

# PET and SPECT of Neurobiological Systems

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Andreas Otte  
Erik F. J. de Vries  
Aren van Waarde  
*Editors*

Adriaan A. Lammertsma  
*Guest Editor*

*Second Edition*

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## Foreword

When in 2014 the editors published the first edition of this three-volume series, dedicated to the use of PET and SPECT in the CNS, a major undertaking saw the light in print. These were significant multi-authored books, providing the most comprehensive review of this challenging field at the time.

Now in 2020, a second edition is launched, demonstrating the success of this initial endeavor. With a further major effort, over 50 % of the numerous chapters are either entirely novel or rewritten anew, by an impressive list of international contributors. The team from Groningen deserve warm congratulations for this achievement.

In 2004, PETMR had just emerged as a novel imaging modality, and the medical applications of machine learning, or artificial intelligence, had hardly surfaced. PET imaging of Tau had just made it.

Now in 2020, amazing progress in the understanding of the brain is being made, and even dedicated brain PET and PETMR imaging instruments are in development (Catana 2019). Yet, with its hundred billion neurons, the brain keeps its mystery and continues to engage and stimulate our enquiry.

Little by little progress is being made, from decoding consciousness (Sohn 2019) to growing neurons from reprogrammed skin fibroblasts (Kofler 2019). We now understand that protein deposits in the brain may precede clinical manifestations by years, great advances have been made in extracting quantitative data from such studies (as shown with PETCT and amyloid and tau numerical data) and that we can intervene effectively, inter alia, in receptor deficiencies and in deep brain stimulation. The brain connectome is unravelling, with it, greater understanding of, among others, pain and drug-induced addictions, the dementias, and the movement disorders. All of this and much more is being critically reviewed by 119 chapters spread into these three major volumes.

It is hence perfect timing that the second edition shines a new light at the significant progress made in PET and SPECT, describing and analyzing latest information from novel biological radionuclide probes, neuroreceptors, and the clinical progress achieved.

Peter Josef Ell  
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## Preface

Neuroscientists of today dispose of a powerful armament for functional, physiological, and molecular imaging that has never made more impressive advances than before, helping to better understand the mechanisms of diseases and to develop and design drug treatment options with a superior efficacy and safety profile. Among these instruments, positron emission tomography (PET) and single-photon emission computed tomography (SPECT) have become forerunners in the *in vivo* imaging arena, and for this reason, the present trilogy, now in its second, completely revised and supplemented edition, is dedicated to PET and SPECT. The volumes of this trilogy are *PET and SPECT in Psychiatry*, *PET and SPECT in Neurology*, and *PET and SPECT in Neurobiological Systems*. In all volumes, we have again assembled the combined expertise of the renowned authors of the first edition and expanded this by some new authors for the second edition, whose dedication to the investigation of psychiatric and neurological disorders or of neurobiological systems through nuclear medicine technology has achieved international recognition.

The editors, who are nuclear medicine specialists, radiochemists, and biologists with a strong exposure to neurosciences, have again invited experts from the psychiatry, neurology, and medical physics fields to enhance the editorial board as guest editors for each volume of the trilogy. For *PET and SPECT in Psychiatry*, this was Iris Sommer, professor of cognitive aspects of neurological and psychiatric disorders; for *PET and SPECT in Neurology*, it remained Klaus (Nico) L. Leenders, emeritus professor of neurology; and for *SPECT in Neurobiological Systems*, it was Adriaan A. Lammertsma, professor of medical physics and positron emission tomography.

We are very happy that our trilogy has become a state-of-the-art compendium with top downloads already for the first edition. This is certainly also due to the production and distribution by one of the premier publishers in the field, guaranteeing a high quality of reproduction and allowing for the inclusion of many color figures, which is essential in the field of neuroimaging. We are intrigued by the enthusiastic response from contributors from all over the world who made this endeavor successful and are confident that the second edition continues to live up to this onus. Finally, we would like to thank Mrs. Gesa Frese from *Springer-Verlag* for her continuous help and support during the development of the second edition of this book series.

We sincerely hope that the new trilogy will still serve as a key tool not only for all physicians in nuclear medicine, psychiatry, neurology, or geriatrics, but also for all professionals working to understand or treat brain disorders. With today's ageing population, PET and SPECT imaging can provide new insights into the processes that may lead to unhealthy ageing of the brain. May this book series serve as a guide towards the present use of PET and SPECT in brain disorders and as a catalyst for future research. The progress achieved by PET and SPECT in the diagnosis of the many facets of diseases disposed in the neurosciences has been astounding. Nevertheless, in line with the Socratic paradox "*I know that I know nothing,*" it seems that we are still at the beginning of understanding the brain. This book hopes to provide a renewed platform to further contribute to this quest at the benefit of patients suffering from neurologic and psychiatric disorders.

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**Part I**  
**Basics**



# Animal Models for Brain Research

1

Debby Van Dam and Peter Paul De Deyn

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## Abstract

Animal models are important experimental tools in neuroscience research since they allow appraisal of selected and specific brain pathogenesis-related questions—often not easily accessible in human patients—in a temporal and spatial pattern. Translational research based on valid animal models may aid in alleviating some of the unmet needs in the current pharmaceutical market. Of primary concern to a neuroscience researcher is the selection of the most relevant animal model to achieve pursued research goals. Researchers are challenged to develop models that recapitulate the disorder in question, but are quite often confronted with the choice between models that reproduce cardinal pathological features of the disorders caused by mechanisms that may not necessarily occur in the patients versus models that are based on known aetiological mechanisms that may not reproduce all clinical features. Besides offering some general concepts concerning the relevance, validity and generalisation of animal models for brain disorders, this chapter focuses in detail on animal models of brain disease, in particular schizophrenia models as examples of animal models of psychiatric disorders and Alzheimer’s disease models as examples of animal models of neurological/neurodegenerative disorders.

## Abbreviations

AChE	Acetylcholinesterase
AD	Alzheimer’s disease
APP	Amyloid precursor protein
A $\beta$	Amyloid $\beta$
BPSD	Behavioural and psychological signs and symptoms of dementia
ChAT	Choline O-acetyltransferase
DISC1	Disrupted in schizophrenia-1 gene
DSM	Diagnostic and Statistical Manual of Mental Disorders
DTNBP1	Dysbindin gene
ERBB4	Neuregulin 1 receptor gene
FRL	Flinders resistant line
FSL	Flinders sensitive line
HAB	High-anxiety-related behaviour Wistar rat line
LAB	Low-anxiety-related behaviour Wistar rat line
LI	Latent inhibition
NFT	Neurofibrillary tangle
NGF	Nerve growth factor
NMDA	N-methyl-D-aspartate
NPS	Neuropsychiatric symptoms
NRG1	Neuregulin 1 gene
PCP	Phencyclidine
PDAPP	Platelet-derived growth factor promoter-driven APP

---

PPI	Prepulse inhibition
PSEN	Presenilin
RDoC	Research Domain Criteria
RELN	Reelin gene
SAM	Senescence-accelerated mouse
SAMP	SAM-prone substrain
SNP	Single nucleotide polymorphism
TDP-43	TAR DNA-binding protein 43

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## 1.1 Introduction to Animal Modelling for Human Brain Disease

Animal models aiming at studying human disorders emerged in the 1800s and experienced a major boost over the last decades. The value of animal experimentation in the advances of human health is exemplified by the list of Nobel Prizes awarded for Physiology or Medicine. Since the beginning of the twentieth century, these prizes have chronicled the world's greatest medical advances. Of the 109 Nobel Prizes awarded for Physiology or Medicine up to 2018, at least 87 were directly dependent on animal-based research or the discovery relied on crucial data obtained from animal studies by other research groups.

Basic neuroscience research on animal models is essential to understand the nature of brain disorders that afflict humans and develop purposeful therapies. In the mid-1960s, the neural circuits containing and utilising the neurotransmitters dopamine and norepinephrine were visualised for the first time in rodent brain (Glowinski et al. 1966; Glowinski and Iversen 1966a, b; Iversen and Glowinski 1966). Further development of other (antibody-based) methods allowed neuroscientists to link synaptic molecules to neural circuits in rodent and nonhuman primate brains, thereby coordinating neuroanatomy, neuropharmacology and neurophysiology into a cohesive view of biochemically coded brain circuitry. Application of these techniques in human brain aided in elucidating the neurotransmitter-based nature of specific brain circuits affected in disease, e.g. the degeneration of cholinergic neurons in the basal forebrain of Alzheimer's disease (AD) patients (Whitehouse et al. 1982). The concept of synaptic communication between nerve cells as the key level of interaction in the brain further culminated with the work of E. Kandel and colleagues, who were awarded the Nobel Prize in 2000. More recently, the 2013 Nobel Prize in Physiology or Medicine was awarded jointly to J.E. Rothman, R.W. Schekman and T.C. Südhof for their discoveries of machinery regulating vesicle traffic, a major transport system in our cells. The synapse has become a key target for pharmaceutical intervention in many brain disorders that are now understood at the level of synaptic circuitry and functional brain organisation, including multiple psychiatric disorders (e.g. schizophrenia, depression), as well as neurodegenerative disorders (e.g. AD, amyotrophic lateral sclerosis, Parkinson's disease).



The starting point for the development of a new animal model for a specific brain disease is mostly the current dominant theory about the disorder. Although a logical first approach, it is essential to broaden the focus of animal models under development based on the increasing knowledge of underlying disease mechanisms.

Animal models of human disease can be classified into spontaneous, induced, negative and orphan models, of which the two latter types are in general not relevant for psychiatric and neurological disorders. Negative models after all are models in which a specific human disorder does not develop, while orphan models display a condition which has never been described in man or other target species. Spontaneous models are presumed to develop their condition without experimental manipulation, but selective breeding is often compulsory to establish and maintain the desired line. Especially for psychiatric and neurological conditions, few spontaneous models exist, and experimentally induced pathology is often required. Such artificial manipulations include surgical procedures, administration or withdrawal of biologically active substances, application of physical or physicochemical stimuli and genetic manipulations.

### 1.1.1 Relevance of Animal Models

Relevance reflects the meaningfulness and usefulness of results obtained with an animal model for a particular scientific and/or clinical purpose (van der Staay 2006). The two most important clinically inspired applications of animal models in neuroscience research are (1) the development and evaluation of mechanistic hypotheses about neurological and psychiatric disorders in general and their neural substrates in particular (i.e. brain–behaviour relation) and (2) the identification and screening of novel therapeutic approaches, most frequently drugs.

Opponents of animal research often question the relevance of animal models to understanding diseases and disease processes in humans. The disputed similarities between human and other species at the structural and functional level of the CNS are often considered a major hurdle for the use of animal models in studies of human neurological and psychiatric conditions.

Neuroscientists are far from reaching consensus about the level of similarity between the brains and ‘minds’ of humans and other species. Comparative neuroscience can provide essential information for the adequate comparison of both structural and functional aspects of human and nonhuman brain and thereby identify research areas in which animal models are likely to be useful, as well as the appropriate species for such studies. Specifically when considering models of psychiatric disease, the issue of similarities between the mental lives of animals and humans is fundamental. Animal consciousness or unconsciousness is difficult, if not impossible, to corroborate. On the other hand, one could argue whether it is necessary for the animal mind to be as complex as the human mind. Behavioural responses and brain mechanisms associated with anxiety, fear, aggression and coping serve such a crucial adaptive function that they must have evolved very early in the development of mammalian species and are probably highly conserved (Steimer 2011).

Other classes of vertebrates and even lower invertebrates may display some form of anxiety and have the capacity to detect danger and react to threat, i.e. display a fight-or-flight response. Components of emotional processes, as well as the biological and neuronal logistic systems essential for their realisation, gain in complexity higher up in the phylum. Species located higher in the phylogenetic tree gain additional capabilities (e.g. autothetic consciousness in humans, orang-utans, chimpanzees and bonobos), but never lose the more primitive abilities shared with lower invertebrates (Belzung and Philippot 2007), thereby opening perspectives for the use of lower species like zebrafish (Gerlai 2010), or even fruit flies (Iliadi 2009), in the studies of emotional responses and specific psychiatric disorders. The further focus of this chapter will be, however, on rodent models given their widespread use and applicability of microimaging techniques in these species.

Although some attributes still appear to be rather unique to the human brain, other aspects seem to be shared with more species than originally believed. Also at the neuroanatomical level, major species differences should be taken into consideration. The complex pattern of human cortical sulci and gyri, as well the extent of the prefrontal cortex, the neural substrate of many higher functions, is beyond compare in the rest of the animal kingdom (Fuster 1980). The prefrontal cortex may be present, however, in more species than initially thought, and the prefrontal to nonprefrontal cortical surface ratio in itself may not be an optimal index of evolution. Improved knowledge of rat prefrontal cortex anatomy has modified strategies to manipulate specific cortical areas to model prefrontal involvement, although reports delineating boundaries between prefrontal cortical subareas of the rat in comparison with the primate homologues show some dissimilarities. Nevertheless, current anatomical and functional data indicate that rats do have a prefrontal cortex, although not as differentiated as the primate prefrontal cortex. Rats have a functionally divided prefrontal cortex that includes not only features of the medial and orbital areas in primates, but also some features of the primate dorsolateral prefrontal cortex (Uylings et al. 2003).

### 1.1.2 Validity of Animal Models

The conclusions drawn from animal models largely depend on the validity of the model in representing the human condition. The more levels of validity a model satisfies, the greater its value, utility and relevance to the human condition. The validity of a model can only be determined in a multidisciplinary approach. A 'perfect' model would account for aetiology, symptomatology, treatment response and physiological basis, as originally proposed by McKinney and Bunney in 1969 (McKinney Jr and Bunney Jr 1969). Animal models in general do not meet all of these criteria. Moreover, in developing and assessing an animal model, it is critical to consider the explicit purpose intended for the model, because the intended purpose determines the criteria that the model must satisfy to establish its validity. The following perspectives are commonly used in the characterisation of a model's validity (Van Dam and De Deyn 2006, 2011a).

### **1.1.2.1 Face Validity**

Face validity refers to the resemblance between the animal model and the situation or process being modelled. It refers to the phenomenological similarity between the model and the human condition. Similarity of symptoms is most commonly used to assess face validity in animal models of behavioural dysfunction. Throughout this chapter, several behavioural paradigms will be mentioned. Writing a complete ‘how-to manual’ for the behavioural phenotyping of a new brain disease model would go beyond the scope of this chapter. For further reading, we would like to refer the readers to excellent manuals on behavioural neuroscience as, e.g. ‘What’s wrong with my mouse?’ (Crawley 2000) and ‘Methods of Behavior Analysis in Neuroscience’ (Buccafusco 2008).

### **1.1.2.2 Construct Validity**

Construct validity refers to the theoretical clarification of what a test measures or a model is supposed to mimic. Because a given condition may manifest itself in different ways in different species, the behaviour used in the animal model may not match that of humans, yet the test or model may still be valid. Construct validation is useful in the incessant process of further developing and refining an animal model.

### **1.1.2.3 Aetiological Validity**

Aetiological validity is closely related to construct validity and refers to identical aetiologies or phenomena in the model and the human condition. Models with high aetiological validity are most valuable in drug development and discovery.

### **1.1.2.4 Predictive Validity**

Predictive validity represents the extent to which the performance of the animal model in a test predicts the performance in the condition being modelled. This level of validation necessitates parallel development of clinical measures for meaningful comparisons between model and man. In a more narrow sense, this term is sometimes used to indicate pharmacological isomorphism, i.e. the model’s ability to identify compounds with potential therapeutic effects in the human condition.

## **1.1.3 Homology, Analogy and Isomorphism**

As a consequence of the different validation levels, validity is highest in the so-called homologous models, where the symptoms displayed and the cause of the condition in the animal are identical to those of the human condition. Especially in neurosciences, homologous models are very rare and mostly limited to well-defined lesion syndromes. Analogous or isomorphic models are more common, but, although they display similar symptoms, the condition is not provoked by the same events as the human condition. In reality, most animal models are partial models, focusing only on restricted aspects of a disease, and modelling the complete condition is often not pursued. Of primary concern to neuroscientists is the selection of

the most relevant animal model to achieve his/her research goals. Researchers are challenged to develop models that recapitulate the disorder in question, which is often not as straightforward as it may seem. Quite often they are confronted with the choice between models that reproduce cardinal pathological features of the disorders caused by mechanisms that may not necessarily occur in the patients versus models that are based on known etiological mechanisms that may not reproduce all clinical features. Although animal models cannot replicate human pathophysiology or psychopathology in every detail, we should envisage them as experimental systems in which selected and specific CNS pathogenesis-related questions—often not easily accessible in human patients—can be studied in a temporal and spatial pattern. Partial models may be of substantial value in the gradual process of building a more complete image of the disease and underlying pathophysiological mechanisms, whereas more complete and complex models are only possible in case of better understood diseases (De Deyn and Van Dam 2011).

### 1.1.4 Generalisation and Extrapolation

The ultimate goal of animal modelling is the generalisation of results and insights to the target species, which in most occasions of course is man. The term extrapolation is also often used to describe how data obtained from animal studies can be reliably used to apply to the human condition. Extrapolation is therefore not performed in its mathematical sense, in which data fit a certain function that may be described graphically and the graph extends beyond the highest or lowest sets of data to describe a situation outside the window of observation. Generalisation across species is intrinsically based on the Darwinian evolutionary concept. Phylogenetic similarity in morphology and physiology between different species allows observations made in animals to be translated to the human condition based on homology. Related to animal models for psychiatric and neurological disorders, this cross-species comparison may focus or be based on the biological, genetic or environmental basis of personality traits (Gosling 2001), or neuroanatomical structures and their function (Striedter 1998). Uncritical generalisation or extrapolation of animal findings to the human condition may lead to unreliable or even dangerous conclusions. As a general rule, extrapolation tends to be most reliable when a plurispecies approach is taken and the condition is studied in a variety of relatively unrelated laboratory animals. Differences in metabolic patterns and speed, as well as several other confounding variables (e.g. age, sex, distress, diet, housing condition, route of administration, rhythmic variations), need to be taken into account, particularly when quantitative extrapolation is intended (De Deyn and Van Dam 2011).

Generalisation may be advanced by the development of translatable preclinical biomarkers, including small animal imaging (Kaimal and McConville 2009; Sakoğlu et al. 2011), and biochemical assays on biological fluids such as plasma and cerebrospinal fluid (Liu and Duff 2008). These biomarkers could be used to characterise the translatability of animal models, determine the translatability of a

novel therapeutic intervention if the same biomarker can be used in a clinical trial, assess target engagement of investigational treatments and monitor biological responses to treatment in real time.

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## 1.2 Animal Models of Psychiatric Disorders

Psychiatric disorders represent a diverse set of conditions, variously affecting all domains of mental function and affecting the most fundamental human attributes, namely, language, thought, perception, mood and sense of self. Many psychiatric symptoms, e.g. hallucinations, delusions, sadness, guilt and suicidal thoughts, are probably uniquely human and cannot be convincingly ascertained in animals. Researchers should also refrain from uncritical anthropomorphising of emotional behaviour in general. Feeling anxious or depressed are subjective emotional experiences that may require a certain minimal level of consciousness that may well be uniquely human.

The evaluation of animal models of neuropsychiatric disorders is further challenged by the use of the Diagnostic and Statistical Manual of Mental Disorders (DSM), criteria in the clinical diagnosis, combined with the lack of knowledge of pathophysiological mechanisms and objective diagnostic tests. Both in the fourth (American Psychiatric Association 2000) and fifth edition (American Psychiatric Association 2013) of the DSM, the diagnosis is based on phenomenology (i.e. signs, symptoms, course of illness), and mostly, a minimal number of symptoms should be present for at least a specified period of time. In the case of schizophrenia, two (or more) of the following symptoms should be present for a significant portion of time during a 1-month period (or less if successfully treated): delusions, hallucinations, disorganised speech, grossly disorganised or catatonic behaviour and negative symptoms, i.e. affective flattening, alogia, or avolition, in addition to social/occupational dysfunction (American Psychiatric Association 2000, 2013). Two individuals with the same DSM diagnosis of schizophrenia may therefore present with extremely heterogeneous symptom combinations. Similar problems exist for most DSM-based diagnoses, thereby further complicating the evaluation and validation process of newly developed mouse models for psychiatric illnesses (Nestler and Hyman 2010).

### 1.2.1 The Endophenotype Concept in Psychiatry

The concept of endophenotypes appears to be unique to the broad field of psychiatry. An endophenotype is a biomarker associated with genetic components, as well as the clinical symptoms of neuropsychiatric disorders. In that way they differ from biomarkers in general that often lack the genetic link. Endophenotypes play an important role to bridge the gap between the microscopic and macroscopic level of neuropsychiatric disorders. It is per definition any hereditary characteristic that is normally associated with some condition but is not a direct symptom of that condition. Criteria useful for the identification of endophenotypes of diseases that display

complex inheritance patterns have been suggested (Gottesman and Gould 2003; Gould and Gottesman 2006):

1. The endophenotype is associated with illness in the population.
2. The endophenotype is heritable.
3. The endophenotype is primarily state-independent (manifests in an individual whether or not illness is active), but age-normed and may need to be elicited by a challenge.
4. Within families, endophenotypes and illness co-segregate.
5. The endophenotype found in affected family members is found in nonaffected family members at a higher rate than in the general population.

Endophenotype analysis can be based on neuroanatomical, neurophysiological, biochemical, endocrinological, cognitive and neuropsychological/behavioural measures (Leboyer et al. 1998). Advanced neuroimaging techniques, including PET and SPECT, have left little doubt that specific structural CNS anomalies can be linked to particular neurological and psychiatric diseases. Neuroimaging endophenotypes are vitally important to the identification of genetic determinants of complex brain-related disorders (Glahn et al. 2007).

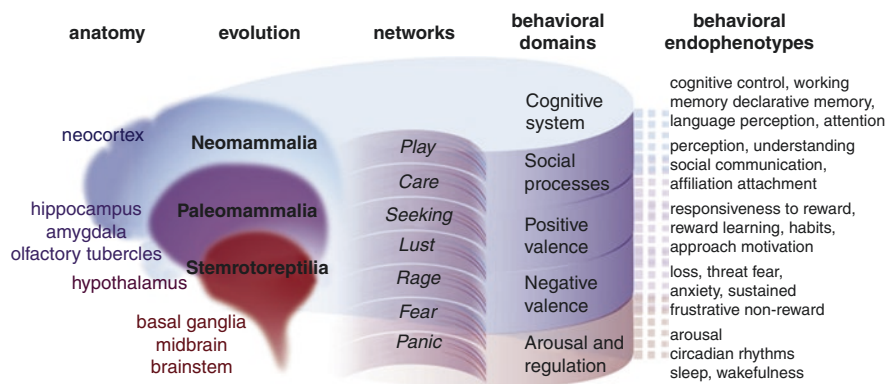
Especially with regard to behavioural measures, the endophenotype approach is considered particularly relevant in the context of the Research Domain Criteria (RDoC) framework from the National Institute of Mental Health for new approaches to investigate psychopathologies by bridging basic neuroscience and mental health and supporting the concept of a normality and pathology continuum. According to the RDoC system, behavioural endophenotypes attribute five main domains, including arousal and regulation, negative valence, positive valence, social processes and cognitive systems. Changes at all neurobiological levels, ranging from molecules to brain networks, are related to the specified behavioural domains (Anderzhanova et al. 2017; Maximino and van der Staay 2019) (Fig. 1.1).

The study of endophenotypes is particularly useful to understanding underlying disease mechanisms in neuropsychiatric disorders, essential for the accurate diagnosis, classification, early detection purposes, the development and evaluation of valid animal models and accelerating the drug discovery pipeline concerned.

## 1.2.2 Approaches to Modelling Psychiatric Disease

### 1.2.2.1 Behavioural Approach

In the absence of knowledge of underlying genetic causes, the first attempts to model psychiatric disturbances in animals consisted of the behavioural approach that defined behavioural assays thought to mimic particular psychiatric symptoms. Animals with specific behavioural deficits were generated through pharmacological, surgical or genetic (e.g. comparison of inbred strains) procedures. Well-known examples are the learned helplessness model of depression, spontaneous alternation indexing working memory in schizophrenia, decreased social interaction linked to



**Fig. 1.1** According to the RDoC system, behavioural endophenotypes attribute five main behavioural domains. These five domains and their respective constructs can be used to comprehensively profile behaviour (i.e. behavioural endophenotypes). They are not only distinct but describe personality as an interacting and dynamic individual trait. The domains are objectively different, determined by particular neuronal networks. Separation of the networks is due to a distinction in the functions they serve (The Affective Neuroscience Scale, Panksepp, 1998). This distinction corresponds well with the concept of a gradual evolution of the brain (the triune brain model of MacLean 1970). Legend: RDoC, Research Domain Criteria. Adapted from Anderzhanova et al. (2017) *Neurobiol Stress* 7:47–56 (Anderzhanova et al. 2017) under the CC-BY license 4.0 (<http://creativecommons.org/licenses/by/4.0>). Copyright © 2017 Anderzhanova, Kirmeier and Wotjak

autism spectrum disorder and approach–avoidance conflict behaviours associated with anxiety disorders (Crawley 2000; Mitchell et al. 2011). The development of valid behavioural models and paradigms is currently still ongoing, but with more attention for the neurobiological and clinical relevance (Baker 2011). A rodent behavioural paradigm has neurobiological construct validity if it isolates and measures a behavioural process with similar environmental regulators and neural mediators as the human process of interest. However, many behavioural assays lack sufficient construct validity, and their use in the development of rodent models for psychiatric disorder is mainly based on face validity (Moore 2010). These translational challenges are—at least partially—tackled by the development of reverse translational tools that take the rodent paradigm as a starting point to develop a human variant that allows appraisal of construct validity. Relevant examples include human real-life or virtual reality analogues of visual spatial learning and memory as tested in a water maze (Laczó et al. 2010; Woolley et al. 2013; Fajnerová et al. 2014), or a tool to assess unconditioned locomotor behaviour similarly as in rodents (Young et al. 2007).

Although brain–behaviour relationships are assumed to be conserved across evolution for many aspects of normal motor and sensory functions, this may not be translated as easily to abnormal behaviour, let alone psychiatric disease. Animal brain circuitry and neurochemical systems involved may differ from the complex human (disease) system. A mental disorder is not characterised by an aberration in one single feature, but is a syndrome with a myriad of deviations in behaviour,

cognition and/or emotion. Behavioural models of psychiatric disease, like the learned helplessness model of depression, are of course partial models that mimic only one single isolated feature of the human condition (Insel 2007).

### 1.2.2.2 Pharmacological Models

The administration of certain chemicals may induce behavioural changes reminiscent of specific psychiatric symptoms based on interference with central neurotransmission pathways. Understanding the mechanisms of these actions might contribute to our fundamental understanding of the illness. Drug-based models include the reserpine model of depression (Hendley and Welch 1975) and N-methyl-D-aspartate (NMDA) receptor antagonist-based models of psychosis (Andiné et al. 1999).

### 1.2.2.3 Genetic Models

Genetic models of psychiatric disorders were initially dominated by inbred strains, often after selection for a relevant psychiatric-like phenotype. Examples include the selectively bred Flinders Sensitive Line and the Flinders Resistant Line of rats (FSL versus FRL), a genetic animal model of depression (Overstreet et al. 1989); selectively bred Wistar rat lines with high- versus low-anxiety-related behaviour (HAB versus LAB) (Liebsch et al. 1998); and the continuous distribution of PPI (see paragraph on “Sensory Discrimination” in Sect. 1.2.3.2) response amongst mouse inbred strains, which indicates PPI to be a polygenetic trait (Willott et al. 1994; Bullock et al. 1997; Paylor and Crawley 1997). The success of inbred and selectively bred strains was closely followed by single-gene models developed through homologous recombination in embryonic stem cells, as illustrated by neurokinin-1 receptor knockout mice that display significantly reduced anxiety and stress-related responses (Santarelli et al. 2001), and the glutathione S-transferase M1-knockout mouse model for autism (Yochum et al. 2010). Based on information obtained from genome-wide association studies, single-gene knockout or mutation models will undoubtedly maintain a stable position in the field of psychiatry models. They aid in assessing the putative link between gene and disorder using increasingly powerful and sophisticated tools to elucidate underlying neurobiology and disease mechanisms from the level of synapses over brain circuitry to behavioural outcome (Mitchell et al. 2011). With the advancement of genetic engineering techniques, the number of genetic mouse—and even rat—models is strongly increasing. The development of genetic rat models is of specific value for the study of psychiatric disorders, since they display more complex and sociable behaviour compared to mice. Genetic possibilities in rodents include full knockouts, spatial and/or temporal knockouts, knock-in of human alleles, and precisely engineered chromosomal rearrangements (e.g. inversions, deletions, translocations or duplications) into syntenic animal genes (van der Weyden and Bradley 2006). Although novel molecular tools, like the bacterial CRISPS/Cas9 system, mainly promote *in vitro* genome editing, the ability of CRISPR/Cas9 to directly target any gene of interest in the embryonic genome holds great promise to faster, less expensively and more reliably generate *in vivo* psychopathology models (Tschaharganeh et al. 2016).



The theory that genes and environment combine to confer susceptibility to the development of diseases surfaced in the early half of the last century, but the use of such a framework for exploring the aetiology of psychiatric disorders is more recent. Given the fact that many—if not all—psychiatric disorders are multifactorial, sophisticated genetically engineered animal models can of course also be used to disentangle complex aetiological mechanisms with the appraisal of gene–environment interactions. For example, the effect of repeated maternal separation or exposure to pesticides early in ontogeny on the reelin-deficient mouse, i.e. the reeler model that has been linked to autism and schizophrenia, has been studied (Laviola et al. 2009).

### 1.2.3 Animal Models of Schizophrenia

#### 1.2.3.1 Aetiology and Symptomatology of Schizophrenia

Schizophrenia is a devastating and common psychiatric disorder that is associated with a high degree of medical morbidity and reduced life span. Both incidence and prevalence estimates vary across the world, and even within countries, and at the local and neighbourhood level. The median lifetime prevalence was estimated at 4/1000 and the lifetime morbid risk at 7.2/1000, while the median incidence was estimated at 15.2/100,000 individuals. Incidence rates differ between sexes with a male to female ratio of 1.4:1 (McGrath et al. 2008). These variations are supportive of the assumption that schizophrenia is a clinical syndrome, perhaps comprising several disease entities.

Symptoms cluster into three categories: positive symptoms (including hallucinations, delusions, thought disorder and conceptual disorganisation), negative symptoms (including emotional blunting, social withdrawal, anhedonia, avolition, poverty of thought and content of speech) and cognitive dysfunction (including impaired executive functioning, working memory and attention) (Andreasen 1995). The onset of some aspects of the disease may be observed from birth onward, but psychotic symptoms generally become manifest in late adolescence and early adulthood in males, with an extended onset period in females (McGrath et al. 2008; Carpenter and Koenig 2008).

The dopamine hypothesis of schizophrenia proposes that dysfunction in dopaminergic neurotransmission, specifically hyperactivity of dopaminergic neurons in the limbic system and striatum, produces the positive symptoms, while underactive mesocortical dopaminergic neurons cause the negative, cognitive and affective symptoms of schizophrenia (Seeman 1987). Dopamine dysfunction and its treatment are not sufficient to explain the psychopathology of schizophrenia and its treatment outcomes (Yang and Tsai 2017; Stepnicki et al. 2018). Besides the dopaminergic hypothesis, dysfunction of the glutamatergic system has been implicated in schizophrenia (Olney and Farber 1995; Tsai and Coyle 2002). Glutamate is the major excitatory neurotransmitter in the CNS. The glutamate hypothesis posits that the function of the NMDA receptor is compromised in this disease. NMDA

receptors mediate slow excitatory postsynaptic potentials, which are considered critical for the proper expression of complex behaviours, such as associative learning, working memory, behavioural flexibility and attention, many of which are impaired in schizophrenia. These neurochemical hypotheses have been further extended to include contributions of other neurotransmitter systems in the pathophysiology of schizophrenia (Yang and Tsai 2017; Stepnicki et al. 2018). The serotonin (5-hydroxytryptamine; 5-HT) hypothesis of schizophrenia is derived from early studies of interactions between two major classes of psychedelic hallucinogens, the indoleamines (e.g., lysergic acid diethylamide, LSD) and phenethylamines (e.g., mescaline) and the serotonergic system, i.e. via 5-HT<sub>2A</sub> receptor agonisms (Aghajanian and Marek 2000). Although no direct evidence of serotonergic dysfunction in the pathogenesis of schizophrenia is yet available, specific 5-HT receptors continue to be a focus of interest in schizophrenia (Yang and Tsai 2017; Stepnicki et al. 2018). Of relevance in this matter is of course the serotonergic inhibitory modulation of dopaminergic neurotransmission in midbrain and forebrain (Kapur and Remington 1996), and the 5-HT-related enhancement of glutamatergic transmission in for example frontal cortex (Aghajanian and Marek 2000). Gamma-aminobutyric acid (GABA) is the main inhibitory neurotransmitter in the CNS. GABAergic dysfunction leads to an imbalance between inhibitory and excitatory activity in cortical areas which has been implemented in the pathophysiology of schizophrenia (Yang and Tsai 2017; Stepnicki et al. 2018; Guidotti et al. 2005). The high prevalence of smoking in schizophrenia, motivated the evaluation of nicotinic acetylcholine receptors, but also muscarinergic receptors are, respectively, potential targets for cognitive symptoms and positive symptoms (Yang and Tsai 2017; Stepnicki et al. 2018; Raedler et al. 2007; Brunzell and McIntosh 2012). In addition, several other systems including the histaminergic, adrenergic and endocannabinoid systems, as well as various neuropeptides have been implicated in the pathomechanisms of schizophrenia (Yang and Tsai 2017; Stepnicki et al. 2018).

### 1.2.3.2 Validating Animal Models of Schizophrenia

When validating animal models of schizophrenia, behavioural and electrophysiological endophenotypes are commonly employed, in addition to alternative endophenotypes focussing on neurochemical alterations, e.g. brain dopamine levels (Winter et al. 2009; Ayhan et al. 2011), or other neurochemical readouts (Corcoba et al. 2015; Puhl 2015), in addition to neuroanatomical changes, like reduction of brain (region) volume (Ayhan et al. 2011; Drew et al. 2011), ventricular enlargement (Ayhan et al. 2011; Corcoba et al. 2015; Puhl 2015; Li et al. 2009), and dendritic spine density (Ayhan et al. 2011; Berlanga et al. 2011; Jaehne et al. 2015; Nakao 2017).

#### Electrophysiological Endophenotypes

Electroencephalography recordings have been widely used to assess sensory processing deficits in schizophrenia. Analogous human and rodent auditory-evoked event-related potentials that are similarly affected by pharmacological interventions

and stimulus manipulations were described. The human auditory-evoked potential waveform shows characteristic positive deflections 50, 200 and 300 ms poststimulus (respectively, P50, P200 and P300) and a negative deflection 100 ms poststimulus (N100). The rodent analogues are P20, N40, P80 and P120 in order of occurrence. When exposed to a paired-click paradigm (S1–500 ms interval—S2), the waveform amplitudes are lower after S2 than after S1. Schizophrenia patients display a gating deficit based on a decreased response to S1 and/or a failure to inhibit S2 that results in similar amplitudes after S1 and S2. Useful endophenotypes with acknowledged heritability include decreased P50 (P20), N100 (N40) and P200 (P80) amplitude and gating, as well as decreased P300 (P120) amplitude. Schizophrenia patients also show a reduced ability to detect changes in the auditory environment, which is measured with a mismatch negativity paradigm, an exaggerated negative voltage deflection following N100 (N40), elicited when the qualitative features of a novel tone fail to match the pattern of a previous series of repetitive tones. Electrophysiological endophenotypes may aid in understanding schizophrenia mechanistically as gene and neurobiological associations have been described (Amann et al. 2010; Rosen et al. 2015; Owens et al. 2016).

### **Cognitive Endophenotypes**

The cognitive phenotype can be evaluated using various protocols and paradigms (Crawley 2000), including a wide range of dry land (e.g. radial arm maze) and water mazes (e.g. Morris water maze) to specifically assess working memory. Contextual and fear conditioning are based on learning the association of a nonaversive context or cue with an aversive stimulus and the evaluation of freezing responses. The novel object recognition task is based on the rodents' tendency to focus attention on novel objects in their surroundings and the evaluation of exploration of a familiar versus a new object. Impairments in different forms of behavioural flexibility are commonly associated with schizophrenia. A number of behavioural paradigms using different sensory modalities and classes of stimuli have been developed to assess behavioural flexibility, in particular reversal learning, in rodents (Floresco et al. 2009).

### **Locomotor Activity**

The majority of early tests for schizophrenia-related behaviours in rodents were based on the antipsychotic activity of dopamine D2 receptor antagonists (Ellenbroek and Cools 1990), with behavioural outcome parameters including locomotor hyperactivity and stereotyped sniffing and/or grooming. Locomotor activity can be scored in an open-field arena equipped with a video tracking system or infrared sensors to detect spontaneous exploration and ambulatory horizontal locomotion over a specific time frame (Crawley 2000). Motor stereotypy is mostly scored in rodents employing the Creese–Iversen stereotypy scale ranging from 0 (mouse either sleeping or inactive) to 6 (mouse engaged in continuous and nonstop route-tracing stereotypic behaviours) (Creese and Iversen 1973). Repeated administration of psychostimulants, including amphetamine, causes sensitisation to the motor-activating properties, as exemplified by enhanced hyperlocomotion and increased neuronal release of dopamine after the second or third administration. Behavioural

sensitisation is quantified by repeatedly testing spontaneous ambulatory locomotion in an open-field arena (Crawley 2000).

### Sensory Discrimination

Abnormalities in the automatic processes of sensory discrimination, orienting and reorienting of attention are evident in the early phases of schizophrenia. Two attentional tasks were developed to be used both in humans and rodents: latent inhibition (LI) and PPI of the startle reflex. LI is the retarded conditioning to a stimulus that is repeatedly presented without reinforcement. In other words, the prior experience that a stimulus does not have a consequence makes it less likely that the brain will form an association with that stimulus later. LI is widely considered to relate to the cognitive abnormalities that characterise schizophrenia because it reflects an organism's ability to ignore irrelevant stimuli. One major strength of the LI task is that it can be applied across (mammalian) species (Amann et al. 2010; Lubow and Gewirtz 1995).

PPI of the startle reflex is a neurophysiological and behavioural measure of sensorimotor gating. One of its primary advantages is its ability to translate between mice and humans, as it is one of the few tests that is largely conserved across all vertebrate species. In rodents, the startle response is typically evoked using either acoustic or tactile stimuli and is characterised by contractions of the major muscles of the body, generally leading to extension of the forepaws and hind paws followed by muscle flexion into a hunched position. PPI is evoked when a weak stimulus (e.g. a 70- to 80-dB tone or a puff of air) inhibits the subsequent startle response to a strong stimulus (e.g. a 120-dB tone or a second puff of air) if presented within 100 ms. The whole-body startle from the rodent, restrained in a cylindrical holder, is quantified with piezoelectric motion sensors. Abnormal sensory inhibition may reflect a deficit in processing and prioritising incoming sensory information (Crawley 2000; Amann et al. 2010). A genetic component underlying PPI was suggested based on inbred mouse strain differences (Paylor and Crawley 1997).

### Negative Symptoms

Certain negative symptoms, like poverty of speech, may be uniquely human, but other negative symptoms can be assessed in rodents using relatively simple behavioural paradigms. Anhedonia is studied with the sucrose preference test, which measures reduction in reward function in rodents (Crawley 2000). Deficits in social functioning is best scrutinised in rodents using a range of behavioural protocols, which may include a social choice paradigm for spontaneous affiliative behaviour, assessments of social recognition–discrimination, play-soliciting behaviour, social grooming, dominance, aggression and social defeat. Other negative symptoms, including avolition and blunted affect, have been studied in rodents using respectively behavioural despair tests (forced swim test and tail suspension test) and general anxiety tests, but interpretation is rather controversial (O'Tuathaigh et al. 2010). The controversy surrounding behavioural despair tests (Reardon 2019; Trunnell 2019), quite recently lead to the ban of the FST by several large pharma companies.

### 1.2.3.3 Neurodevelopmental Schizophrenia Models

Epidemiological evidence indicates gestational or perinatal exposure to adverse environmental events, like maternal stress, malnutrition and infection, and obstetric complication (e.g. hypoxia) increases the risk of developing schizophrenia (Lewis and Levitt 2002). Exposure to early-life adverse events combined with genetic predisposition is hypothesised to trigger deviating neuronal development and connectivity leading to the development of schizophrenia. When developing animal models based on this neurodevelopmental hypothesis, the timing of the exposure to a certain adverse event during the sensitive perinatal period is critical. Various adverse events have been applied, including postweaning social isolation (Fone and Porkess 2008), sometimes combined with physical stressors like e.g. alternating warm and cold temperature exposures (Hohmann et al. 2012), gestational exposure to the DNA-alkylating agent methylazoxymethanol acetate (Lodge and Grace 2008; Gulchina et al. 2017), exposure to bacterial or viral infection of pregnant rats (Gayle et al. 2004; Buka et al. 2008), or maternal exposure to stress (Martínez-Télez et al. 2009; Markham et al. 2010). Depending on the chosen procedure and timing, these models may display, amongst others, disrupted sensory gating, decreased social interaction, cognitive deficits, enhanced drug-induced locomotion, neuroanatomical, neurostructural and neurochemical alterations.

### 1.2.3.4 Drug-Induced Schizophrenia Models

The development of an animal model is often based on the observation that certain drugs produce abnormal behaviour in healthy people. Usually the following steps are included in the process of developing an animal model based on pharmacological compounds: (1) the drug in question induces behavioural effects in healthy people that resemble human psychiatric pathology; (2) the drug induces specific biochemical alterations in animals; (3) the drug produces similar behavioural effects in animals, allowing for any species differences in behaviour; (4) the biochemical observations in animals provide insight into the behavioural effects of the drug in humans; and (5) if the behavioural effect of the drug in humans has the same characteristics as pathological behaviour, then the biochemical changes produced by the drug in animals may also provide data relevant for the understanding of the abnormal human behaviour. The vast majority of schizophrenia-related models have been based on the induction of abnormal behaviour by psychotomimetics, including both psychostimulants and hallucinogens. These drug-induced states, however, resemble the early stages of a range of psychotic disorders, and not necessarily the diagnostic syndrome of schizophrenia.

#### Psychostimulant Models

The most widely studied class of drug-induced animal models of schizophrenia is based on behavioural effects of psychostimulant drugs, such as amphetamine. Amphetamine-induced psychosis has been used as a model to support the dopaminergic hypothesis of schizophrenia (Seeman 1987). A hypodopaminergic state in the frontal-cortical terminal fields of mesocortical dopaminergic neurons has also been proposed to be the basis of negative symptoms (Dworkin and Opler 1992), but

these are relatively rare in amphetamine-induced psychosis, due to the hyperdopaminergic nature of this model. One of the primary consequences of amphetamine treatment is the release of catecholamines from nerve terminals. In addition to its effects on release, amphetamine also blocks the reuptake of catecholamines from the synaptic cleft, and in high dosages, it is a potent monoamine oxidase inhibitor (Kokkinidis and Anisman 1981).

In rodents, chronic amphetamine administration induces a persistent sensitisation that, both behaviourally and neurochemically, mirrors several features linked to the positive symptoms of schizophrenia. Behavioural alterations include locomotor hyperactive and/or stereotypy or perseverative behaviours, reduced PPI and disrupted LI (Featherstone et al. 2007). While amphetamine sensitisation does not produce memory impairments similar to those seen in schizophrenia, it does produce strong impairments in set-shifting, suggesting schizophrenia-like changes in prefrontal functioning (Featherstone et al. 2007, 2008). Interestingly, rhesus monkeys displayed behavioural alterations reminiscent of schizophrenia-like hallucination, including tracking, grasping 'at thin air', manipulating nonapparent stimuli (e.g. picking at imaginary parasites) and hypervigilance after amphetamine administration (Castner and Goldman-Rakic 1999, 2003).

### Hallucinogen Models

NMDA receptor antagonists, such as phencyclidine (PCP, 'angel dust') and ketamine, produce psychotic symptoms and cognitive disturbances reminiscent of schizophrenia (Cohen et al. 1962; Krystal et al. 1994), findings that have contributed to the hypoglutamatergic hypothesis of schizophrenia. As opposed to amphetamine-induced psychosis, hallucinogen-induced psychosis encompasses negative symptoms (e.g. emotional withdrawal) as well. Positive schizophrenia-like symptoms induced in laboratory animals include hyperlocomotion, stereotyped movements, circling and ataxia, while deficits in social behaviour are reminiscent of negative symptoms typical for schizophrenia. PCP and PCP-like drugs impair sensorimotor discrimination (LI and PPI) and affect performance in several cognitive paradigms, both in rats and monkeys (Javitt and Zukin 1991; Jentsch and Roth 1999). Also the serotonin 5-HT<sub>2A</sub> receptor agonist LSD has been applied to develop a rodent model with a schizophrenia-related behavioural state entailing e.g. psychosis, hyperactivity and hyperirritability, increased locomotor activity, anhedonia, cognitive and/or impairment in social interaction (Marona-Lewicka et al. 2011; Moreno et al. 2013).

### 1.2.3.5 Lesion-Induced Schizophrenia Models

To address some of the issues surrounding progressive neurodevelopmental or neurodegenerative changes in schizophrenia, a number of targeted lesion animal models have been developed. Besides electrolytic and aspiration lesions, the majority of models is based on the application of excitotoxic agents (Marcotte et al. 2001). Three major sites of actions were chosen based on their presumed involvement in schizophrenia, i.e. the prefrontal cortex, hippocampus and thalamus. The prefrontal cortex is involved in higher cognitive functions, including attention, working

memory, emotional expression and social interaction. Executive function allows us to interact with the world in a purposive, goal-directed manner. It relies on several cognitive control operations that are mediated by different regions of the prefrontal cortex. The hippocampus modulates activity of the prefrontal cortex and thus exerts direct control over the schizophrenia-linked mesolimbic dopaminergic system. The thalamus is the brain's relay station that filters and gates sensory information and is therefore involved in PPI. Despite reasonable claims of predictive and face validity for many adult lesion models, the required size and temporal nature of these lesions limits their construct validity as animal models of schizophrenia. The neonatal ventral hippocampus lesion model is the best characterised model to test neurodevelopmental hypotheses of schizophrenia. This excitotoxicity model leads to dopamine-related behavioural alterations in adolescence or early adulthood, which include hyperlocomotion in response to stress, sensory gating deficits (PPI and LI), disturbed working memory and reduced social contacts, as well as psychotomimetic-induced hyperactivity, apomorphine-induced stereotypies and reduced catalepsy in response to haloperidol. Aberrant development of the prefrontal cortex, illustrated by molecular changes in this region, in response to neonatal hippocampal damage may be a critical factor in the onset of symptoms (Lipska and Weinberger 2000; Lipska 2004). Based on the neonatal disconnection of the ventral hippocampus presumably altering the development and plasticity of prefrontal cortical circuitry, the neonatal rat excitotoxic brain lesion model was also adapted to the ventral thalamus (Wolf et al. 2010, 2018), and the prefrontal cortex (Klein et al. 2008; Lazar et al. 2008).

### 1.2.3.6 Genetic Schizophrenia Models

As exemplified by twin studies, schizophrenia has a substantial genetic component (Cardno et al. 1999). An important shift can be observed in the way animal models are being used for the genetic dissection of psychiatric disease. Initially, inbred strains, often selectively bred for a disease-specific trait, dominated the literature, followed by single-gene knockout or mutation models developed with homologous recombination in embryonic stem cells. Whole-genome association studies are presently expected to provide strong evidence for the involvement of susceptibility genes that will lead to attempts to model multiple susceptibility genes. In addition, the development of microarray and proteomic technology has enabled global description of gene expression in schizophrenia, although it may be difficult to distinguish between primary aetiology and secondary pathology, confounding influences or compensatory mechanisms. Changes in gene expression with a primary etiopathological role may arise from sequence variants in regulatory regions of genomic DNA, epigenetic/stochastic variation or environmental influences, while secondary changes may reflect points of convergence in the action of individual susceptibility genes (Bray 2008). Given the complex aetiological nature of schizophrenia, genetic models expressing risk genes may form the basis to study the effect of environmental manipulation on such mutant phenotypes at specific developmental stages and may help in defining the trajectory, relative contribution of and

interaction between genes and environmental factors in the emergence of the schizophrenia phenotype (Desbonnet et al. 2009).

### **Inbred and Selectively Bred Rodent Strains**

Remarkable variability in outcome of specific cognitive and behavioural tasks is evident in different rodent strains (Crawley 2000; van der Staay and Blokland 1996; Crawley et al. 1997). Genetic factors determine sensory and sensorimotor gating in rodents, and consequently, strain differences in baseline startle response and PPI have been described (Paylor and Crawley 1997; Palmer et al. 2000; Willott et al. 2003; Pietropaolo and Crusio 2009). Moreover, strain and substrain differences for pharmacological (e.g. amphetamine-based) disruption of sensorimotor gating were observed (Kinney et al. 1999; Swerdlow et al. 2000, 2012; Varty et al. 2001). Interestingly, strain-dependent changes across adolescence have been observed, indicating that genetic factors and the early adolescent phase are critically important considerations in the design of animal models of neuropsychiatric disturbances (Pietropaolo and Crusio 2009). Variations between strains and substrains, whether or not accomplished through selective breeding, form a good basis to study the underlying biological mechanisms and hence, therapeutic strategies in schizophrenia (Swerdlow et al. 2005, 2012; Dieckmann et al. 2007; Gogos et al. 2008; Flood et al. 2011). Selective breeding also forms an excellent basis for the development of highly relevant multiple-hit models (Kekesi et al. 2015; Horvath et al. 2016; Büki et al. 2019).

### **Genetically Modified Models**

Candidate genes for schizophrenia are dysbindin (*DTNBP1*) (Williams et al. 2005); neuregulin 1 (*NRG1*) and its receptor *ERBB4* (Banerjee et al. 2010); *DISC1* (i.e. disrupted in schizophrenia-1) (Muir et al. 2008); reelin (*RELN*) (Grayson et al. 2006); components of the Akt-GSK3 $\beta$  signalling pathway, including AKT1 (Freyberg et al. 2010); and genes located in chromosomal region  $\Delta 22q11.2$ , which has been implicated in schizophrenia based on the 22q11.2 deletion syndrome (Paylor and Lindsay 2006). For an updated list of published genetic associations studies for schizophrenia readers are referred to the comprehensive, multi-omics online SZGR 2.0 database (<https://bioinfo.uth.edu/SZGR/>). Besides genetics studies that reported variants and genes potentially associated with schizophrenia, SZGR 2.0 also compiles transcriptome and epigenetic information (Peilin et al. 2017).

Several spontaneous mutant rodent models have been studied as presumed phenocopies of schizophrenia. The *sandy* mouse has a deletion of two of the exons of the dysbindin gene and is therefore a naturally occurring dysbindin-knockout model. *Sandy* mice display some behavioural phenotypes with relevance to schizophrenia, including increased anxiety and deficits in social interaction (Hattori et al. 2008). After the discovery of *DISC1* as a potential schizophrenia candidate gene, some of the earliest animal studies on its potential role included the detection of



natural mutations in different mouse strains. All 129 mouse substrains carry a 25-bp deletion in the *mDISC1*, which modulates working memory (Clapcote and Roder 2006; Koike et al. 2006). Reeler is an autosomal recessive mutant mouse with reduced cerebellar size, disruption of the laminar organisation in several brain regions and inversion of neocortical cellular layers (Falconer 1951; Caviness Jr 1976). The reelin-haploinsufficient heterozygous reeler mouse has been used as an animal model for schizophrenia based on several neuropathological and behavioural abnormalities homologous to schizophrenia. Heterozygous reeler mice exhibit alterations in sensorimotor gating (Barr et al. 2008), deficits in working memory (Brigman et al. 2006), impaired social interaction (Podhorna and Didriksen 2004), but also neurochemical, neuroanatomical and protein expression alterations relevant to schizophrenia have been reported (Schmitt et al. 2013; Nullmeier et al. 2014; Varela et al. 2015).

DNA variation that affects expression of candidate disease genes can take various forms, including single nucleotide polymorphisms (SNPs), deletions, insertions, variable repeat sequences, rearrangements or duplications (copy number variations). Downregulation of individual susceptibility genes can be mimicked via gene knockout or via RNA interference, while upregulation can be achieved with gene knock-in techniques or RNA activation.

Multiple genes (perhaps thousands) of small effect are thought to exist for the illness, given its status as a complex genetic illness (Peilin et al. 2017; Gejman et al. 2011). Nonetheless, tissue effects in animals with risk mutations are presumed to be associated in some way with schizophrenia pathophysiology. The biological characterisation of animals with risk genes is highly relevant to ultimately discovering disease pathophysiology, even though every aspect of the biology may not be critical. For the currently most promising risk genes, few transgenic mouse models will be briefly discussed.

Many mouse models with manipulated expression levels of the different NRG1 isoforms have been created. Mutants with heterozygous deletion of the transmembrane domain exhibit hyperactivity, a deficit in PPI, selective impairment in social novelty preference and altered patterns of social interaction. Studies focusing on deletion of specific NRG1 isoforms have shown that type III NRG1 mutant mice display even more pronounced PPI deficits in addition to working memory dysfunction (Desbonnet et al. 2009). Homozygous knockout of *NRG1* is developmentally lethal in mice, but viable heterozygous, hypomorphic/conditional knockouts that can modulate neuregulin–ErbB4 signalling have been developed, all with distinct ‘schizophrenia-like’ alterations, including—depending on the genetic models studied—hyperactivity, PPI, LI and social interaction deficits, impairment in contextual fear conditioning and mismatch negativity (Harrison and Law 2006; Mei and Xiong 2008; Jones et al. 2011). At least seven different strains of transgenic mice containing inducible and/or partial *DISC1* gene mutations resulting in a (partial) loss of DISC-1 function have been created (Jaaro-Peled 2009). These mice exhibit enlarged lateral ventricles and reduced cortical thickness and brain volume, mimicking some characteristics of schizophrenia. Some mutants display reductions in hippocampal dendritic complexity, structure and density. Some *DISC1* mice display PPI deficits

that are attenuated with antipsychotics, while hyperactivity, reduced sociability, working memory and executive function impairments have been described as well.

Over the past decades a multitude of genetically modified schizophrenia models has been developed. Discussing all currently developed genetic models of schizophrenia would go beyond the scope of this chapter. Readers are referred to several excellent review papers (Jones et al. 2011; Leung and Jia 2016; Moran et al. 2016; St Clair and Johnstone 2018).

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### 1.3 Animal Models of Neurological Disorders

The discovery of new therapies for neurological disorders is predicated on the use of animal models both to identify new therapeutic targets and to perform preclinical trials of drugs prior to clinical application. In both cases, the challenge is to develop models that recapitulate the disorder. The starting point for the development of a new animal model for a specific neurological condition is often the current dominant theory about the disease. Although a logical first approach, it is essential to broaden the focus of animal models under development based on the increasing knowledge of underlying disease mechanisms. The development of animal models for Alzheimer's disease (AD) serves an excellent example of such a strategy.

Advances in biochemical pathology and human genetics have yielded striking progress in our understanding of molecular mechanisms underlying nervous system diseases such as AD, Parkinson's and Huntington's disease. Of great importance is the pivotal concept that certain normally soluble neuronal proteins can misfold and aggregate into oligomers and fibrils, which can confer profound cytotoxicity. Perhaps the foremost example, both in terms of its societal impact and how far knowledge has moved toward the clinic, is that of AD, hence the focus of the following paragraphs.

#### 1.3.1 Approaches to Modelling Neurological Disorders

In contrast to psychiatric disorders, approaches to model neurological conditions are not easily categorised. The approach largely depends on the aetiology of the human neurological condition, which may be acquired or degenerative. Specific approaches when developing new animal models should be applied depending on the aetiology, which may be amongst others infectious (e.g. Creutzfeldt–Jakob disease, HIV dementia), immune-related (e.g. multiple sclerosis), genetic (e.g. Charcot–Marie–Tooth disease; leukodystrophies), lesion-based (e.g. head trauma), and environmental/toxicity-related (e.g. alcohol-related dementia, toxic-metabolic encephalopathy).

## 1.3.2 Animal Models of Alzheimer's Disease

### 1.3.2.1 Aetiology and Symptomatology of Alzheimer's Disease

As the prototype of cortical dementias, AD is characterised by prominent cognitive deficits. Patients initially exhibit limited forgetfulness with disturbance of memory imprinting, which further evolves to short-term memory disruption and, ultimately, to long-term memory deficits. At more advanced stages, patients display executive dysfunctioning leading to advanced helplessness (Selkoe 2000). Besides cognitive deterioration, patients demonstrate behavioural and psychological signs and symptoms of dementia (BPSD), including paranoid and delusional ideation, hallucinations, activity disturbances, aggressiveness, diurnal rhythm disturbances, affective disturbances, anxieties and phobias (Reisberg et al. 1987). BPSD are nowadays also often referred to as neuropsychiatric symptoms or NPS (Lyketsos et al. 2011; Van Dam et al. 2016). The histopathological hallmarks of AD brain are extracellular amyloid- $\beta$  (A $\beta$ ) plaques and intracellular neurofibrillary tangles (NFT), accompanied by decreased synaptic density, which eventually leads to widespread neurodegeneration and failure of neurotransmitter pathways, particularly those of the basal forebrain cholinergic system (Van Dam et al. 2016; Selkoe 2001). The number of affected individuals is likely to grow in the decades to come as a result of demographic changes and rising life expectancy. It is forecast that the worldwide number of elderly people suffering from dementia will rise to 63 million in 2030 and to 114 million in 2050 (Wimo et al. 2003).

### 1.3.2.2 Validating Animal Models of Alzheimer's Disease

#### Cognitive Symptoms

A variety of paradigms and protocols has been developed to assess cognitive functions in rodents (Crawley 2000; Buccafusco 2008). Ideally, several paradigms requiring distinct sensory and motor abilities are chosen when phenotyping a new dementia model. Protocols distinguishing explicit versus implicit memory can be chosen, as well as designs assessing short-term versus long-term memory (Tulving 1987). The Morris water maze is presently the most widely used paradigm for the evaluation of hippocampus-dependent visual-spatial learning and memory skills in rodents, which represents the highest cognitive level appreciable in rodents (D'Hooge and De Deyn 2001). Besides the Morris water maze, several other mazes could be employed, which are all based on the same principle; successfully passing through the maze is rewarded by escaping from the water in wet mazes or by food as a positive reinforcer in dry-land mazes. Examples include the plus-shaped water maze, radial arm mazes, multiple Y mazes and the Barnes maze. Complex nonspatial hippocampus-dependent tasks like the odour paired-associates task (Bunsey and Eichenbaum 1996), are analogous to the verbal paired-association task for humans, but the long duration of the protocol, which requires several months, makes it unsuitable for the assessment of memory in progressing phenotypes (Van Dijk et al. 2008). The same disadvantage may arise when using schedule-induced

operant tasks employing operant conditioning or Skinner boxes. Both active and passive avoidance learning protocols are widely used to assess cognitive function in a short time frame. However, procedural components of the task are not easily distinguishable from declarative memory components as is the case in the Morris water maze or in cued and contextual learning (Crawley 2000). Associative learning (cued and contextual conditioning) requires a different set of sensory and motor functions, so that the procedural components of associative tasks do not overlap. Other potentially useful learning and memory tasks include novel object recognition, conditioned taste aversion, social recognition and discrimination learning (Crawley 2000).

### **BPSD-Related Symptoms**

In accordance with the increased clinical focus on BPSD, major efforts have been made to mimic specific behavioural alterations in animal models and to develop useful tools (Crawley 2000; Buccafusco 2008), to evaluate new psychopharmacological strategies to replace atypical antipsychotics or classic neuroleptics, which display only modest effect size and are frequently associated with significant side effects (De Deyn et al. 2005).

Activity and circadian rhythm disturbances can be easily screened using infrared sensors surrounding the animal's home cage. The number of beam interruptions over a specific time interval is recorded as a measure of activity comparable to actigraphic measurements in the clinical setting (Vloeberghs et al. 2004). Aggressive behaviour in male rodents can be provoked using a variety of behavioural protocols based on dominance hierarchy, as is the case in the commonly used isolation-induced resident-intruder protocol (Valzelli 1973). Anxiety and fear-related behaviours can be assessed using both conditioned (i.e. conflict tests like the Vogel's lick-suppression test) and unconditioned response tests, which are based on the quantification of fear-related responses like freezing, defecation and thigmotaxis, and the conflict between the innate trait of nocturnal animal like mice and rats to prefer narrow, dark enclosures and the tendency to explore new environments (Crawley 2000). Appraisal of depression-related symptoms in rodents is based on learned helplessness and behavioural despair phenomena, in which animals are exposed to uncontrollable and inescapable stress, e.g. the Porsolt forced swim test, the tail suspension test and inescapable shock paradigms. Failure to try to escape from this type of aversive stimulus in rodents is considered to model a depression-like state (Chourbaji et al. 2005). Anhedonia, a core symptom of clinical depression, which is defined as the loss of sensitivity to reward, can be assessed in rodents with a sucrose preference test during which the consumption of a 0.8% sucrose solution is compared to the simultaneous consumption of tap water (Sanchis-Segura et al. 2005). Phenotyping can also focus on ingestive behaviour. The intake of food and water can be screened using metabolic cages. More detailed analysis of food intake is possible in operant conditioning boxes equipped with pellet dispensers and optical lick-o-meters that allow appraisal of the circadian rhythm of ingestive behaviour (Vloeberghs et al. 2008).

## Pathological Alterations

The neuropathological hallmarks of AD include “positive” lesions such as amyloid plaques, cerebral amyloid angiopathy, neurofibrillary tangles, and neuroinflammatory responses, and “negative” lesions, such as neuronal and synaptic dysfunction and loss (Van Dam et al. 2016). Amyloid structure is most commonly stained using congo red, thioflavin S or A $\beta$ -specific antibodies (Castellani et al. 2007; Vidal and Ghetti 2011). Amyloid plaque burden can be evaluated by employing stereological methods, which allow a three-dimensional geometrical interpretation of structures based on observations made on two-dimensional sections. Levels of A $\beta$  peptides and other APP-derived molecules can be quantified using ELISA or Western blotting techniques. While ELISA is a very sensitive method to detect and quantify total A $\beta$  levels, commercially available kits provide no or very little information on the possible aggregation state and structure of A $\beta$ . Western blotting, on the other hand, is less suited for quantification but makes it possible to detect aggregates and roughly estimate their size. However, it still only provides very limited information about the structure and composition of the aggregates. Researchers more interested in the molecular structure, composition and formation of the oligomers turned to mass spectrometry techniques, which allow very accurate determination of protein identity, amino acid sequence and presence of modifications, or even—when using non-denaturing approaches—the macromolecular structure (Ashcroft 2010; Bleiholder et al. 2011). Tau pathology can be assessed at various levels, including a quantitative assessment using Western blot, biochemical and immunohistochemical determination of different phosphorylation states and sites, as well as the immunohistochemical assessment (e.g. with the ALZ-50 monoclonal antibody) of the spatiotemporal progression of tau pathology (Rankin and Gamblin 2008). Monitoring inflammatory processes can be based on the use of histopathological and immunohistochemical markers of astrocyte and microglia activation, as well as on various biochemical assays of inflammatory markers in body fluids (McGeer and McGeer 2003). Potential biomarkers of oxidative stress, including an imbalance between free radicals (reactive oxygen and nitrogen species) and the antioxidant response (free radical scavengers and antioxidant enzymes), as well as the subsequent damage to macromolecules (lipids, proteins, nucleic acids and sugars), can be determined in body fluids or brain using a wide range of biochemical techniques, including spectrophotometric, fluorometric, chromatographic and immunohistochemical assays (Migliore et al. 2005). Degenerative processes (e.g. neuritic spheroids) can be visualised by silver impregnation, neurofilament or ubiquitin immunostaining (Castellani et al. 2007).

## Neurochemical Alterations

The cholinergic hypothesis of AD is based on the fact that degeneration of cholinergic neurons in the nucleus basalis of Meynert, situated in the basal forebrain and primarily projecting to the neocortex, occurs early in the course of the disease and leads to a marked decline in the activities of choline O-acetyltransferase (ChAT) and acetylcholinesterase (AChE) (Contestabile 2011). In rodent models, the presence of cholinergic deficits can be scrutinised with the determination of ChAT or AChE enzyme activities based on, respectively, a radiochemical (Fonnum 1975), and a spectrophotometric protocol (Ellman et al. 1961). Microdialysis extraneuronal

sampling coupled with a high-sensitivity HPLC detection and quantification method has been often used for assessing the cholinergic impairment and its recovery in many AD models (Pepeu and Giovannini 2007). Moreover, immunohistochemical visualisation and quantification of ChAT- or AChE-positive neurons and fibres has been employed, as well as determination of high-affinity choline uptake, considered a biochemical marker of the localisation and integrity of the cholinergic nerve endings and a measure of the activity of the cholinergic neurons (Pepeu and Rosi 2011).

Neurochemical alterations observed in AD brain, however, are not confined to the cholinergic system. Moreover, many data indicate that the neurochemical alterations underlying cognitive deterioration and related disturbances in cortical processing implicate more widespread neurodegeneration that cannot be attributed solely to the cholinergic system (Dringenberg 2000). Neuronal loss and, inherently, alterations in the concentration of neurotransmitters and metabolites of the noradrenergic, adrenergic, dopaminergic and serotonergic system, as well as neurotransmitter amino acids, have been described in AD (Van Dam et al. 2016; Vermeiren et al. 2014a, b; Gsell et al. 2004). In rodent models, chromatography-based determination of neurotransmitter (and metabolite) levels in microdialysis samples or brain (region) homogenates can be employed to assess neurotransmitter alterations. Deficits in peptidergic neurotransmission (or co-transmission) have been implicated in AD. Classically, neuropeptides are mainly studied by radioimmunoassay and immunohistochemistry, but recently researchers have shifted to mass spectrometry-based methodologies (Van Dam et al. 2013).

### 1.3.2.3 Spontaneous and Selectively Bred Alzheimer's Disease Models

Various species across the phylogenetic tree, ranging from invertebrates like *Drosophila melanogaster* and *Caenorhabditis elegans* (Mhatre et al. 2013), up to non-human primates have been applied in AD-related research (Van Dam and De Deyn 2017). Some species, including dogs (Cummings et al. 1993, 1996; Rofina et al. 2006), cats (Head et al. 2005; Gunn-Moore et al. 2006), bears (Cork et al. 1988; Tekirian et al. 1996; Uchida et al. 1995), goats and sheep (Braak et al. 1994), wolverine (Roertgen et al. 1996), as well as several non-human primate species (Van Dam and De Deyn 2017; Bons et al. 1994; Gearing et al. 1994, 1997; Lane 2000; Geula et al. 2002; Kimura et al. 2003; Sani et al. 2003; Lemere et al. 2004, 2008), spontaneously develop plaque pathology and some species even display tauopathies. These histopathological changes can be accompanied by cognitive decline (Cummings et al. 1996; Rofina et al. 2006; Gunn-Moore et al. 2006; Voytko and Tinkler 2004). The use of these species for experimental research is however limited by availability, economical (based on long lifespan) and/or ethical reasons (Van Dam and De Deyn 2017).

Rodents are unequivocally the front runners as model organisms in this field. Rodents are easier and cheaper to house and maintain than larger mammals, and they display a fast generation time with large numbers of offspring. Moreover, rodents and humans do share approximately 95% of their genome, and standardized breeding gives researchers access to various inbred strains contributing to a less variable phenotype (Van Dam and De Deyn 2006, 2011a). Ageing rodents do not spontaneously develop histopathological AD-like hallmarks, but do display

senescence-related cognitive decline and behavioural alterations associated with neurochemical and morphological alterations (Erickson and Barnes 2003), including age-associated cholinergic hypofunction (Sherman and Friedman 1990). In addition, they aid in uncovering the boundary between normal and pathological ageing, allowing in-depth investigation of basic neural mechanisms underlying brain ageing.

Selective breeding from a genetic pool of AKR/J mice has led to the development of the senescence-accelerated mouse (SAM) model, which includes nine major SAM-prone (SAMP) substrains and three major SAM-resistant substrains. The SAMP8 substrain is of particular interest given the development of age-associated learning and memory deficits in association with A $\beta$  deposition (Butterfield and Poon 2005; Ren $\acute{a}$  and Butterfield 2011).

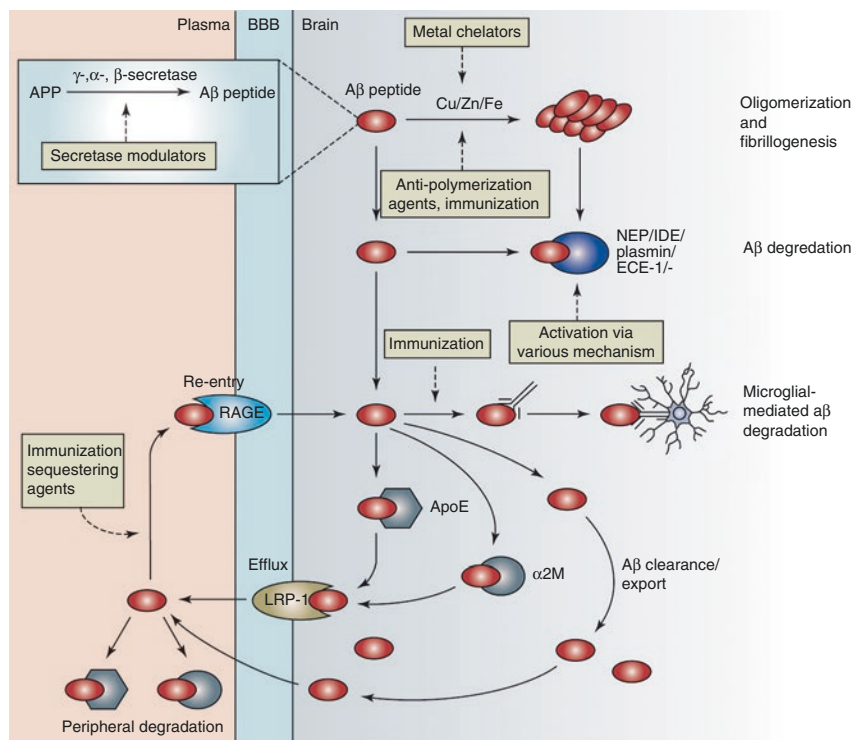
### **Pharmacological, Chemical and Lesion-Induced Rodent Models of Alzheimer's Disease**

Disruption of multiple neurotransmitter systems underlies the cognitive and behavioural disturbances associated with AD. The majority of animal models that fall within this category are based on the cholinergic hypothesis of AD, which states that degeneration of cholinergic neurons in the nucleus basalis of Meynert, located in the basal forebrain and projecting to the neocortex, occurs early in the course of the disease (Whitehouse et al. 1982; Davies and Maloney 1976). The most widely used pharmacological model is scopolamine-induced amnesia (Sunderland et al. 1986; Ebert and Kirch 1998). The application of this muscarinic antagonist has increased our knowledge of the role of the cholinergic system in cognition and allows pre-clinical evaluation of symptomatic efficacy of cholinomimetics. The applicability of this model is however limited by the fact that cholinergic hypofunction is not associated with the development of AD-typical pathology, and the lack of disease progression both at the cholinergic and cognitive level. Blockade of nicotinic receptors by mecamylamine also induces learning impairment (Moran 1993; Estapé and Steckler 2002). Since AD brain shows both reduced muscarinic and nicotinic acetylcholine receptor densities (Whitehouse and Au 1986; Nordberg et al. 1989), blockade of both receptors may offer a better animal model for AD-related amnesia (Levin et al. 1990; Riekkinen Jr et al. 1990).

In addition to scopolamine-induced amnesia, cortical cholinergic involution has been replicated in lesion models that focus on specific cholinergic centres of the basal forebrain, as well as more general lesions of all basal forebrain cholinergic neurons. Focal lesions are most often directed at the nucleus basalis magnocellularis (Lescaudron and Stein 1999; Vale-Martínez et al. 2002), the rodent analogue of the nucleus basalis of Meynert, the septal area (Mulder et al. 2005), or include fimbria/fornix transection leading to septo-hippocampal cholinergic denervation (He et al. 1992; Alonso 1996). Lesions can be induced by mechanical, i.e. knife cut, or electrolytic procedures and intraparenchymal or intracerebroventricular microinjections of neurotoxic substances, such as quinolic, kainic, ibotenic and quisqualic acids, NMDA, the cholinotoxin AF64, and the immunotoxin 192 IgG-saporin (Toledana and Alvarez 2011). Such lesion models increase our understanding of the role of cholinergic innervations in the aetiology and treatment of cognitive disorders.

AD-related memory deficits can also be (partially) reproduced by lesioning brain structures or pathways essential for different aspects of learning and memory, such as the hippocampus, striatal or cortical regions (Gray and McNaughton 1983; Glenn et al. 2003; Sloan et al. 2006; Castañé et al. 2010). These models increase our knowledge of the neural mechanisms underlying memory dysfunction, but of course lack disease progression and do not develop AD-like pathological hallmarks. Moreover, selected lesions are compared with the more global disease process of AD.

Certain chemically induced models focus on one specific pathophysiological pathway thought to underlie AD, such as neuroinflammation or glucose/energy metabolism impairment, and their effects on neurodegeneration. Neuroinflammation develops upon the infusion of endotoxins, like lipopolysaccharide (Hauss-Wegrzyniak et al. 1998), or proinflammatory cytokines (Wenk et al. 2003). Brain metabolism can be disrupted through interference with mitochondrial metabolic pathways (Szabados et al. 2004), or neuronal insulin signal transduction (Ishrat et al. 2009).



**Fig. 1.2** This figure represents the major processing pathways of amyloid precursor protein (APP) and amyloid  $\beta$  ( $A\beta$ ) peptides, including the potential therapeutic targets and therapeutic interventions. *APP* amyloid precursor protein, *CTF* C-terminal fragment, *NTF* N-terminal fragment, *A $\beta$*  amyloid  $\beta$  peptide, *ApoE* apolipoprotein E, *BBB* blood-brain barrier, *CSF* cerebrospinal fluid, *ECE-1* endothelin-converting enzyme-1, *IDE* insulin-degrading enzyme, *LRP-1* low-density lipoprotein receptor-related protein-1, *NEP* neprilysin, *RAGE* receptor for advanced glycation end products,  *$\alpha$ 2M*  $\alpha$ -2-macroglobulin. Repinted with permission from Van Dam and De Deyn (2006)



### 1.3.2.4 Amyloid- $\beta$ Infusion Rodent Models of Alzheimer's Disease

The amyloid cascade hypothesis states that cerebral accumulation and aggregation of A $\beta$  peptides into amyloid plaques is the primary culprit in AD (Selkoe 2000; Hardy and Selkoe 2002). More recently, a key role was assigned to soluble A $\beta$  oligomers (Gong et al. 2003; Lacor et al. 2004; Walsh and Selkoe 2007; Selkoe 2008) (Fig. 1.2).

Aspects of AD can be modelled by acute or repetitive intracerebral or intracerebroventricular infusion of A $\beta$  peptides in rodent brain (Lawlor and Young 2011). To better mimic the progressive nature of AD, chronic and continuous administration is accomplished by connecting an implanted cannula to an osmotic mini-pump (Nakamura et al. 2001; Olariu et al. 2002), or a micro-infusion pump (Nag et al. 1999), or with microdialysis (Harkany et al. 2000). In addition to measurable adverse effects on memory and behaviour (Harkany et al. 1998; Yamada et al. 2005; Sipos et al. 2007), exogenous administration of A $\beta$  peptides can lead to AD-like neuropathological alterations (Sipos et al. 2007; Frautschy et al. 1996), although the full complexity of the human pathology is not reproduced and pathology is not widespread as in patients. Accumulation of A $\beta$  deposits can be associated with, e.g. inflammation and microglial activation, oxidative stress and local cell loss (Weldon et al. 1998). More specifically, disruption of cholinergic function was reported (Harkany et al. 1998; Yamada et al. 2005).

CNS A $\beta$  infusion animal models provide insight into the mechanisms and secondary effects of A $\beta$  toxicity and allow preclinical evaluation of drugs targeting A $\beta$ , as well as evaluation of protective efficacy of pharmacological modulation of microglial signalling. Rodent infusion models have some advantages over transgenic amyloid-related models. Infusion models can deliver the desired (pathological) outcome within a time frame of a few weeks (Frautschy et al. 1996), versus several months or years in transgenic models. Moreover, they allow researchers to administer defined amounts of a specific A $\beta$  species of known sequence and length or to introduce controlled cofactors related to plaque development, while transgenic overexpression of APP results not only in increased production of A $\beta$ 1–40 and/or A $\beta$ 1–42, but in elevated levels of other APP fragments, which can have neuroprotective, neurotoxic or signalling functions and influence learning and memory. On the other hand, major caveats of infusion models are the fact that administered A $\beta$  concentrations are much higher than A $\beta$  levels found in the brain or cerebrospinal fluid of AD patients (Vickers et al. 2000), and that effects of ageing on AD progression are lacking. Moreover, the invasive nature of A $\beta$  infusion inevitably brings about brain injury, which—in addition to the potential neurotoxic effects of vehicles used—may contribute to the induction of inflammation observed in these models. These potentially confounding effects should be controlled for by including proper sham and/or scrambled peptide groups.

### 1.3.2.5 Genetically Modified Alzheimer's Disease Mouse Models

Modelling of AD in transgenic mouse models became reality in the mid-1990s with the development of the platelet-derived growth factor promoter-driven APP (PDAPP) model (Games et al. 1995; Basak and Holtzman 2011), followed in

subsequent years by the Tg2576 (Hsiao et al. 1996; Deacon 2011), and the APP23 mouse model (Sturchler-Pierrat et al. 1997; Van Dam and De Deyn 2011b), currently the most widely used amyloidosis models in AD-related research. The choice of *APP* as the incorporated transgene was based on two lines of evidence: APP is the precursor of A $\beta$  and it is the target of the first described early-onset AD mutations. The PDAPP model expresses human *APP* carrying the Indiana familial AD mutation driven by the platelet-derived growth factor- $\beta$  promoter, whereas both the Tg2576 and APP23 model express human APP with the Swedish mutation driven by the hamster prion protein and murine Thy-1 promoter, respectively. The discovery of familial AD mutations in the presenilin (PSEN) genes, which influence APP processing, opened the path for PSEN1 and PSEN2 transgenic mouse models and double-cross APP/PSEN models. Although early-onset AD accounts for only a small proportion of AD cases, models expressing pathogenic mutations of human genes have become crucial tools in unravelling disease processes and have boosted drug discovery and development.

Several major genetic risk factors for developing late-onset AD have been identified, among which apolipoprotein E (ApoE) and triggering receptor expressed on myeloid cells 2 (TREM2). Mice expressing a human ApoE isoform, such as ApoE4 (Huber et al. 2000), and TREM2 (Kleinberger et al. 2017) have been developed to study the pathophysiological link between, respectively, ApoE and TREM2, and AD.

Genetically modified mouse models based on other aetiological hypotheses that are considered relevant to the preclinical AD field include, e.g. mutated human  $\alpha$ -synuclein models (Freichel et al. 2007), TAR DNA-binding protein 43 (TDP-43)-related models (Wu et al. 2010; Wils et al. 2012), mutated human cyclooxygenase-2 overexpression models (Andreasson et al. 2001), anti-nerve growth factor (NGF) mice (in which NGF activity is neutralised using antibodies) (Capsoni et al. 2000), neprilysin (Iwata et al. 2001), insulin-degrading enzyme knockout mice (Farris et al. 2003), and crosses between mutated human APP overexpression strains and nitric oxide synthase 2 knockouts (Colton et al. 2008). For more in-depth information and latest updates, readers are referred to the Alzheimer Research Forum website (<http://www.alzforum.org>), which includes an online compendium of relevant mouse models for AD and related disorders. Most of these models robustly mimic a subset of AD features, including A $\beta$  deposition, plaque formation and cerebral amyloid angiopathy, as well as neurodegeneration, synaptic dysfunction and inflammation, in addition to cognitive decline.

The major limitation of the above-mentioned models, i.e. the lack of NFT formation, was partially counterbalanced with the development of several (mutated) tau models, and the crossing of tau and amyloidosis models, which featured enhanced amyloid deposition accompanied by tau phosphorylation, NFT-like formation and overt neuronal loss (Götz et al. 2004; Ribé et al. 2005). It has proven challenging to develop mice in which both histopathological hallmarks occur in AD-relevant brain regions, such as the hippocampus and cortex. However, the development of triple transgenic (3  $\times$  Tg) mice seems to have overcome this hurdle (Oddo et al. 2003; Sy et al. 2011). Rather than crossing independent mutant mouse lines, two transgenes

(mutant APP and tau) were microinjected into single-cell embryos from homozygous mutant PSEN1 mice. The major advantages of this approach are the integration of APP and tau at the same genetic locus (which prevents segregation in subsequent generations), more cost-effective breeding requirements and the fact that there is no mixing or altering of the genetic background. In accordance with the amyloid cascade theory, these 3 × Tg mice develop A $\beta$  plaques prior to NFT pathology with a temporal and spatial profile equivalent to AD. Studies using the 3 × Tg-AD mice have revealed a strong interaction between A $\beta$  and tau, which synergistically drive the pathogenesis in the brain. The first signs of cognitive decline manifest at age 4 months as a deficit in long-term retention and correlate with accumulation of intraneuronal A $\beta$  in the hippocampus and amygdala (Billings et al. 2005). Similarly, the 5 × FAD model was developed that expresses human APP and PSEN1 transgenes with a total of five AD-linked mutations: the Swedish (K670N/M671L), Florida (I716V), and London (V717I) mutations in APP, and the M146L and L286V mutations in PSEN1. The Swedish mutation results in higher levels of total A $\beta$ , whereas the other mutations specifically increase the production of A $\beta$ <sub>x-42</sub>. Already at age 2 months, extracellular amyloid plaques are detected by immunohistochemistry in hippocampus, cortex, and thalamus (Oakley et al. 2006). At the age of 1 year, 5 × FAD mice display a 40% loss of layer V pyramidal neurons (Oakley et al. 2006; Neuman et al. 2015). The model develops age-dependent cognitive (spatial, associative and working) decline from the age of 3 months onwards (Girard et al. 2013), as well as non-cognitive behavioural abnormalities (Jawhar et al. 2012).

Of interest is the development of genetically modified rat models for AD, which may serve better tools than mouse models for microPET/SPECT applications. TgF344-AD rats for example express human APP with the Swedish mutation and human PSEN1 with the  $\Delta$  exon 9 mutation. The rat model shows an age-dependent accumulation of amyloid plaques between 6 and 26 months of age, with plaque-associated neuritic dystrophies, cerebral amyloidosis, tauopathy, gliosis, apoptotic loss of neurons in the cerebral cortex and hippocampus, and cognitive disturbances (Cohen et al. 2013), as well as enhanced anxiety-like behaviour (Pentkowski et al. 2018).

More recently, the use of viral vector gene transfer technology has allowed the development of ‘somatic transgenic’ models, whereby genes putatively involved in AD pathogenesis can be selectively overexpressed in specific brain regions involved in AD (Lawlor and Young 2011; Jaworski et al. 2010). Viral transgenesis also allows investigators to explore the role of newly discovered genetic linkages in a more efficient and cost effective manner, indeed, viral technologies can be used to introduce many genetic changes at once for the screening of many individual factors (and their interactions) simultaneously (Platt et al. 2013). The CRISPR/Cas9 system further improves the biological toolkit to develop genetically modified animals (Tschaharganeh et al. 2016). For an updated list of genetically modified AD-related models, readers are referred to the website of the Alzheimer Research Forum (<http://www.alzforum.org>).

## 1.4 Imaging in Rodent Models of Brain Disease

This chapter does not aim at giving an overview of neuroimaging involving PET and SPECT in models of brain disease, but rather serves as an introduction to models of brain disorders with a selected overview of modelling approaches and methodological aspects with regard to phenotyping. We will briefly describe some relevant imaging techniques and show a couple of examples related to schizophrenia and AD.

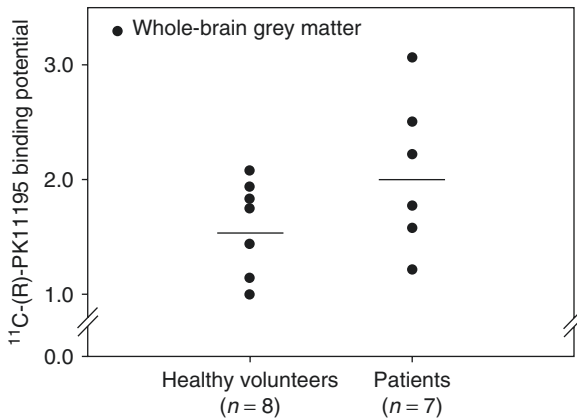
With the development of more sophisticated techniques, neuroimaging has become an essential component of studying brain structure and function in the diseased state. In drug development research, utilising methodology and results that can easily be translated from preclinical to clinical platforms will facilitate the identification and the assessment of efficacy of neuroprotective therapeutic compounds to treat CNS diseases. Translational neuroimaging can provide information on brain morphology and function in preclinical models of CNS diseases, healthy human subjects and patients. Amongst the variety of imaging methods used for understanding and treating brain diseases are PET, SPECT, X-ray/computed tomography, near-infrared optical imaging and magnetic resonance imaging.

Normal laboratory rodents may be used for *in vivo* biodistribution experiments focussing on CNS penetration and washout of radiotracers, while (transgenic) disease models may form the basis for *ex vivo* autoradiography to assess binding of probes to the targeted molecule on brain slices, or of course the *in vivo* evaluation of tracer retention in brain.

### 1.4.1 Imaging in Schizophrenia

Advances in the development and applications of neurochemical brain imaging have improved the ability to study the neurochemistry of the living brain in psychiatric disorders. In particular, PET and SPECT have been used to determine neurochemical substrates of schizophrenia and to uncover the mechanism of action of antipsychotic medications. The growing availability of radiotracers for monoaminergic neurotransmitter synthesis, transporters and receptors has enabled the evaluation of hypotheses regarding neurotransmitter function in schizophrenia derived from preclinical and clinical observations. Clinical schizophrenia PET-/SPECT-based trials study (1) abnormalities in indices of neurotransmitter systems with an established role in schizophrenia in patients versus controls, (2) new tools to study other neurotransmitter systems or potential biomarkers (e.g. inflammation markers) (Doorduyn et al. 2009, 2010) (Figs. 1.3 and 1.4) and (3) characterisation of target occupancy by antipsychotic drugs, as well as its relationship to efficacy and side effects (Urban and Abi-Dargham 2010; Kirino et al. 2017); see also <http://www.clinicaltrials.gov>.

A variety of PET tracers has been used to identify brain abnormalities in schizophrenia based on kinetic modelling. The most consistent early findings showed a difference in the dopamine content in the prefrontal cortex, anterior cingulate gyrus

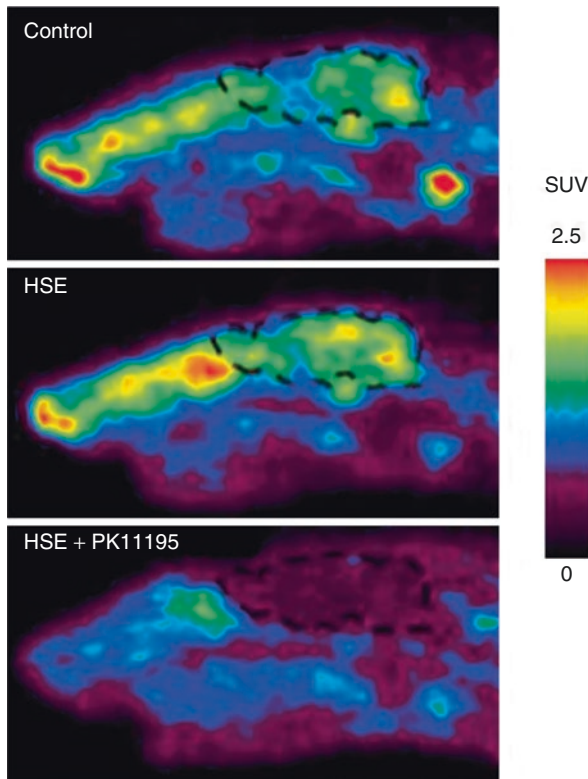


**Fig. 1.3** Binding potential of the PET marker (*R*)-[<sup>11</sup>C]-PK11195 that binds to translocator protein (TSPO), a biomarker to study activated microglia cells, in healthy volunteers and schizophrenia patients. All schizophrenia patients met the DSM-IV criteria of schizophrenia spectrum disorders. Each dot represents an individual subject, and horizontal lines represent mean values. Whole-brain grey matter binding potential was 30% higher in patients versus healthy controls, although this difference did not reach statistical significance (Reprinted from (Doorduyn et al. 2009); Doorduyn et al. (2009) with permission. Copyright © 2009 Society of Nuclear Medicine and Molecular Imaging)

and hippocampus between healthy controls and schizophrenia patients. In addition, a higher density of D2 receptors in the striatum and neural brain dysconnectivity was observed (Patel et al. 2010), while [<sup>11</sup>C]SCH23390 and [<sup>11</sup>C]N-methylspiperone to map D1/D2 receptors identified reduced D1R densities in the prefrontal cortex (Okubo et al. 1997). Dopaminergic abnormalities have been further substantiated in a large number of PET/SPECT-based studies in recent years (Kambeitz et al. 2014; Weinstein et al. 2017; Kim 2018; Avram et al. 2019).

In addition, other neurotransmitter systems, including the serotonergic (Milak et al. 2008; Huang et al. 2010; Herth et al. 2016; Petersen et al. 2016), and the cholinergic system (Horti et al. 2010; Wong 2018), are being targeted in more recent years by PET/SPECT imaging in line with the elaboration of the neurochemical hypothesis in schizophrenia as discussed in Sect. 1.2.3.1. Aetiology and Symptomatology of Schizophrenia.

There have been some PET studies of the behavioural sensitisation to psychostimulant drugs in dogs (Nakamura et al. 1997), rhesus monkeys (Castner 2000), and rats (Kilbourn and Domino 2011). Given the presumed contribution of excitatory NMDA-linked neurotransmission to schizophrenia, PET/SPECT has scarcely been used for investigations of NMDA perturbations in experimental animals, including rats (Reith et al. 1998), cats (Vollenweider et al. 2000), and monkeys (Tsukada et al. 2000; van Berckel et al. 2006). Albeit the application of microPET/SPECT in schizophrenia remains currently still limited, insight in disease processes underlying schizophrenia, with of course an important focus on the dopaminergic aspects, lay at the basis of the development of novel genetically modified rodent



**Fig. 1.4** Many neurological and psychiatric disorders are associated with neuroinflammation, which can be studied using PET markers like (*R*)-[<sup>11</sup>C]-PK11195 that binds to translocator protein (TSPO), a biomarker to study activated microglia cells. The sensitivity of (*R*)-[<sup>11</sup>C]-PK11195 may however be insufficient to visualise mild neuroinflammation. [<sup>11</sup>C]-N-(2,5-dimethoxybenzyl)-N-(4-fluoro-2-phenoxyphenyl)-acetamide, [<sup>11</sup>C]-DAA1106, proposed as a potentially more sensitive neuroinflammation tracer, was evaluated in a rat model of herpes encephalitis (*HSE*). Sagittal view of the head of a control rat (*control*), a rat with *HSE* and a *HSE* rat pretreated with 5 mg/kg of unlabeled PK11195 (*HSE + PK11195*), in which the brain is delineated by a dashed line. The images represent tracer uptake between 30 and 60 min after injection of [<sup>11</sup>C]-DAA1106 ( $44 \pm 14$  MBq). Images show a high uptake of [<sup>11</sup>C]-DAA1106 in the brainstem. Pretreatment with unlabeled PK11195 resulted in effective blocking of [<sup>11</sup>C]-DAA1106 uptake. *In vivo* uptake, however, did not significantly differ between control and *HSE* rats. [<sup>11</sup>C]-DAA1106 may not be an ideal tracer to perform longitudinal PET studies in small animals to study the role of neuroinflammation in neurological and psychiatric disorders (Reprinted from (Doorduyn et al. 2010); Doorduyn et al. (2010) with permission. Copyright©2010 Elsevier)

models. In an interesting bedside-to-bench approach, the enhanced D2 stimulation in the dorsal striatum formed the basis for the development of the D2R-OE mouse model designed to overexpress dopaminergic D2 receptors in dorsal striatum during development, thereby modelling the temporal and regional-specific state of D2 hyperstimulation in schizophrenia (Abi-Dargham 2020). Interestingly, in a non-human primate model of maternal immune activation showing schizophrenia-like

aberrant behavior, PET imaging indicated increased striatal dopamine in late adolescence—a hallmark molecular biomarker of schizophrenia (Bauman et al. 2019).

This recent moving into gear with the development of various relevant schizophrenia animal models will undoubtedly boost valuable insights into disease-causative molecular and cellular pathology of schizophrenia and other psychiatric disorders, as well as the application of these models for the evaluation of novel PET/SPECT tracers.

### 1.4.2 Imaging in AD

A $\beta$  plaque-bearing mice are obviously a potentially interesting tool for the development of PET/SPECT tracers aiming at the visualisation of amyloid plaques and/or other AD-typical pathological alterations. With the development of a specific tracer that reliably visualises AD pathology in an amyloidosis mouse model, small animal imaging techniques may be used to assess treatment efficacy of anti-A $\beta$  leads in the AD drug discovery pipeline. More in general, the impact of PET/SPECT imaging on drug discovery can also be situated at various levels, including validation of mechanisms of drug localisation, establishing transport efficiency of a drug to the target, establishing drug occupancy of saturable receptor sites and determining half-time of occupancy of the drug (Eckelman 2003).

2-Deoxy-2-(<sup>18</sup>F)fluoro-D-glucose-based PET, [<sup>18</sup>F]FDG-PET, is a sensitive surrogate marker for AD diagnosis. A typical activity pattern shows decreased metabolism in the posterior cingulate and temporoparietal and prefrontal association cortex, while metabolism is maintained in the primary sensorimotor cortex and basal ganglia. Changes in [<sup>18</sup>F]FDG-PET are evident prior to the onset of cognitive symptoms in AD-related mutation carriers. Moreover, [<sup>18</sup>F]FDG-PET has predictive power for conversion of MCI to AD (Zhang et al. 2012). However, given the lack of specificity of [<sup>18</sup>F]FDG-PET for AD, much efforts are directed at the development of A $\beta$ -specific radioligands. [N-methyl-11C]2-(4'-methylaminophenyl)-6-hydroxybenzothiazole, a fluorescent analogue of thioflavin T better known as Pittsburgh compound B, [<sup>11</sup>C]PiB-PET, emerged as the lead A $\beta$ -specific PET ligand to proceed to clinical development. [<sup>11</sup>C]PiB-PET studies have demonstrated significant regional retention of radioactive signal in CNS areas with extensive plaque burden in AD brain, including the frontal, parietal and lateral temporal cortex and the striatum (Klunk et al. 2004; Price et al. 2005). [<sup>18</sup>F]FDDNP (2-(1-{6-[(2-[fluorine-18]fluoroethyl)(methyl)amino]-2-naphthyl}-ethylidene)malononitrile) is the only currently available PET radiotracer to image neurofibrillary tangles, besides A $\beta$  aggregates, and it is also the only radiotracer to visualise AD pathology in the hippocampal region of living humans (Shin et al. 2011).

Mid-2012, [<sup>18</sup>F]AV-45 (Amyvid™) became the first FDA-approved PET ligand. The search for other A $\beta$ -binding radiopharmaceuticals is still ongoing, and several [<sup>18</sup>F]- and [<sup>11</sup>C]-labelled tracers have been developed that underwent or are still undergoing extensive clinical evaluation. Besides amyloid-targeting tracers, also tau PET as a biomarker of tau pathology in AD is being scrutinized (Okamura et al. 2014; Maass et al. 2017). More information concerning current clinical trials in AD

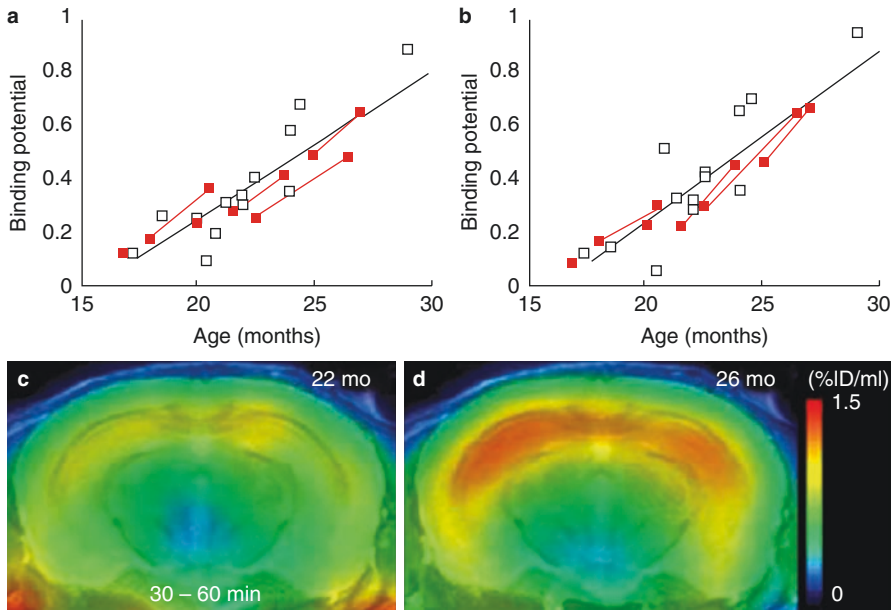
with PET and/or SPECT imaging can be obtained at <http://www.clinicaltrials.gov>, while some excellent reviews summarize the state-of-the-art in radioligand-based imaging in AD (Koric et al. 2016; Cistaro et al. 2018; Femminella et al. 2018), as well as in several interesting chapters included in this book series.

Based on often disappointing PET imaging results in AD mouse models (Klunk et al. 2005; Toyama et al. 2005; Kuntner et al. 2009; Luo et al. 2012), the usefulness of currently available transgenic mouse models for the identification and optimisation of imaging tracers is somewhat controversial. The spatial resolution of PET is often insufficient to allow reliable estimation of A $\beta$ -specific tracer binding. Repeated PET imaging in the same mouse, however, may be applied to study the evolution of plaque burden and may be useful to complement *ex vivo* autoradiographic radionuclide imaging of brain slices, which provides sufficient spatial resolution for assessment of specific tracer binding but of course lacks the possibility of time-linked observations (Wirhns et al. 2009; Teipel et al. 2011). Substantial differences in pharmacokinetics and pharmacodynamics are often observed between animals and humans for CNS targeted molecules (Carpenter Jr et al. 2009). Moreover, the detection of amyloid plaques often depends on the presence of one specific A $\beta$  subtype that may not be present in the same proportion in mouse versus man, or in different transgenic amyloidosis mouse models, as is the case for A $\beta$ N3-pyroglutamate recognised by [<sup>11</sup>C]PiB (Maeda et al. 2007). In addition, the presence of high-affinity binding sites may differ between species, as well as between the sources of A $\beta$  even within the same species (Klunk et al. 2005; Rabinovici and Jagust 2009; Rosen et al. 2011). Latter hurdle may be overcome, however, by increasing the specific activity of [<sup>11</sup>C]PiB to afford an A $\beta$ -specific binding signal in transgenic mouse brain that could be quantified with small animal PET (Maeda et al. 2007) (Fig. 1.5).

Despite differences between humans and animals, the development of new radioligands for AD brain lesions still strongly relies on preclinical tests in transgenic amyloidosis mice. For example, the [<sup>18</sup>F]AV-45 ligand, which allows significant discrimination between AD patients and healthy control individuals (Wong et al. 2010), has first been successfully tested in transgenic amyloidosis mice (Choi et al. 2009). Noteworthy is the fact that given the (relative) similarity in brain structure, as well as important protein homology (Van Dam and De Deyn 2017; Podlisny et al. 1991), non-human primates still also play an important role in the preclinical evolution of radiopharmaceuticals with regard to safety, kinetics, selectivity and specificity.

The most important advantage of (micro)PET imaging over (micro)SPECT is that of exhibiting a much higher sensitivity, while (micro)SPECT offers the possibility to widen the observational time window (based on the longer half-life of single-photon emitters) thus allowing the observation of biological processes *in vivo* several hours or days after administration of the labelled compound. By using different energy radioisotopes conjugated to different molecular targets, (micro)SPECT has the advantage over (micro)PET in being able to image several molecular events simultaneously. *In vivo* evaluation of novel SPECT tracers in AD rodent models is however limited. Most studies evaluate the *in vivo* biodistribution and brain penetration of newly developed SPECT tracers in normal mice and A $\beta$ -specific

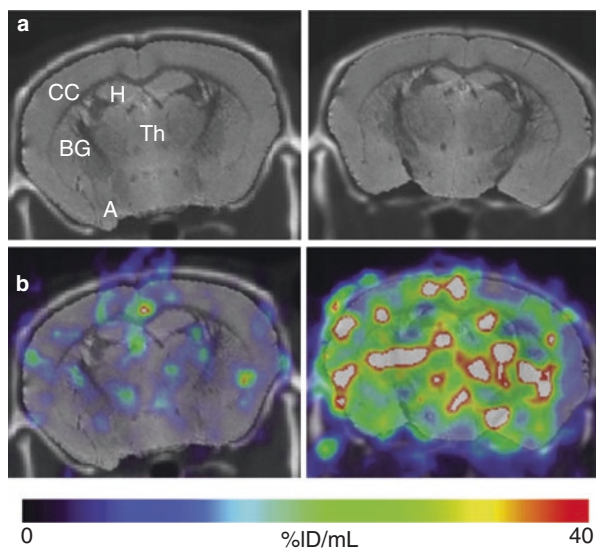




**Fig. 1.5** Ability of mouse PET to capture age-dependent evolution of brain amyloidosis. (a, b) Represent scatter plots of binding potential of N- $^{11}\text{C}$ methyl-2-(4-methylaminophenyl)-6-hydroxybenzothiazole (or  $^{11}\text{C}$ PIB for ‘Pittsburgh compound B’) versus age in the hippocampus (a) and neocortex (b) of Tg2576 mice, with the correlation of the two parameters being statistically significant in both regions. Black lines represent regression, and red elements indicate longitudinal analysis of the same individuals ( $n = 5$ ). (c, d) Represent PET images to illustrate  $^{11}\text{C}$ PIB’s substantial increase in tracer binding in the hippocampal formations of the same animal at the age of 22 (c) and 26 months (d), indicating progression of amyloid accumulation. Images were generated by averaging dynamic scan data at 30–60 min after tracer injection and were merged onto an MRI template. *mo* month (Reprinted from (Maeda et al. 2007); Maeda et al. (2007) with permission. Copyright©2007 Society for Neuroscience)

binding via *ex vivo* autoradiography on brain slices or brain homogenates of plaque-bearing rodents, often in addition to post-mortem brain tissue (extracts) of AD patients (Cui et al. 2011; Cheng et al. 2012; Martins et al. 2013; Fuchigami 2014; Molavipordanjani et al. 2019). Few published studies, however, also compared tracer retention in plaque-bearing versus control animals (Opazo et al. 2006; Ono et al. 2013; Chen et al. 2015; Kiritsis et al. 2017; Sagnou et al. 2019), and few examples of tau pathology-targeting SPECT tracer development based on tau rodent models (*ex vivo* or *in vivo*) have been published up to date (Ono 2016; Watanabe 2017).

MicroSPECT imaging is not only useful for tracer development and evaluation, but may also be used to characterize novel biological targets, with the aim to elevate the accuracy of AD diagnosis. Since amyloid or tau-related neuropathological changes may also occur in cognitively normal individuals (Johnson et al. 2013), there is a need for additional and more accurate biomarkers, or biomarker combinations. As butyrylcholinesterase (BuChE) becomes associated with amyloid and



**Fig. 1.6** Comparison of BuChE activity at mid coronal level in specific brain regions of WT (left) and 5× FAD (right) mice detected with radiotracer SPECT analysis. **(a)** CT with coregistered MR. **(b)** SPECT images acquired at 4-minute post-injection with coregistered CT/MR and ROIs. Marked retention in the 5 × FAD cerebral cortex is evident in the 5 × FAD brain compared to WT with less difference in retention evident in amygdala, hippocampus, basal ganglia, and thalamus. Image intensities expressed as %ID/mL and set to a common scale of 0%ID/mL–40%ID/mL. Abbreviations: A, amygdala; BuChE, butyrylcholinesterase; BG, basal ganglia; CC, cerebral cortex; CT, computed tomography; H, hippocampal formation; MR, magnetic resonance; ROIs, regions of interest; SPECT, single photon emission computed tomography; Th, thalamus; WT, wild-type; 5 × FAD, B6SJL-Tg(APP<sup>S</sup>SwFLon, PSEN1\*<sup>M146 L</sup>\*<sup>L286 V</sup>) 6799Vas/Mmjax mouse strain. (Adapted from (DeBay et al. 2017); DeBay et al. (2017) *Alzheimers Dement (N Y)* 3:166–176 under the CC-BY license 4.0 (<http://creativecommons.org/licenses/by/4.0>). Copyright © 2017 DeBay, Reid, Pottie, Martin, Bowen and Darvesh)

tau deposits in AD, BuChE pathology in the cerebral cortex, a region of scant BuChE activity in healthy brain, was targeted by N-methylpiperidin-4-yl 4-[<sup>123</sup>I]iodobenzoate, a BuChE radiotracer, in the 5 × FAD mouse model. Tracer retention was consistently higher in the 5 × FAD mouse versus healthy control animals, and differences were particularly evident in the cerebral cortex, paving the way for further investigation of BuChE-specific radiotracers as AD diagnostic agents (DeBay et al. 2017) (Fig. 1.6).

## 1.5 Conclusion

Patho(physio)logical alterations in rodent models of human disease are usually studied using invasive techniques, requiring sacrificing of the animals. This approach implies, amongst other, the use of a large number of animals to reach statistical significance, high associated (labour) costs and follow-up or longitudinal studies are impossible in the same animal. Recent improvements in molecular imaging

technologies employed to image small animal models, at least partially, overcome these limitations. Small animal imaging is increasingly recognised as an important facet of preclinical and translational (neuroscience) research. Perhaps most significant amongst the clear advantages of imaging experimental animals is that physiology, pathology and novel phenotypes can be understood in the most relevant milieu—in an intact, living system. Moreover, longitudinal imaging studies will allow researchers to gain more insight into the (developmental) time course of the modelled endophenotypes. The nuclear imaging modalities PET and SPECT can provide the sensitivities required to obtain the same physiological imaging acuity in small animals as can be obtained in humans, which will greatly facilitate the translation of preclinical studies to applications in the clinic. Valid animal models provide essential tools for the preclinical development of novel PET/SPECT tracers, be it via *ex vivo* autoradiography or *in vivo* tracer biodistribution and brain penetration. Besides in the framework of tracer development and evaluation, microPET/SPECT imaging support the identification of novel biological targets to elevate the accuracy of AD diagnosis.

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# The Use of Small Animal Molecular Imaging ( $\mu$ PET) Exemplified in a Neurobiological Pathology

# 2

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## Abstract

Positron emission tomography (PET) imaging is a molecular imaging modality with an extreme high versatility. Theoretically, all biological processes can be visualized longitudinally and quantified in vivo with high sensitivity, once a radioli-

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gand for a certain target of interest is developed. This chapter introduces the principle of PET and how relevant biological parameters can be extracted from PET images using pharmacokinetic modelling. In addition, we discuss awake PET imaging in small animal ( $\mu$ PET) studies. The novelty of point source tracking enables to follow tracer distribution in the brain of rodents during the conduct of certain (pathological) behavior. As a consequence, anesthesia-related factors are eliminated which continue to be an obstacle for bench-to-bedside translation. Lastly, this chapter emphasizes the value of  $\mu$ PET in the exploration and evaluation of disease mechanisms, as illustrated by a pharmacological and genetic rodent model for obsessive compulsive disorder (OCD). Longitudinal brain PET studies can be set up to investigate whether behavioral alterations parallel changes in the binding of radioligands for multiple targets. Altogether,  $\mu$ PET imaging provides unique whole brain information on underlying disease mechanisms and potential treatment targets, especially when the PET data are collected in awake animals.

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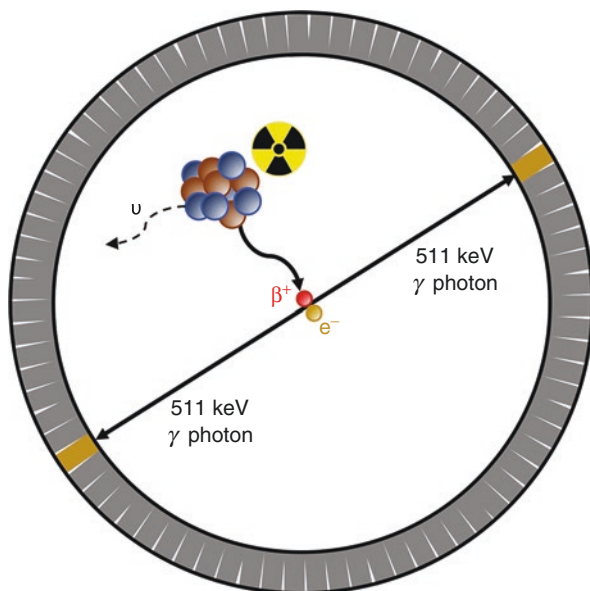
## 2.1 Introduction on $\mu$ PET

Positron emission tomography (PET) is a noninvasive imaging technique which detects the distribution of radionuclide-labelled molecules to visualize and quantify biochemical and physiological processes on a molecular level. While the first human PET cameras date back to the 1970s (Phelps et al. 1975), the application of PET imaging in rodents started about 15 years later (Ingvar et al. 1991; Hume et al. 1992). This preceded the introduction of miniaturized systems with higher spatial resolution given the smaller brain size compared to humans (Watanabe et al. 1992; Shuping et al. 1994; Bloomfield et al. 1995; Lecomte et al. 1996; Cherry et al. 1997). MPET imaging has a moderate spatial resolution (typically 1.0–1.5 mm); however, its ability to detect picomolar changes makes PET imaging the most sensitive noninvasive functional imaging technique available. PET imaging can be theoretically applied to image all biological processes once a radioligand for the target of interest has been developed, indicating the extremely high versatility of the technique. In addition, PET imaging has the potential to assess target engagement through quantitative measurements of receptor occupancy when a radioligand shares the same binding site as a drug of interest, allowing to estimate the optimal treatment dosage in vivo. Most importantly, given its noninvasive character and high sensitivity, it is well suited for longitudinal study designs and allows relatively small sample sizes to perform adequately powered experiments.

### 2.1.1 Physics and Principle of PET

PET imaging relies on the use of radioisotopes, which are unstable isotopes undergoing radioactive decay in order to form a more stable element. The radioactivity decays exponentially with a half-life,  $t_{1/2}$ , which is constant for a certain

**Fig. 2.1** Principle of PET. The radioisotope of the radioligand emits a positron ( $\beta^+$ ), which annihilates with an electron ( $e^-$ ) present in the surrounding medium and generates two antiparallel gamma ( $\gamma$ )-ray photons with an energy of 511 keV each. In the process of positron emission, also a neutrino ( $\nu$ ) is released. The photons are detected in coincidence by the PET ring detector as one event. Image reconstruction algorithms use the detected events to generate a PET image



radioisotope. Radioisotopes suited for PET imaging decay through positron ( $\beta^+$ ) emission, a positron being the antiparticle to the electron ( $e^-$ ). Once emitted, the positron loses energy and collides with an electron from the surrounding medium, a process known as the annihilation reaction (Fig. 2.1). This results in the generation of two antiparallel  $\gamma$ -ray photons with an energy of 511 keV each. The coincident detection of two photons at opposite sides of the PET ring detector (composed of scintillation crystals) generates an event. The collection of these events, followed by the application of image reconstruction algorithms, informs about the position of the radioligand in the tissue.

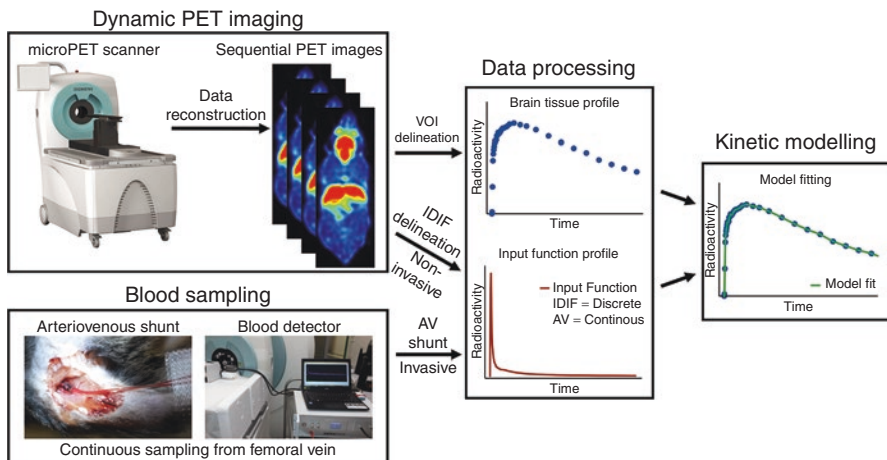
The selection of a specific radioisotope mainly depends on the atomic structure of the ligand, the possible radiolabelling reaction, and the half-life. The two most commonly employed radioisotopes are [ $^{18}\text{F}$ ] and [ $^{11}\text{C}$ ]. The former features a relatively long half-life ( $t_{1/2} = 109.8$  minutes); thus, it is generally preferred as it allows to perform acquisitions of multiple subjects with the same radiosynthesis and it can be transported to imaging centers without an on-site cyclotron. However, a fluor atom in the structure of the ligand is required, but radiofluorination is not always a viable option. On the other hand, Carbon-11 has the main advantage that C atoms are present in all molecular structures; therefore, [ $^{11}\text{C}$ ] labelling can virtually always be applied. However, this latter radioisotope is characterized by a shorter half-life ( $t_{1/2} = 20.39$  minutes), which requires a higher number of radiosyntheses, and it cannot be used by imaging centers lacking an on-site cyclotron. Nevertheless, a benefit of its short half-life is the possibility of performing repeated PET imaging studies in the same subject in a short time frame as well as the low radiation burden for the subjects.

## 2.1.2 Pharmacokinetic Modelling

Following systemic injection of a radioligand, dynamic PET imaging measures the concentration of radioactivity as a function of time. As a result, serial PET images depicting sequential time windows (frames) are created. Thus, dynamic acquisition allows the generation of a time-activity curve (TAC) of the radioligand in the region of interest over time. By combining the tissue TAC with the quantitative measurement of the plasma radioactivity (input function), it is possible to derive quantitative information about, for instance, the blood flow, enzymatic activity, and protein densities (Fig. 2.2).

A large number of factors and variables come into play when characterizing the dynamic biodistribution of a radioligand. Mathematical models simplifying the biological system can be helpful in summarizing the main relevant variables in a limited number of parameters.

Radioligand dynamics can be modelled in order to allow a quantitative prediction of the parameter of interest. The selection of an appropriate pharmacokinetic model, describing the underlying biology of the tissue, is crucial to obtain accurate parameter estimates. Since irreversible binding kinetics are not commonly used in PET imaging, only reversible binding kinetics are described. Overall, kinetic models can be divided into two main categories: compartment models, when no reference tissue is available, and reference tissue models, when a reference tissue is available.



**Fig. 2.2** Principle of dynamic PET. The radioligand is injected in the vascular system (via tail vein for rodents) and distributes with specific in vivo kinetics. The profile of the radioligand in the volumes of interest (VOI) and the amount of radioactivity in the blood (input function) can both be measured as a function of time. The input function can be obtained with either invasive (arteriovenous, AV shunt) or noninvasive (image-derived input function (IDIF)) approaches. Pharmacokinetic modelling can be performed in order to fit a specific model to the data and to derive the parameter estimations

### 2.1.3 Compartment Models

Compartment models are applied when no reference tissue is available. These models require knowledge of an input function, which can be measured with invasive or noninvasive approaches as later described. The physiological system is decomposed into a number of compartments, each contributing to the overall PET signal. Each compartment represents a part of the physiological system and is characterized by a pool of homogenous radioligand that changes over time as a consequence of the exchange of radioligand between compartments. According to compartment models, the radioactivity measured in a region of interest is composed of the radioligand distributed in different compartments at any given time. The fitting of a model to the measured radioligand profile allows to generate a model estimate of the radioligand's steady-state parameters such as the volume of distribution (described in the next section), which is related to receptor density and affinity (Gunn et al. 2001). Thus, if affinity of the radioligand toward the target is not altered, the model estimate informs about changes in receptor density.

#### 2.1.3.1 Volume of Distribution

For receptor radioligands with no reference tissue, the gold standard outcome parameter to estimate is the volume of distribution ( $V_T$ ).  $V_T$  is defined as the ratio of radioligand concentration in the tissue of interest ( $C_T$ ) (measured in kBq/cm<sup>3</sup>) to the concentration of intact radioligand in plasma ( $C_P$ ) (measured as kBq/mL) *at equilibrium* (Innis et al. 2007). It is measured in mL/cm<sup>3</sup> and can be expressed as follows:

$$V_T = \frac{C_T}{C_P} \quad (2.1)$$

This can be visually depicted when infusion of radioligand is performed. After a certain period of time, the TAC of the radioligand will reach *equilibrium*, and its measurement could provide the  $V_T$  (Fig. 2.3). For instance, a  $V_T$  equal to five means that the radioligand is concentrated in tissue by five times compared to the blood.

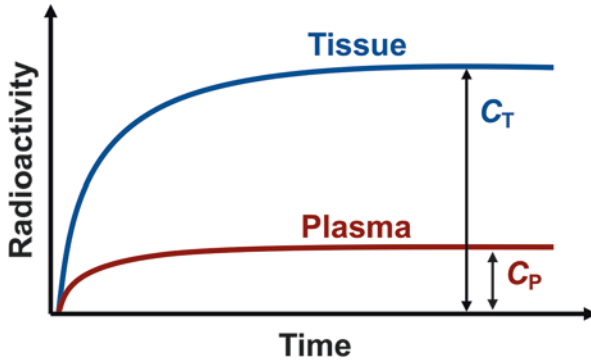
The  $C_T$  component of the  $V_T$  measurement is composed of three parameters in total:

$$C_T = C_{FT} + C_{NS} + C_S \quad (2.2)$$

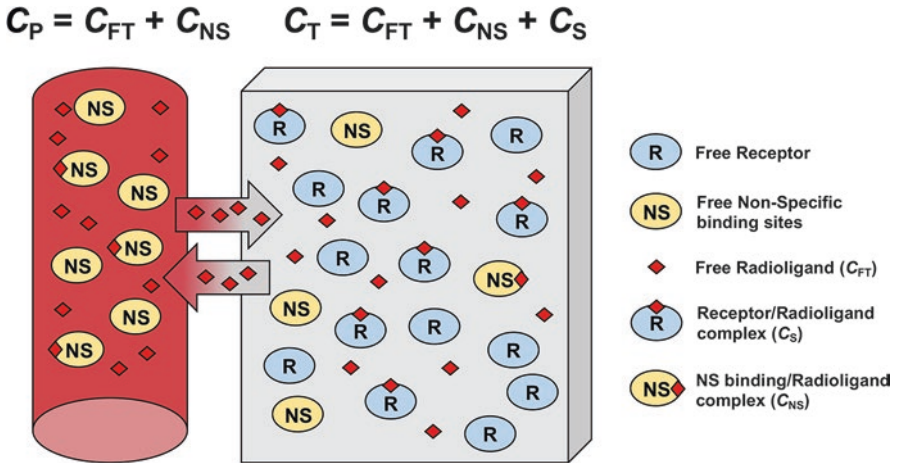
where  $C_{FT}$  represents the concentration of free radioligand in the tissue,  $C_{NS}$  the nonspecifically bound radioligand, and  $C_S$  the radioligand bound to the receptor of interest (Fig. 2.4).  $C_{FT}$  and  $C_{NS}$  can be combined as a single measurement, as they represent the total non-displaceable ( $C_{ND}$ ) radioligand uptake. This component is independent of the presence of the receptor in the tissue of interest. Thus, Eq. (2.1) can be substituted as follows:

$$V_T = V_S = V_{ND} \quad (2.3)$$

where the total volume of distribution ( $V_T$ ) is a combination of the specific ( $V_S$ ) and non-displaceable ( $V_{ND}$ ) volume of distribution.



**Fig. 2.3** Schematic representation of the radioactivity in tissue and plasma *at equilibrium*. Following radioligand infusion, it is possible to reach *equilibrium* after a certain amount of time, as shown here. *At equilibrium*, the  $V_T$  can be measured as the ratio of the concentration in the tissue ( $C_T$ ) to the concentration in plasma ( $C_P$ ) as described in Eq. (2.1)



**Fig. 2.4** Schematic representation of the components contributing to the radioligand signal in the tissue and the plasma. The concentration of radioligand in the plasma ( $C_P$ ) is the summation of the concentration of free radioligand ( $C_{FT}$ ) and the concentration of nonspecifically bound radioligand ( $C_{NS}$ ). The concentration of radioligand in the target tissue ( $C_T$ ) is constituted by  $C_{FT}$  and  $C_{NS}$ , which represent the non-displaceable component ( $C_{ND}$ ), with the addition of the concentration of specifically bound radioligand to the receptor ( $C_S$ )

$V_s$  and thus  $V_T$  are linked to the receptor density in the region of interest and the binding affinity and thus to the binding potential, which will be described later.

**2.1.3.2 Input Function**

In order to perform accurate quantification of the PET data, an input or reference tissue is required to provide information of the amount of available radioligand. A plasma input function is used for compartment models, while a brain region TAC is



employed in reference tissue models. The selection of one over the other depends on the distribution of the radioligand's target in the brain. If a region devoid of a target is present in the brain, that region can be used as reference tissue; otherwise, a plasma input function is required. The use of a brain region as reference tissue will be discussed in Sect. 2.1.4.

The input function describes the concentration of intact radioligand in arterial plasma as a function of time and represents the delivery of the radioligand to the brain. Subject cannulation for continuous arterial blood sampling with correction for radiometabolites in plasma represents the gold standard in quantitative PET data analysis. However, this procedure is laborious and invasive for patients (or small animals) (Jons et al. 1997), including many challenges and possible biases related to blood sampling, peripheral metabolism, and plasma protein binding. In particular, arterial blood sampling is specifically required for accurate measurement of the plasma radioactivity; however, arterial cannulation is invasive and, while it may be well tolerated in healthy subjects, is often a concern in ill or aged subjects with arteriosclerosis, especially for longitudinal studies. Venous blood sampling is less invasive and laborious, improving the comfort of the patient. However, radioligand venous concentration is never consistent with the arterial concentration since the venous concentration is dependent on the clearance of the compound from the tissue, thus the sampling site (Chiou 1989; Gumbleton et al. 1994; Zanotti-Fregonara et al. 2011).

Alternatively, blood sampling at a few precise moments during the scan is a valid alternative but still requires the arterial cannulation of the subject. Although blood sampling is also possible in rodents (Weber et al. 2002), it often represents a terminal procedure. Therefore, it cannot be applied for longitudinal studies. In addition, in mice, blood sampling has further limitations because the amount of available blood is extremely limited (usually 1.5–2 mL, around 58.5 mL/kg of body weight).

A possible alternative is the noninvasive extraction of the blood radioactivity data using the dynamic  $\mu$ PET data, an approach referred to as image-derived input function (IDIF). IDIF application is relevant in small animal imaging due to the difficulty of regular blood sampling without altering the physiological state (Laforest et al. 2005), especially in longitudinal study paradigms that require serial PET acquisitions. The majority of IDIF approaches use the signal from a large blood vessel or the left cardiac ventricle from the dynamic PET acquisition. However, IDIF approaches have many challenges (Zanotti-Fregonara et al. 2011) since the quality of the IDIF is affected by several factors: the measurement of whole blood, the accumulation of radiometabolites, the cardiac or large vessel region in the  $\mu$ PET scanner field of view (FOV), the sampling rate (PET frames), the cardiac rhythm, the sensitivity of the  $\mu$ PET scanner, and the quality of the reconstruction software. As many of these limitations are related to the individual  $\mu$ PET scanner and/or reconstruction software, it is usually recommended to validate an IDIF protocol in each laboratory independently. An important drawback of IDIFs is the presence of blood radiometabolites, which cannot be distinguished from the signal provided by the intact radioligand. A possible approach to overcome this restriction is the application of a population-based input function. For this approach, in vivo radioligand

metabolism is calculated at different times postinjection in an animal cohort representative of the overall population, and the average profile is calculated. However, this method has the limitation of requiring extremely highly comparable metabolic profiles of the radioligand across subjects, which, if not the case, may introduce under-corrections or overcorrections in the estimation of the parent radioligand contribution, thus in the  $V_T$ .

### 2.1.3.3 One-Tissue Compartment Model

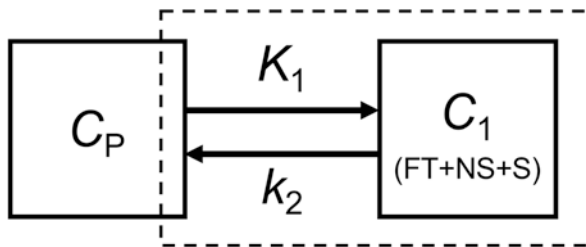
The one-tissue compartment model (ITCM) is composed of two compartments (Fig. 2.5). As for all compartment models, the first compartment is described by the input function  $C_p$ . The concentration of radioligand in the region of interest is defined by a single compartment, and thus,  $C_{FT}$ ,  $C_{NS}$ , and  $C_S$  are lumped together. Exchange of radioligand between the two compartments is represented by the rate constants  $K_1$  and  $k_2$  (Koeppel et al. 1991; Lassen 1992), and the change of activity in the tissue compartment is given by the following differential equation:

$$\frac{dC_1(t)}{dt} = K_1 C_p(t) - k_2 C_1(t) \quad (2.4)$$

When applying the ITCM, the volume of distribution can be determined as follows *at equilibrium*, hence the first order derivative being zero:

$$V_T = \frac{K_1}{k_2} \quad (2.5)$$

The two rate constants  $K_1$  and  $k_2$  represent the transport of radioligand from and to the plasma compartment, respectively.  $K_1$  denotes the unidirectional transport of the radioligand from plasma to the tissue. Its value depends on perfusion and plasma extraction, which is a combination of capillary permeability and capillary surface area. Thus,  $K_1$  is representative of the delivery rate in the region of interest. For radioligands featuring high brain permeability, it nearly equals perfusion.  $K_1$  is measured as mL/cm<sup>3</sup> per minute, while  $k_2$  describes the fractional rate constant for the washout of free radioligand transferred to plasma per unit time, measured as 1/minute.



**Fig. 2.5** Schematic overview of the one-tissue compartment model. Following the injection into the vascular system ( $C_p$ ), the radioligand is transported from the plasma to the tissue ( $C_1$ ) at a certain delivery rate ( $K_1$ ), with the free radioligand component ( $C_{FT}$ ) transferred back to plasma at a fractional rate constant ( $k_2$ ). The dashed box indicates the total component present in the tissue of interest, as a vascular component is within the brain region and cannot be separated

### 2.1.3.4 Two-Tissue Compartment Model

The two-tissue compartment model (2TCM) is composed of three compartments, as depicted in Fig. 2.6. Analogous to 1TCM, the first compartment is described by the input function  $C_p$ . In the 2TCM, the concentration of the radioligand in the region of interest is defined by two compartments: the first compartment ( $C_1$ ) is the non-displaceable contribution ( $C_{FT} + C_{NS}$ ), while the second component ( $C_2$ ) represents the specific binding to the receptor ( $C_S$ ).

Exchange of radioligand between the three compartments is described by the rate constants  $K_1$ ,  $k_2$ ,  $k_3$ , and  $k_4$  (Koepppe et al. 1991; Lassen 1992) with the following differential equations:

$$\frac{dC_1(t)}{dt} = K_1 C_p(t) - (k_2 + k_3)C_1(t) + k_4 C_2(t) \quad (2.6)$$

$$\frac{dC_2(t)}{dt} = k_3 C_1(t) - k_4 C_2(t) \quad (2.7)$$

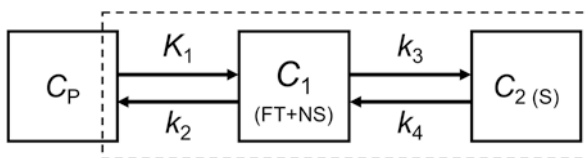
When applying the 2TCM, the volume of distribution can be determined as follows:

$$V_T = \frac{K_1}{k_2} \left( 1 + \frac{k_3}{k_4} \right) \quad (2.8)$$

Compared to 1TCM, 2TCM includes two additional rate constants.  $k_3$  represents the association rate constant to the specific binding site, while  $k_4$  describes the dissociation rate constant of the receptor-radioligand complex. Analogous to  $k_2$ , both  $k_3$  and  $k_4$  are reported per unit time (1/minute).

Although 2TCM has a larger number of parameters to fit, thus it is more sensitive to noise, and it has the advantage of allowing estimation of more complex kinetics, including approximation of  $V_{ND}$  and  $V_S$ .  $V_{ND}$  can be seen as the  $V_T$  of the first compartment  $C_1$ , and thus, it is described by a 1TCM equation as follows:

$$V_{ND} = \frac{K_1}{k_2} \quad (2.9)$$



**Fig. 2.6** Schematic overview of the two-tissue compartment model. Following the radioligand injection into the vascular system ( $C_p$ ), the tracer is transported from the plasma to the tissue ( $C_1$ ) at a certain delivery rate ( $K_1$ ), with the free radioligand component ( $C_{FT}$ ) transferred back to plasma at a fractional rate constant ( $k_2$ ). The radioligand binds specifically to the receptor ( $C_2$ ) at an association rate ( $k_3$ ) and dissociates from the receptor at a precise rate ( $k_4$ ). The dashed box indicates the total component present in the tissue of interest, as a vascular component is within the brain region and cannot be separated

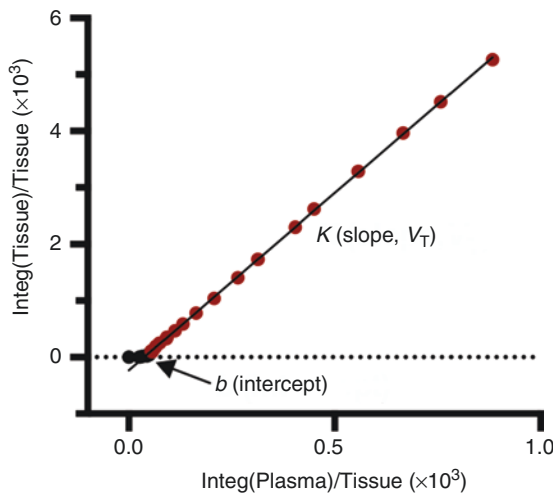
In addition, as the main interest relies on specific binding of the radioligand to the receptor, 2TCM allows to estimate the  $V_s$  as the difference between  $V_T$  and  $V_{ND}$  based on Eq. (2.3).

### 2.1.3.5 Logan Plot

One- and two-tissue compartment models, as full kinetic models, are very informative and provide estimations for the microparameters  $K_1$ ,  $k_2$ ,  $k_3$ , and  $k_4$ . However, due to the number of variables that needs to be fitted, they are sensitive to noise, and in certain situations, the accuracy of the estimation might be compromised. Therefore, the Logan plot (Logan et al. 1990) represents a robust alternative for radioligands with reversible kinetics. It is a “graphical” method that normalizes the measurements using a mathematical transformation, which simplifies the differential equations of the more complex compartment models to a linearized model:

$$\frac{\int_0^t C_T(\tau) d\tau}{C_T(t)} = K \frac{\int_0^t C_P(\tau) d\tau}{C_T(t)} + b \quad (2.10)$$

As a result, the estimation of the  $V_T$  (slope  $K$ ) using a Logan plot is easier compared to using compartment models (Fig. 2.7). However, also the Logan plot is susceptible to noise in the data for low activities, which can result in an underestimation of the  $V_T$  (Logan et al. 2001; Varga and Szabo 2002). For this reason, before application of this linearization approach, it is always recommended to verify its stability over the compartment models.



**Fig. 2.7** Representative Logan plot fitting. As described in Eq. (2.10), when applying the Logan plot graphical analysis, the integral of the tissue activity over the tissue activity is plotted versus the integral of the input function over the tissue activity. The result of the fitting is a straight line (black line) starting after an equilibration period defined as  $t^*$ . The slope of the fitting,  $K$ , depicts the  $V_T$  value. The tissue activity values considered for the linearization are depicted in red, while values before  $t^*$  are depicted in black

### 2.1.4 Reference Tissue Models

Unlike compartment models, reference tissue models have the advantage of not requiring a vascular input function. Thus, all challenges and possible biases related to blood sampling, peripheral metabolism, and plasma protein binding are absent. These models provide an estimate of the binding potential (BP) of a radioligand-receptor complex. The concept of BP is described in the next section.

To apply reference tissue models, there are two conditions that need to be met: (i) A reference tissue completely devoid of target, i.e., with no (or negligible) specific binding of the radioligand, is required (Lammertsma and Hume 1996). (ii) The  $V_{ND}$  of the radioligand should be similar in all investigated regions, i.e., the ratio  $K_1$  over  $k_2$  is comparable. Thus:

$$\frac{K_1}{k_2} = \frac{K_1'}{k_2'} \quad (2.11)$$

where  $K_1'$  and  $k_2'$  are the microparameters of the reference tissue.

Based on the equation above, it is possible to estimate the delivery ratio  $R_1$ :

$$R_1 = \frac{K_1}{K_1'} = \frac{k_2}{k_2'} \quad (2.12)$$

which describes the ratio of relative influx in the region of interest and in the reference tissue. As a ratio,  $R_1$  is unitless.

#### 2.1.4.1 Binding Potential

When applying reference tissue models, the gold standard is represented by the binding potential (BP). BP is the ratio of the total density of receptors in the tissue (receptor concentration) ( $B_{max}$ ) to the equilibrium dissociation constant of the radioligand ( $K_D$ ) (Mintun et al. 1984). Thus, BP is given by a combination of receptor density and affinity of the radioligand toward the receptor as follows:

$$BP = \frac{B_{max}}{K_D} = \text{receptor density} \times \text{affinity} \quad (2.13)$$

When performing PET imaging, BP can be determined in vivo if a region with only non-displaceable uptake exists, in order to estimate the so-called non-displaceable BP ( $BP_{ND}$ ) (Innis et al. 2007).  $BP_{ND}$  quantifies the equilibrium concentration of specific binding as compared to a region with only non-displaceable binding (reference tissue) and can be calculated as follows:

$$BP_{ND} = \frac{(V_T - V_{ND})}{V_{ND}} = \frac{V_T}{V_{ND}} - 1 = \frac{k_3}{k_4} \quad (2.14)$$

The ratio  $V_T$  to  $V_{ND}$  is referred to as distribution volume ratio (DVR) (Logan et al. 1996), and it can be determined using the compartment models if a reference tissue region is available for calculation of  $V_{ND}$ .  $BP_{ND}$  is a ratio of two volumes of distribution and is therefore unitless.

### 2.1.4.2 Simplified Reference Tissue Model

The simplified reference tissue model (SRTM) is a simplification of the full reference tissue model in which no distinction is made between the non-displaceable and specific components of the target region, as depicted in Fig. 2.8. For this reason, it performs the best when the kinetic of the radioligand could be described using 1TCM (Lammertsma and Hume 1996). In the SRTM, a total of three components can be identified: the input function ( $C_P$ ), the region of interest ( $C_{ROI}$ ), and the reference tissue ( $C_{REF}$ ). As the input function  $C_P$  is the same for both  $C_{ROI}$  and  $C_{REF}$ , its measurement is not required if the delivery rates ( $K_1$  and  $K_1'$ ) are similar. In addition, the  $C_{REF}$  needs to be devoid of specific binding (Lammertsma and Hume 1996).

Exchange of radioligand for SRTM can then be described by the following differential equation:

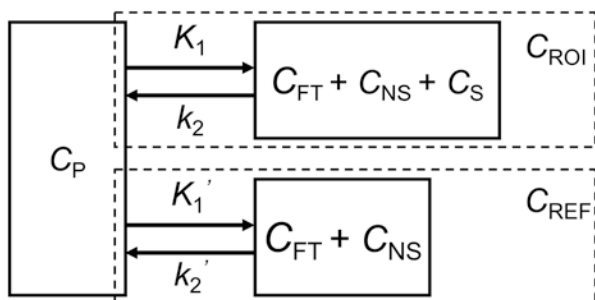
$$\frac{dC_{ROI}(t)}{dt} = R_1 \frac{dC_{REF}(t)}{dt} + k_2 C_{REF}(t) - \frac{k_2}{1 + BP_{ND}} C_{ROI}(t) \quad (2.15)$$

### 2.1.5 Receptor Occupancy and Displacement

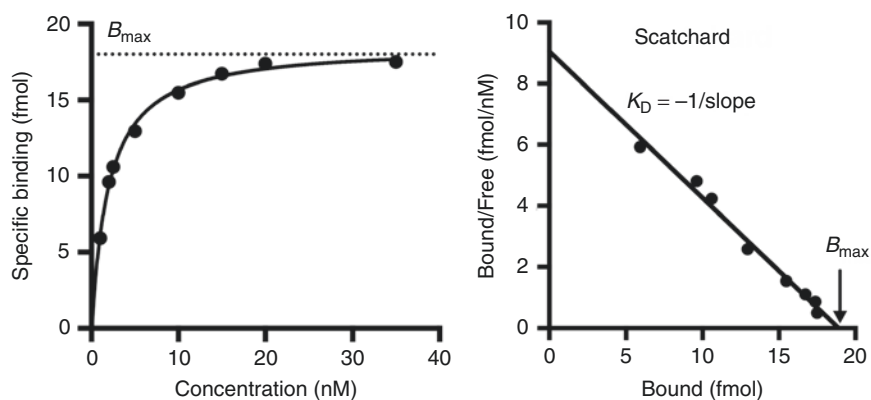
In addition to quantification of receptor density, PET imaging has the potential to assess target engagement through quantitative measurements of receptor occupancy (RO) (Lassen et al. 1995; Cunningham et al. 2010). RO of the radioligand binding site by a treatment drug can be described as follows:

$$RO = \left( 1 - \frac{B_{available}(\text{treatment})}{B_{available}(\text{baseline})} \right) \times 100\% \quad (2.16)$$

where  $B_{available}(\text{treatment})$  and  $B_{available}(\text{baseline})$  represent the density of available receptors for the radioligand under the drug condition and during baseline,



**Fig. 2.8** Schematic overview of the simplified reference tissue model. Following the injection into the vascular system ( $C_P$ ), the radioligand is transported from the plasma to the tissue of interest ( $C_{ROI}$ ) and the reference ( $C_{REF}$ ) at a similar delivery rate ( $K_1$  and  $K_1'$  prime), with the free radioligand component ( $C_{FT}$ ) transferred back to the plasma at a fractional rate constant ( $k_2$  and  $k_2'$ ). The non-displaceable binding ( $C_{FT} + C_{NS}$ ) is comparable between the tissue of interest ( $C_{ROI}$ ) and the reference ( $C_{REF}$ ). The dashed box indicates the total component present in the tissue regions



**Fig. 2.9** Representative saturation study to estimate  $B_{max}$  and  $K_D$ . The receptor density ( $B_{max}$ ) and equilibrium dissociation constant ( $K_D$ ) can be estimated by generating a saturation curve where the specific binding of a compound is measured at different doses (concentrations of compound) (left). The estimation can be linearized by transforming the data of the saturation curve in a Scatchard plot (right), where the intercept with the  $x$ -axis represented the  $B_{max}$  and the  $K_D$  is  $-1/\text{slope}$

respectively (Innis et al. 2007). For instance, this paradigm is extremely useful in the determination of drug dosage to determine the desired in vivo target occupancy (Cunningham et al. 2005). Additionally,  $B_{max}$  and  $K_D$  can be estimated by performing a saturation binding study to generate a curve where the specific binding of the compound of interest is measured at different doses (concentrations of compound) (Fig. 2.9). This estimation can also be represented linearly by transforming the data of the saturation curve in a Scatchard plot, where the intercept with the  $X$ -axis represented the  $B_{max}$  and the  $K_D$  is  $-1/\text{slope}$  (Fig. 2.9). Notably, by performing serial PET scans with increasing amounts of cold ligand, thus with cumulative target saturation studies, an estimation of the in vivo  $B_{max}$  and  $K_D$  can be determined. Alternatively, PET scans can be performed using a competitive ligand to block or displace the receptor. This approach has the potential to confirm the specificity of the binding of novel radioligands, to provide an estimate of the amount of non-displaceable binding contributing to the PET signal in the tissue of interest, and to investigate whether a reference tissue region devoid of specific binding exists.

### 2.1.6 Semiquantitative Approach

When a radioligand presents slow and stable (reversible or irreversible) kinetics, the quantitative pharmacokinetic modeling based on dynamic imaging may be replaced by a simpler semiquantitative static approach. The most common static measurement is the standardized uptake value (SUV). Unlike dynamic imaging, SUV is based on a static acquisition starting when, following radioligand system injection,

the kinetics of the radiotracer have stabilized. SUV typically lasts for a time window of 15–20 minutes, and it is measured as follows:

$$\text{SUV} = \frac{C_t}{\text{ID} / \text{BW}} \quad (2.17)$$

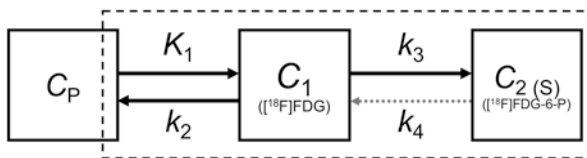
where  $C_t$  represents the concentration of activity in the tissue of interest, ID is the dose injected into the animal, and BW is the body weight of the subject.

The most common radioligand measured with semiquantitative approaches is [ $^{18}\text{F}$ ]2-fluoro-2-deoxy-D-glucose ([ $^{18}\text{F}$ ]FDG), thanks to its semi-irreversible kinetics. [ $^{18}\text{F}$ ]FDG is a glucose analogue; therefore, it can be used to measure glucose metabolism.

Following system injection, [ $^{18}\text{F}$ ]FDG enters the brain via glucose transporters. Due to its similarity to glucose, [ $^{18}\text{F}$ ]FDG directly competes with glucose for the binding to the glucose transporters in the brain. Because of this competition, animals are generally fasted several hours before [ $^{18}\text{F}$ ]FDG administration in order to guarantee low blood glucose levels and, consequently, high cerebral [ $^{18}\text{F}$ ]FDG uptake (Deleye et al. 2014). Once in the brain, free [ $^{18}\text{F}$ ]FDG can either be transported back to the vascular system ( $C_p$ ) or be phosphorylated by hexokinase to [ $^{18}\text{F}$ ]FDG-6-phosphate ([ $^{18}\text{F}$ ]FDG-6-P). The latter cannot be further metabolized through metabolic pathways and remains trapped into the tissue. Thus, as depicted in Fig. 2.10, [ $^{18}\text{F}$ ]FDG kinetics for glucose transport and phosphorylation in the brain resemble the 2TCM, as described above. Although the [ $^{18}\text{F}$ ]FDG-6-P uptake is not fully irreversible, its efflux constant ( $k_4$ ) is much slower than the metabolic rate ( $k_3$ ); therefore,  $k_4$  can be assumed to be equal to zero.

### 2.1.7 Radioligands

Given the versatility of PET imaging, this technique can be theoretically applied to image all biological processes if a radioligand for the target of interest is available.



**Fig. 2.10** Schematic overview of the 2TCM applied to [ $^{18}\text{F}$ ]FDG. Following [ $^{18}\text{F}$ ]FDG injection into the vascular system ( $C_p$ ), the radiotracer is transported from the plasma to the tissue ( $C_1$ ) at a certain delivery rate ( $K_1$ ), with the free radioligand component ( $C_{\text{FT}}$ ) transferred back to the plasma at a fractional rate constant ( $k_2$ ). [ $^{18}\text{F}$ ]FDG is phosphorylated into [ $^{18}\text{F}$ ]FDG-6-P ( $C_2$ ) at a certain metabolic rate ( $k_3$ ), and its efflux from the tissue ( $k_4$ ) occurs at a much slower rate; thus, it is assumed to be negligible. The dashed box indicates the total component present in the tissue of interest, as a vascular component is within the brain region and cannot be separated



Nonetheless, reversibly binding radioligands feature a series of requirements that need to be met in order to be suitable for receptor imaging:

- *Availability of the target*: The amount of target available for the radioligand depends on the specific target. For receptor imaging, this amount is generally in the nanomolar range; thus, the amount of radioligand injected into the animal has to be consistent (and low) to avoid any target saturation or pharmacological effect.
- *Specificity toward the target*: In order to investigate the pharmacokinetics, the process of reversible binding of the radioligand from the target should be the main, if not the only, process to occur in the region of interest. This means that the presence of labelled metabolites should be minimal or absent and a low specific binding toward any other molecule is desired. For instance, the radioligands for the dopaminergic  $D_1$  receptor, namely, [ $^{11}C$ ]SCH23390 and [ $^{11}C$ ]NNC-112, feature affinity toward serotonin 2A receptors (Ekelund et al. 2007).
- *Radiometabolites*: Following injection into the vascular system, radioligands undergo metabolism, which results in the generation of radiometabolites as a consequence of xenobiotic-degrading enzymes in the blood, liver, and kidneys. For brain imaging, this does not represent a major drawback as long as the radiometabolites generated in plasma are polar and cannot penetrate the blood-brain barrier (BBB) and no metabolism occurs in the brain.
- *Low nonspecific binding*: Another variable that might hamper the sensitivity of a radioligand is the amount of nonspecific binding in the tissue of interest. Ideally, a good radioligand washes out from a region quickly if no target is present. However, radioligands might have nonspecific binding toward other undefined molecules causing longer retention. This represents the major contributor to the non-displaceable binding component and cannot be avoided.
- *Trace amount*: The radioligand introduced to the system needs to be in trace amount in order to avoid perturbation of the process meant to be measured. Accordingly, less than 5% of receptors should be occupied by the radioligand (Hume et al. 1998). However, this is challenging in small animals because of the low amount of total target available, which demands extremely high molar activities. Molar activity is a measure of the amount of radioactivity in a given number of moles of ligand. Thus, the higher the molar activity, the less non-labelled molecules of the radioligand are injected into the system and compete for the receptor (Elsinga 2002). This is particularly relevant for small animal imaging where the amount of injected radioligand is often above the 5% occupancy (trace amount) and a high-affinity radioligand is used. Thus, the calculated injected mass of each animal should always be corrected for the molar activity of the specific radiosynthesis in order to avoid changes in receptor occupancy across subjects. This erroneous methodology can generate artificial changes in radioligand binding.

- *Low plasma protein binding*: Following injection into the vascular system, a portion of the total radioligand binds to plasma proteins. The binding to plasma proteins means less free fraction of radioligand available to reach the region of interest, and high protein binding leads to prolonged blood circulation. Plasma protein binding is generally related to the lipophilicity of the molecule.
- *Target accessibility*: The major factor avoiding target accessibility in brain imaging is the BBB. While radiometabolites should not cross the BBB, high penetration by the radioligand is critical to allow in vivo pharmacokinetic analysis. It is poorly understood which factors contribute to membrane permeability, with the exception of lipophilicity.
- *Appropriate in vivo pharmacokinetics*: In order to perform dynamic PET imaging and obtain accurate parameter estimations for a radioligand, it is critical that the in vivo pharmacokinetic profile is in the time window of PET scan acquisition time and isotope decay. For instance, PET imaging is sensitive to noise; thus, the temporal sampling rate of the kinetic data has some restrictions. However, if radioligand efflux from the region of interest is extremely fast, the PET scanner cannot detect it accurately. Conversely, if the influx and efflux of the radioligand labelled with C-11 ( $t_{1/2} = 20.39$  minutes) into the region of interest is too slow, thus requiring a longer acquisition time, the kinetic data may become too noisy for accurate quantification.

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## 2.2 PET Motion Correction: State of the Art

One technological innovation in small animal PET imaging that is of particular interest to imaging of neurobiological systems is PET imaging of awake unconstrained animal. As such, an approach has the potential to circumvent the, often unknown, impact of anesthesia on the PET reading of the brain function. Since the first demonstration of awake animal imaging, now already 10 years ago, a few research groups have worked on further improving the technology (Kyme et al. 2008). Our group has mainly focused on making the method practically feasible in a routine small animal PET imaging setting. As this technology is now becoming available, we can not only avoid the impact of anesthesia on the PET outcome, but we can also implement previously unfeasible experimental paradigms where we can monitor and/or alter behavior during PET imaging. Indeed, this approach has great potential to dynamically monitor brain uptake of a radioligand during the occurrence of, for example, compulsive-like behaviors in animal models for obsessive compulsive disorder, as discussed in Sect. 2.3. In addition, the method also allows to image interacting animals in larger imaging systems (Miranda et al. 2019b). The next paragraphs introduce the problem of anesthesia, give a brief overview of several approaches to awake animal imaging, and finally demonstrate the technique in a typical small animal brain PET study with excessive animal motion due to pharmacological stimulation.

### 2.2.1 The Need for Awake PET: Anesthesia, Stress, and Its Impact on Small Animal PET

A major difference between preclinical brain PET research in small animals and human brain PET is the use of anesthesia to immobilize the animal during the study to avoid image blurring caused by animal motion. In contrast, human subjects are usually scanned in a conscious state. Several studies have investigated to which degree anesthesia affects the animal's brain response for specific PET radiotracers as reviewed by Alstrup and colleagues (Alstrup et al. 2013). For example, in line with previous *ex vivo* studies using 2- $^{14}\text{C}$ -deoxyglucose (Sokoloff et al. 1977; Lenz et al. 1998), an overall lower  $^{18}\text{F}$ FDG brain uptake was observed in anesthetized animals (Toyama et al. 2004; Mizuma et al. 2010). However, for certain radiotracers, the effect of anesthesia on the PET uptake is still unknown.

Injectable and inhalant anesthetic agents are used to immobilize small animals to perform imaging scans. Due to its practicality, inhalant anesthetic agents are more commonly used.

Since scanning of awake animals under physical restraint has been proposed to avoid the effects of anesthesia, studies have also been performed to assess the impact of restraining stress on the radiotracer brain uptake. Sung et al. (2009) performed  $^{18}\text{F}$ FDG scans of rats that underwent restraining stress 1 and 2 h prior to a PET scan. It was shown that restraining stress, induced by fixing the rat limbs, modifies the  $^{18}\text{F}$ FDG brain uptake in comparison with unrestrained rats. Even after a 1 h recovery from restraining, the change in  $^{18}\text{F}$ FDG brain uptake in comparison with control rats endured.

The effect of anesthesia and stress on the animal's brain response have motivated research in performing brain PET scans of awake and unrestrained small animals. A method proposed by Schulz et al. (2011) makes use of a specially designed miniaturized PET scanner attached to a rat's head. A mechanical system is needed to counterbalance the weight of the PET scanner and allow motion of the rat. Another approach proposed by Balasse et al. (2015) measures the activity in small regions of the awake rat brain using a beta-probe. A portable wireless measurement system for the beta-probe was built inside a backpack that is placed on the rat. These methods require specially designed devices and surgical implantation of the device in the skull of the rat.

A different approach that makes use of typical preclinical PET scanners to perform awake rat PET scans was proposed by Kyme et al. (2008). In this method, the motion of the rat head is measured while it is scanned, and the PET data is corrected for motion using the tracking data. Awake mouse brain scanning has also been investigated (Baba et al. 2013), although the problem is more challenging than rat awake scanning due to the smaller size of the animal.

Finally, awake brain scanning not only avoids undesired effects on the animal's brain response from anesthesia and stress; it would also allow to expand the experimental design to setups where the conscious response of the subject to external sensorial stimuli could be investigated, similar to what has been done for several years in human brain PET and MRI studies (Frith et al. 1991; Reiman et al. 1997; Crespo-Facorro et al. 2001).

## 2.2.2 Motion Correction in PET Reconstruction

In order to perform brain PET scans in awake animals, methods for human brain rigid motion correction can be adapted to be used in preclinical PET scanners. One of these methods consists of measuring the six degrees of freedom (DOF) motion of the human head and relocating each line-of-response (LOR) back to a reference pose (translation and rotation) according to the measured motion with (Menke et al. 1996) or without (Rahmim et al. 2004) additional rebinning to a sinogram. This approach is convenient to use when continuous motion is present and is therefore the usual method for small animal PET brain motion correction. The method requires data acquisition in list-mode format, a condition fulfilled in most modern PET scanners. In the next section, methods to perform the motion tracking of the animal head are detailed.

## 2.2.3 Head Motion Tracking for Awake Small Animal PET

The challenges in awake small animal brain PET scanning are considerably greater than in human brain PET motion correction. The motion of freely running small animals is considerably larger than the motion of a human head inside a scanner. In addition, the tracking devices need to be able to track inside smaller bore size scanners than in human motion tracking. Also, the tracking accuracy needs to be higher to avoid image degradation in preclinical scanners with finer spatial resolution. Moreover, different correction approaches in the motion correction algorithm need to be implemented or developed to obtain reconstructions with quantitative information.

### 2.2.3.1 State of the Art

Several approaches have been proposed to track the animal head motion. One of the earliest methods is the one proposed by Kyme et al. (2008). In this method the marker-based optical tracking camera MicronTracker (Claron Technology Inc., Toronto, Canada) is used to measure the motion of the rat's head during the PET scan. A small rectangular marker is pasted on the head to measure its motion. This method performs well if the motion of the rat is restricted to face toward the tracking camera, normally placed in front of the scanner. If the rat turns around or the marker is occluded by the scanner bore, the tracking information is lost. With this method, motion-corrected images are obtained by repositioning of the LORs and performing LOR rebinning or list-mode motion correction.

Kyme et al. (2014) also developed a markerless rat head tracking method. By acquiring different views from the rat head and matching face features in the different views, the pose of the head is determined. Although in theory natural features of the rat head can be used (e.g., eyes), in this work, it was found that painting a pattern on the rat fur is necessary to increase the accuracy of the motion tracking data.

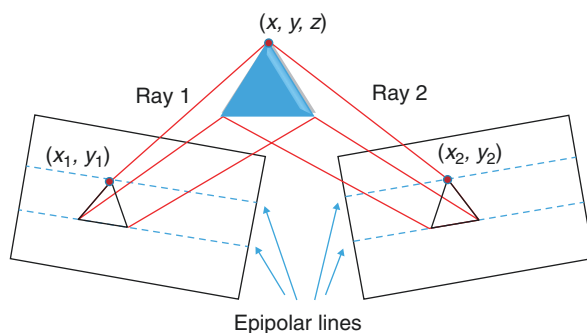
### 2.2.3.2 Optical Tracking Methods

Optical tracking methods work with the principle of stereovision. For these methods, features in different views of the space that needs to be mapped are used to determine the 3D location of points in this space. In addition, since the tracking data is measured with respect to the camera coordinate system and time, a spatial and temporal calibration needs to be performed to transform the camera tracking information to the coordinate system and time of the PET scanner.

#### Stereovision

Two 2D images acquired from two different views of the same scene can be used to calculate the 3D position of points in the scene. A 2D image can be seen as a planar projection of a 3D scene on which each point in the 2D image plane is the image of a 3D point on the scene which intersects a 3D ray (line) coming from the camera that forms the image. By triangulation of two of these rays that intersect the same point (from different views), the 3D coordinate of the point can be calculated (Fig. 2.11). If all points (up to the image resolution) on the two different images can be related to a point in the other image, a disparity map can be calculated by triangulating all points. The disparity map is then a 3D surface mapped space.

There also exist other devices that could be used for motion tracking using 3D vision. These devices have different working principles. For example, time-of-flight (TOF) cameras measure the distance from several points on the surface of objects to the camera by measuring the distance travelled from light projected and detected back to the camera. This information renders a 3D map of the object. Most of these cameras are used for broad-field applications, such as geological surface mapping



**Fig. 2.11** By determining that the point  $(x_1, y_1)$  on image 1 corresponds to the image of the point  $(x_2, y_2)$  on the second image, e.g., using image correlation, and with knowledge of the camera intrinsic parameters, the 3D point  $(x, y, z)$  can be calculated. The intrinsic parameters of the camera serve to determine the equations of ray 1 and 2. The 3D point  $(x, y, z)$  can be found by triangulation of these two rays. The search for the corresponding 2D point in the other image can be simplified by calculating the epipolar lines. The epipolar line in one image will contain all points of the corresponding epipolar line of the other image (if the points are within the corresponding field of view of the camera)

and atmospheric mapping. To our knowledge, no high precision TOF camera for small-scale applications with high frame rate, such as that required for PET motion tracking, is commercially available.

### 2.2.3.3 PET-Based Motion Tracking

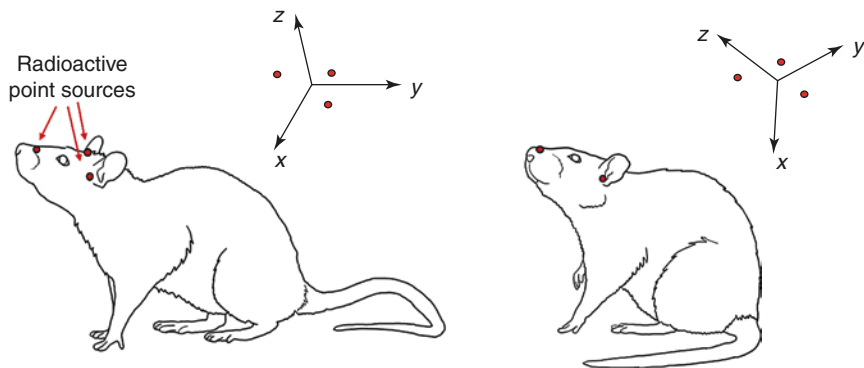
As the PET image provides the 3D location in space of the radiotracer, in theory, this information could be used to determine the pose of the object containing the radiotracer. However, short time-frame PET images, required to determine fast motion, present high noise, thereby hindering the possibility of accurate image registration. In addition, the radiotracer distribution changes over time, making it difficult to define a unique image to perform the registration.

The problem of high image noise in the short time frames and changing tracer distribution can be solved by using radioactive point sources as fiducial markers. The point sources can be accurately reconstructed in short time frames, and their activity distribution does not change over time.

#### Point Source Tracking

For this method, radioactive point sources are used as markers attached to the object to be tracked. By detecting the position of at least three noncollinear points in 3D space arranged in an asymmetric cluster, the six DOF of the point configuration (and of the object they are attached to) can be uniquely defined. This method performs the motion tracking without additional hardware, and therefore, a spatial and temporal synchronizations are not necessary, which avoid any associated calibration errors.

The point sources used as markers are prepared using sodium polyacrylate grains with sizes of less than 1 mm. The grains are repeatedly soaked in  $[^{18}\text{F}]\text{FDG}$  for a few seconds until an activity between 148 and 222 kBq is reached. The radioactive point sources are attached to the object prior to the PET scan. At least three point sources must be attached to the head of the rat or mouse (Fig. 2.12).



**Fig. 2.12** By attaching radioactive point sources to the head of the rat (or mouse), its rigid motion can be determined by defining a rigid coordinate system fixed with respect to the point source geometry

The tracking algorithm works by identifying the position of point sources in the image space. In order to track high-speed motion, the PET data is divided in short time frames, and these frames are reconstructed. The duration of every frame is 32 ms, i.e., a tracking frame rate of 31.2 Hz.

### 2.2.4 Awake [ $^{18}\text{F}$ ]FDG PET Imaging of Memantine-Induced Brain Activation and Test-Retest in Freely Running Mice

The point source tracking method has been used to perform a [ $^{18}\text{F}$ ]FDG brain uptake experiment in freely running mice (Miranda et al. 2019a). A test-retest and challenge study using the stimulant drug memantine was performed. For these experiments, point sources were attached to the head of the mouse to track their motion during the PET scan. Awake mice were placed on a cylindrical holder inside the PET scanner and scanned during 20 minutes. A test, a retest, and a memantine challenge scan were performed in 8 mice (24 scans in total). Additionally, eight mice were scanned in the same three conditions under anesthesia to obtain a motion-free image for comparison. Since an awake [ $^{18}\text{F}$ ]FDG uptake period of 20/30 minutes was allowed before the beginning of the PET scan for both the awake and the anesthesia scans, [ $^{18}\text{F}$ ]FDG brain uptake in the anesthesia and the awake state should be similar (Matsumura et al. 2003).

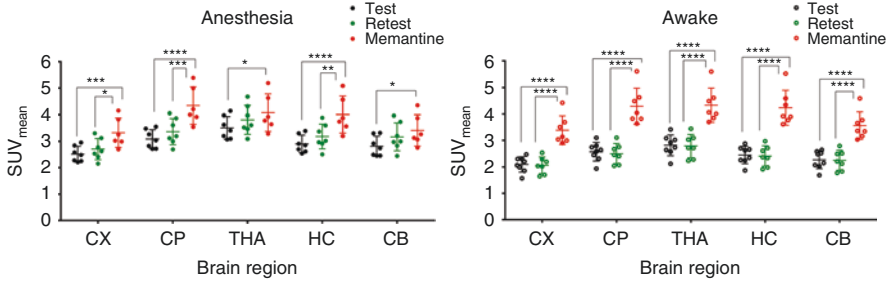
#### 2.2.4.1 Mouse Behavior in Awake Scans

The average speed of the mice during the test-retest scans was 2.09 cm/s and more than doubled during the memantine scans to 4.25 cm/s. The travelled distance during the memantine scans (18.2 m) was significantly higher compared to the test-retest conditions (6.49 m). During test-retest scans, mice moved sporadically and remained in the same location for long periods of time. On the other hand, during memantine scans, mice moved constantly during the entire scan duration with short resting periods.

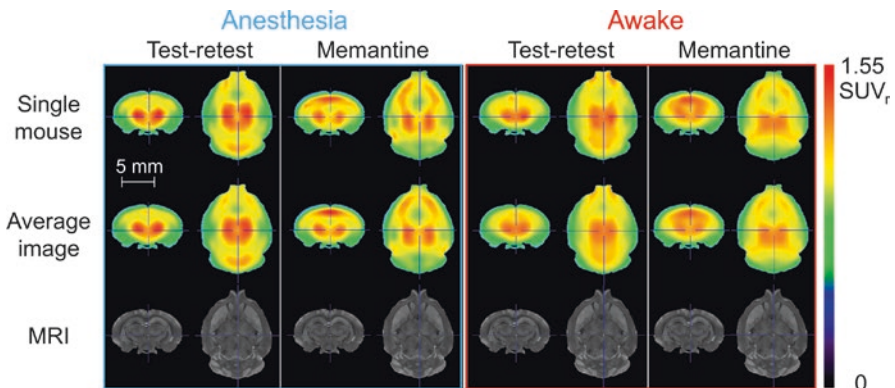
#### 2.2.4.2 Brain Regional Quantification

Figure 2.13 shows the regional brain [ $^{18}\text{F}$ ]FDG uptake in the test, retest, and memantine conditions, in both awake and anesthetized mice. The uptake in the cortex, the caudate putamen, the thalamus, the hippocampus, and the cerebellum was investigated. In both the anesthesia and the awake state, there is a significant difference ( $p < 0.05$ ) in regional brain uptake between test-retest and memantine conditions, except in the thalamus and cerebellum in the anesthesia state. Differences between test-retest and memantine conditions were more pronounced in the awake state in comparison with the anesthesia state. This might be caused by the use of anesthesia before (awake) [ $^{18}\text{F}$ ]FDG injection to attach the point sources in awake mice. In both the awake and the anesthesia state, there was no significant difference between the test and the retest conditions.

When comparing the uptake in the anesthesia and awake groups, the test and retest uptake tends to be lower in the awake state. On the other hand, there was no significant difference in uptake when comparing anesthetized with awake animals in the



**Fig. 2.13** Scatter plot of [ $^{18}\text{F}$ ]FDG standard uptake values (SUV) for different brain regions in the test, the retest, and the memantine challenge conditions for both the anesthesia and the awake group. (Significance level \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ , \*\*\*\* $p < 0.0001$ . CX cortex, CP caudate putamen, TH thalamus, HC hippocampus, CB cerebellum. This research was originally published in *JNM*. Miranda A, et al. Awake  $^{18}\text{F}$ -FDG PET Imaging of Memantine-Induced Brain Activation and Test–Retest in Freely Running Mice. *J Nucl Med*. 2019;60:844–850. © SNMMI)



**Fig. 2.14** The anesthesia and awake [ $^{18}\text{F}$ ]FDG images for a single mouse and averaged over all mice from each group. MRI template images are included as an anatomical reference. The test-retest and the memantine challenge conditions are displayed for each group. Image values are normalized to whole brain uptake for visualization purposes. (SUV ratio (SUV<sub>r</sub>); This research was originally published in *JNM*. Miranda A, et al. Awake  $^{18}\text{F}$ -FDG PET Imaging of Memantine-Induced Brain Activation and Test–Retest in Freely Running Mice. *J Nucl Med*. 2019;60:844–850. © SNMMI)

memantine condition. This might be caused by the longer waiting period between the administration of the anesthesia to paste the point sources and the [ $^{18}\text{F}$ ]FDG injection in the memantine scans (40 minutes), compared to the test-retest scans (10 minutes).

Figure 2.14 shows the brain reconstructions for all conditions in the anesthetized and the awake mice. The brain uptake pattern is similar between the anesthesia and the awake states, but awake reconstructions show a lower spatial resolution. There is a change in the memantine condition brain uptake pattern compared to the test-retest condition. This change is observable in both the anesthesia and the awake states. Compared to the test-retest uptake, the memantine brain uptake is reduced in the cerebellum and increased in the hippocampus, relative to other brain regions.



## 2.3 Illustration of PET Applications Exemplified for a Neuropsychiatric Disorder

Previously, *ex vivo* techniques are predominated in basic research to obtain information on pathological processes in animal models of disease. The development of dedicated small animal imaging equipment enabled the opportunity to collect such information *in vivo*. With  $\mu$ PET, interactions between a radioligand and its target can be quantified on a molecular level within a picomolar range and with high sensitivity. In this way,  $\mu$ PET has acquired an important role in preclinical research. This modality creates the opportunity to perform longitudinal study designs, wherein each animal can function as its own control. This allows noninvasive monitoring of biological processes and disease progression but also therapy response, thereby improving the scientific value of the obtained data using paired observations. Also, fewer animals can be used. In addition, exploring (patho)physiology in small animals using  $\mu$ PET is associated with a high potential for translation to a clinical setting. The same radioligands can be used for both animal and human research.

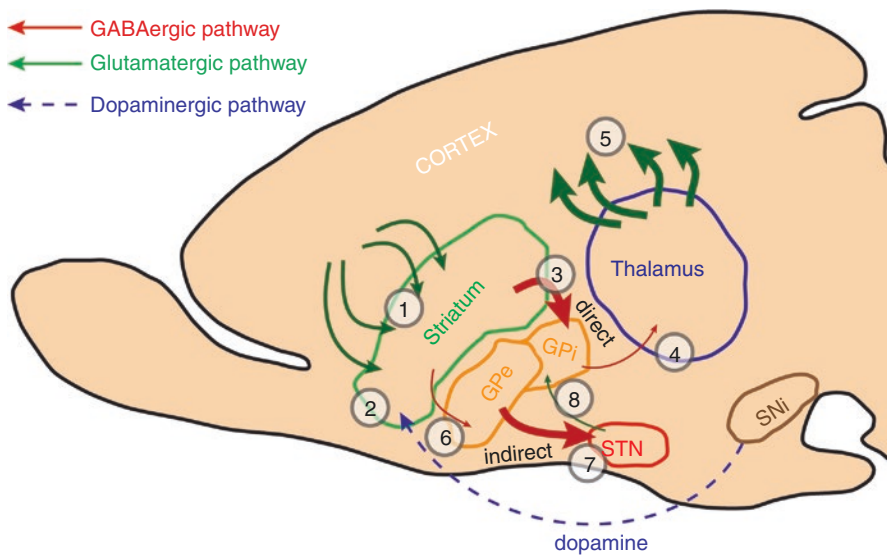
In disease model studies, for example, the animal's behavior can function as a readout for the presence of a certain pathology. Longitudinal brain PET studies can be designed to explore whether characteristic behavioral changes parallel alterations in a relevant target. Using PET, a wide range of molecular targets and entities such as brain receptors can be visualized in the same animal. Hence, information about, for example, different neuroreceptors belonging to separate neurotransmitter systems can be integrated. This provides us with unique information on underlying disease mechanisms. In this section, we provide an illustration of the application of  $\mu$ PET in obsessive compulsive disorder (OCD) animal models and how this imaging modality can provide new insights in the complex pathophysiology of OCD.

### 2.3.1 Obsessive Compulsive Disorder

OCD is a chronic disabling psychiatric disease with a lifetime prevalence of 1–3% (Ruscio et al. 2010). The two hallmark symptoms include unwanted obsessions and compulsions (Abramowitz et al. 2009). Obsessions are recurrent and persistent thoughts (e.g., of contamination), urges (e.g., to stab someone), or images (e.g., of violent or horrific scenes) causing anxiety or distress. Also, individuals attempt to suppress or ignore these thoughts, urges, or images or to neutralize these with another thought or action (compulsion). These compulsions include repetitive behaviors (e.g., washing, checking) and mental acts (e.g., counting, repeating words silently), which are driven by an uncontrollable need to be performed in response to the obsession. Clearly, this condition causes serious impairment in daily functioning, which drastically reduces the patient's quality of life. OCD often remains underdiagnosed, resulting in a treatment gap of 57.3% (Kohn et al. 2004). The first-line treatment consists of selective serotonin reuptake inhibitors (SSRIs) and cognitive behavioral therapy (CBT) and was shown to reduce symptoms with circa 50% in half of the patients (Denys et al. 2010). A substantial part of patients remains

treatment refractory (Denys 2006). This emphasizes the need for an improved treatment approach to tackle this disease.

Despite the extensive research focusing on OCD in recent years, the exact mechanism of action is not known. One of the more consistent findings has been the identification of alterations in cortico-striato-thalamocortical (CSTC) circuitry (Pauls et al. 2014), which was mainly established via a range of imaging studies (Bandelow et al. 2016), although additional brain regions are likely to be involved in the pathogenesis as well (Milad and Rauch 2012; Lapidus et al. 2014). Furthermore, it is hypothesized that an altered local neurotransmission lies at the basis of a behavioral disinhibition (Fineberg et al. 2011), which is not fully characterized yet. More specifically, as illustrated in Fig. 2.15, when dopamine (DA) from nigral projections is released in the striatum, a direct pathway is activated. This results in a disinhibition of the thalamus with activation of excitatory projections returning to the cortex. Alternatively, when striatal DA is absent, an indirect



**Fig. 2.15** Simplified scheme of the basal ganglia circuitry in the presence of dopamine. The striatum receives (1) glutamatergic projections from the entire neocortex, which is involved in higher functioning, and (2) dopaminergic projections originating from the substantia nigra (SNi). Dependent on the dopamine (DA) levels in the striatum, a direct or indirect pathway is chosen: (i) When dopamine (DA) is present, GABAergic projections (3) from the striatum will inhibit the globus pallidus interna (GPi) through the direct pathway, leading to less inhibition of the thalamus (4). The thalamus then projects back to the cortex completing the loop (5). Furthermore, the inhibitory projections (6) from the indirect pathway to the globus pallidus externa (GPe) are inhibited themselves due to binding of DA on the dopamine D2 receptor (D2R). This will lead to inhibition (7) of the subthalamic nucleus (STN), followed by less activation of the GPi (8), again resulting in less inhibition of the thalamus (4). (ii) However, the opposite occurs when less DA is present. The inhibitory projections from the striatum to the GPe will become active, leading to less inhibition of the subthalamic nucleus (STN), excitation of the GPi, and inhibition of the thalamus

pathway is activated, thereby inhibiting the thalamus resulting in less signalling to the cortex (Milad and Rauch 2012). OCD is hypothesized to have an excessive relative activation of this aforementioned direct pathway. Other neurotransmitters, such as serotonin and glutamate, likely play a role in the manifestation of this complex pathology as well (Kariuki-Nyuthe et al. 2014; Pauls et al. 2014). Also, new evidence points out synaptic pathology “synaptopathies” as a potential cornerstone in the development of OCD (Bienvenu et al. 2009; Züchner et al. 2009; Ting et al. 2012; Stewart et al. 2013; Mattheisen et al. 2015).

Taken together, OCD is a complex disease involving alterations in multiple brain regions and neurotransmitter systems. Therefore, PET is a promising tool to use in the context of OCD research. PET enables whole brain visualization of multiple targets, e.g., dopamine and glutamate receptors can be visualized in the same animal (model). Also, the use of PET enables regional quantification of radioligand binding. Hence, it can be demonstrated whether or not a certain brain region is involved in the origin and/or progression of OCD or OCD-like behavior(s) in an animal model.

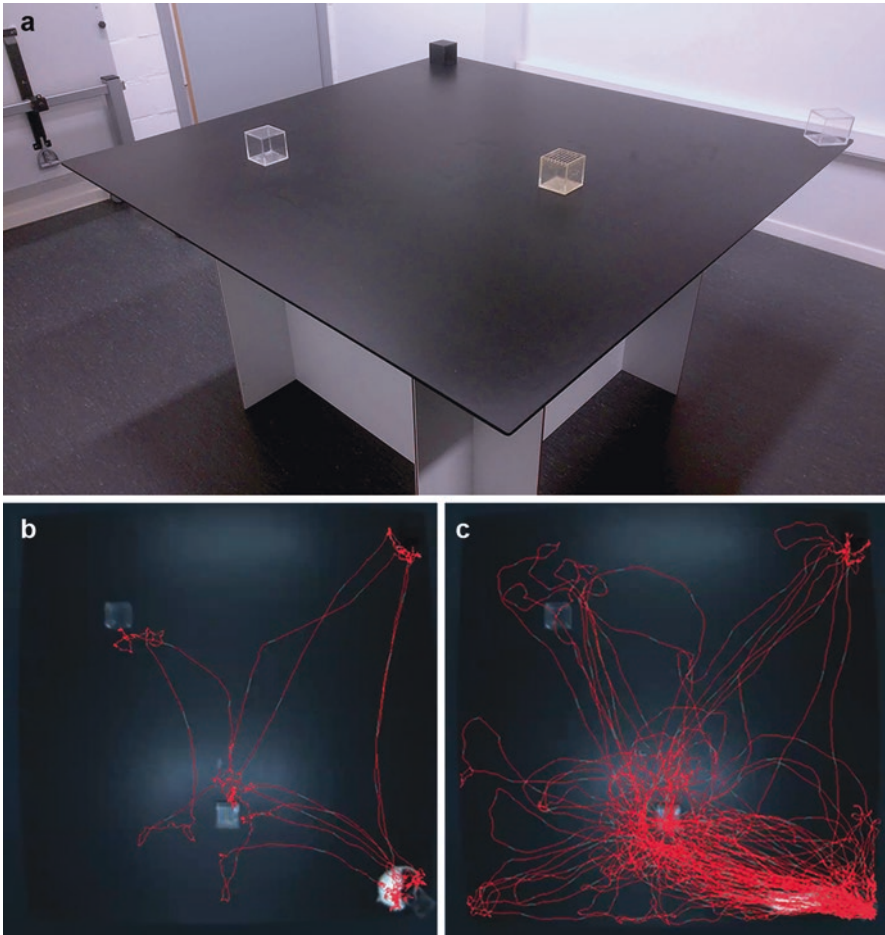
### 2.3.2 Animal Models for Obsessive Compulsive Disorder

The heterogeneous OCD patient population and the presence of comorbidities hamper clinical research. Also, the majority of patients are already receiving treatment, thereby potentially modifying existing disease processes. This further complicates the investigation of underlying pathological mechanisms. Therefore, animal models serve as a valuable tool to dissect mechanisms underlying pathological behavior such as excessive self-grooming and compulsive-like checking behavior. They allow to investigate circuit dysfunction on an experimental level, which is not possible in clinical OCD research. Besides the classic methods such as electrophysiology, the use of *in vivo* methods like optogenetics and molecular imaging is indispensable to obtain a more in-depth comprehension of underlying neuropathology at a molecular, cellular, and circuit level.

Here, we focus on two common OCD disease models to illustrate the value and applicability of  $\mu$ PET imaging within this field.

#### 2.3.2.1 A Pharmacological Rat Model

The pharmacological-induced quinpirole (QP) model belongs to the most used models for OCD and was already extensively validated (Stuchlik et al. 2016) and proven to be effective in a wide variety of studies. In this model, rats receive a biweekly dose of QP for a total duration of 10 weeks. QP is a dopamine D2 receptor (D2R) agonist, causing a sensitization effect on the receptor (Janikova et al. 2019). Activation of D2R, being an inhibitory receptor, results in an inhibition of the inhibitory indirect pathway, thus activating the downstream striatal circuit (Fig. 2.15). This results in ritual-like repetitive checking behavior combined with hyperlocomotion (Szechtman et al. 1998). Fifteen minutes after the QP injection, the animal is placed on a large open field table with four different cubes for a duration of 30 minutes to evaluate such behavior (Fig. 2.16a). This results in a characteristic pattern of locomotion that differs from the pattern of saline-treated control animals, as illustrated in Fig. 2.16b,

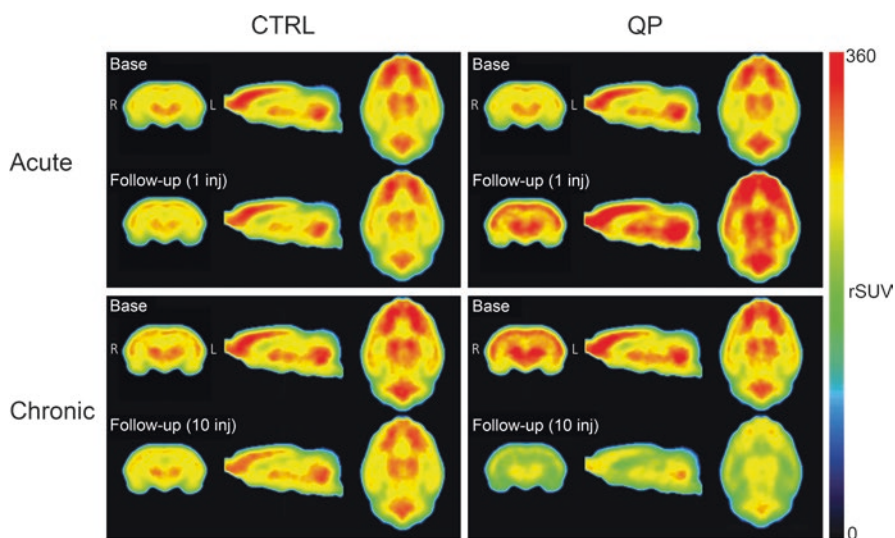


**Fig. 2.16** Behavioral testing. The open field table (OFT) that was used to setup the model is shown here (a). In total, 4 objects are placed on the surface of the table at the specified locations. Panel (b) and (c) show a 30-minute exemplary trajectory of (b) a control animal and (c) a quinpirole animal with a visible contrast between the two trajectories with regard to compulsive-like checking and hyperlocomotion

c. Despite the extensive amount of behavioral studies done with the QP model (Zadicario et al. 2007; Winter et al. 2008; De Carolis et al. 2011; De Haas et al. 2011; Eagle et al. 2014; Hatalova et al. 2014), no in vivo longitudinal data was collected in this model, and the currently available in vivo imaging data is sparse. This is where  $\mu$ PET enters the picture, because it could provide new insights on regional brain metabolism and the interplay between neurotransmitter systems in this model. For example, a  $\mu$ PET study permits to visualize what happens to the D2R within the interval at which the pathological behavior occurs. In the future, awake PET imaging could be applied in this context (cf. Sect. 2.2).

### Imaging of Cerebral Glucose Consumption

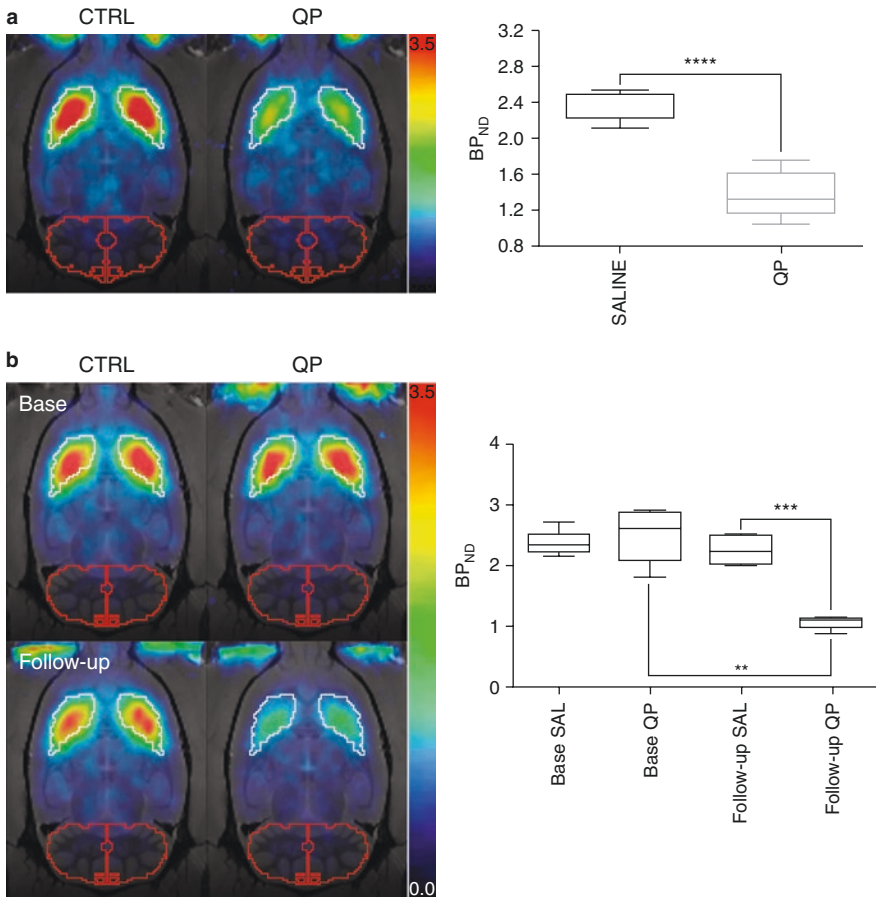
We reported on a first imaging study in the QP model, using the glucose analogue [ $^{18}\text{F}$ ]FDG as a marker for cerebral glucose consumption expressed via the rodent standardized uptake value (rSUV) parameter (cf. Sect. 2.1.6 semiquantitative approach). A baseline [ $^{18}\text{F}$ ]FDG  $\mu$ PET scan was acquired of both the acute and chronic groups prior to (i) an acute QP or saline injection or (ii) twice weekly injections of either QP or saline, each time paired with an open field test. Further, follow-up scans to stage the pathology, after the first injection for the acute group and after the tenth injection (in week 5) for the chronic group, were acquired. This study established significant decreases in [ $^{18}\text{F}$ ]FDG uptake in cortical and striatal regions of chronic QP-treated rats (Fig. 2.17) (Servaes et al. 2016), consistent with the hypothesis stating involvement of the CSTC circuit in the pathophysiology of this OCD model and the process of inefficient thalamic gating (Del Casale et al. 2011). Furthermore, they showed alterations in other key structures such as the amygdala, the hippocampus, and the entorhinal cortex, which are directly involved in memory and fear processing, pointing toward a more extensive disease mechanism. These findings further hint toward a dysregulation of a neural feedback circuit that includes not only the cortical and striatal structures but also the (para)limbic regions in this model.



**Fig. 2.17** Rodent standardized uptake value (SUV<sub>r</sub>) averaged [ $^{18}\text{F}$ ]FDG images at baseline and follow-up for both the acute and chronic quinpirole (QP) group in both conditions (saline control and QP). Acute exposure increased while chronic exposure to quinpirole decreased the total glucose consumption in the brain. Strong reductions were seen within the cortico-striato-thalamocortical (CSTC) circuit after chronic QP exposure. Additional alterations in the (para)limbic regions suggest a more extensive disease mechanism. (CTRL control, inj injection, QP quinpirole, SUV<sub>r</sub> rodent standardized uptake value; acute group,  $n = 8/\text{group}$ ; chronic group,  $n = 7/\text{group}$ )

### Dopamine D2 Receptor Imaging

A second  $\mu$ PET study used [ $^{11}\text{C}$ ]raclopride to visualize the striatal dopamine D2 receptor availability in QP rats with compulsive-like checking behavior (Servaes et al. 2017). This study clearly showed that animals receiving acute and chronic QP had a reduction in D2R availability in the striatum (expressed as the  $BP_{\text{ND}}$  cf. Sect. 2.1.4.2 SRTM), which was more pronounced after chronic injections (Fig. 2.18). Since QP is a D2R agonist, it most likely competes with [ $^{11}\text{C}$ ]raclopride for the same binding site. Hence, a reduced uptake was to be expected. However, rats receiving chronic QP had significantly lower striatal  $BP_{\text{ND}}$  levels compared to rats



**Fig. 2.18** Panel (a) displays the [ $^{11}\text{C}$ ]raclopride  $BP_{\text{ND}}$  maps (left) and the corresponding striatal  $BP_{\text{ND}}$  values (right) after acute administration of quinpirole and saline. Panel (b) shows the [ $^{11}\text{C}$ ]raclopride  $BP_{\text{ND}}$  maps (left) and the corresponding striatal  $BP_{\text{ND}}$  values (right) for the chronic groups at baseline and after chronic administration of quinpirole and saline. ( $BP_{\text{ND}}$  non-displaceable binding potential, CTRL control, QP quinpirole;  $n = 8/\text{group}$ ;  $p$ -values  $t$  tests (unpaired and paired): \*\* $<0.01$ , \*\*\* $<0.001$ , \*\*\*\* $<0.0001$ )

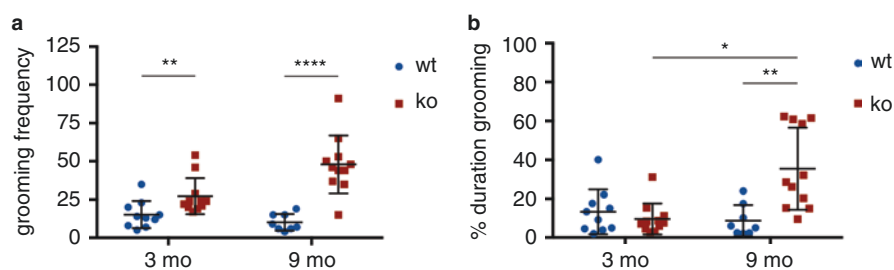
receiving only one QP injection (Fig. 2.18). Therefore, we hypothesized that repeated administration of QP causes a decreased number of available binding sites due to receptor internalization (Goggi et al. 2007), or tolerance development (Lomanowska et al. 2004), while also influencing extracellular DA levels (De Haas et al. 2011).

### 2.3.2.2 A Genetic Mouse Model

As mentioned before, repeated  $\mu$ PET acquisitions can be combined with the setup of a pharmacological model to provide information on alterations linked to the provoked behavior. Furthermore, similar study designs allow us to follow pathological changes in genetic models during the entire lifespan of the animal. For example,  $\mu$ PET acquisitions can be performed in the same animal prior and during the expression and/or progression of certain pathological behavior. This approach enables imaging of secondary effects to a primum movens. In the next example, the primum movens is the loss of a synaptic protein.

Mice with a genetic deletion of the SAP-90/PSD95-associated protein 3 (Sapap3) exhibit an OCD-like phenotype with pathological self-grooming and increased anxiety (Welch et al. 2007). Sapap3 is a scaffolding protein localized in the postsynaptic density (PSD) of excitatory synapses with a high expression in the striatum (Welch et al. 2004). The PSD is a dense multiprotein complex consisting of neurotransmitter receptors, scaffolding proteins, adhesion molecules, signalling enzymes, and cytoskeletal components (Kim and Sheng 2004; Bourgeron 2015). Spatial and temporal organization of these components plays a central role in neurotransmission and synaptic plasticity (Zhu et al. 2016). This widely used model was previously characterized with an abnormal striatal output based on an imbalance between direct and indirect pathway medium spiny neurons located within the CSTC “OCD” circuit (Fig. 2.15) (Ade et al. 2016).

In this model, the grooming phenotype was shown to worsen with age with the development of skin lesions (Fig. 2.19, Glorie et al. 2020). Furthermore, young adult Sapap3 ko mice already have significantly higher grooming frequencies in the



**Fig. 2.19** Progression of compulsive-like grooming in Sapap3 ko mice reflected by behavioral parameters (a) grooming frequency and (b) % duration grooming, compared to wt controls. Both parameters were collected from a 30-minute video recording. (mo, months; wt, wild types; ko, knockouts; animal numbers, wt 3 mo  $n = 10$ , ko 3 mo  $n = 11$ , wt 9 mo  $n = 8$ , ko 9 mo  $n = 11$ ;  $p$ -values Mann-Whitney and Wilcoxon matched pairs signed rank test: \* $<0.05$ , \*\* $<0.01$ , \*\*\*\* $<0.0001$ )

absence of visible skin lesions. Longitudinal in vivo  $\mu$ PET imaging provides the opportunity to further investigate changes in relevant molecular targets such as glutamate receptors and other synaptic targets related to Sapap3, parallel to the aggravation of this phenotype.

### Metabotropic Glutamate Receptor 5 PET Imaging

Sapap3 belongs to a group of scaffolding proteins that links metabotropic glutamate receptors (mGluRs i.a. mGluR5) to ionotropic glutamate receptors (iGluRs i.a. NMDAR and AMPAR) anatomically and functionally (Zhu et al. 2016). Sapap3 ko mice are characterized by glutamatergic abnormalities: (i) an altered N-methyl-D-aspartate receptor (NMDAR) composition (Welch et al. 2007), (ii) a reduction in corticostriatal synaptic strength driven by metabotropic glutamate receptor 5 (mGluR5)-dependent  $\alpha$ -amino-3-hydroxy-5-methyl-4-isoxazolepropionic acid receptor (AMPA) endocytosis (Wan et al. 2011), and (iii) an elevated mGluR5 signalling driving OCD-relevant behavior (Ade et al. 2016). These findings substantiate the application of in vivo [ $^{11}\text{C}$ ]ABP688  $\mu$ PET imaging to longitudinally visualize mGluR5 availability (Ametamey et al. 2006) in the brain of Sapap3 ko mice, expressed as the non-displaceable binding potential  $\text{BP}_{\text{ND}}$  (cf. Sect. 2.1.4.2 SRTM) (Lammertsma and Hume 1996).

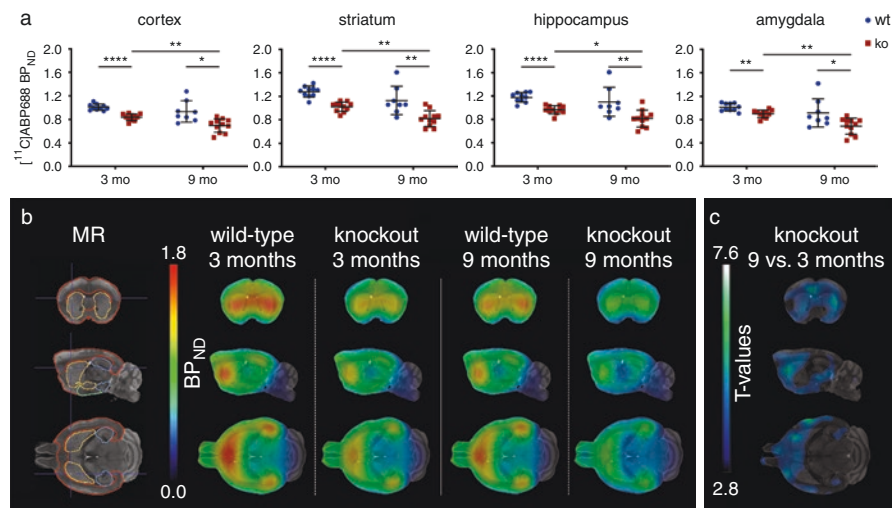
The results of this study emphasize the potential of  $\mu$ PET to follow disease progression in vivo. We established a significant progressive decline in the availability of the mGluR5 allosteric site (expressed via [ $^{11}\text{C}$ ]ABP688  $\text{BP}_{\text{ND}}$ ) in both striatal and extra-striatal brain regions of Sapap3 ko mice (Fig. 2.20). This parallels an increase in compulsive-like grooming behavior (Fig. 2.19) (Glorie et al. 2020). In line with previous findings (Ade et al. 2016), our data, showing dynamic changes in mGluR5 availability during the lifespan of Sapap3 ko mice, support the hypothesis of excessive mGluR5 activation as a driver of the OCD-like phenotype in this model. Previously, the mGluR5 was shown to be constitutively activated, not via the conventional binding of its orthosteric ligand glutamate. Lower mGluR5 availability for [ $^{11}\text{C}$ ]ABP688 in ko mice could be related to this aberrant activation pattern. Altogether, these findings suggest a promising role for [ $^{11}\text{C}$ ]ABP688 as a biomarker for constitutive mGluR5 activation and to study disease progression in this model. Also, this study points out the potential of [ $^{11}\text{C}$ ]ABP688  $\mu$ PET to evaluate drugs targeting mGluR5 together with the drug's effect on the behavioral readout.

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## 2.4 Summary/Conclusion

This chapter highlights the added value of PET imaging in the preclinical research setting, with its high versatility and its ability to collect in vivo molecular information during the lifespan of a laboratory animal. Indeed, an illustration of the use of  $\mu$ PET with multiple radiotracers in OCD animal models confirms these characteristics. Beside the pharmacokinetic modelling methods to extract biologically relevant





**Fig. 2.20** Sapap3 ko mice show cross-sectional differences and a longitudinal decline in mGluR5 availability for  $^{11}\text{C}$ ABP688. (a) Averaged  $\text{BP}_{\text{ND}}$  values showing a progressive decline in  $\text{BP}_{\text{ND}}$  in the brain of Sapap3 ko mice in the cortex, striatum, hippocampus, and amygdala. Cross-sectionally, the averaged  $\text{BP}_{\text{ND}}$  value of Sapap3 ko mice is significantly lower in all four analyzed regions compared to their wt counterparts both at the age of 3 and 9 months. (b) Corresponding averaged  $^{11}\text{C}$ ABP688  $\text{BP}_{\text{ND}}$  images superimposed on a mouse MR template at both time points (3 and 9 months) and for both groups separately (wt versus ko). (c) Hypo T-map (voxel-based statistical parametric mapping (SPM) analysis) showing clusters of voxels (voxel threshold = 100) with a significantly ( $p < 0.01$ ) decreased  $\text{BP}_{\text{ND}}$  at 9 mo, when compared to the 3 mo time point in Sapap3 ko mice. (MR, magnetic resonance; MR delineation, red cortex, yellow striatum, blue hippocampus, green amygdala; wt, wild type; ko, knockout; animal numbers, wt 3 mo  $n = 10$ , ko 3 mo  $n = 11$ , wt 9 mo  $n = 8$ , ko 9 mo  $n = 11$ ; statistical analysis, cross-sectional data with a two-way ANOVA with post hoc Bonferroni correction for multiple comparisons and longitudinal data with a mixed effects model;  $p$ -values, \* $<0.05$ , \*\* $<0.01$ , \*\*\* $<0.001$ , \*\*\*\* $<0.0001$ )

information from PET images, we describe the new and promising method of point source tracking with the possibility to perform high-throughput awake  $\mu$ PET in a routine small animal setting. Further work should focus on combining awake  $\mu$ PET imaging paradigms with animal models of brain disease, thereby collecting unique information on underlying pathological mechanisms and eliminating anesthesia-related disturbances. In this context, the use of multiple tracers within one animal enables the study of the interplay between different targets of interest during the conduct of behavior. With this progress, an important problem associated with translation to a clinical setting will be tackled.

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Charlotte Thyssen and Stefaan Vandenberghe

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## Abstract

Since the introduction of positron emission tomography (PET) in the mid-1970s, multiple attempts were made to build a scanner able to cover a large portion of the body, but none led to a clinical research system. However, many research centers remained interested in the idea, and with the help of simulation studies, the potential of total-body PET was shown. The building of a system was started only 2 years ago, and the first clinical studies were performed at UC Davis in 2019. The reason for the success of total-body PET is the increased angle coverage of the subject. The larger this angle, the more coincidences are potentially detected by the system. Studies showed that a gain in sensitivity up to a factor 40

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can be obtained with this new scanner design. This allows a reduction in administered dose as well as acquisition time, paving the way for dynamic PET imaging and long-term follow-up of patients. Although the main application of PET is in oncology, other disciplines like neurology, pediatrics, and pharmaceutical (drug) development can take advantage of the increased sensitivity.

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### 3.1 Introduction

It is safe to say that since the introduction of positron emission tomography (PET) in the mid-1970s (Phelps et al. 1975), nothing fundamentally changed to the original idea. We inject the patient with a short living radiotracer, and we attempt to trace its biodistribution by detecting the annihilation photons that are produced inside the body. However, the sensitive detectors that surround the patient have changed in several ways. Already in the early 1990s, Terry Jones came up with the idea of a PET system that covers the whole body of a human subject (Vandenberghe et al. 2020). The following years, only prototypes of such systems were produced, and the idea never led to a scanner for clinical research. In 2015, a consortium called EXPLORER was founded, committed to build a long field-of-view (FOV) PET scanner, and this resulted in the first publication of a human study with a total-body PET system in 2019 (Badawi et al. 2019). In order to understand the added value of total-body imaging, the present chapter starts with a discussion of the importance of coincidence detection in PET imaging and how this is influenced by scanner length.

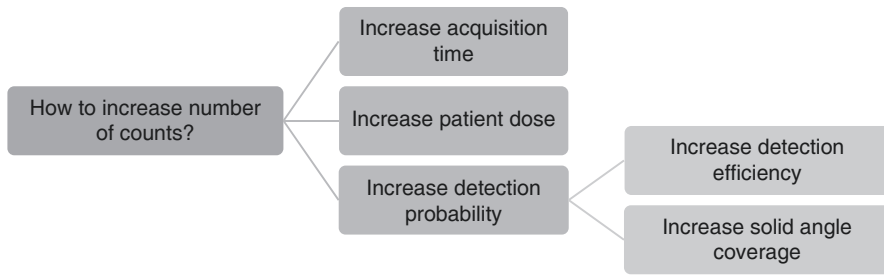
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### 3.2 The Importance of Detection

In medical imaging, very good spatial resolution is important in order to detect small lesions and abnormalities. Spatial resolution is the ability to distinguish two objects (like lesions) from their surrounding tissue. However, the spatial resolution cannot be improved infinitely without simultaneously increasing the signal-to-noise ratio (SNR) (Nesterets and Gureyev 2015; Gureyev et al. 2014). In order to explain this, one could draw the analogy with photography where enough light exposure on the sensor is necessary in order to see the image appear against the black background. The longer we expose the camera to the object we want to capture, the better it appears on the resulting image. Reverting back to medical imaging, multiple sources of noise exist that each individually adds to this background signal. In the specific case of PET, these sources include, among others, the scintillation process, the movement of charge carriers in the p-n junction of the silicon photomultiplier (SiPM), the electronic noise from the readout circuit, and the random process of radioactive decay of the radioisotopes.

Detection of this radioactive decay is a perfect example of a Poisson distribution. The Poisson distribution is a statistical distribution that exists for situations in which an experiment is repeated a large number of times with a low probability of success, as is photon detection in PET imaging. The Poisson distribution predicts a noise





**Fig. 3.1** Schematic overview of three main strategies for increasing the number of counts

level that equals the square root of the number of detected photons. This implies that the SNR of measured data is equal to (Tsui et al. 1981):

$$\text{SNR} = \frac{n}{\sqrt{n}} = \sqrt{n} \quad (3.1)$$

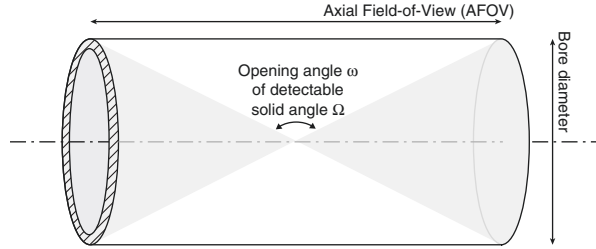
In words, the square of the SNR scales linearly with the detected number of photons. In its turn, better SNR implies a better ability to distinguish the signal from the background. The schematic overview in Fig. 3.1 shows how the number of detected photons can be increased in PET. We distinguish three main strategies. The number of detections could be increased by increasing (i) the acquisition time, (ii) the amount of radioactivity administered to the patient, and (iii) the sensitivity—the ability to detect photons—of the set of detectors surrounding the patient. However, increasing the acquisition time (of each static image, strategy 1) will decrease the image quality due to movement artifacts. This is not favorable, and the acquisition time should therefore be as short as possible. Also, increasing the radioactive dose (strategy 2) increases the risk of radiation-induced problems. Therefore, the only sustainable solution is to improve the probability of detection of photons by the PET ring (strategy 3). This can be done either on the detector level, by increasing detection probability when a photon hits the sensitive ring, or on the system level, by increasing the probability of a gamma photon hitting the ring.

### 3.3 The Effect of the Scanner Configuration on the Sensitivity

The sensitivity of a PET scanner is a measure for the counting efficiency of the system. It is defined as the count rate of incoming coincidences, in counts per second (cps), divided by the activity, often expressed in Becquerel (Bq). Since both count rate and activity are essentially expressed in the unit 1/s, the sensitivity can be thought of as a unitless number expressing the fraction of radioactive decays, captured by the system. As stated by Habte et al. (2007), the sensitivity ( $S$ ) can be estimated as:

$$S = E_g \times E_i \quad (3.2)$$

**Fig. 3.2** Definition of the opening angle of the detectable solid angle of a PET scanner



The sensitivity can thus be written as the product of the geometric efficiency,  $E_g$ , depending on the geometry of the system, and the intrinsic detection efficiency,  $E_i$ , depending among other things on the atomic number of the detecting material. The geometric detection efficiency,  $E_g$ , equals the (fractional) solid angle  $\Omega$  that is covered by the PET system. This solid angle expresses what portion of an imaginary sphere in the center of the scanner is covered by detectors. The magnitude of  $\Omega$  will vary with the opening angle  $\omega$  of the triangle, as defined in Fig. 3.2, and is also referred to as the detectable solid angle or solid angle coverage. The magnitude of this angle is defined as:

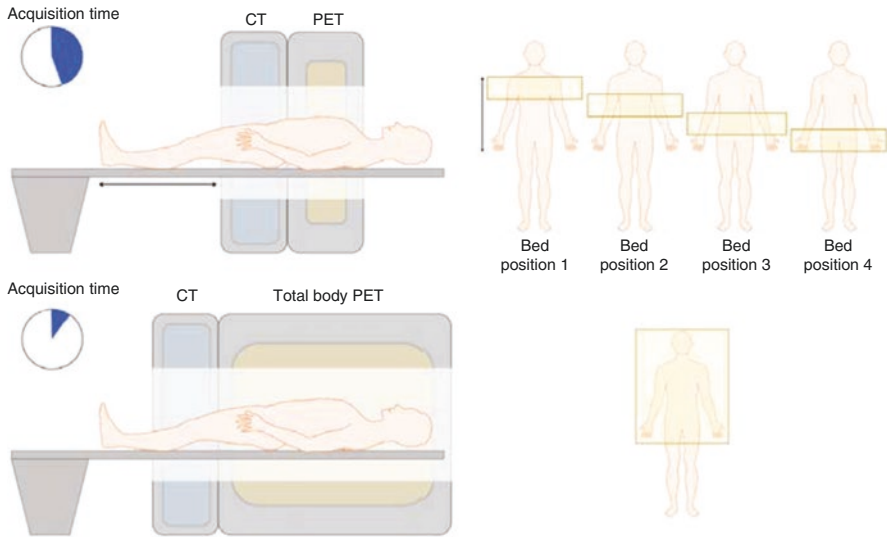
$$\tan(\omega / 2) = \frac{\text{AFOV}}{\text{Diameter}} \quad (3.3)$$

Since all emission directions falling inside  $\omega$  will possibly lead to a coincidence on the scanner, the number of coincidences will increase according to the magnitude of  $\omega$ . When the axial length (also called axial field of view (AFOV)) is increased, Eq. (3.3) proves that the solid angle coverage will increase accordingly. The detected count rate per unit of activity will thus be higher, which implies a higher geometric sensitivity of the system.

Since only  $\sim 1.6\%$  of all decays inside the body are captured in state-of-the-art PET systems (Van Sluis et al. 2019), there is clearly a lot of space for further improvement in terms of sensitivity. As sensitivity is dependent on geometrical and intrinsic sensitivity, it can be improved in two ways (Fig. 3.1).

1. Increasing detection efficiency
2. Increasing solid angle coverage

Increased solid angle coverage can be achieved by increasing the AFOV of the scanner or reducing the diameter. Figure 3.3 sketches the concept of total-body PET and highlights the differences between conventional and total-body PET systems. On the one hand, the geometric efficiency is increased, and on the other, a larger portion of the body is imaged at once which makes translation (i.e., stepwise horizontal movement of the human subject) in the system redundant (Vandenberghe et al. 2020).



**Fig. 3.3** Comparison between normal PET as we find it today in clinic and total-body PET. In the first situation, the patient has to move through the scanner while this is not the case in total-body PET. Here, the patient is scanned at once

The idea of drastically increasing the scanner length as a solution to the limited sensitivity has recently (re)gained a lot of attention from the PET community. The idea was already considered in the early 1990s but never became reality. This was mainly due to limitations in the performance of the scintillators and detectors to do fully 3D acquisitions.

Also, today there are obstacles on the road toward a system able to cover the complete human body. The major obstacle here is the cost of the system, since cost increases linearly with length of the system. Also, component choices affect the price significantly. Among others, the choice of scintillator material is a decisive factor for total-body PET designs since it forms the largest component cost of the system; the choice of photosensor and electronics has a large impact on the total price as well.

Apart from the financial obstacles that will arise, there are some other, more practical, problems that will have to be overcome. One of these is the claustrophobic feeling that might be evoked from the long and narrow tunnel in which the patient will have to lie during image acquisition. It might be necessary to incorporate screens inside the scanner bore or to go for an adaptive design, for example, in which there exist gaps between the detectors. Via such gaps, the patient is able to see outside the scanner which will counteract the feeling of claustrophobia.

### 3.4 State of the Art in Total-Body PET

#### 3.4.1 The Road Toward Total-Body PET

Although total-body PET scanners are a new member of the nuclear imaging family, the idea to extend the axial FOV had already been suggested in the early 1900s by Terry Jones (Vandenberghe et al. 2020). Ever since, it has fascinated scientists around the world, and it led to several research projects (Watanabe et al. 2003; Wong et al. 2007; Poon et al. 2012; Freedenberg et al. 2014; Mikhaylova et al. 2017; Schmall et al. 2016; Crosetto 2003; Conti et al. 2006). In 2003, a 68.5-cm-long prototype was built by Hamamatsu with 96 rings and pixelated BGO crystals read out by flat panel PS-PMTs (Watanabe et al. 2003). However, the detector's energy resolution—36% on average—was not sufficient to obtain qualitative images.

#### 3.4.2 The EXPLORER Project

In 2015, a research consortium, called EXPLORER, was founded by the University of California (UC) at Davis, University of Pennsylvania (Upenn), and Lawrence Berkeley National Laboratory based on the idea of scanning the entire body of a patient at once. This consortium led to the construction of a total-body PET system by United Imaging under the control of UC Davis. This system is called uEXPLORER and has an AFOV of 194 cm. First images were taken mid-2018 (Reardon 2019) followed by a study on healthy volunteers (Badawi et al. 2019). Since the acquisition time per image can be reduced to a few seconds or even less, it becomes possible to image the distribution of a radiotracer in the patient as a function of time. A second system, the PennPET Explorer, is under construction at the University of Pennsylvania. This system is scalable since it comprises multiple ring segments. Recently, a 64-cm-long prototype of the PennPET Explorer consisting of three ring segments has been completed and tested on human subjects. This system also allows dynamic imaging. Table 3.1 gives an overview of the performance parameters of both systems in the EXPLORER project.

**Table 3.1** Overview of the performance parameters of both scanners in the EXPLORER project

System	uEXPLORER (Badawi et al. 2019)	PennPET Explorer (Karp et al. 2019)
Length AFOV	194 cm	64 cm <sup>a</sup>
Bore diameter	78.6 cm	76.4 cm
Spatial resolution center (FBP)	~ 2.9 mm	~ 4.0 mm
Energy resolution	~ 11.7%	~ 12%
Time resolution	~ 430 ps	~ 256 ps
Sensitivity NEMA	~ 191.5 kcps/MBq	~ 55 kcps/MBq

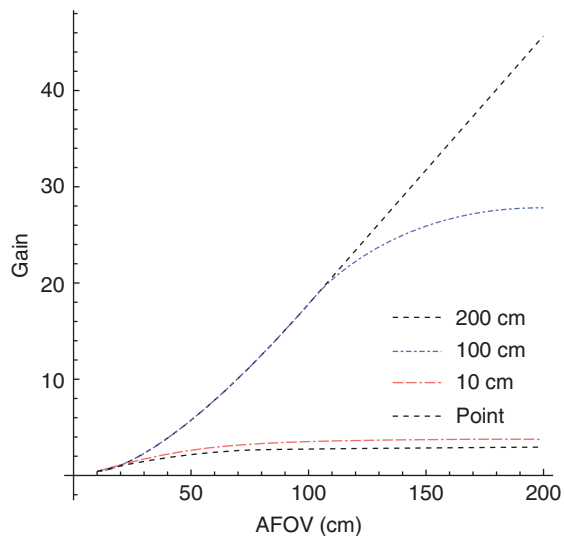
<sup>a</sup>This is the active AFOV of the current prototype comprising three ring segments. The scanner is scalable in length and in the future more rings will be added (Karp et al. 2019)

### 3.4.3 Gain Versus System Length

As mentioned before, moving to a scanner with a long bore affects the sensitivity of the scanner. The actual gain in image quality depends not only on the geometry of the system but also on the object under study. Figure 3.4 gives a graphical overview of how the predicted sensitivity of the scanner improves with increasing length for different line sources. In case of single organ examinations, the sensitivity gain is limited to a factor between 3 and 4 depending on the final size of the system, and the primary gain in sensitivity is already in the first 50 cm of the system. This is also visible in Fig. 3.4, where the sensitivity is going up in the beginning of the curve in case of a point source and a 10-cm-long source but stabilizes after the 50-cm point. Both sources can act as surrogate for a single organ examination. However, the real added value of a total-body system is not in single organ examinations but in the ability to image a large portion of a tall object at once. This comes forward in both blue and black curves in Fig. 3.4. Gain for a 2-m source is predicted to range between 15, for a 1-m-long scanner, and 40, for a 2-m-long scanner (Vandenberghe et al. 2020).

A side note to these results can be drawn when one looks at a nationwide study in the Netherlands, the country known to have the tallest average inhabitant (Risk and Collaboration F 2016). This study shows that 99% of male adults have a sitting height (torso and head) below 104 cm (Fredriks et al. 2005). Since most relevant information is confined in head and torso, a 100-cm source is an accurate estimate for a patient under study. From Fig. 3.4, we see that for a 1-m source, only limited extra sensitivity gain is expected between 100-cm and 200-cm AFOV. This supports the idea of limiting the AFOV of a total-body PET system to around 1 m in order to reduce cost and make this new technology more accessible for research centers and clinics.

**Fig. 3.4** Evolution of the sensitivity gain as a function of system length for different line sources. This figure was originally published in EJNMMI Physics. Vandenberghe S, Moskal P, Karp J. State of the art in Total Body PET. EJNMMI Phys (Vandenberghe et al. 2020)



### 3.5 Total-Body PET Applications

It is a well-known fact that early detection of diseases is very important in order to improve the quality of life of the patient and to prolong his/her lifespan. In many diseases, like cancer and dementia, changes happen to the physiology (e.g., metabolism and signaling pathways) long before anatomical changes (e.g., tumor formation and neuron degradation) occur. Therefore, in order to diagnose a patient in the early disease stage, focus on physiology over anatomy is important. For this purpose, PET is the modality of choice. In anno 2020, it is the most powerful molecular imaging modality with main applications in oncology. Early detection is not only important for the patient, but it also makes expensive and specific treatments redundant. The follow-up of patient treatments is one step further in the direction of personalized medicine, where therapy can be adapted to the patient and revised through the treatment cycle.

The gain in sensitivity in total-body PET systems has a big impact not only on the image quality but also on the possibilities and applications of molecular imaging. Basically, the advantages can be split up into four categories: signal-to-noise ratio (SNR), dynamic imaging, injected dose, and acquisition time.

- (a) **Better SNR:** When the dose and the acquisition time are kept constant, the increased sensitivity can be used to visualize tracers in tissues or lesions with only small uptake or with tracers that could not be visualized before because they only send out a very small number of positrons. One particular example of this is visualization of therapeutic isotopes like  $^{90}\text{Y}$  that do not have positrons as primary emission source. Moreover, the increased sensitivity allows to visualize the radiotracer distribution at later time points—e.g., the next day—when they have already decayed for several half-lives.
- (b) **Dynamic image:** The increased sensitivity of a large system can also be applied to reduce the length of the acquisition frames (to an order of magnitude of 1 s). Thus, the tracer can be followed from the moment of administration while it spreads in the patient's body.
- (c) **Lower dose:** Since a larger portion of the emitted photons are captured in total-body PET, another possibility is the reduction of the administered dose to the patient. Especially for children who are much more likely to develop radiation-induced tumors, this is very important. The uEXPLORER claims to have a sensitivity gain of  $\sim 20$  for children (Berg et al. 2018). An extra advantage is the ability to scan a larger number of patients with the same amount of tracer. This makes rare tracers accessible for more patients, and it reduces the need for an expensive on-site cyclotron.
- (d) **Faster:** Instead of using the decreased acquisition time for dynamic imaging, one could also use this to reduce the overall acquisition time per patient without decreasing image quality. This allows to image patients faster and to increase scanner throughput. Especially in regions with limited availability of PET systems – e.g., in regions with a high population density – this is very desirable.

### 3.5.1 Applications in Oncology

Since the invention of positron emission tomography in 1975, it lasted until the early 2000s before PET established itself in clinical routine. Ever since, PET has been used mostly for diagnosis, staging, and follow-up of cancer patients. In order to diagnose and stage a patient, it is typically necessary to scan a large portion of the patient's body in order not to miss metastases that have spread to other body parts far from the primary tumor. However, conventional systems only allow to scan a small portion of the body at once. By the time an image of a certain body portion is acquired, the uptake of a rapidly changing radioactive tracer might already have altered, leading to the possible loss of detection of metastatic lesions. By scanning the whole body at once and in a dynamic way, no lesions will be missed in this way.

Already at the turn of the century, it was shown that malignant carcinomas have more tracer uptake at later time points (Kubota et al. 2001; Cherry et al. 2017) and normal tissue uptake decreases at later time points (Cheng et al. 2013). However, at late time points, the radioactivity of the tracer will have decayed already significantly, and visibility is poor with conventional scanners. The higher sensitivity of a total-body PET system makes it possible to acquire a delayed image of the patient with sufficient image quality for lesion detection.

### 3.5.2 Applications in Neurology

Many diseases that were thought to be confined to one single organ in the past now appear to be much more complex and involve interactions between organs. Total-body PET imaging, i.e., simultaneous imaging of the brain and torso, allows the medical doctor to look at these interactions during a single examination. This interaction of peripheral organ pathologies and the brain or the interaction between the body and the brain is an interesting topic of novel neurological studies that could lead to new information and a more thorough understanding on the origin of CNS diseases. One example of this is the gut-brain axis or the relationship between the brain and the microbiota in the gastrointestinal tract. It has been shown that changes in the gut microbiota induce activation of certain pathways in the central nervous system and may even lead to conditions like anxiety and depression (Foster and McVey Neufeld 2013; Cryan and O'Mahony 2011). The introduction of total-body PET and the ability to visualize the whole body and the dynamics of the tracer at once can lead to breakthroughs in neurological research.

### 3.5.3 Research on New Radiopharmaceuticals

One of the main driving factors for the realization of total-body PET is its potential use for research. The first research advantage is the ability to visualize the complete body of the patient at once and in a dynamic way. Thus, the effect and the

time-dependent distribution of a drug can be studied in the entire body. A second reason is the ability to scan patients with a radioactive dose lower than the monthly background radiation. This makes it safe for healthy subjects to participate in clinical trials, which will make new radiotracers more rapidly available. One example of a new tracer under investigation is the use of radioactively labeled nanobodies for the detection of breast carcinoma (Keyaerts et al. 2016).

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### 3.6 Conclusion

Since PET physics is based on electronic collimation by means of coincidence detection, its sensitivity is highly restricted by the geometry of the system. Because of the limited length of clinically available system (~20 cm), they are only able to detect less than 2% of all emitted activity from inside the patient. For this reason, the patient has to be injected with a high dose of radioactivity and lie still in the scanner for a long time. The first requirement is associated with an increased risk for radiation-induced cancers, while the latter may introduce artefacts due to small movements of the patient. The only way to drastically increase the fraction of detected photons is by increasing the length of the scanner and making it cover a larger portion of the patient's body. This is exactly what was done in the EXPLORER project. This project resulted in the construction of two total-body PET systems with lengths of 1 and 2 m and up to 15 and 40× higher sensitivity, respectively. The highest gain is in applications where the full length of the scanner is used, meaning where the whole body is under examination. A typical application of PET where this is the case is in oncology where PET is used for diagnosis, staging, and follow-up of patients. Although the main application of PET is in oncology, other disciplines like neurology, pediatrics and pharmacology can take advantage of the increased sensitivity.

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# Cerebral Glucose Metabolism

# 4

Wolf-Dieter Heiss and Olivier Zaro-Weber

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## Abstract

Glucose is the main substrate for energy metabolism of the brain, and the regional cerebral metabolic rate is directly related to regional brain activity. Therefore, the measurement of regional glucose metabolism is of great importance for the assessment of regional normal function and of pathological changes. Quantitation of glucose metabolism by PET is based on the 2-deoxyglucose method developed by Sokoloff and colleagues: F18-labelled deoxyglucose (FDG) is transported into the brain and phosphorylated, but cannot be further metabolised and therefore is accumulated intracellularly. The concentration of the tracer can be

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measured three dimensionally by PET, and together with the arterial tracer concentration, the kinetics of glucose uptake can be assessed, and the regional cerebral metabolic rates for glucose (rCMRGlc) can be calculated.

rCMRGlc is high in cortex and grey matter structures and low in white matter, but there are significant differences among various regions. Metabolic rate is slightly reduced with ageing and changed by sleep, dream, and functional activation. CMRGlc is significantly affected in pathological states, and the regional and global changes are important for assessing the severity of disorders and for differential diagnosis of diseases of the brain. Therefore, FDG-PET has still great importance in brain research and many applications in clinical neurology.

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## 4.1 Energy Requirements of Brain Tissue

The energy demand of the nervous tissue is very high, and therefore sufficient blood supply to the brain must be maintained consistently. A normal adult male's brain containing approximately 130 billion neurons (21.5 billion in the neocortex) (Pakkenberg and Gundersen 1997) comprises only 2% of the total body mass yet consumes at rest approximately 20% of the body's total basal oxygen consumption supplied by 16% of the cardiac blood output. The brain's oxygen consumption is almost entirely for the oxidative metabolism of glucose, which in normal physiological conditions is the almost exclusive substrate for the brain's energy metabolism (Clarke and Sokoloff 1999). It must be kept in mind that the glucose metabolised in neuronal cell bodies is mainly to support cellular vegetative and housekeeping functions, e.g. axonal transport, biosynthesis of nucleic acids, proteins, lipids, as well as other energy-consuming processes not related directly to action potentials. Therefore, the rate of glucose consumption of neuronal cell bodies is essentially unaffected by neuronal functional activation. Increases in glucose consumption (and regional blood flow) evoked by functional activation are confined to synapse-rich regions, i.e. neuropil which contains axonal terminals, dendritic processes, and also astrocytic processes that envelope the synapses. The magnitudes of these increases are linearly related to the frequency of action potentials in the afferent pathways, and increases in the projection zones occur regardless of whether the pathway is excitatory or inhibitory. Only at the next downstream projection zones glucose utilisation (and, as a consequence, blood supply) is depressed in inhibited neurons and increased in excited neurons. Energy metabolism by functional activation is due mostly to the stimulation of the  $\text{Na} + \text{K} + \text{ATPase}$  activity to restore the ionic gradients across the cell membrane and the membrane potentials that were degraded by the spike activity and is rather high compared to the demand of neuronal cell bodies (Sokoloff 1999). Overall, 87% of the total energy consumed is required by signalling, mainly action potential propagation and postsynaptic ion fluxes, and only 13% is expended in maintaining membrane resting potential (Attwell and Laughlin 2001).

## 4.2 Brain Energy Metabolism

Glucose is the obligatory energy substrate for the brain, and it is almost entirely oxidised to  $\text{CO}_2$  and  $\text{H}_2\text{O}$ . Although the brain represents only 2% of the body weight, it receives 15% of the cardiac output, 20% of the total body oxygen consumption, and 25% of the total body glucose utilisation (Villien et al. 2014). With a global blood flow of 57 mL/100 g min, the brain extracts approximately 50% of oxygen and 10% of glucose from the arterial blood. Hence, the glucose utilisation of the brain, as assessed by measuring the arterial-venous difference (Kety and Schmidt 1948), is 31 mmol/100 g min. Oxygen consumption is 160 mmol/100 g min; because  $\text{CO}_2$  production is almost identical, the respiratory quotient (RQ) of the brain is nearly 1, indicating that carbohydrates are the substrates for oxidative metabolism (Sokoloff 1989). Given a theoretical stoichiometry of 6 mmol of oxygen consumed for each mmole of glucose, glucose utilisation by the brain should in theory be 26.6 mmol/100 g min. As indicated earlier, the measured glucose utilisation is 31 mmol/100 g min, indicating that an excess of 4.4 mmol/100 g min of glucose follows other metabolic fates. Glucose can produce metabolic intermediates, such as lactate and pyruvate, which do not enter necessarily in the tricarboxylic acid cycle, but rather can be released and removed by circulation. Glucose can be incorporated into lipids, proteins, and glycogen, and it is also the precursor of certain neurotransmitters such as gamma-aminobutyric acid (GABA), glutamate, and acetylcholine (Sokoloff 1989).

Of molecules that could substitute for glucose as an alternative substrate for brain energy metabolism, mannose is the only one that can sustain normal brain function in the absence of glucose. Lactate and pyruvate can sustain synaptic activity *in vitro*. Because of their limited permeability across the blood–brain barrier, they cannot substitute for plasma glucose to maintain brain function (Pardridge and Oldendorf 1977). However, if formed inside the brain parenchyma, they are useful metabolic substrates for neural cells (review in Magistretti and Allaman 2015).

Whole-organ studies, which allowed the determination of the substrate requirements for the brain, failed to provide the appropriate level of resolution to appreciate two major features of brain energy metabolism: (a) its regional heterogeneity and (b) its tight relationship with the functional activation of specific pathways. The autoradiographic 2-deoxyglucose (2-DG) method developed by Sokoloff and colleagues afforded a sensitive means to measure local cerebral metabolic rates of glucose (LCMRGlc) with a spatial resolution of approximately 50–100  $\mu\text{m}$  (Sokoloff et al. 1977). The method is based on the fact that tracer amounts of radioactive 2-DG are taken up by glucose transporters and phosphorylated by hexokinase with kinetics that are similar to those for glucose; however, unlike glucose-6-phosphate, 2-deoxyglucose-6-phosphate cannot be metabolised further and therefore accumulates intracellularly, thus providing, after appropriate corrections (Sokoloff et al. 1977), an accurate measurement of the amount of glucose utilised. Using this method, LCMRGlc have been determined in virtually all morphologically and

functionally defined brain structures in various physiological and pathological states including sleep, seizures, and dehydration and following a variety of pharmacological treatments. Furthermore, the increase in glucose utilisation following activation of pathways subserving specific modalities, such as visual, auditory, olfactory, or somatosensory stimulations, as well as during motor activity, has been revealed in the pertinent brain structures.

Basal glucose utilisation of the grey matter as determined by the 2-DG technique varies, depending on the brain structure, between 50 and 150  $\mu\text{mol}/100\text{ g wet weight}/\text{min}$  in the rat (Sokoloff et al. 1977). Physiological activation of specific pathways results in a 1.5–3-fold increase in LCMRGlc as determined by the 2-DG technique.

With the advent of PET and the use of positron-emitting isotopes such as  $^{18}\text{F}$ , local glucose utilisation has been studied in humans with 2- $(^{18}\text{F})$ fluoro-2-deoxyglucose (Reivich et al. 1979). Changes in local brain energy metabolism can now be studied in humans with PET by monitoring alterations in glucose utilisation, oxygen consumption, and blood flow during activation of specific areas. Studies in which these three parameters have been analysed during activation of a given modality have yielded an uncoupling between glucose uptake and oxygen consumption during activation, since the increase in blood flow and in glucose utilisation in the activated cortical area was not matched by an equivalent increase in oxygen consumption (Fox et al. 1988; Madsen et al. 1995; Blomqvist et al. 1994). This observation raises the puzzling possibility that, at least during the early stages of activation, the increased energy demand is met by glycolysis rather than by oxidative phosphorylation (Vaishnavi et al. 2010; Raichle and Mintun 2006).

### 4.2.1 Glycolysis and Oxidative Phosphorylation

Glycolysis (Embden–Meyerhof pathway) is the metabolism of glucose to pyruvate and lactate. It results in the net production of only 2 mol of adenosine triphosphate (ATP)/mol of glucose as well as in the regeneration of reducing equivalents (the oxidised form of nicotinamide adenine dinucleotide ( $\text{NAD}^+$ )) through the conversion of pyruvate into lactate. Alternatively, pyruvate can enter the tricarboxylic acid (TCA) cycle (or the Krebs cycle) and produce 30 mol of ATP/mol of glucose via the mitochondrial oxidative phosphorylation cascade. The energetic value of oxidative phosphorylation over glycolysis is thus obvious. The respiratory quotient of the brain is virtually 1; PET studies indicate an uncoupling between glucose uptake and oxygen consumption during activation (Fox et al. 1988; Madsen et al. 1995), and rises in lactate have been monitored. During activation, lactate may normally be taken up by neurons as an energy fuel. It should be remembered that after conversion to pyruvate, lactate can enter the TCA cycle with the potential to generate a total of 36 mol of ATP/mol of glucose. Activation-induced glycolysis may provide ATP to fuel energy-dependent ion transport, in particular the  $\text{Na}^+/\text{K}^+ - \text{ATPase}$ , which represents the main energy-consuming process in neural cells (Siesjo 1978).

### 4.2.2 Determination of the Regional Cerebral Metabolic Rate for Glucose (rCMRGlc)

The study of glucose metabolism with  $^{18}\text{F}$ FDG is a direct application of the autoradiographic technique of Sokoloff et al. (1977) with  $[^{14}\text{C}]$ deoxyglucose. The model developed by Sokoloff et al. can be applied directly because the fluorodeoxyglucose labelled at point 2 behaves in the same way as deoxyglucose. It is transported into the cell in the same way as glucose and, with the aid of hexokinase, is phosphorylated to  $[^{18}\text{F}]$ deoxyglucose-6-phosphate. Deoxyglucose-6-phosphate, however, cannot be further converted to fructose-6-phosphate and degraded to  $\text{CO}_2$  and  $\text{H}_2\text{O}$ , but accumulates in the cell. The kinetics of the back reaction (phosphatase) to deoxyglucose is much slower, and the deoxyglucose-6-phosphate can penetrate through the cell membrane only in small amounts. The kinetics of the accumulation of deoxyglucose-6-phosphate can be described with the transport and enzyme constants of a three-compartment model (Fig. 4.1). The corresponding complex formula (Reivich et al. 1979) for calculating the regional cerebral metabolic rate of glucose (rCMRGlc) (Fig. 4.2) can be simplified to the following form (Phelps et al. 1979):

$$\text{rCMRGlc} = \frac{(\text{GI})}{\text{LC}} \cdot \frac{C(^{18}\text{F}) - C(\text{FDG})}{A_b}$$

where  $C(^{18}\text{F})$  is the total fluorine activity measured in the tissue, which is determined directly in the PET, and  $C(\text{FDG})$  is the concentration of free FDG in the tissue, calculated from the plasma concentration up to time point  $T$  with the aid of the

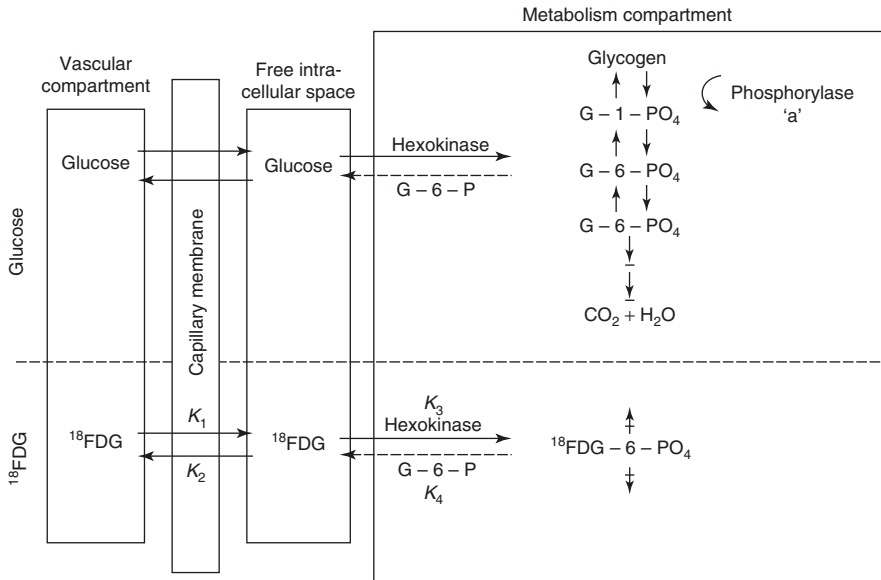
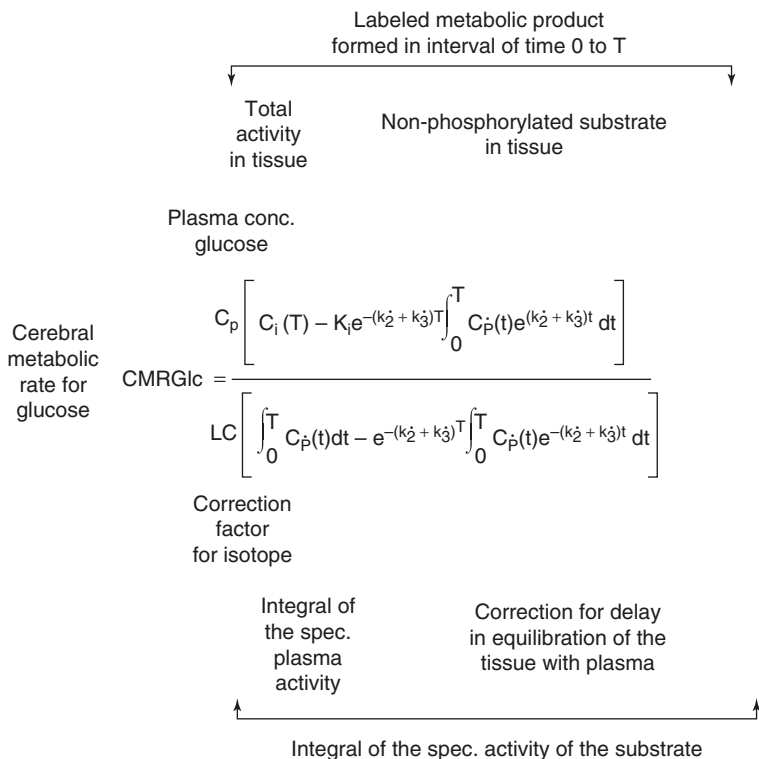


Fig. 4.1 Transport and metabolism of glucose and  $^{18}\text{F}$ -FDG in brain tissue (see text for details)



Operational equation for calculation of rCMRGlc

**Fig. 4.2** Equation for calculation of regional cerebral metabolic rate of glucose (rCMRGlc) in the brain

constants of the model. The difference between these two values gives the local tissue concentration of FDG-6-phosphate;  $A_b$  is the total quantity of FDG released into the tissue and is calculated from the plasma FDG concentration curve up to time point  $T$ , decreased by the delay in the tissue equilibration using the corresponding model constants. The quotient therefore gives the proportional phosphorylation rate of FDG. Multiplication by the plasma concentration of glucose (Glc) would yield the rate of glucose phosphorylation if it behaved in the same way as FDG. Since the arterial-venous extraction of glucose is not identical with that of FDG, the value must be corrected with an experimentally determined constant ( $LC =$  lumped constant). For the measurement of the regional glucose consumption in the brain, therefore, after i.v. administration of 3–6 mCi  $^{18}\text{F}$  FDG, the plasma curve of  $^{18}\text{F}$  FDG from the injection to the measurement time point (usually determined in arterialised venous blood), the glucose value in plasma, and the regional  $^{18}\text{F}$  activity in the brain must be determined.

Errors resulting with this model from widely diverging kinetic constants in pathological tissue (Hawkins et al. 1981) can be reduced by dynamic PET. For this purpose, the tissue activity is determined at short time intervals from the time point of injection. By variation of the values for the kinetic constants, the curve deriving from the model equation is adapted to the measured regional activity time curve. The kinetic constants thus determined correspond best to the activity uptake in the corresponding tissue segment. They allow the regional metabolic rate for glucose to be directly calculated (Wienhard et al. 1985). It is now only necessary to assume a known value for the LC.

There remains some uncertainty as to the exact value of the normal “lumped constant” (LC) for FDG. The initial value empirically derived by Phelps et al. (1979) was 0.42. It has been directly measured by Reivich et al. (1985), who found a value of 0.52, assuming  $k_4 = 0$ . But LC could be even as high as 0.65 (Wu et al. 2003). However, regional changes in LC are small. For the sake of uniformity, the actual value used should be quoted in all publications. This will allow direct comparison of numerical values because the LC is a linear scaling factor in the operational equations. With rate constants measured by dynamic curve fitting or by integration techniques, the equation for calculation of CMR<sub>glc</sub> is:

$$\text{CMR}_{\text{glc}} = C_a / \text{LC} \times (k_1 k_3 / (k_2 + k_3))$$

The term  $k_1 k_3 / (k_2 + k_3)$ , representing the metabolic rate of FDG, can be substituted by the influx rate constant  $k_i$  determined with the linear approximation of Patlak et al. (1983). There have also been modifications that avoid the assumption of a fixed LC and refer instead to the Michaelis–Menten equation to account for the relations between enzyme affinities for FDG and glucose (Kuwabara et al. 1990).

Determination of individual rate constants is not very practical in many clinical applications, and methods are preferred that can be done with a single scan, a situation similar to the original development of the method for autoradiography. Then the deviation from population average CMR<sub>glc</sub> (given by the average rate constants) is estimated from a single scan. Actual measured FDG activity is compared with the activity that would have been expected at the time of the scan with the individual’s blood activity time course and average rate constants (Wienhard et al. 1985).

To avoid the conversion factor needed with the analogue tracer FDG, native glucose labelled with <sup>11</sup>C in the 1-position (1-<sup>11</sup>C-d-glucose) has also been used for quantitation of CMR<sub>glc</sub> (Raichle et al. 1975). Modelling is based on the same two-tissue compartment model as with FDG, but an additional term is necessary to account for labelled metabolites (mainly lactate and other monocarboxylic acids and CO<sub>2</sub>). Metabolites occur in the plasma and in the brain, and loss of labelled CO<sub>2</sub> from the brain is dependent on CBF. Data indicate that there is a rapid loss of labelled lactate from the brain, suggesting that it represents a significant nonoxidative part of glucose metabolism in the brain (approximately 10% of the total CMR<sub>glc</sub>) (Blomqvist et al. 1990).

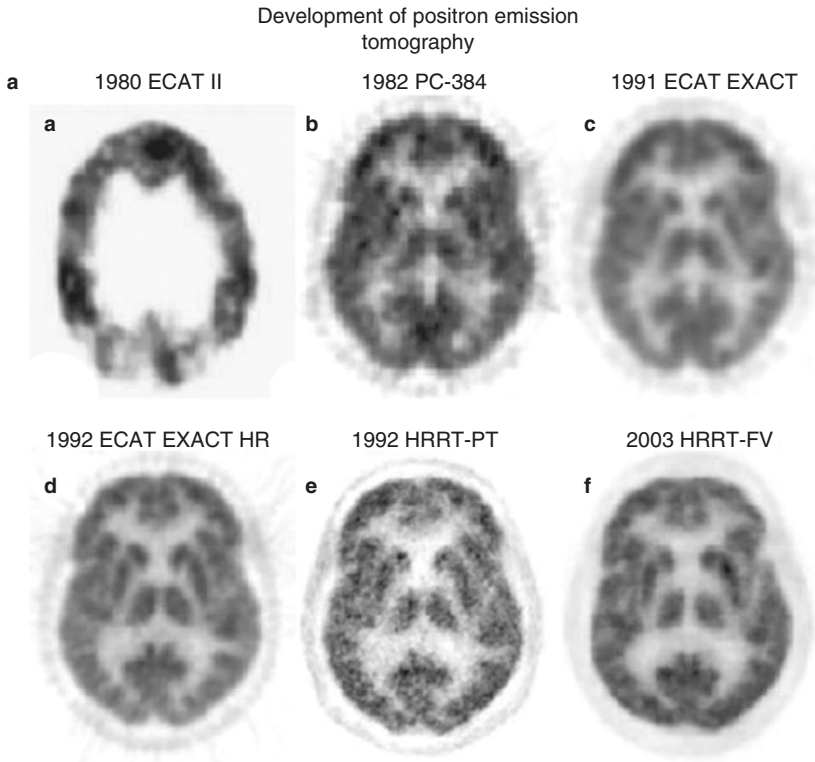


### 4.2.3 Normal Glucose Consumption of the Brain

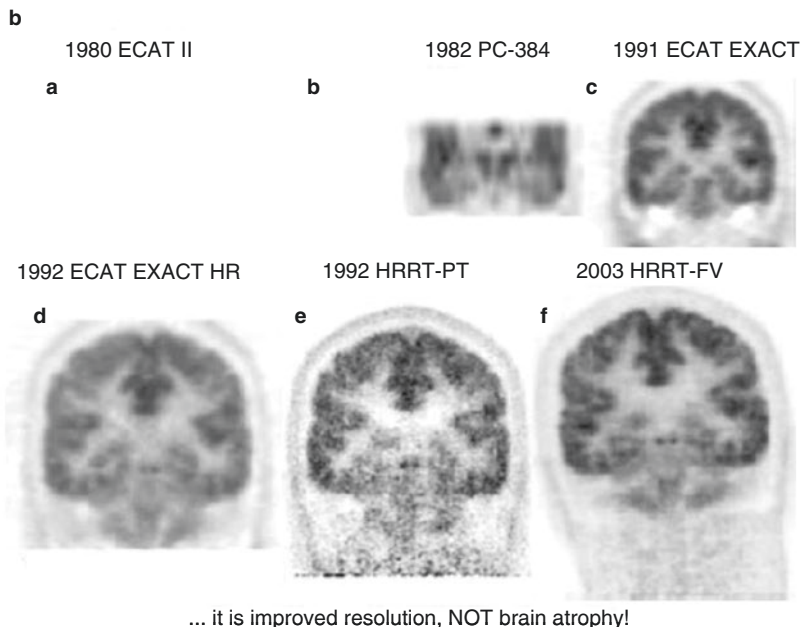
In healthy volunteers, a mean glucose consumption of 29–32  $\mu\text{mol}/100\text{ g}/\text{min}$  was found by means of FDG and PET (Reivich et al. 1979; Heiss et al. 1984), which correspond well to whole-brain metabolic rates provided by the Kety–Schmidt method. Under controlled conditions (darkened laboratory and steady noise from fans of equipment cooling systems), the functional anatomy of the brain is reflected in the metabolic activity of the individual regions. However, reliable regional values for cerebral metabolic rate of glucose (rCMRGlc) can only be obtained by equipment permitting high 3D resolution of tracer concentration in the brain tissue (Heiss et al. 2004). This progressively improved spatial resolution of PET is documented in Fig. 4.3 showing FDG images of the brain in the same volunteer assessed with different tomographs over the years (Heiss 2009a, b). Typical resting state grey matter CMRGlc values are in the range of 40–60  $\mu\text{mol}/100\text{ g}/\text{min}$ , and the corresponding level in the white matter is about 15  $\mu\text{mol}/100\text{ g}/\text{min}$ . There are significant differences among regions with highest values in the basal ganglia, primary visual cortex, and cingulate and frontal cortex (42–50  $\mu\text{mol}/100\text{ g}/\text{min}$ ) and lower values in other cortical and subcortical areas (35–42  $\mu\text{mol}/100\text{ g}/\text{min}$ ) and in the structures of the brain stem (25–30  $\mu\text{mol}/100\text{ g}/\text{min}$ ) and the cerebellum (33  $\mu\text{mol}/100\text{ g}/\text{min}$ ). There exist also significant asymmetries with largely right hemispheric predominance (Pawlik and Heiss 1989), review in Silverman and Melega (2004). The resting regional metabolism and its asymmetry are highly dependent on the state of resting wakefulness (e.g. apprehensive or relaxed) and background conditions (e.g. laboratory noise).

Local CMRGlc measured with PET is influenced by age: glucose metabolism of various grey matter structures was low at birth (13–25  $\mu\text{mol}/100\text{ g}/\text{min}$ ), reached a level of 19–33  $\mu\text{mol}/100\text{ g}/\text{min}$  by 2 years and continued to rise until age 3–4 years, and was maintained at a high level (49–55  $\mu\text{mol}/100\text{ g}/\text{min}$ ) until age of 10 years (Chugani et al. 1987). At about 10 years, CMRGlc began to decline with a rather uniform decrease by 26% in all investigated brain regions of 40 healthy resting subjects between the ages of 18 and 78 years (Kuhl et al. 1982). However, these age-dependent changes were not observed in all studies (Duara et al. 1984). In our own study on 42 normal subjects aged 15–85 years, a small (0.65  $\mu\text{mol}/100\text{ g}/\text{min}$  per decade/ $p < 0.05$ ) age-dependent decrease in global CMRGlc was found (Fig. 4.4). However, as demonstrated in Fig. 4.4, the various regions contributed differently to this overall effect: decreases of 16.6–11.3% in cingulate, frontal, parietal, insular, temporal, and sensorimotor cortex, virtually no change in the primary visual cortex and cerebellum (Pawlik and Heiss 1989). Similar age-dependent changes of rCMRGlc were described in further studies (Kalpouzos et al. 2009; Hsieh et al. 2012; Chetelat et al. 2013; Berti et al. 2014; Shen et al. 2012; Bonte et al. 2017; Jiang et al. 2018).

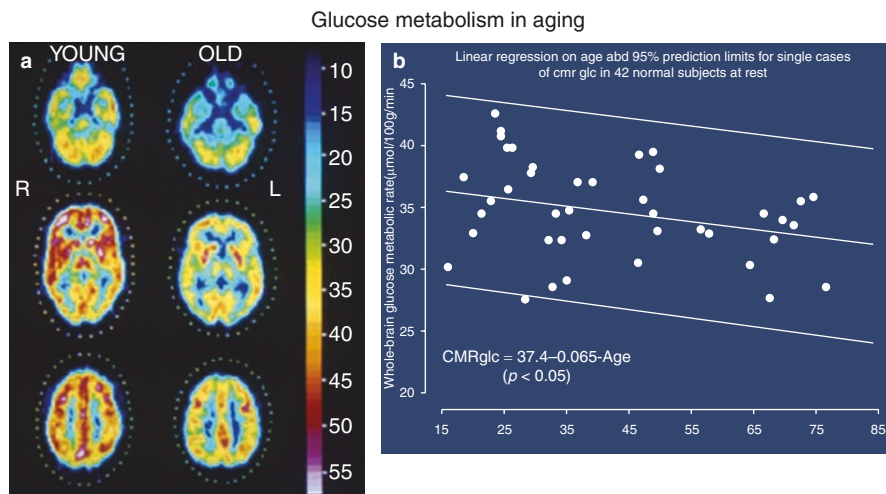
Hardly any normal functional state is as regularly associated with as dramatic changes of general behaviour and shifting of attention as is sleep. The conclusion of no effect of sleep on human cerebral haemodynamics and metabolism, derived from early Kety–Schmidt studies (Mangold et al. 1955), could be disproved with PET (Heiss et al. 1985). As shown in Fig. 4.5, during stages II–IV sleep, a significant ( $P < 0.001$ ) global decrease of brain functional activity was observed, with the



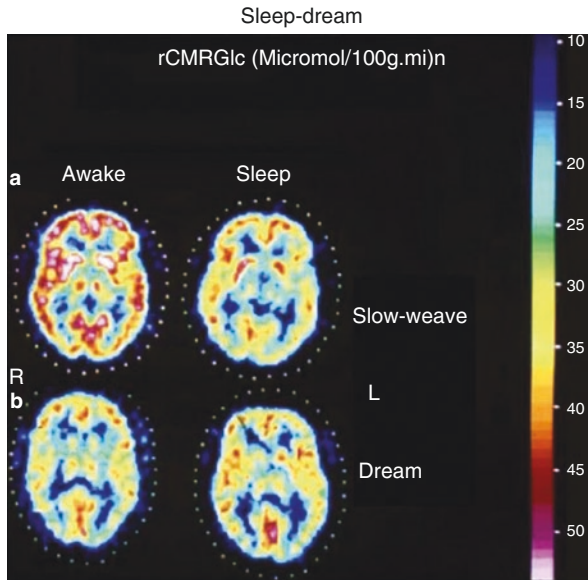
**Fig. 4.3** (a) Development of PET: a horizontal slice. Various PET systems over the years demonstrate improvement in image quality and spatial resolution. (a) ECAT II, spatial resolution 15 mm. (b) PC-384, FWHM 8.4 mm. (c) ECAT EXACT, FWHM 6.5–7 mm. (d) ECAT EXACT HR, FWHM 3.6–4.5 mm. (e) f HRRT PET: e prototype. (f) final version (HRRT-FV), FWHM 2.3–3.2 mm. Images of glucose metabolism were acquired for 20 min of steady-state starting 30 min after tracer administration. (b) Coronal views of glucose consumption of the brain in a volunteer acquired with various PET systems over the years demonstrate improvement in axial resolution due to decreased slice thickness and advances in image reconstruction. (a) ECAT II (1980) was a single-ring camera; axial reconstruction was therefore not feasible. (b) PC-384, slice thickness 12 mm. (c) ECAT EXACT, axial FWHM 5–8 mm. (d) ECAT EXACT HR, axial FWHM 4.0–6.7 mm. (e) f HRRT PET: e prototype. (f) final version (HRRT-FV), axial FWHM 2.5–3.4 mm



**Fig. 4.3** (continued)



**Fig. 4.4** (a) PET scans of glucose metabolism ( $\mu\text{mol}/100 \text{ g}/\text{min}$  according to scale) in cerebral sections at the level of the cerebellum, basal ganglia, thalamus, and semioval centre in young (23 years) and old (67 years) healthy subjects. The individual brain structures can be differentiated according to different metabolic rates; metabolism decreases slightly in all regions in older patients. (b) Decrease of mean global glucose metabolic rate in 42 healthy subjects with increasing age. The regression line shows a significant relationship despite the large range of variation



**Fig. 4.5** (a) Corresponding images of the local cerebral metabolic rate for glucose determined by FDG-PET in a characteristic horizontal brain slice across the basal ganglia of a healthy 37-year-old male subject representative of the non-dreamers group, showing nonselective decrease in glucose utilisation from wakefulness (W) to sleep (S). Frontal poles are on top, occipital poles at bottom, sides as marked (R, L). Values on reference scale are in  $\mu\text{mol}/100\text{ g}\cdot\text{min}$ . (b) Corresponding metabolic maps of a 28-year-old normal volunteer's brain slice recorded by FDG-PET while the subject was awake (W) and asleep dreaming (S). A generalised activation, most marked in the insular regions, visual cortex, and hippocampal formations, is clearly demonstrated during sleep with dreaming

largest declines in the orbitofrontal cortex and in the thalamus. In dream sleep, by contrast, both a general metabolic increase and conspicuous regional activations of superior frontal, insular, inferior parietal, hippocampal, and visual association cortex (Heiss et al. 1985) were found. Increased CMRGlc during REM sleep has also been observed in limbic and paralimbic regions including hypothalamus, amygdala, orbitofrontal cingulate, entorhinal, and insular cortices (Nofzinger et al. 1997).

#### 4.2.4 Coupling of Neuronal Activity to Metabolism and Flow

The activation of  $\text{Na}^+$ ,  $\text{K}^+$ -ATPase represents the coupling mechanism between the increase in glucose utilisation and functional activity of the nervous tissue. The activation-induced increase in glucose uptake is visualised in the neuropil, that is, where synapses ensheathed by astrocytes are present, not at the level of the neuronal perikarya. Glucose, taken up by astrocytic processes, is metabolised glycolytically to lactate and pyruvate, which are then released as substrates for oxidative

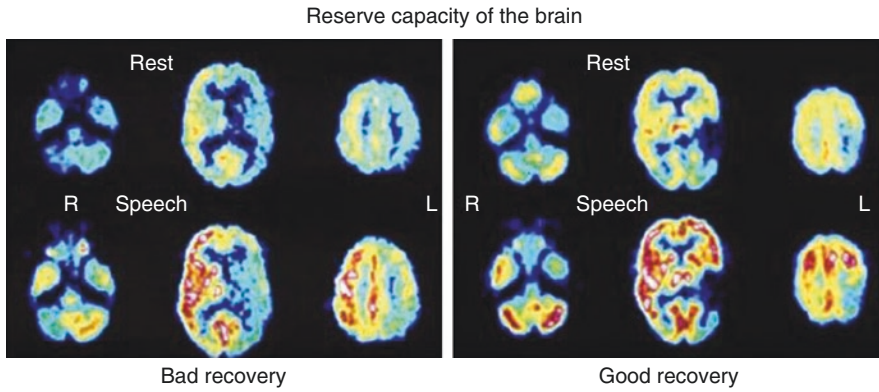
phosphorylation in neurons (Wyss et al. 2011; Juaristi et al. 2019). Mapping of neuronal activity in the brain can be primarily achieved by quantitation of the regional cerebral metabolic rate for glucose (rCMRGlc), as introduced for autoradiographic experimental studies by Sokoloff (1977) and adapted for positron emission tomography (PET) in humans (Reivich et al. 1979). Functional mapping, as it is widely used now, relies primarily on the hemodynamic response assuming a close association between energy metabolism and blood flow. While it is well documented that increases in blood flow and glucose consumption are closely coupled during neuronal activation, the increase in oxygen consumption is considerably delayed leading to a decreased oxygen extraction fraction (OEF) during activation (Villien et al. 2014; Mintun et al. 2001). PET detects and, if required, can quantify changes in CBF and CMRGlc accompanying different activation states of the brain tissue. The regional values of CBF or CMRGlc represent the brain activity due to a specific state, task, or stimulus, in comparison to the resting condition, and colour-coded maps can be analysed or coregistered to morphologic images.

Due to the radioactivity of the necessary tracers, activation studies with PET are limited to a maximum of 12 doses of  $^{15}\text{O}$ -labelled tracers, e.g. 12 flow scans, or 2 doses of  $^{18}\text{F}$ -labelled tracers, e.g. 2 metabolic scans. Especially for studies of glucose consumption, the time to metabolic equilibrium (20–40 min) as well as the time interval between measurements required for isotope decay (HT for  $^{18}\text{F}$  108 min, for  $^{15}\text{O}$  2 min) must be taken into consideration. FDG-PET was the leading method to investigate functional activation in humans in the 1980s (Pawlik and Heiss 1989; Phelps et al. 1981). PET-FDG activation studies assess task-induced CMRGlc changes either by performing the bolus method, with one or two separate PET scans, or as described recently by constant infusion of FDG during the entire scan for rest and activation condition (Villien et al. 2014; Hahn et al. 2016). FDG activation studies can also be applied in patients with functional disorders due to localised brain damage, e.g. by stroke and tumour (review in Chiaravalloti et al. (2019)), and has found broad application to patients with aphasia (Heiss 2009a, b). An example is given in Fig. 4.6 showing different activation patterns in poststroke aphasia which are related to prognosis and recovery of language function.

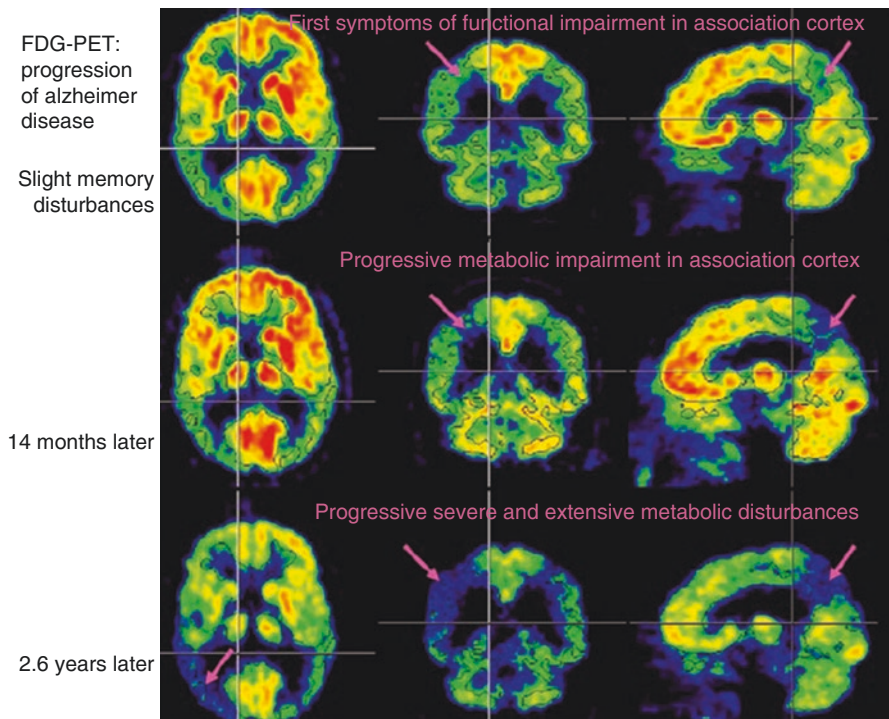
#### 4.2.5 Clinical Applications of FDG-PET

Since its introduction, FDG-PET has been applied for studying the pathophysiology and for differential diagnosis of several neurological and psychiatric disorders (Chiaravalloti et al. 2019; Herholz et al. 2013; Jones et al. 2012). These applications will be described in the special clinical chapters of this book series. Some examples where FDG-PET has gained special importance are shown here.

In dementias, FDG-PET has attained a special role to detect progression of regional functional disturbance related to severity of cognitive and memory impairment (Fig. 4.7) and for differential diagnosis to other degenerative disorders

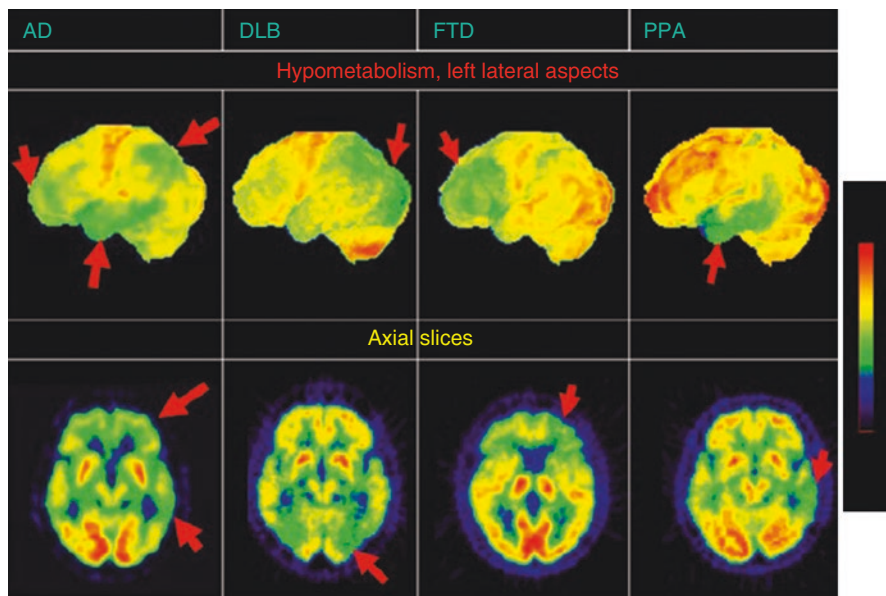


**Fig. 4.6** Resting and speech-activated regional glucose metabolism in two patients with aphasia after ischemic stroke: if only contralateral regions are activated by speech, the prognosis is poor. If activation takes also place in homolateral periinfarct regions, prognosis is better, and speech performance shows satisfactory recovery



**Fig. 4.7** Decline of cerebral metabolic rate (glucose) (CMRGlc) in association areas with progression of AD from the clinical stage of mild cognitive impairment (MCI) to mild dementia (three follow-up FDG-PET scans, each showing the same orthogonal slices at position marked by crosshairs)

## Differential diagnosis of dementias by FDG-PET



**Fig. 4.8** FDG-PET in the differential diagnosis of various degenerative dementias. *Upper row*: typical transaxial slices. *Lower row*: reconstructed surface view. *AD* Alzheimer dementia, characterised by the decreases in temporoparietal and temporal association and in cingulate cortex; *FTD* frontotemporal dementia, the metabolic decrease is most severe in the anterior frontal and temporal regions; *DLB* dementia with Lewy body (Parkinson's disease), the metabolic disturbance also affects the visual cortex; *PPA* primary progressive aphasia, the disturbance is most accentuated in the temporal (Wernicke) area. *Arrows* indicate most prominent changes

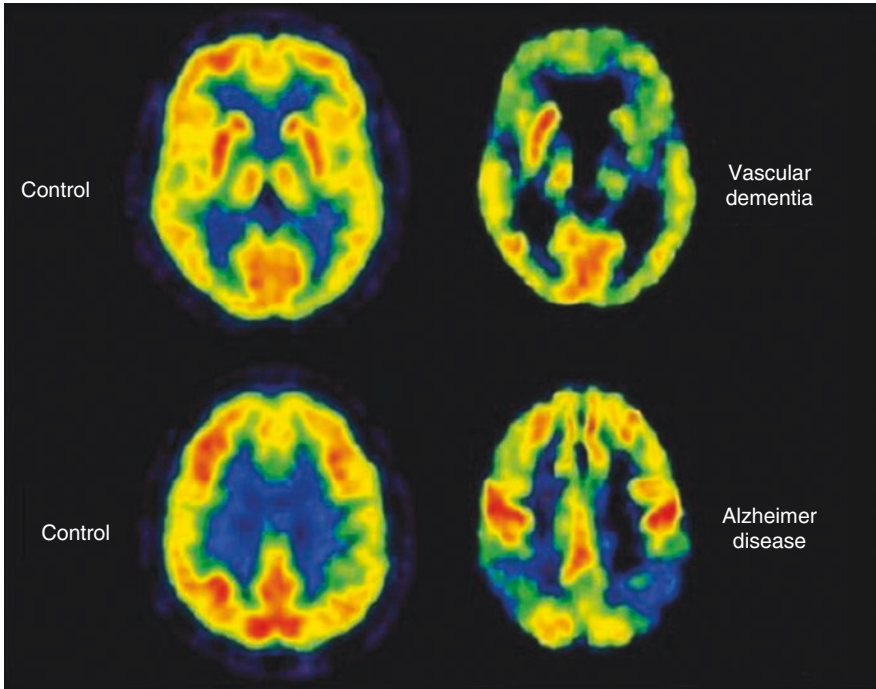
(Fig. 4.8) and to vascular dementia (Fig. 4.9) (Drzezga 2009; Drzezga et al. 2018; Heiss and Zimmermann-Meinzingen 2012; Heiss 2018; Bohnen et al. 2012; Choo et al. 2013; Garibotto et al. 2017).

In brain tumours, FDG-PET has been successful in differentiating between necrosis and recurrent tumour (Fig. 4.10) and has value for grading of gliomas and for assessing the effect of chemotherapy (Heiss et al. 2011; Chierichetti and Pizzolato 2012; Herholz 2017; Herholz et al. 2012).

FDG-PET is the most common tracer used in epilepsy since epileptogenic foci are hypometabolic on interictal imaging, and FDG imaging is a commonly used tool in presurgical assessment of epilepsies (von Oertzen 2018; Broski et al. 2018; Lotan et al. 2020).

FDG-PET imaging has also been extensively performed in Parkinson's disease and is able to distinguish between several other movement disorders (Meyer et al. 2017; Meles et al. 2020).

Metabolic pattern in dementia:  
VaD ns AD



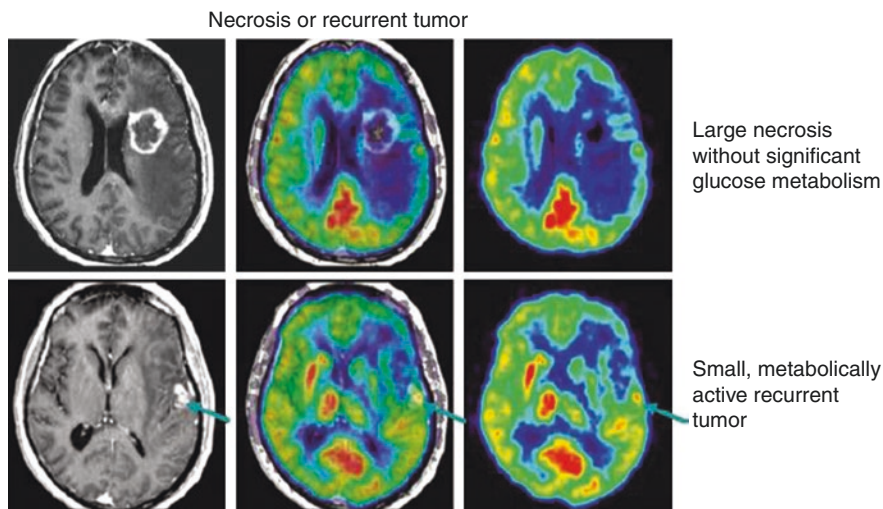
**Fig. 4.9** Glucose metabolism in a normal control, in a patient with vascular dementia and a patient with Alzheimer's disease. The severity of dementia was comparable; the pattern of pathological changes differentiated these two cases: patchy metabolic defects in VaD in the frontal lobe, basal ganglia, and thalamus and hypometabolism in AD bilateral in parieto-temporal cortex and to a lesser degree in the frontal association areas, whereas primary cortical regions are spared

Neuroimaging of sleep disorders such as narcolepsy and primary hypersomnias with FDG-PET combined with other MR-based measures has given insight into the neural basis and pathogenesis of narcolepsy and primary or idiopathic hypersomnias (for review, see Cavaliere et al. 2020).

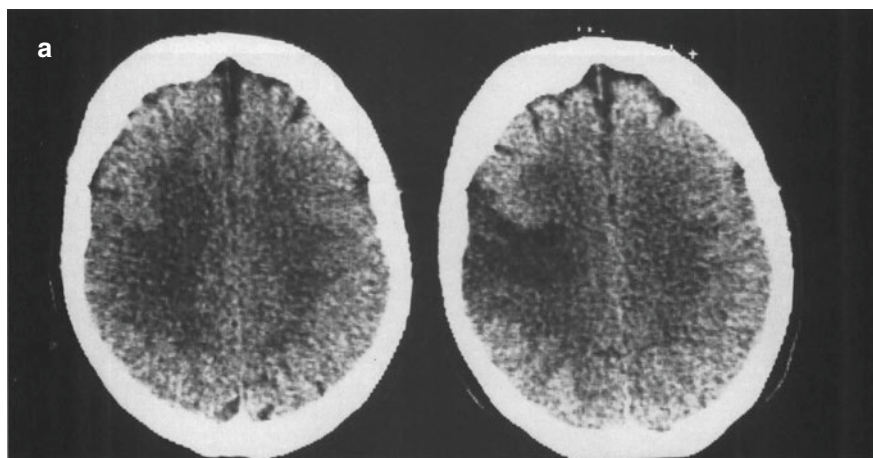
In ischemic stroke, 18FDG-PET has a role in discriminating recoverable ischemic brain tissue (penumbra) from infarcted tissue (infarct core) to predict along with 15O-PET tissue fate in acute and subacute ischemic stroke (Fig. 4.11) (Heiss et al. 1992; Nasu et al. 2002; Bunevicius et al. 2013).

Since plaque inflammation contributes to stroke and FDG identifies carotid plaque inflammation-related metabolism, FDG-PET is suitable to independently predict future recurrent stroke which may improve patient selection for revascularisation therapies as well as anti-inflammatory therapy (Marnane et al. 2012; Kelly et al. 2019).

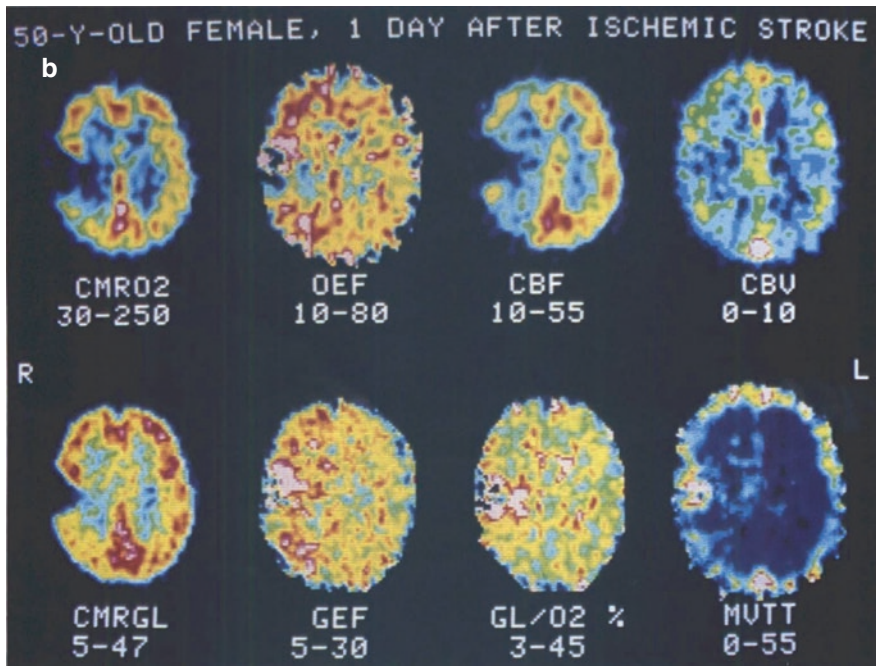




**Fig. 4.10** FDG-PET and coregistered MRI in a patient with a large contrast-enhancing radiation necrosis (*top row*) and a small recurrent active carcinoma metastasis (*bottom row*)



**Fig. 4.11** CT and multitracer PET study of a patient 24 h after ischemic stroke in the territory of the right middle cerebral artery. CT and PET images of a brain slice 55 mm above the canthomeatal plane are presented. **(a)** While the initial CT is inconclusive, the ischemic infarct is clearly demarcated on the CT 4 days later. **(b)** PET images of the measured variables clearly demonstrate the flow defect (on CBF image) and the metabolic disturbance (CMRO<sub>2</sub> and CMRGl<sub>c</sub>): the flow defect, however, is larger than the CMRO<sub>2</sub> and CMRGl<sub>c</sub> defect, leaving border zone regions with preserved oxygen and glucose consumption and therefore increased oxygen extraction fraction (OEF) and glucose extraction fraction (GEF). This anterior portion is preserved at the later CT. In the posterior rim, however, CMRO<sub>2</sub> is more severely impaired than CMRGl<sub>c</sub> leading to an increase in the ratio of glucose to oxygen consumption indicative of anaerobic glycolysis. On later CT, this area is infarcted. In the infarct, CBV is increased in relation to CBF leading to an increased transit time (TT). Due to the occlusion of the right internal carotid artery, CBF in the ipsilateral hemisphere outside the infarct is reduced without effect on CMRO<sub>2</sub> and CMRGl<sub>c</sub> since OEF and GEF are increased



**Fig. 4.11** (continued)

The impact of PET has further increased by the advent of integrated MRI-PET facilities (Broski et al. 2018; Catana et al. 2012; Portnow et al. 2013; Tondo et al. 2019; Shepherd and Nayak 2019; Shiyam Sundar et al. 2020).

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# Cerebral Blood Flow Measurement with Oxygen-15 Water Positron Emission Tomography

# 5

Henryk Barthel, Vilia Zeisig, Björn Nitzsche, Marianne Patt, Jörg Patt, Georg Becker, Antje Dreyer, Johannes Boltze, and Osama Sabri

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## Abstract

The human brain receives approximately 15% of the cardiac output and therefore is the most demanding organ in respect to blood flow supply. This fact emphasizes the importance of perfusion as a key factor in a variety of cerebrovascular and other diseases including stroke, migraine, and brain tumors. Today, numerous imaging techniques are able to visualize brain perfusion, but only few of them provide quantitative information. In the field of modern in vivo imaging techniques, positron emission tomography (PET) is considered to be the gold standard to give reliable results about major aspects of cerebral physiology. [ $^{15}\text{O}$ ]H $_2$ O allows for quantitative cerebral blood flow (CBF) measurement within a few minutes, and subsequent  $^{15}\text{O}$ [O $_2$ ] imaging can provide precise information on oxygen metabolism like cerebral oxygen metabolism and oxygen extraction fraction. As a result, PET has become an extremely useful research tool for defining cerebral blood flow and physiology. However, complex methodological logistics and a limited availability of the imaging system hamper the widespread use of CBF PET in clinical routine. The chapter aims at summarizing the radio-synthesis, data acquisition, and analysis, as well as major preclinical and clinical applications of [ $^{15}\text{O}$ ]H $_2$ O PET.

## Abbreviations

$c_a(t)$	Arterial input function (arterial activity concentration over time)
CBF	Cerebral blood flow
CBV	Cerebral blood volume
CMRO $_2$	Cerebral metabolic rate of oxygen
CT	Computed tomography
$c_v(t)$	Activity concentration in venous blood over time
$f_a$	Arterial blood flow
GBq	Gigabecquerel
GM	Gray matter
IAP	Iodoantipyrine
iNO	Inhaled nitric oxide
$k$	Washout constant
kBq	Kilobecquerel
MBq	Megabecquerel
MRI	Magnetic resonance imaging
NIRS	Near-infrared spectroscopy
NO	Nitric oxide
OEF	Oxygen extraction fraction
PET	Positron emission tomography
rCBF	Regional cerebral blood flow
ROI	Region of interest
S/N	Signal-to-noise
SPECT	Single-photon emission computed tomography



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TSE	Turbo spin echo
$V_d$	Partition coefficient
$V_{\text{tissue}}$	Tissue volume
WM	White matter
Xe	Xenon

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## 5.1 Introduction

Although the adult human brain assumes only about 2% of the total body weight, it receives nearly 15% of the resting cardiac output and 20% of the total body oxygen consumption. This high metabolic rate reveals the need to ensure a constant delivery of oxygen and energy-providing substrates at the capillary level and to remove the waste products of metabolism. Owing to a complex autoregulatory mechanism, the cerebral perfusion is maintained relatively constant over a wider range of mean arterial pressures. A complete interruption of brain–blood supply, however, leads immediately to neuronal impairments because of the limited availability of nutrition and energy reserves in the brain. Hence, perfusion parameters are important key factors involved in major cerebrovascular and other brain diseases. Important information about cerebral characteristics is given by measurement of the cerebral blood flow (CBF) which describes the rate of blood delivery to the brain parenchyma. Traditionally, the respective literature reports CBF units in mL blood/100 g of tissue per minute. Other authors use mL instead of gram tissue to describe the volumetric character especially with imaging techniques. Due to the fact that 1 g of brain tissue nearly corresponds to 1 mL, both values can be more or less used interchangeably.

Global average CBF values in middle-aged healthy human subjects are around 55 mL/100 g/min (Kety and Schmidt 1945). However, CBF values vary regionally: In cortical gray matter, the CBF is 60–100 mL/100 g/min (Slosman et al. 2001; Sokoloff et al. 1957) and around 20 mL/100 g/min for white matter (Law et al. 2000). It has been demonstrated that neuronal activity as well as CBF is closely coupled to brain metabolism (Pantano et al. 1984; Roy and Sherrington 1890). This may explain the generally higher blood flow in younger subjects which is typically exceeding values that are 50–85% higher than those of adults (age: 6–7 years, Chiron et al. 1992). This age dependency of CBF is probably caused, at least in parts, by an age effect on the gray-to-white matter ratio. Additionally, a decline of CBF with age is described (Brody 1955).

Kety and Schmidt (1945) were the first who introduced a method for quantitative CBF measurement in the unanesthetized human. This method is based on the Fick principle. During the inhalation of the freely diffusible nitrous oxide, the brain perfusion was calculated by measuring the amount of gas removed from the blood by the brain per minute and dividing this by the arteriovenous difference of  $\text{N}_2\text{O}$ . This method was an important step to study brain function in humans and has contributed fundamentally to our understanding of physiological and pathological cerebral processes. However, this first technique lacked the possibility to measure CBF in different brain subregions. This limitation has led to further developments in perfusion

imaging techniques. As important examples, radioactive agents such as  $^{133}\text{Xe}$  and  $^{85}\text{Kr}$  were later used to measure their washout with external radiation detectors (Lassen and Ingvar 1961; Veall and Mallett 1967; Bruce et al. 1973; Obrist et al. 1975). Today, a wide variety of brain perfusion imaging techniques is available for preclinical and clinical research settings as well as clinical routine applications. CBF measurement using positron emission tomography (PET) became available in the 1970s, followed by other *in vivo* imaging techniques, like single-photon emission computed tomography (SPECT), magnetic resonance imaging (MRI), and transmission computed tomography (CT), to obtain CBF readouts (Hoeffner 2005). This chapter will focus on CBF PET with  $^{15}\text{O}$  as this represents the gold standard technique for *in vivo* CBF measurements (Carroll et al. 2002; Hoeffner 2005). One possible application for its usage is studying CBF in brain activation studies to localize brain segments involved in processing certain tasks. More important today, however, is the diagnostic potential of CBF PET imaging mainly in cerebrovascular disorders and oncology. Hence alterations of cerebral perfusion can theoretically be detected by quantitative CBF measurement leading to optimal therapy and providing estimates of the potentially salvable part of the affected brain.

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## 5.2 Radiochemistry of $^{15}\text{O}$

Oxygen-15 can be produced by different nuclear reactions using both high- and low-energy protons as well as deuterons as projectiles. The  $^{14}\text{N}(\text{d}, \text{n})^{15}\text{O}$  nuclear reaction is the most efficient and economic production pathway and is therefore applied most frequently. For cyclotrons that do not have the possibility of accelerating deuterons, either the  $^{15}\text{N}(\text{p}, \text{n})^{15}\text{O}$  (Powell and O'Neil 2006) for low-energy protons (>3.7 MeV) or the  $^{16}\text{O}(\text{p}, \text{pn})^{15}\text{O}$  (Beaver et al. 1976; Krohn et al. 1986) for high-energy protons (>16.6 MeV) nuclear reactions can be used as an alternative.

Two methods are available for conversion of the  $^{15}\text{O}$  to  $^{15}\text{O}$ , the in-target production method and the out-of-target external conversion method. In case of the in-target production method,  $^{15}\text{O}$  is produced by either addition of small amounts of hydrogen to the target gas or direct irradiation of  $\text{H}_2^{16}\text{O}$ . For the first approach, the  $^{15}\text{N}(\text{p}, \text{n})^{15}\text{O}$  nuclear reaction is used (Powell and O'Neil 2006), and the trace amounts of  $\text{H}_2$  in the target lead to the formation of  $^{15}\text{O}$  by radiolytic reactions which can be trapped in a cooled stainless steel loop. The possible radioactive impurities such as  $^{11}\text{CH}_4$  and  $^{13}\text{NH}_3$  (resulting from trace amounts of N-14 and O-16 in the target gas) are directed to the exhaust. By heating the loop,  $^{15}\text{O}$  vapor is released and can be trapped again in a sterile water or saline solution. The major drawbacks are the high costs for enriched  $^{15}\text{N}_2$  which is used as target material. An alternative approach for  $^{15}\text{O}$  in-target production is the direct irradiation of  $\text{H}_2^{16}\text{O}$  in a liquid target (Mulholland et al. 1990; van Naemen et al. 1996). As mentioned above this method is feasible only with high-energy protons, a restriction that excludes this approach for commonly used cyclotrons in a clinical setting.

As an alternative to the in-target production of  $^{15}\text{O}$ , external out-of-target tracer production by catalytic conversion of a mixture of  $\text{H}_2$  and  $^{15}\text{O}_2$  has been

demonstrated for all abovementioned nuclear reactions, i.e.,  $^{16}\text{O}(p, pn)^{15}\text{O}$  (Krohn et al. 1986),  $^{15}\text{N}(p, n)^{15}\text{O}$  (Explora<sup>®</sup>H<sub>2</sub>O module, Siemens Healthcare, Erlangen, Germany), and  $^{14}\text{N}(d, n)^{15}\text{O}$  (Clark et al. 1987; Clark and Tochon-Danguy 1991; Sajjad et al. 2000). For the  $^{14}\text{N}(d, n)^{15}\text{O}$  nuclear reaction, N<sub>2</sub> with up to 4% O<sub>2</sub> is irradiated with deuterons ( $\approx 7$  MeV). In most cases, Pd is used as the catalyst; however, if the same target is used as well for production of C<sup>15</sup>O, it might be necessary to reduce the amount of O<sub>2</sub> in order to minimize the production of toxic carrier CO. Under these conditions it has been shown that Pt as catalyst is superior (Berridge et al. 1990). The most simple production and application approach consists of an H<sub>2</sub> supply that is connected to the target line via a T connector. The target gas/H<sub>2</sub> mixture (2–10%, Berridge et al. 1990; Sajjad et al. 2000) is then directed over the heated catalyst ( $\approx 170$  °C, temperatures up to 450 °C have been reported as well (Berridge et al. 1990)), and the resulting [<sup>15</sup>O]H<sub>2</sub>O vapor is bubbled into a sterile reservoir containing water, saline, or preferably buffer since the catalyst might as well generate trace amounts of ammonia which un-buffered may lead to elevated pH values. The [<sup>15</sup>O]H<sub>2</sub>O-containing solution is then drawn up into a syringe and manually applied to the investigated subject. However, due to the relatively high radiation exposure of the medical personnel, it might be worth to automate the injection procedure. In a relatively simple setup, the injection is performed by means of two infusion pumps and a 4-port valve (Sajjad et al. 2000). A more sophisticated method uses a dialysis membrane to enable exchange of [<sup>15</sup>O]H<sub>2</sub>O with sterile water (Clark and Tochon-Danguy 1991) together with an infusion pump and several valves to enable automatic injection. This system is commercially available (Veenstra Instruments, Joure, Netherlands).

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### 5.3 [<sup>15</sup>O]H<sub>2</sub>O Brain PET Data Generation

Over the years, different methods to perform [<sup>15</sup>O]H<sub>2</sub>O CBF measurements have been developed and applied. Early users employed a technique originally developed by Kety and coworkers to determine rCBF with tissue autoradiography in laboratory animals (Kety 1951). Later on, groups used the steady-state technique developed by Jones and colleagues in which [<sup>15</sup>O]CO<sub>2</sub> needs to be continuously inhaled over the examination time (Jones et al. 1976).

For the dynamic PET scanning method which is the current standard approach, the emission scan is, after a transmission scan for attenuation correction, acquired preferentially in 3D mode and initiated immediately before tracer administration. The tracer is injected as a fast bolus followed by a flush of inert saline solution. The amount of injected tracer in humans typically ranges between 550 and 1000 MBq for an adult subject, but studies using up to 2.2 GBq were also found in the literature (Heiss et al. 2000). For this bolus injection method, Kanno et al. (1991) investigated an optimal scan time for [<sup>15</sup>O]H<sub>2</sub>O to improve image quality and signal-to-noise ratios. A minimum scan duration of 90 s was recommended. For most applications, however, scan durations of 2–5 min are chosen. A typical protocol for a 5-min dynamic data acquisition is shown in Table 5.1.

**Table 5.1** Acquisition protocol for a 5-min PET scan with [ $^{15}\text{O}$ ]H $_2$ O

Frame duration [s]	5	10	30
Number of frames	24	12	2

The advantageous short half-life of the tracer (122 s) enables the performance of multiple image acquisition scans in rapid sequence. Inter-scan intervals should, however, not be shorter than 15 min to allow isotope decay. Due to the fast radioactive decay, an on-site cyclotron or linear accelerator for isotope production becomes necessary. Further data analyses generally include the absolute quantification of CBF. For this purpose, the experimental setting implies arterial blood sampling in parallel to the PET acquisition. This is preferentially performed using an automated sampling system (e.g., ALLOGG AB blood sampler; Allogg Mariefred, Sweden) with a peripheral artery, e.g., the radial artery. Using such a device, arterial blood samples are continuously drawn at a constant speed with activity measurements for every 0.5 or 1 s. The blood sampler needs to be cross-calibrated to the PET scanner, allowing the decay-corrected blood data to be used as input function for kinetic modeling.

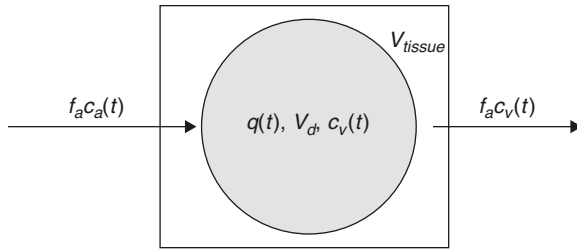
Obtaining arterial blood samples via the placement of an arterial catheter is accepted as the gold standard method for CBF quantification. However, there are some limitations, like invasiveness, complications for the patient, and sensitivity to errors (Hall 1971; Machleder et al. 1972). Aiming to replace the arterial canalization and corresponding input function, alternatives were suggested, like image-derived input functions (Zanotti-Fregonara et al. 2011) and methods without the need of any input function (Lammertsma 1994; Watabe et al. 1996). Another possibility to avoid arterial canalization is the use of arterialized venous blood, which is an often used method also with other PET tracers. With this technique, the hand of the patient or volunteer is heated well above 37 °C to achieve a shunting of arterial blood to the venous system (Wakita et al. 2000). Nevertheless, all alternative techniques have their drawbacks, and the CBF values obtained by them need to be handled with caution.

As for all brain PET tracers, the resulting PET data require a correlation with structural information from MRI (preferably 3D-T1 data). In this regard, the use of combined PET/MRI scanners might improve this situation in the future. Combined PET/MR imaging gives the opportunity for accurate registration and exact correlation of PET functional aspects with anatomical information from MRI. This will result in better image quality because of the comparably lower spatial resolution of PET in contrast to MRI and also shortens imaging times for the patients/anesthetized research animals.

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## 5.4 Kinetic Modeling of CBF

The first method to measure CBF in humans was proposed by Kety and Schmidt in 1945 who utilized nitrous oxide for CBF detection. Later, other tracers like  $^{133}\text{Xenon}$  (Veall and Mallett 1967) were applied within this concept which was based on the



**Fig. 5.1** Schematic description of a single-tissue compartment.  $V_{\text{tissue}}$  is the anatomical tissue volume ( $1 \text{ cm}^3$ ), and  $V_d$  is the volume of the tissue compartment accessible to the tracer. The products of flow and concentration describe the amount of tracer which enters or leaves the tissue compartment per time unit. Please refer to the main text for further explanation of the other symbols

Fick principle and a single-tissue compartment model. It states that the amount of a metabolically inert and freely diffusible gas that is taken up by a tissue per unit of time is equal to the product of the blood flow through that tissue and the difference between the amount of gas entering it via the arterial blood and the gas leaving in the venous blood.

The single-tissue compartment model describes the behavior of a freely diffusible tracer like  $[^{15}\text{O}]\text{H}_2\text{O}$  in tissue as shown in Fig. 5.1 and can be used to determine the local arterial blood flow in the brain (on a region of interest (ROI) or voxel basis). The model consists of two parameters that have to be estimated from the data of a dynamic PET scan ( $q(t)$ ) and from the measured arterial input function  $c_a(t)$ .

Assuming that the transport of tracer from the vessel into the tissue compartment is fast (high permeability surface area product) compared to the delivery by the arterial blood flow, the tracer dynamics can be described by a one-tissue compartment model with one input function. The mass balance for the tracer (Fick principle) yields the differential equation

$$\frac{dq(t)}{dt} = f_a c_a(t) - f_a c_v(t) \quad (5.1)$$

where  $q(t)$  is the quantity of tracer per unit volume of tissue ( $\text{kBq cm}^{-3}$ ),  $f_a$  is the local arterial blood flow per unit volume ( $\text{mL min}^{-1} \text{cm}^{-3}$ ), and  $c_a(t)$  and  $c_v(t)$  are the tracer concentrations in arterial and venous blood ( $\text{kBq cm}^{-3}$ ).

The tracer concentration in the venous blood ( $c_v(t)$ ) is related to the tracer concentration in the tissue space ( $q(t)$ ) through the relative volume of distribution ( $V_d$  ( $\text{mL cm}^{-3}$ )) by the Kety–Schmidt assumption:

$$q(t) = V_d c_v(t) \quad (5.2)$$

reflecting the assumption that the concentrations in the water spaces of venous blood and tissue are always equilibrated (this assumption is not generally valid, because a diffusion limitation exists for  $[^{15}\text{O}]\text{H}_2\text{O}$  at low CBF rates).

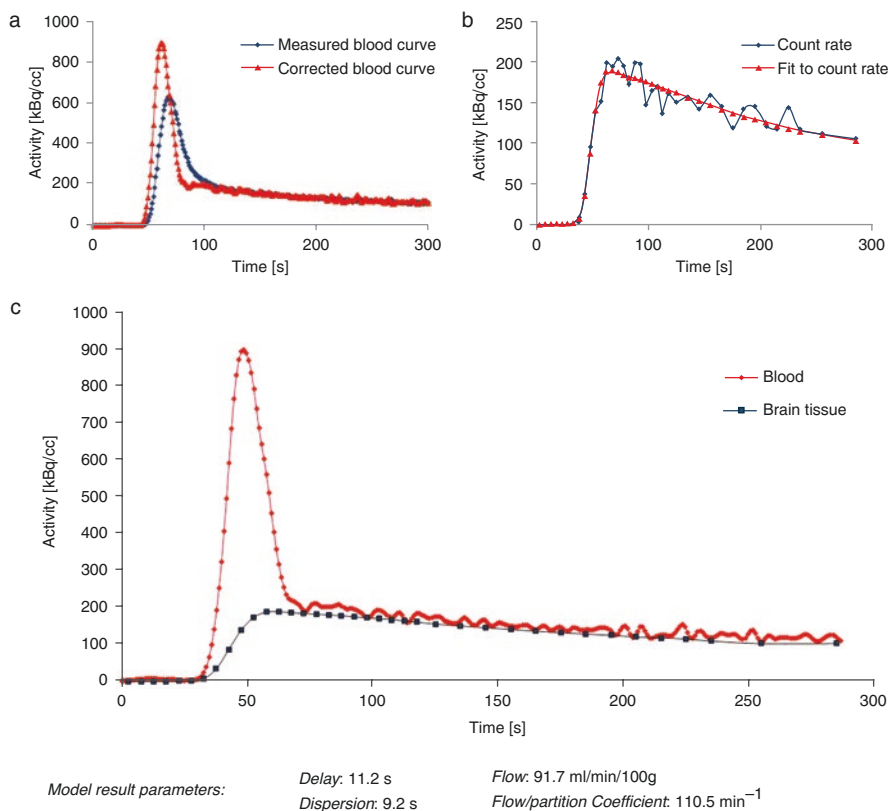
From Eqs. (5.1) and (5.2), the differential equation

$$\frac{dq(t)}{dt} = f_a c_a(t) - kq(t) \quad (5.3)$$

is obtained with the washout constant  $k$  ( $\text{min}^{-1}$ ) defined to be

$$k = \frac{f_a}{V_d}. \quad (5.4)$$

By estimating values of  $f_a$  and  $k$ , the partition coefficient of the tracer can be calculated for every tissue voxel. An example of parameter estimation for a cortical region of interest based on the arterial input function is shown in Fig. 5.2.



**Fig. 5.2** Model parameter estimation. The arterial input function (a) and corresponding tissue response in a cortical ROI (b) after bolus injection of [<sup>15</sup>O]H<sub>2</sub>O are shown. Blood activity data were determined by taking arterial blood samples with a dedicated sampling device. The input function and tissue response are corrected for tracer arrival times and bolus dispersion. Further kinetic modeling is based on the resulting corrected curves (c)

Equation (5.3) has the solution

$$q(t) = f_a e^{-kt} \otimes c_a(t) \quad (5.5)$$

where  $\otimes$  describes the convolution of an exponential function with the arterial input function  $c_a(t)$ :

$$e^{-kt} \otimes c_a(t) = \int_0^t e^{-k\tau} c_a(t-\tau) d\tau \quad (5.6)$$

The experimental design is further specified in a way that the tissue response,  $q(t)$ , as well as the arterial,  $c_a(t)$ , input is measured and thus known for the duration of the experiment. Additionally,  $q(0) = 0$ .

[<sup>15</sup>O]H<sub>2</sub>O is the most commonly used tracer for brain perfusion imaging and CBF quantification with PET. However, there are some notable restrictions concerning the diffusion limitation of [<sup>15</sup>O]H<sub>2</sub>O. A study of Eichling et al. investigated the cerebral behavior of [<sup>15</sup>O]H<sub>2</sub>O after administration to rhesus monkeys. They found that only about 90% of the injected tracer freely exchanges with the brain tissue, with even lower rates at higher flow rates. This incomplete first-pass extraction (80–90% in gray matter structures) resulted in a slightly underestimation of CBF especially in high-flow regions (Eichling et al. 1974; Bolwig and Lassen 1975; Raichle et al. 1983).

In this respect, lipophilic gaseous tracers (like nitrous oxide and xenon) behave superior for CBF measurements as compared with [<sup>15</sup>O]H<sub>2</sub>O.

Further consideration should be done concerning partial volume effect that among others derive from relatively low spatial resolution of typically 6–10 mm in PET. The resulting spread out of signal is a phenomenon that introduces distortion effects in the targeted region and adjacent tissue (Links et al. 1996). The partial volume effect primarily leads to an increased bias for small anatomical structures (like vessels) in the brain (Rousset et al. 1998).

Since the early 1980s, several methods have been described to calculate CBF from PET measurements with [<sup>15</sup>O]H<sub>2</sub>O, including simplified techniques that include parameter fixation, e.g., fixation of the partition coefficient (Watabe et al. 1996). The partition coefficient of [<sup>15</sup>O]H<sub>2</sub>O is a parameter that has been investigated in many studies and varies from 0.77 to 1.05 mL/mL (Herscovitch and Raichle 1985; Iida et al. 1993; Kanno et al. 1991). A good approximation for the whole brain was considered by Herscovitch and Raichle to be 0.9 mL/mL (Herscovitch and Raichle 1985). However, CBF calculation methods with fixed values for  $V_a$  are only applicable to identify global blood flow changes. In presence of regional blood flow deficits, CBF calculation results in incorrect values.

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## 5.5 Role of PET for CBF Measurements

### 5.5.1 General Principles for CBF Measurements

A general distinction must be made between the behaviors of different CBF tracers. One class of techniques utilizes agents which are restricted to the intravascular space and do not interact with the nonvascular space. These can be, in cases of an

intact blood–brain barrier, referred to as nondiffusible CBF tracers. Many brain imaging techniques, such as contrast-enhanced MR and CT, use these intravascular tracers to calculate CBF on the basis of the indicator dilution theory (Meier and Zierler 1954; Zierler 1962). In contrast, a direct tracer exchange from the arterial vascularity to the parenchyma occurs in case of freely diffusible CBF tracers and enables to give a direct measurement of parenchymal blood flow, as originally described by Kety (1951). This concept is utilized, for example, with [ $^{15}\text{O}$ ]H $_2$ O PET, Xenon CT, and Xenon SPECT. Some basic characteristics appear to be necessary for a useful CBF measurement method: Ideally, the incorporated indicator should be early and completely mixed with blood and must stay identifiable for position and concentration in time of image acquisition. Further, the indicator should be metabolically inert and rapidly eliminated.

## 5.5.2 Advantages and Disadvantages of Perfusion Imaging Methods

Various modalities have been developed to obtain hemodynamic parameters in research and clinical settings. These include the older  $^{133}\text{Xe}$  inhalation method, PET, SPECT, X-ray computed tomography methods, and several MRI techniques. However, each technique has its own advantages and drawbacks. It depends on the study subject and the targeted question which method to choose appropriately. A review from Wintermark et al. (2005) gives a comparative overview of current brain perfusion measurement techniques and their clinical relevance. At this point, a brief overview is given on the role of [ $^{15}\text{O}$ ]H $_2$ O PET in this context.

### 5.5.2.1 Nuclear Medicine Methods

CBF can be measured after incorporation of radioactive agents which are detected outside the investigated subject with dedicated scintillation detectors. Based on this principle, tracers like [ $^{99\text{m}}\text{Tc}$ ]HMPAO or [ $^{99\text{m}}\text{Tc}$ ]ECD and  $^{133}\text{Xe}$  were commonly used for CBF measurements with SPECT (Barthel et al. 2001; Lass et al. 1998; Sakai et al. 1987). In comparison with the coincidence method in PET imaging, the detection of single photons is less sensitive. SPECT imaging with [ $^{99\text{m}}\text{Tc}$ ]HMPAO or [ $^{99\text{m}}\text{Tc}$ ]ECD only allows for semiquantitative CBF estimation (Markus 2004). In contrast, the  $^{133}\text{Xe}$  SPECT method relies on the Kety–Schmidt model (Kety and Schmidt 1945) and is considered to give quantitative measures of CBF (Wintermark et al. 2005). In several studies, however, a systematic CBF overestimation was reported in low-flow areas, as well as an underestimation of cortical CBF (Matsuda et al. 1996; Skyhøj Olsen et al. 1981).

However, the coincidence PET technique is commonly accepted to be the reference standard for CBF imaging. With PET, the tissue perfusion can be directly measured by using the diffusible radiotracer [ $^{15}\text{O}$ ]H $_2$ O. This method is well validated and combines several favorable properties. The tracer is easy to produce, and the fast acquisition time permits repetitive measurements with whole brain coverage. Additionally, the major advantage of CBF measurement with PET is the high



accuracy for assessing quantitative parameter maps as well as a high reproducibility (Carroll et al. 2002; Matthew et al. 1993).

Additionally, PET imaging with  $^{15}\text{O}$ -labeled compounds is of special interest for studying cerebrovascular diseases and if a comprehensive view on brain hemodynamic is demanded. In addition to the determination of CBF with  $[^{15}\text{O}]\text{H}_2\text{O}$ , a successive  $^{15}\text{O}[\text{O}_2]$  inhalation allows for quantitative determination of essential parameters of hemodynamics and energy metabolism like oxygen consumption (cerebral metabolic rate of oxygen;  $\text{CMRO}_2$ ) and oxygen extraction fraction (OEF) (Frackowiak et al. 1980; Ibaraki et al. 2004). Further, because of its binding to hemoglobin in red blood cells (Martin et al. 1987),  $[^{11}\text{C}]\text{CO}$  is used as an intravascular tracer to measure the cerebral blood volume (CBV). Compared to other modalities, PET is the only technique which is able to gain all of these different functional parameters noninvasively and in 3D for the entire brain, a fact which allowed PET imaging to become the gold standard method in the field of brain circulation physiology and pathophysiology imaging (Hoeffner 2005).

In comparison with SPECT, which is widely accessible and a routine perfusion imaging tool, PET imaging is technically more demanding, and its availability is limited by complex logistics. Not only a PET scanner but also the constant access to a cyclotron producing the radiopharmaceutical online is required, a fact limiting the application of the method especially in emergency settings. A further restriction occurs in patients who will receive a thrombolytic therapy. Because quantitative PET preferentially requires invasive arterial blood sampling in order to obtain an input function for kinetic CBF modeling, this procedure is not applicable in these patients. As a consequence,  $[^{15}\text{O}]\text{H}_2\text{O}$  PET did not manage to become a clinical routine imaging tool in the acute stroke situation. Clinical applications instead mainly refer to chronic cerebrovascular disorders, brain tumors, and brain activation studies. Moreover, due to the wide acceptance of PET as standard for CBF visualization and quantification, the method is used as the reference to validate other brain perfusion imaging techniques, like perfusion-weighted or arterial spin labeling MRI (Zaro-Weber et al. 2010a, b; Chen et al. 2008).

Apart from  $[^{15}\text{O}]\text{H}_2\text{O}$  PET,  $[^{11}\text{C}]\text{butanol}$  has been suggested to be an alternative tracer for CBF measurements with PET. In comparison with radiolabeled water, butanol has the advantage of being permeable through the blood–brain barrier to 100%. However, this tracer is not used in routine practice because of its complex and radiochemical synthesis (Herscovitch et al. 1987).

$[^{14}\text{C}]\text{iodoantipyrine}$  (IAP) autoradiography is another nuclear medicine perfusion imaging method used in preclinical research (Hatakeyama et al. 1992; Jay et al. 1988). As with butanol and  $\text{H}_2\text{O}$ , IAP is also able to freely cross the blood–brain barrier. It is not metabolized and as such accumulates in the brain tissue depending on the regional CBF. After tracer application, the animals need to be sacrificed to prevent tracer diffusion and to autoradiographically determine the CBF at the time point of tracer injection. Sequential arterial blood samples can be used for absolute CBF quantification. This autoradiographic *ex vivo* method provides accurate and high-resolution quantitative CBF values at a specific time point and is therefore mainly used in small animal studies.

### 5.5.2.2 Computed Tomography Methods

The physical principle that underlies the CT technique is based on tissue-specific attenuation of X-rays that are directed to the body. The image contrast then resulted from variations in attenuation depending on tissue density. Due to similar densities in white and gray matter structures, this technique is not the ideal tool to image anatomical brain structures (Griffiths et al. 2001). However, with a bolus injection of a contrast agent, such as iodine, most prerequisites of the abovementioned indicator dilution theory are satisfied to measure blood flow in the brain. However, due to different acquisition hardware, acquisition protocols, varying post-processing protocols, and differences in the interpretation of perfusion CT data, a reliable CBF quantification remains challenging and varies widely between centers (Kudo et al. 2010). A further approach for CBF measurement uses inhaled Xenon to detect concentration changes of the substance (Pindzola and Yonas 1998). The lipophilic gas is soluble in water, and its X-ray attenuation is similar to that of iodine. As in  $^{133}\text{Xe}$  SPECT, the Xe-CT technique also utilizes the Kety-Schmidt method to calculate quantitative CBF maps with sufficient accuracy (Wintermark et al. 2005). Although newer CT scanners are able to achieve whole brain coverage, a main limitation of commonly used CT scanners derives from the limited anatomical coverage, which is restricted to few brain slices.

### 5.5.2.3 Magnetic Resonance Methods

Several methods for CBF estimation by means of MR have been developed. The most commonly used method for neuroimaging studies is the dynamic susceptibility contrast (DSC) MRI method. It relies on changes in relaxation time on T2\*-weighted images. With the bolus of a paramagnetic contrast agent (e.g., gadolinium-DTPA) passing through the vascular system, a detectable signal loss occurs in T2\*-weighted sequences. Mathematical conclusions were then drawn from the signal reductions to further calculate several perfusion or perfusion-related parameters including mean transit time (MTT), time to peak (TTP), relative cerebral blood volume (rCBV), and relative CBF (Ostergaard et al. 1996a, b). Contrast agents for MR imaging are not radioactive and relatively inexpensive as compared with PET and SPECT tracers. A further advantage of this method is the short acquisition time that enables to visualize perfusion-weighted measurements within a few minutes. However, the absolute quantification of CBF remains unsolved (Wintermark et al. 2005). The use of a local internal input function (Calamante et al. 2004) is necessary to receive parametric maps. As such, the detection of a plausible AIF which is influenced by numerous factors, such as partial volume effects, is important for reproducible and reliable perfusion values (van Osch et al. 2001). Another respective challenge is the localization of the intracranial region of interest for the AIF calculation (Zaro-Weber et al. 2012). Thus, most of the calculated parameter maps in perfusion MR are named as “relative” (Griffiths et al. 2001; Jezard 1998). Nevertheless, perfusion MRI is employed for diagnostic purposes, for instance, in acute stroke and in clinical settings regarding the combination of the various readout parameters. In comparison with gadolinium-based MRI, the arterial spin labeling approach is another promising MR perfusion

technique. Here, magnetically labeled water protons are used as endogenous tracer. However, problems with image interpretation may occur due to a limited signal-to-noise ratio and in the presence of prolonged blood transit times, like in patients with stroke or atherosclerosis (Petersen et al. 2006). In latter cases (e.g., stroke patients), the labeled water spins did not reach the target brain tissue within a given time, with the consequence of underestimating the real blood flow values (Jezzard 1998; Kimura et al. 2005).

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## 5.6 Applications for CBF PET

### 5.6.1 Acute Cerebral Ischemia

The increasing incidence of vascular diseases, like atherosclerosis, is associated with a worldwide increasing number of ischemic attacks (Feigin et al. 2009). Interruptions in brain–blood supply rapidly leads to ischemic cell damage that results in necrotic tissue if no sufficient therapy or spontaneous reperfusion becomes available. Current therapies aim to restore perfusion in the ischemic, salvageable brain tissue. In order to determine this hypoperfused yet viable tissue—the so-called ischemic penumbra (Astrup et al. 1981)—in acute stroke patients, perfusion imaging techniques were developed to identify this “tissue at risk” and to separate it from the already necrotic infarction core. The transition from reversible to irreversible damage is a function of ischemia duration as well as of CBF. A hemodynamic determination of the ischemic penumbra is used for many research studies in laboratory animals and humans, and CBF thresholds to characterize different tissue states were proposed: While normal human CBF is in the range of 50–80 mL/100 g/min, reversible ischemia (“ischemic penumbra”) is evident when CBF drops below values around 22 mL/100 g/min, and neuronal cell death occurs below a CBF of 8 mL/100 g/min (Baron 2001). However, penumbra detection, based on CBF thresholds, is highly dependent on a reliable and accurate quantitative imaging method in an acute stroke diagnostic setting.

The opportunity to investigate different parameters of brain function, like CBF, CBV, OEF, and CMRO<sub>2</sub>, within one PET imaging session is a further advantage in the investigation of cerebrovascular diseases such as ischemic stroke. A first example for the use of the steady-state [<sup>15</sup>O]CO<sub>2</sub> inhalation technique to determine CBF (together with most of the abovementioned parameters) in acute stroke patients was provided by Wise et al. (1983).

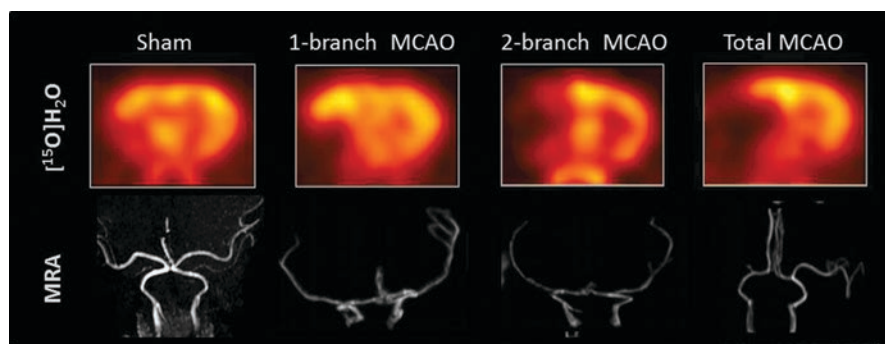
In general, it is essential to reliably separate primary perfusion deficits from events of decreased metabolic demand. This is as CBF decreases may not only appear in the surrounding tissue of the stenotic vessel but sometimes also distant from the obviously damaged part of the brain. This well-recognized phenomenon is called “diaschisis.” One example is the so-called crossed cerebellar diaschisis in which a CBF reduction in the cerebellum contralateral to the stroke-affected brain hemisphere occurs as a result of crossed functional deafferentiation (Baron et al. 1981; Feeney and Baron 1986).

### 5.6.1.1 PET Perfusion Imaging in Preclinical Stroke Research

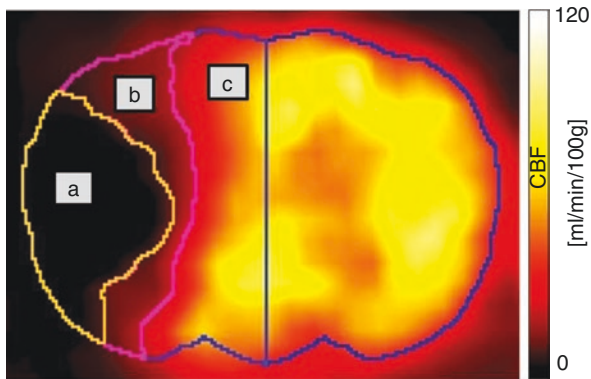
This paragraph will handle the employment of [ $^{15}\text{O}$ ]H $_2\text{O}$  PET in translational research studies of experimental stroke.

In 2008, our group proposed a new large animal model that is applicable for acute and chronic stroke induction (Boltze et al. 2008) and is highly suitable to reflect the human brain pathophysiology. Due to a similar cerebral anatomy and the favorable ovine brain size, brain imaging protocols, scanners, and data analysis techniques as used in clinical routine become feasible. This enables us to perform studies while meeting main conditions for translational research. Like in humans, in sheep, the middle cerebral artery (MCA) usually gives rise to three arterial branches. Different stroke sizes can be induced by permanent transcranial occlusion of one, two, or all three MCA branches, with the latter being referred to as permanent MCA occlusion (pMCAO). Figure 5.3 shows examples of different occlusion types for experimental pMCAO in sheep. One, two, and three branch (total) occlusions of the MCA could clearly be visualized by magnetic resonance angiography (MRA), together with the resulting CBF defects in [ $^{15}\text{O}$ ]H $_2\text{O}$  PET. In addition, slight CBF decreases, probably due to the transcranial surgery, were also detected in the sham-operated animals. Further, it was possible to demonstrate that the ischemic strokes induced by the transcranial pMCAO lead to reproducible CBF deficits, which remain stable over time and eventually leading to necrotic brain tissue.

The major advantage of [ $^{15}\text{O}$ ]H $_2\text{O}$  PET is the option to perform serial scans in a short time period due to the short half-life of the  $^{15}\text{O}$  (122 s). As an example, CBF PET measurements were applied for a controlled preclinical study that aimed to test inhaled nitrous oxide (iNO) for its potential to protect the ischemic tissue in the penumbra in acute ischemic stroke. All animals were subjected to repeated PET scans at 110, 150, 175, and 210 min following pMCAO (a total of four PET scans within 100 min). In the treatment group, 50 ppm iNO were applied from 120 to 180 min after pMCAO. By using kinetic modeling, parametric CBF maps were



**Fig. 5.3** Differential CBF deficits dependent on extent of experimental permanent middle cerebral artery occlusion in sheep. The extent of the post-pMCAO CBF deficit is clearly visualized with [ $^{15}\text{O}$ ]H $_2\text{O}$  PET. Corresponding to angiographic MRI findings, the CBF deficit increases in the order sham >1-branch pMCAO >2-branch pMCAO > total pMCAO. *MCAO* middle cerebral artery occlusion, *MRA* magnetic resonance angiography (Modified from Boltze et al. (2008))

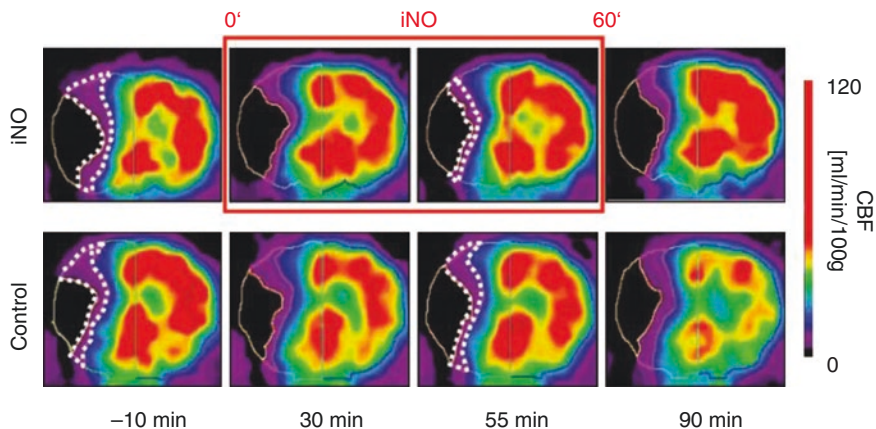


**Fig. 5.4** Parametric CBF map as obtained by  $[^{15}\text{O}]\text{H}_2\text{O}$  PET in the Leipzig permanent middle cerebral artery occlusion sheep stroke model. For volume of interest analysis, the stroke-related regions were defined as follows: infarction core (a),  $<8$  mL/100 g/min; ischemic penumbra, (b) 8–22 mL/100 g/min; normal brain tissue (c),  $>22$  mL/100 g/min

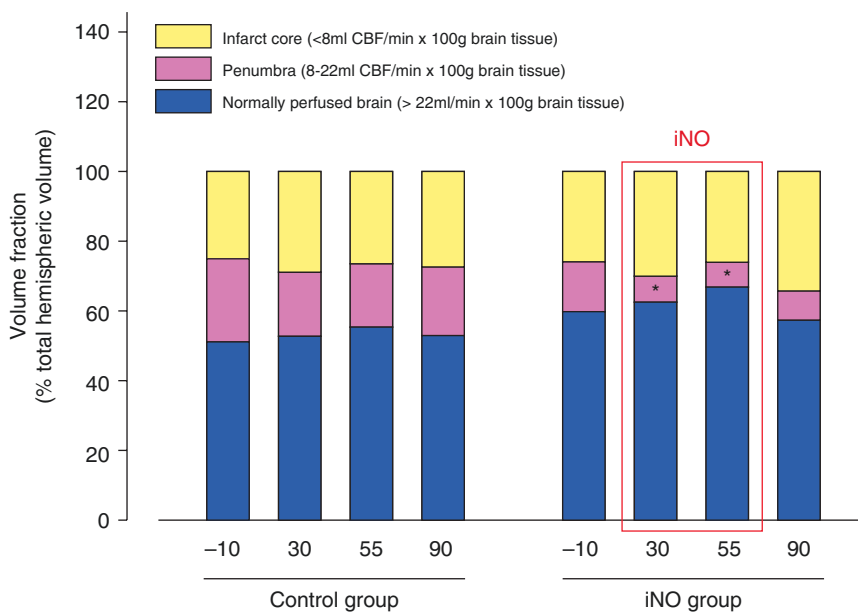
created. Based on the abovementioned commonly accepted CBF thresholds (Baron 2001), operator-independent brain volumes of interest were defined for penumbra, infarction core, and remaining normal brain tissue (Fig. 5.4). Our experiments showed that iNO selectively restores CBF in the ischemic penumbra. While the volume of the necrotic core was not affected, the volume of the penumbra decreased by up to 50% turning into normally perfused tissue ( $>22$  mL/100 g/min) under iNO application but remained unchanged in the untreated control animals ( $p < 0.05$  vs. baseline and vs. control; Figs. 5.5 and 5.6; Terpolilli et al. 2012).

To give an outlook for further projects on CBF PET imaging in the Leipzig sheep pMCAO stroke model, Fig. 5.7 shows first images acquired by a simultaneous PET/3T-MRI system (Biograph mMR, Siemens). Comparative PET and MR imaging studies in acute stroke setting will greatly benefit from the new possibility to acquire data of both modalities simultaneously. This will significantly improve the investigation of the very fast pathophysiological processes in early ischemia. The first experience with this new simultaneous imaging approach, however, triggers great enthusiasm to employ this new technique for further preclinical and clinical research in the acute stroke situation.

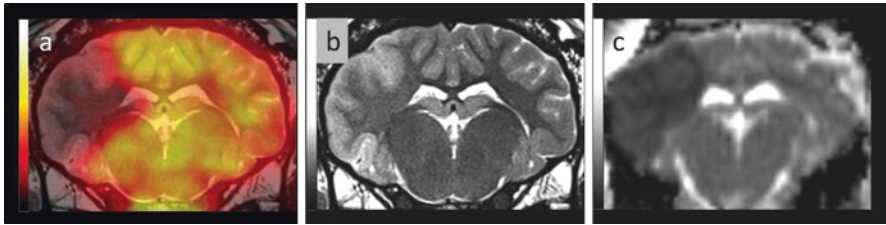
With regard to the abovementioned possibility of PET imaging to provide multi-parameter readouts (CBF, OEF,  $\text{CMRO}_2$ , CBV), in experimental ischemic stroke, so far mainly the brains of monkeys (Kuge et al. 2001; Pappata et al. 1993), pigs (Sakoh et al. 2000a), and felines (Heiss et al. 1994) were investigated by using  $[^{15}\text{O}]\text{H}_2\text{O}$  for CBF measurements. With examinations of different time points after stroke, the studies by Heiss et al. aimed at monitoring important parameters of the temporal transversion of penumbral tissue to the final infarct core. The ischemic penumbra is characterized by an initial increase of OEF and CBV and preserved values of  $\text{CMRO}_2$ . In progress of the infarct and decreasing blood flow supply,  $\text{CMRO}_2$  declines to values of 25% of baseline within the first hour after stroke,



**Fig. 5.5** Serial [<sup>15</sup>O]H<sub>2</sub>O PET scans show therapeutic effect of iNO in the Leipzig permanent middle cerebral artery occlusion sheep stroke model. Stroke-related tissue regions were defined on the basis of quantitative CBF maps. Volumetric analysis showed a significant decrease of the penumbra volume under iNO therapy in favor of the normal brain tissue compartment. This beneficial effect was not detected in control animals (Terpolilli et al. 2012)



**Fig. 5.6** iNO improves penumbral blood flow after permanent middle cerebral artery occlusion in sheep. Quantification of normally perfused, ischemic, and penumbral tissue volumes revealed that penumbral volume decreased significantly during NO inhalation ( $n = 3$  per group;  $*p < 0.05$  vs. control and vs. baseline PET at  $t - 10$  min) in favor of the normally perfused brain tissue (Terpolilli et al. 2012)



**Fig. 5.7** Multimodal PET–MRI imaging in the Leipzig sheep stroke model. Four hours after permanent middle cerebral artery occlusion, (a) CBF was determined by  $[^{15}\text{O}]\text{H}_2\text{O}$  PET (shown here in overlay with anatomical T2-MRI) demonstrating the typical post-stroke deficit. (b) T2-MRI acquired to exclude hemorrhage. (c) Diffusion-weighted MRI with deficit similar to the CBF abnormality. The image data were acquired simultaneously using a PET/3 T-MR (Siemens mMR) system

while the initial increase of OEF becomes less prominent. In the final stage, the lack of CBF is associated with a decrease in OEF and the occurrence of brain tissue necrosis. This progress from still viable to necrotic tissue was defined as a function of duration and severity of stroke (Heiss et al. 1994, 1997; Heiss and Rosner 1983). In further experiments from the Cologne group, postischemic hemodynamic and metabolic processes were investigated after temporary MCAO (from 30 to 120 min of duration) in anesthetized cats (Heiss et al. 1997). CBF PET was performed immediately after MCAO and was repeated at 30-min intervals. After reopening the vessel, a distinct hyperperfusion was found in all animals, depending on the duration of the MCAO. After 30 min of ischemia, the reactive CBF increase was found to be transient with a fast normalization to pre-occlusion CBF values and without major tissue necrosis. By comparison, the reperfusion period after 60 or 120 min of ischemia was associated with severe hyperperfusion (CBF increase up to 300% compared to basal levels) and irreversible tissue damage, depending on the severity of prior ischemia. Approximately 50% of animals died after the prolonged ischemic period and suffered from higher CBF than the surviving animals from the same group. One explanation might be the hyperperfusion processes with massive cerebral edema that resulted in malignant brain swelling. These observations let to assume a relatively high tolerance against moderate hyperemia, whereas extensive hyperperfusion leads to an increased mortality (Heiss et al. 1997).

Other large animal experiments utilized  $[^{15}\text{O}]\text{H}_2\text{O}$  (Sakoh et al. 2000a, b) in combination with  $[^{15}\text{O}]\text{CO}$  PET to study regional correlations of CBF and CBV after acute stroke and in cases of reperfusion (Sakoh et al. 2000a). For that purpose, 13 pigs underwent PET, and the results were compared to MRI measurements of the same parameters. A good agreement was found between the values of both modalities in the ischemic tissue. After MCAO, a significant correlation between CBF reduction and CBV increase was observed. However, the decrease of CBF below 60% of the contralateral side was found to induce a reduction in CBV. In contrast, both parameters were less correlated in cases of reperfusion (Sakoh et al. 2000a).

### 5.6.2 Chronic Cerebral Ischemia

Earliest human work on CBF visualization and quantification by PET in chronic cerebrovascular disease was carried out by Gibbs and coworkers (1984). They successfully imaged, by using the steady-state [ $^{15}\text{O}$ ]CO<sub>2</sub> inhalation technique, the consequences of carotid artery occlusion on brain hemodynamics in their patients (Gibbs et al. 1984). Currently, [ $^{15}\text{O}$ ]H<sub>2</sub>O PET imaging is used as a diagnostic standard tool in specialized centers handling patients with Moyamoya disease (Taki et al. 1988). In this chronic occlusive cerebrovascular disorder of unknown etiology which is most prevalent in East Asia, cerebral angiography and cerebral MRI are often accompanied by testing the hemodynamic consequences of the eminent arterial occlusion on the brain parenchyma by PET (Zhang et al. 2019).

### 5.6.3 Brain Activation Studies

Perfusion parameters are closely coupled with changes in neuronal activity (Roy and Sherrington 1890; Fox et al. 1988; Villringer and Dirnagl 1995). In the last decades, numerous neuronal activation studies have reported this effect, which, nevertheless, is not completely understood (Peterson et al. 2011). The possibility to perform sequential measurements within a certain time window made CBF PET an attractive tool to study perfusion during cognitive, motor, or sensomotor tasks in humans. During local brain activation, cerebral areas that are involved in task performance were identified by CBF changes (Feng et al. 2004; Worsley et al. 1992). It is also worth noting that many tools to analyze brain activation studies, like the Statistical Parametric Mapping (SPM) toolbox (Friston et al. 1991), which are nowadays intensively employed to analyze functional MRI data, were originally developed to process [ $^{15}\text{O}$ ]H<sub>2</sub>O brain PET studies.

### 5.6.4 Other Applications for CBF PET

In general, PET is a useful tool to study perfusion and oxygen metabolism in any cerebral disease, such as dementia (Yao et al. 1990; Cohen et al. 1997), schizophrenia (Ragland et al. 2001), and migraine. Latter approaches addressed hemodynamic changes and oxygen metabolism during and between acute attacks. An increase in CBF was found in cortical areas and in the brainstem which appeared to be pain related (Andersson et al. 1997; Cutrer et al. 2000). Similar to these migraine studies, other preclinical and clinical investigations focus on epilepsy to determine perfusion changes during and between epileptic seizures (Szabo et al. 2007; Kahane 1999; Gaillard et al. 1995). A few studies also used CBF PET (mostly in combination with other PET readouts, like oxygen metabolism from  $^{15}\text{O}$  inhalation) for detailed characterization of brain tumors (Lammertsma et al. 1985; Leenders 1994; Hino et al. 1990; Tomura et al. 1993). Due to the newly formed vessels which often



appear in the tumor parenchyma or its surrounding tissue, a regional CBF increase could be detected and may provide additionally useful information about the progression of the brain tumors.

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## 5.7 Simplification/Improvement of CBF Quantification by [ $^{15}\text{O}$ ]H $_2$ O PET

There are ongoing efforts to simplify and/or improve CBF quantification by [ $^{15}\text{O}$ ]H $_2$ O PET. For instance, ways to replace invasive blood sampling to derive the required arterial input function have been suggested. Here, Jochimsen et al. (2016) and Khalighi et al. (2018) proposed to generate image-derived input functions on hybrid PET/MRI systems. This was realized by the help of arterial vessel segmentation on time of flight MR angiography images perfectly aligned to the PET data, with convincing results as compared to the arterial sampling gold standard. Nevertheless, Koopman et al. (2019) recently compared different non-invasive [ $^{15}\text{O}$ ]H $_2$ O PET CBF quantification methods and pointed out that absolute CBF quantification is limited using these approaches regarding accuracy, while relative—in relation to a reference region—CBF determination (CBF pattern determination as used in most clinical routine applications) is possible with sufficient reliability. Contributing to an improvement of the quality of kinetic modeling of [ $^{15}\text{O}$ ]H $_2$ O PET data, Kudomi et al. (2013) proposed to directly determine the appearance time of the tracer to the brain which is also called the delay time. This parameter which is important for accurate modeling would be determined separately for different brain pixels in an image-based manner.

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## 5.8 Future Alternatives to [ $^{15}\text{O}$ ]H $_2$ O PET Imaging in Determining CBF

Different strategies were tested over the last years to potentially substitute the challenging [ $^{15}\text{O}$ ]H $_2$ O PET approach by more convenient imaging techniques. This search was significantly stimulated by the appearance of the hybrid PET/MRI technology by which it is now possible to cross-evaluate certain MR techniques which were developed to deliver CBF surrogates simultaneously against the gold standard [ $^{15}\text{O}$ ]H $_2$ O PET. Also, the future possibility to obtain CBF information via MRI in parallel to other (for instance, neuropathology or neurotransmitter) PET readouts is seen by many as promising, not only to improve brain research but also different neuropsychiatric diagnoses. Following this line, several versions of the arterial spin labeling MRI technique (van Golen et al. 2014; Schmid et al. 2015; Heijtel et al. 2016), phase contrast mapping MRI (Puig et al. 2018), the classical contrast medium-based perfusion-weighted MRI (Werner et al. 2015), and blood oxygenation level-dependent functional MRI (Fierstra et al. 2018) were cross-evaluated against [ $^{15}\text{O}$ ]H $_2$ O PET over the last years with success for most techniques, at least

for the clinically relevant relative CBF determination. Another technique with potential for delivering CBF surrogate readouts is near-infrared spectroscopy (NIRS). Recently, Polinder-Bos et al. (2018) cross-evaluated NIRS against [ $^{15}\text{O}$ ]H $_2$ O PET. As a drawback, a CBF underestimation was reported by the authors. Thus, more work is required to establish this non-radioactive technology for reliable CBF determination. Finally, there is the concept of substituting CBF imaging with [ $^{15}\text{O}$ ]H $_2$ O PET by blood flow surrogate information as obtained by dynamic data acquisition, early after administration of certain brain PET tracers, with amyloid PET tracers currently being in the main focus. The clinical concept here is that it would be possible to obtain both histopathology (amyloid load) and neurodegeneration (blood flow surrogate) biomarker information in dementia disorders after a single tracer administration. To test this approach, three recent studies head-to-head compared [ $^{15}\text{O}$ ]H $_2$ O PET against different blood flow or blood flow surrogate readouts obtained from amyloid PET imaging (Sojkova et al. 2015; Ottoy et al. 2019; Bilgel et al. 2019). All these studies concluded that the data obtained after amyloid tracer administration is of sufficient quality to obtain clinically relevant blood flow pattern information.

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## 5.9 Summary and Conclusions

PET imaging with [ $^{15}\text{O}$ ]H $_2$ O represents the gold standard to visualize and quantify CBF in vivo. As such, this method plays a relevant role mainly for the investigation of cerebrovascular diseases. Despite advantageous characteristics of the tracer, like the possibility to monitor CBF changes over time in sequential scans, [ $^{15}\text{O}$ ]H $_2$ O PET imaging remains a technological and infrastructural challenge, preventing it from gaining a wider acceptance in a clinical routine setting. In contrast to clinical routine, [ $^{15}\text{O}$ ]H $_2$ O PET plays an important role in preclinical and clinical research, for instance, in (1) clarifying the pathophysiology of ischemic stroke and other cerebrovascular disorders, (2) cross-evaluation of alternative imaging methods to estimate CBF, and (3) the testing of new stroke treatment concepts.

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# The Impact of Genetic Polymorphisms on Neuroreceptor Binding: Results from PET and SPECT Neuroreceptor Imaging Studies

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## Abstract

Major psychiatric disorders are highly heritable. Nevertheless, more than three decades of candidate gene studies and genome-wide association studies have yielded few, if any, unambiguous and replicable results that would be strong enough as to direct research toward new pharmacological targets. This is in part due to the complex non-Mendelian inheritance patterns and difficult-to-define phenotypes of psychiatric disorders. In addition, the relationship between genetic risk and phenotypic expression is blurred by the strong contribution of environmental factors and epigenetic modification. A research strategy that has successfully been pursued over the last years with magnetic resonance-based methods is imaging of endophenotypes. However, these techniques have the drawback that their results are not easily transferable to a molecular level potentially accessible to therapeutic drugs. Neuroreceptor imaging methods such as positron emission tomography (PET) and single photon emission computer tomography (SPECT) make it possible to explore the impact of genetic variation on neuroreceptor binding and function of transporters and other molecules that play a central role in neuropsychiatric disorders, besides certain aspects of molecules that play a central role in our understanding and treatment of psychiatric disorders. This chapter tries to cover the current state of knowledge about the impact of genetic variation on the behavior of PET and SPECT radioligands in the living human brain.

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## 6.1 Introduction

Since more than a century, diagnoses in psychiatry are syndrome based, that is, the diagnostic process relies on the presence (or absence) of a bundle of more or less specific symptoms that are either reported by the patient or observed by the physician. The diagnostic classification is the ground for prognosis and treatment decision. However, diagnosis cannot be related to putative neurochemical alterations thought to be the pathogenic substrate of major psychiatric disorders. The high heritability of disorders such as schizophrenia or affective illness, together with enormous advances in molecular genetics made in the last decades, has incited the hope that molecular genetics might be able to contribute substantially to our knowledge on molecular alterations associated with specific disorders. Moreover, it is anticipated that these methods would help identify biomarkers that would make the diagnostic process more reliable and objective.

Until recently, case-control studies comparing frequencies of putative risk alleles in patient groups and healthy subjects were the predominant research strategy in psychiatric genetics. Candidate genes were selected a priori on grounds of hypotheses on the pathogenesis and available knowledge on genetic variation in these

targets (so-called biological candidate genes). In keeping with the predominant pathogenetic hypotheses, many of the investigated polymorphisms (polymorphisms are frequent genetic variants) were in one or the other way related to brain monoamine neurotransmission. However, it is now increasingly evident that the genetic mechanisms associated with heritability of psychiatric disorders are far more complex than initially thought and that the variance introduced by single genetic variants is rarely strong enough for having relevant influence on a specific disease risk. Results of recent genome-wide association studies (GWAS) rather suggest that the genetic risk for disorders such as schizophrenia is due to a combination of common low-risk and rare high-risk variants (Purcell et al. 2009; Shi et al. 2009; Stefansson et al. 2009). One of the major advantages of the methodology employed in GWAS is that it does not rely on *a priori* hypotheses. Thus, results surviving the extremely rigid statistical thresholds merit further scientific attention. Functional gene variants derived from GWAS have been termed “positional candidate genes.” In contrast to biological candidate genes, the function and biological role of positional candidate genes is frequently unknown and needs to be determined in sometimes complicated follow-up studies. A possible way to relate the function of biological and positional candidate genes to the actual body of knowledge on pathogenesis and neuropharmacology of psychiatric disorders is certainly imaging genetics.

Imaging genetics uses magnetic resonance imaging (MRI)-based methods such as functional MRI (fMRI) or voxel-based morphometry (VBM) and radiation-based methods such as single photon emission computer tomography (SPECT) and positron emission tomography (PET) to study the effects of genetic variants in the living brain. While MRI-based methods can provide information on brain structure, regional brain metabolism, and functional or anatomical connectivity between brain regions, radiation-based methods allow investigating distribution and, in part, function of selected molecules in the brain.

In this chapter, we will review the scientific evidence on the influence of genetic variation upon neuroreceptor PET and SPECT radioligand binding in the brain.

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## 6.2 Serotonin

Serotonin is a phylogenetically ancient modulatory neurotransmitter that is synthesized in the brain and periphery from the essential amino acid tryptophan. The rate-limiting step of serotonin synthesis is hydroxylation of tryptophan by the enzyme tryptophan hydroxylase (TPH). Two isoenzymes are known in humans: TPH-1 catalyzes the reaction in the periphery, whereas TPH-2 (Walther et al. 2003; Zill et al. 2004) is found in the central nervous system (CNS). The serotonin system is involved in the regulation of circadian and seasonal rhythms, mood, sleep, feeding behavior, and energy homeostasis and consequently in a number of psychiatric problems and disorders, notably depression, anxiety, obsessive-compulsive disorder, impulsivity, self-harm, and suicide.

### 6.2.1 The Serotonin Transporter

The presynaptic serotonin transporter (5-HTT) is a key regulator of serotonin signaling and a major target for antidepressant medications and psychostimulants. The 5-HTT takes up serotonin after its release into the extracellular space, thereby controlling the magnitude and duration of serotonergic responses (Bel and Artigas 1992; Shen et al. 2004; Jennings et al. 2006). High presynaptic 5-HTT density is associated with low extracellular serotonin (Jennings et al. 2006), and raising extracellular serotonin by pharmacological blocking 5-HTT is the main mechanism of action of many antidepressant drugs.

### 6.2.2 Serotonin Transporter Gene-Linked Polymorphic Region (5-HTTLPR)

An insertion-deletion polymorphism in the promoter region of the 5-HTT transporter gene, the serotonin transporter-linked polymorphic region (5-HTTLPR; Heils et al. 1996), has been extensively studied in psychiatry. In human cell lines, the 5-HTTLPR long (or *l*-allele) is associated with 30–40% higher 5-HTT expression than the short (or *s*-allele) (Heils et al. 1996; Lesch et al. 1996). Initially, 5-HTTLPR was found to be associated with anxiety-related traits: individuals carrying the short allele had higher neuroticism scores than individuals homozygous for the long allele (Lesch et al. 1996). Several studies were able to find associations between 5-HTTLPR and neuropsychiatric illness (Collier et al. 1996; Kunugi et al. 1997; Malhotra et al. 1998; Rosenthal et al. 1998; Anguelova et al. 2003; Bondy et al. 2006; Grunblatt et al. 2006; Premi et al. 2015) However, there are many studies who failed to replicate earlier results (Seretti et al. 1999; Johansson et al. 2003; Willeit et al. 2003). An intensively discussed finding was the association with the 5-HTTLPR *s*-allele and risk for depression in interaction with traumatic life events (Caspi et al. 2003, 2010). Although this finding has been replicated several times (see, e.g., Zalsman et al. 2006), meta-analyses of this finding are conflicting (Risch et al. 2009; Karg et al. 2011).

In light of inconsistent and contradicting results, it is a rational research strategy to investigate biological endophenotypes associated with the 5-HTTLPR genotype, such as amygdala reactivity (Hariri et al. 2002; Hariri and Holmes 2006) or the influence of 5-HTTLPR on the interaction between brain areas relevant for psychiatric disorders (Pezawas et al. 2005). Today, the *s*-allele is not seen any more as a variant leading to deficient function of brain proteins or brain circuits. Interestingly, *s*-allele carriers were found to outperform subjects carrying the long allele in a number of cognitive and social tasks (for review, see Homberg and Lesch 2010). Hyperactivity in corticolimbic structures associated with the *s*-allele resulting in hypervigilance in the sense of increased sensitivity to both aversive and rewarding stimuli and possibly greater mood reactivity (Willeit et al. 2003) may have beneficial or negative consequences, depending on the environmental circumstances (Homberg and Lesch 2010).

### 6.2.3 Effects of 5-HTTLPR on Serotonin Transporter Binding

First studies in human platelets investigated the effects of 5-HTTLPR on serotonin transporter expression and function. The amount of transporter protein was not found to be altered, but an influence of 5-HTTLPR on maximal serotonin uptake velocity was described in some (Greenberg et al. 1999; Nobile et al. 1999; Patkar et al. 2004) but not all (Willeit et al. 2008) studies. Two of these studies also showed a significant effect of season on functional 5-HTT parameters (Greenberg et al. 1999; Nobile et al. 1999), and one study also found a significant interaction between 5-HTTLPR genotype and seasonal change in serotonin uptake velocity (Greenberg et al. 1999).

Studies on the effects of 5-HTTLPR on 5-HTT availability in the human brain using the radiolabeled cocaine-analog on [<sup>123</sup>I]-2-β-carbomethoxy-3-b-(4-iodophenyl)tropane ([<sup>123</sup>I]β-CIT; Neumeyer et al. 1991) and SPECT (Table 6.1) found higher (Heinz et al. 2000b) or unchanged (Jacobsen et al. 2000; Willeit et al. 2001) binding in carriers of the high-expressing *l*-allele, while other studies found higher binding in subjects homozygous for the low-expressing *s*-allele (Van Dyck et al. 2004). However, rather than being selective for the 5-HTT, [<sup>123</sup>I]β-CIT is a nonspecific monoamine transporter ligand. Moreover, [<sup>123</sup>I]β-CIT binding shows a clear age-dependent decrease (Pirker et al. 2000), and a major limitation of the van Dyck et al. study is that *s/s* homozygous subjects were significantly younger than *ll* and *l/s*-carriers. A more recent study using the β-CIT derivative [<sup>123</sup>I]nor-β-CIT described slightly higher [<sup>123</sup>I]nor-β-CIT binding in the thalamus of *s*-allele carriers (Kauppila et al. 2013).

The effects of 5-HTTLPR or the so-called triallelic 5-HTTLPR – a functional single-nucleotide variant within the 5-HTTLPR *l*-allele termed *LA* and *LG*, with only *LA* being associated with high levels of 5-HTT mRNA, whereas *LG* is more similar to the *s*-allele (Nakamura et al. 2000; Kraft et al. 2005)—have also been studied with two PET ligands that are selective for the 5-HTT, [<sup>11</sup>C]-(+)-6beta-(4-methylthiophenyl)-1,2,3,5,6alpha,10beta-hexahydropyrrolo[2,1-*a*]isoquinoline ([<sup>11</sup>C]McN 5652; Suehiro et al. 1993) and [<sup>11</sup>C]-3-amino-4-(2-dimethylaminomethylphenylthio)benzotrile ([<sup>11</sup>C]DASB; Wilson and Houle 1999; Houle et al. 2000; Wilson et al. 2000). [<sup>11</sup>C]DASB is currently the method of choice for PET imaging of the 5-HTT, and it has been shown to be superior to [<sup>11</sup>C]McN 5652 in several aspects (Huang et al. 2002; Szabo et al. 2002; Frankle et al. 2004).

In contrast to studies using [<sup>11</sup>C]McN 5652 (Shioe et al. 2003; Parsey et al. 2006a), three studies using [<sup>11</sup>C]DASB were able to detect higher serotonin transporter binding in *LA/LA* homozygous subjects (Praschak-Rieder et al. 2007; Reimold et al. 2007; Kalbitzer et al. 2009). The three studies disagree on the location of higher [<sup>11</sup>C]DASB binding in *LA/LA* homozygous subjects in the brain: Praschak-Rieder et al. found significant 5-HTTLPR effects in the putamen, Reimold et al. in the midbrain, and Kalbitzer et al. in the caudate nucleus. What these regions have in common is that they contain large amounts of serotonin transporter protein, and it is inherent to PET methodology that accuracy of measurements is greater in

**Table 6.1** Effects of serotonin transporter-linked polymorphic region (5-HTTLPR), brain-derived neurotrophic factor (BDNF) Val66Met, and 5-HT<sub>2A</sub> receptor polymorphisms on serotonin transporter binding

	Polymorphism	Number of subjects <sup>a</sup>	Radioligand	Results	Region	Comment
Heinz et al. (2000b)	5-HTTLPR	22 (14 AD) <sup>b</sup>	[ <sup>125</sup> I]β-CIT	<i>ll</i> > <i>s</i> -carriers (healthy subjects)	Brainstem	Opposite genotype effects in AD <sup>b</sup>
Willeit et al. (2001)	5-HTTLPR	11	[ <sup>125</sup> I]β-CIT	No effect	Midbrain	
Van Dyck et al. (2004)	5-HTTLPR	96	[ <sup>125</sup> I]β-CIT	<i>ss</i> > <i>ll</i> > <i>ls</i>	Midbrain	<i>s/s</i> Homozygotes younger than other genotypes
Shioe et al. (2003)	5-HTTLPR	27	[ <sup>11</sup> C]MeN 5652	No effect	Midbrain	
Parsey et al. (2006a)	Triallelic 5-HTTLPR	67 (25 MDD) <sup>c</sup>	[ <sup>11</sup> C]MeN 5652	No effect	Serotonin-rich regions	
Praschak-Rieder et al. (2007)	Triallelic 5-HTTLPR	43	[ <sup>11</sup> C]DASB	<i>LalLa</i> > <i>s</i> -carriers	Putamen	Results clearer when covarying for ethnicity and effects of season
Reimold et al. (2007)	Triallelic 5-HTTLPR	19	[ <sup>11</sup> C]DASB	<i>LalLa</i> > <i>s</i> -carriers	Midbrain	
Kalbitzer et al. (2009)	Triallelic 5-HTTLPR	50	[ <sup>11</sup> C]DASB	<i>LalLa</i> > <i>s</i> -carriers	Caudate	
Murthy et al. (2010)	Triallelic 5-HTTLPR	63	[ <sup>11</sup> C]DASB	No effect	Serotonin-rich regions	Only male Caucasians, no effects of daylight
Kaupilla et al. (2013)	5-HTTLPR	30	[ <sup>123</sup> I]nor-β-CIT	<i>s</i> -carriers > <i>ll</i>	Thalamus	No effect in other regions
Premi et al. (2015)	5-HTTLPR	76 <sup>d</sup>	99mTc-ECD	<i>Blood flow ll</i> < <i>s/l</i> < <i>ss</i>	Frontal lobes	Additive effect of CR and <i>ll</i>
Laje et al. (2010)	5-HT <sub>1A</sub>	43	[ <sup>11</sup> C]DASB	rs7333412 AA < AG < GG	Thalamus	rs7333412, rs997012, rs977003
Henningsson et al. (2009)	BDNF Val66Met	25	[ <sup>11</sup> C]MADAM	Val/Val > Met+	Serotonin-rich regions	Finding significant in males only

Klein et al. (2010)	BDNF Val66Met	49	[ <sup>11</sup> C]DASB	No effect	Caudate, putamen, thalamus	Also no effect of plasma BDNF levels
Peciña et al. (2014)	BDNF Val66Met	49	[ <sup>11</sup> C]raclopride	Met > Val/Val	VTA, NaC	Met: no reaction to monetary gains, attenuated dopamine response to analgesic placebo in NAc
Kraus et al. (2014)	BDNF Val66Met	41 (16 MDD) <sup>a</sup> ; 51	[ <sup>11</sup> C]DASB; [ <sup>11</sup> C]WAY 100635	No effect	Serotonin-rich regions	
Fisher et al. (2015)	BDNF Val66Met; 5-HTTLPR	68	[ <sup>11</sup> C]SB207145	Val/Val < Met+; s-carriers < ll	Neocortex, subcortical structures	Effects of 5-HTTLPR in neocortex
Fisher et al. (2017)	BDNF Val66Met; 5-HTTLPR	144	[ <sup>11</sup> C]DASB	Val/Val < Met+; No effect	Subcortical structures	No BDNF Val66Met effect in neocortex; no significant effect of 5-HTTLPR

Adapted from Willeit and Praschak-Rieder (2010)

<sup>a</sup>Healthy subjects if not otherwise indicated

<sup>b</sup>AD Patients with alcohol dependence

<sup>c</sup>MDD Major depressive disorder

<sup>d</sup>FTD Frontotemporal dementia

regions with high density of the target molecule than in those who express the target scarcely (Willeit and Praschak-Rieder 2010). Kalbitzer et al. also showed that light-dependent changes in serotonin transporter binding (Praschak-Rieder et al. 2008; Ruhe et al. 2009) were influenced by 5-HTTLPR genotype, with *LA/LA* carriers showing greater seasonal fluctuation in serotonin transporter binding. However, another recent study did not find any relevant effects of 5-HTTLPR on [<sup>11</sup>C]DASB binding in the human brain (Murthy et al. 2010).

In summary, the literature reports contradictory results on the influence of 5-HTTLPR genotype on serotonin transporter binding in the living human brain. Apart from the natural variation in individual 5-HTT density, one major factor may be the quality of the imaging methods applied. [<sup>11</sup>C]DASB is clearly superior to [<sup>11</sup>C]McN 5652 or [<sup>123</sup>I]β-CIT for imaging the 5-HTT in vivo (Huang et al. 2002; Szabo et al. 2002; Frankle et al. 2004). Moreover, nongenetic aspects have been shown to influence or associate with 5-HTT binding. These include ethnicity (Praschak-Rieder et al. 2007), personality (Kalbitzer et al. 2009), biography (Miller et al. 2009), and psychiatric diagnoses (Malison et al. 1998; Willeit et al. 2000; Meyer et al. 2004; Selvaraj et al. 2009; for review, see Stockmeier 2003; Meyer 2007) and the intensity of environmental light at the time of the year when scanning takes place (Buchert et al. 2006; Praschak-Rieder et al. 2008; Ruhe et al. 2009; Kalbitzer et al. 2010). Studies on the effects of season show that 5-HTT binding is higher in fall and winter as compared to spring and summer, and covarying for season in the Praschak-Rieder et al. sample (Praschak-Rieder et al. 2007) increased the significance of the effects of 5-HTTLPR on [<sup>11</sup>C]DASB binding (Praschak-Rieder et al. 2008). Thus, three of four PET imaging studies using [<sup>11</sup>C]DASB support the notion that 5-HTTLPR is indeed associated with 5-HTT binding in the living human brain. However, in contrast to restricted systems like human cell lines, where a 30–40% difference in 5-HTT binding between high- and low-expressing genotypes has been observed, genotype effects of 5-HTTLPR, if any, were found to be quite small.

#### 6.2.4 Serotonin 2A Receptor Polymorphisms

A relatively large study has investigated the effects of several single-nucleotide polymorphisms (SNPs) in the serotonin 2A (5-HT<sub>2A</sub>) receptor gene on [<sup>11</sup>C]DASB binding in the brain (Laje et al. 2010). Genetic variation in this region had previously been associated with antidepressant treatment response to citalopram (McMahon et al. 2006). An association between 5-HTT binding in thalamus and three SNPs (rs7333412, rs7997012, rs977003) was found, and results were suggestive for a gene-dose effect (rs7333412 AA < AG < GG). This study may thus help to explain differences in treatment response to SSRIs due to genetic variation in the 5-HT<sub>2A</sub> receptor gene. This result is among the few ones in the literature strong enough as to survive correction for multiple testing.



### 6.2.5 Brain-Derived Neurotrophic Factor (BDNF) Polymorphisms (Val66Met)

Brain-derived neurotrophic factor (BDNF) is a relatively well-characterized neurotrophic protein that supports survival, sprouting, and synaptogenesis of neurons. It is encoded on chromosome 11p13–14 (Jones and Reichardt 1990; Maisonpierre et al. 1991). Polymorphisms within the BDNF gene have been shown to influence brain structure and function (Egan et al. 2003; Hariri et al. 2003; Pezawas et al. 2004, 2008) and have been associated with affective disorders, antidepressant treatment response, substance abuse, eating disorders, and schizophrenia (Neves-Pereira et al. 2002; Sklar et al. 2002; Castren 2004; Gratacos et al. 2007; Brunoni et al. 2008). Experimental and clinical evidence suggests that there is a close interaction between BDNF and brain serotonin transmission (Neumeister et al. 2005; Brunoni et al. 2008; Maynard et al. 2016).

Two PET studies have investigated a possible association of the BDNF Val66Met polymorphism and binding to brain 5-HT<sub>2A</sub> receptors (Klein et al. 2010) and to 5-HT<sub>1A</sub> receptors and the serotonin transporter (Henningsson et al. 2009). While there was no effect of the BDNF Val66Met polymorphism on 5-HT<sub>1A</sub> or 5-HT<sub>2A</sub> receptor binding, one study (Henningsson et al. 2009) found significantly higher binding of the serotonin transporter radioligand [<sup>11</sup>C]MADAM in male subjects. This finding was replicated in the same study at a trend level in second cohort using the radioligand [<sup>123</sup>I]β-CIT and SPECT. However, the studies by Klein et al. (2010) and Kraus et al. (2014) using the radioligand [<sup>11</sup>C]DASB did not find any effect of the BDNF Val66Met polymorphism or BDNF plasma levels (Klein et al. (2010)) on brain serotonin transporter binding in larger samples. As part of the same study, Kraus et al. (2014) also performed [<sup>11</sup>C]WAY 100635 PET scans but did not demonstrate effects of BDNF Val66Met on 5-HT<sub>1A</sub> binding either.

Effects on 5-HT<sub>4</sub> receptor binding were reported in a PET study using [<sup>11</sup>C]SB207145 that demonstrated increased BP<sub>ND</sub> for Val66met met carriers compared with Val/Val carriers across regions (Fisher et al. 2015). The group also predicted latent variable model (LVM) for the serotonin transporter-linked polymorphic region (5-HTTLPR) and showed lower neocortical binding for s-carriers relative to LL homozygotes.

The same group further researched effects of 5-HTTLPR and BDNF Val66met on 5-HTT binding ([<sup>11</sup>C]DASB PET) in 144 healthy humans and demonstrated 2–7% higher subcortical 5-HTT binding for met carriers compared with Val/Val individuals of BDNF Val66met on 5-HTT (Fisher et al. 2017). No significant effect was shown for 5-HTTLPR or BDNF Val66met in neocortex (Table 6.1).

### 6.2.6 The Serotonin 1A Receptor

The serotonin 1A (5-HT<sub>1A</sub>) receptor is a G-protein-coupled transmembrane protein whose main signaling effect is a reduction in the formation of cyclic adenosine monophosphate (cAMP). 5-HT<sub>1A</sub> receptors are abundantly expressed in the frontal and

temporal cortex, hippocampus, septum, amygdala, and the raphe nuclei, and lower levels are found in basal ganglia and the thalamus (Ito et al. 1999; de Almeida and Mengod 2008; Storvik et al. 2009). 5-HT<sub>1A</sub> receptors are considered inhibitory as, upon activation, they reduce the probability of neuronal discharge. Cortical and subcortical 5-HT<sub>1A</sub> receptors are mostly postsynaptic receptors located on glutamatergic and GABAergic neurons (Freund et al. 1990; Palchadhuri and Flugge 2005; de Almeida and Mengod 2008), while 5-HT<sub>1A</sub> receptors in the raphe nuclei, the origin of brain serotonin innervation, are somatodendritic autoreceptors that reduce serotonergic output from the raphe into the respective target areas (Sprouse and Aghajanian 1988).

Reductions in brain 5-HT<sub>1A</sub> receptor binding have been described in anxiety disorders (Neumeister et al. 2004; Lanzenberger et al. 2007; for review, see Akimova et al. 2009), depression (Drevets et al. 1999, 2000; Meltzer et al. 2004; Hirvonen et al. 2008, but see Parsey et al. 2006b) (for review, see Drevets et al. 2007; Savitz et al. 2009), and schizophrenia (Tauscher et al. 2002; but see Frankle et al. 2006). Results on the effects of antidepressant treatment with selective serotonin reuptake inhibitors or electroconvulsive therapy on 5-HT<sub>1A</sub> binding are mixed (Moses-Kolko et al. 2007; Spindelegger et al. 2009; Saijo et al. 2010; Lanzenberger et al. 2012a). Increased 5-HT<sub>1A</sub> receptor binding has been observed in anorexia nervosa (Bailer et al. 2005, 2007) and post-traumatic stress disorder (Sullivan et al. 2013).

Several polymorphisms in 5-HT<sub>1A</sub> receptor gene (located on chromosome 5q11.2-q13; Fargin et al. 1989; Albert et al. 1990) have been studied for association with psychiatric disorders (for review, see Drago et al. 2008; Le Francois et al. 2008). The best studied polymorphism is a functional SNP in a palindromic region within the 5-HT<sub>1A</sub> promoter region (C(-1019)G; rs6295; Wu and Comings 1999) that influences gene expression by modulating the interaction of the promoter with transcription factors. The C(-1019)G *C*-allele has been associated with reduced 5-HT<sub>1A</sub> expression (Pernhorst et al. 2013) and was found to be associated with affective disorders in a recent meta-analysis (Kishi et al. 2012).

There is only limited evidence on the *in vivo* influence of the C(-1019)G polymorphism on brain 5-HT<sub>1A</sub> receptor binding. One study did not show any effect of the C(-1019)G polymorphism on brain 5-HT<sub>1A</sub> receptor binding but instead reduced 5-HT<sub>1A</sub> binding in carriers of the 5-HTTLPR short allele (David et al. 2005). Another study (Parsey et al. 2006b, c) showed increased binding of the selective 5-HT<sub>1A</sub> receptor radioligand [<sup>11</sup>C]WAY100635 in carriers of the C(-1019)G *G*-allele. This study also showed higher 5-HT<sub>1A</sub> receptor binding in antidepressant-naïve subjects. *GG*-homozygous subjects were overrepresented in the group of depressed patients in this study; however, the majority of studies find the low-expressing *C*-allele of C(-1019)G to be overrepresented in major depression (Kishi et al. 2012). A recent study in a relatively large collective of healthy subjects (Lanzenberger et al. 2012b) did not show any effect of C(-1019)G on brain [<sup>11</sup>C]WAY100635 binding. Kautzky et al. (2017) also researched the C(-1019)G polymorphism with [<sup>11</sup>C]WAY100635 PET in 19 patients with MDD and 62 healthy volunteers. Receptor binding was calculated for ROIs, and a mixed model was applied that showed an interaction effect of seven ROIs (hippocampus, parahippocampus, medial and lateral

occipitotemporal, inferior frontal occipital and rectal gyrus, and supplementary motor area) and genotype in patients but not healthy volunteers. Further, machine learning was used to classify genotypes but didn't result with significant predictions.

The literature also reports some studies that investigated the influence of the 5-HTTLPR polymorphism on 5-HT<sub>1A</sub> receptor binding in the brain (Table 6.2). Two studies (one of them was performed in a relatively large sample of nonhuman primates) describe lower 5-HT<sub>1A</sub> receptor binding in carriers of the low-expressing 5-HTTLPR *s*-allele (David et al. 2005; Christian et al. 2013). Two further studies describe higher 5-HT<sub>1A</sub> receptor binding in female carriers of the 5-HTTLPR *l*-allele (Lee et al. 2005; Borg et al. 2009), and a third study does not detect any effect of 5-HTTLPR on 5-HT<sub>1A</sub> receptor binding in a relatively large sample of healthy subjects (Borg et al. 2009). Finally, a fourth study (Kautzky et al. 2019) researched epistasis of risk alleles in both C(-1019)G (5-HT<sub>1A</sub> gene) and Val66Met (BDNF gene) via [*carbonyl*-<sup>11</sup>C]WAY-100635 PET imaging and observed an average of 17% increase in 5-HT<sub>1A</sub> receptor binding within risk genotypes. In sum, the evidence concerning the impact of genetic polymorphisms on 5-HT<sub>1A</sub> receptor binding in the brain is, as of yet, inconclusive.

## 6.2.7 The Serotonin 4 Receptor

The 5-HT<sub>4</sub> receptor is abundant in serotonin-rich areas of the brain like the basal ganglia and hippocampus (Waeber et al. 1996; Varnas et al. 2003) and plays a role in anxiety- and depression-like behavior. Studies in mice using autoradiography have shown that modification of 5-HTT gene expression was associated with alterations in 5-HT<sub>4</sub> density: 5-HTT knockout mice showed reduced 5-HT<sub>4</sub> density, and mice overexpressing 5-HTT showed increased 5-HT<sub>4</sub> density (Jennings et al. 2011). 5-HT<sub>4</sub> receptor expression and function seem to be influenced by extracellular serotonin levels, as increases in extracellular serotonin induced by paroxetine given for several weeks are associated with low 5-HT<sub>4</sub> levels (Licht et al. 2009; Vidal et al. 2009).

The PET radioligand [<sup>11</sup>C]SB207145 is a newly developed selective tracer for imaging 5-HT<sub>4</sub> receptors in the human brain (Marner et al. 2009, 2010; Paterson et al. 2010). Since a study in healthy volunteers suggested that acute administration of citalopram does not alter binding of [<sup>11</sup>C]SB207145 (Marner et al. 2010), it was speculated that [<sup>11</sup>C]SB207145 PET could be a useful tool for indirectly measuring chronic, but not acute, changes in extracellular serotonin levels. Fisher et al. thus conducted a study on the effects of 5-HTTLPR on binding of the selective 5-HT<sub>4</sub> radioligand [<sup>11</sup>C]SB207145 in healthy subjects (Fisher et al. 2012). 5-HT<sub>4</sub> receptor binding in the neocortex was 9% lower in *s*-allele carriers compared to *l/l* homozygous subjects, providing further evidence that 5-HTTLPR genotype may influence serotonin transmission in humans. No effect of season or season-by-5-HTTLPR interaction on [<sup>11</sup>C]SB207145 binding was found in this sample. Limitations of this study are that *l/l* subjects were older than *s*-allele carriers and that datasets were collected on two different PET scanners with a significant effect of scanner type.

**Table 6.2** Effects of polymorphisms on serotonin 1A (5-HT1A), serotonin 2A (5-HT2A), and serotonin 4 (5-HT4) receptor binding

	Polymorphism	Number of subjects <sup>a</sup>	Radioligand	Results	Region	Comment
David et al. (2005)	5HT1A C(-1019)G	35	[ <sup>11</sup> C]WAY 100635 (5-HT1A)	No effect	5HT1A receptor-rich regions	Subjects also genotyped for 5-HTTLPR (see below)
Parsey et al. (2006b)	5HT1A C(-1019)G	71 (28 MDD) <sup>b</sup>	[ <sup>11</sup> C]WAY 100635 (5-HT1A)	GG > CG > CC	Raphe nuclei	
Lanzenberger et al. (2012b)	5HT1A C(-1019)G	50	[ <sup>11</sup> C]WAY 100635 (5-HT1A)	No effect	5HT1A receptor-rich regions	
Kautzky et al. (2017)	5HT1A C(-1019)G	81 (19 MDD) <sup>b</sup>	[ <sup>11</sup> C]WAY 100635 (5-HT1A)	BP <sub>ND</sub> GG < C	Dorsal raphe projections	No effect in healthy subjects
Kautzky et al. (2019)	5HT1A C(-1019)G/BDNF Val66Met	46	[ <sup>11</sup> C]WAY 100635 (5-HT1A)	Increased BP <sub>ND</sub> in GG/AA	Cortical surface	
David et al. (2005)	5-HTTLPR	35	[ <sup>11</sup> C]WAY 100635 (5-HT1A)	II > I s/ss	Widespread	Subjects also genotyped for 5-HT1A C(-1019)G (see above)
Lee et al. (2005)	5-HTTLPR	16	[ <sup>11</sup> C]WAY 100635 (5-HT1A)	II < I s/ss	Anterior cingulum	All female study sample
Borg et al. (2009)	5-HTTLPR	54	[ <sup>11</sup> C]WAY 100635 (5-HT1A)	No effect	-	5-HTTLPR effects on cognitive performance
Lothe et al. (2009)	Triallelic 5-HTTLPR	38	[ <sup>18</sup> F]MPPF (5-HT1A)	L <sub>a</sub> carriers < s <sub>s</sub> , females only	Widespread except raphe nuclei	No genotype effect in males ( <i>n</i> = 13)
Fisher et al. (2012)	Triallelic 5-HTTLPR	47	[ <sup>11</sup> C]SB207145 (5-HT4)	II > I s/ss	Neocortex	II Subjects older than I s/ss
Henningsson et al. (2009)	BDNF Val66Met	53	[ <sup>11</sup> C]WAY 100635 (5-HT1A)	No effect	Cortex and pons	
Klein et al. (2010)	BDNF Val66Met	133	[ <sup>18</sup> F]altanserin (5-HT2A)	No effect	Cortex	

Adapted from Willeit and Praschak-Rieder (2010)

<sup>a</sup>Healthy subjects if not otherwise indicated<sup>b</sup>MDD major depressive disorder

## 6.2.8 The Serotonin 2C Receptor

Serotonin binding to the 5-HT<sub>2C</sub> receptor causes inhibition of dopamine release in specific areas of the brain, such as the striatum, prefrontal cortex, nucleus accumbens, and the hippocampus (Alex et al. 2005). The receptor has therefore been implemented in the regulation of mood, anxiety, stress, and pain. An abundant missense SNP in the 5-HT<sub>2C</sub> receptor gene (HTR2C) (rs6318, Cys23Ser) has previously been linked to an *in vitro* activity change and also with mood disorders in humans. A group (Mickey et al. 2012) studied 54 healthy adults using PET and [<sup>11</sup>C]raclopride and detected increased dopamine release in the nucleus accumbens, caudate nucleus, and putamen in Ser23 carriers, after controlling for sex, age, and ethnicity, but there was no connection of Cys23Ser with baseline BPND.

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## 6.3 Dopamine

Due to their preeminent role in addiction and psychotic disorders, brain dopamine systems have been extensively studied with PET and SPECT imaging techniques in psychiatric populations (see Cumming 2009 for in-depth review of dopamine imaging). Dopamine neurons in the brain originate almost exclusively in the brainstem substantia nigra and the more medially located ventral tegmental area. Important projection areas for dopamine neurons are the frontal cortex and subcortical structures, especially the striatum. The best studied dopaminergic region in the brain is the striatum, since it has a high density of dopamine nerve terminals and is thus a region that is particularly well suited for PET and SPECT imaging. Frequently targeted molecules are G-protein-coupled postsynaptic dopamine receptors and the dopamine transporter (DAT).

### 6.3.1 Dopamine D<sub>2/3</sub> Receptors

According to their effects on postsynaptic levels of cyclic adenosine monophosphate (cAMP), dopamine receptors are grouped into two families, the D<sub>1</sub> receptor family (dopamine D<sub>1</sub> and D<sub>5</sub> receptors) and the D<sub>2</sub> receptor family (dopamine D<sub>2</sub>, D<sub>3</sub>, and D<sub>4</sub> receptors). As the main target of antipsychotic medication, dopamine D<sub>2/3</sub> receptors have been of particular interest for imaging studies in schizophrenia. Moreover, D<sub>2/3</sub> receptor imaging techniques have successfully been used to demonstrate alterations in brain dopamine transmission in addictive disorders (Volkow et al. 2009; Urban and Martinez 2012). While current evidence speaks against a relevant alteration in the quantity of D<sub>2/3</sub> receptors that are accessible to PET imaging in schizophrenia (Howes et al. 2012), several studies have shown a reduction of D<sub>2/3</sub> binding in addictive disorders (Volkow et al. 2009; Urban and Martinez 2012).

There are several studies on the influence of genetic variation on D<sub>2/3</sub> receptor binding in the literature (Table 6.4; for review, see Willeit and Praschak-Rieder 2010). To our knowledge, the only replicated finding as of yet is a reduction in

binding of the  $D_{2/3}$  radioligand [ $^{11}\text{C}$ ]raclopride in carriers of the Taq1 A1 allele as compared to A2 homozygotes (Pohjalainen et al. 1998; Jonsson et al. 1999; Joutsa et al. 2014). However, a study of this polymorphism with SPECT and the radioligand [ $^{123}\text{I}$ ]IBZM did not show differences between the genotypic groups (Laruelle et al. 1998). Other polymorphisms in the D2 receptor gene showed in part significant effects but still warrant replication (Table 6.3). A recent study investigated a possible association between polymorphisms in the Period2 (PER2) gene and dopamine  $D_{2/3}$  receptor binding as measured using [ $^{11}\text{C}$ ]raclopride and PET (Shumay et al. 2012a). PER2 is part of the clock-gene family and involved in the reaction of organisms to circadian and circannual changes in lighting conditions and day length. Dopamine transmission shows circadian and seasonal changes in rodents and humans (Castaneda et al. 2004; Tsai et al. 2011; Eisenberg et al. 2010). The authors report on a newly identified VNTR polymorphism in the human PER2 gene that was associated with cocaine craving and striatal dopamine  $D_{2/3}$  receptor binding. This finding is so far very exciting as it brings together several research strings on the physiology of dopamine function and its implication in addiction (see also Praschak-Rieder and Willeit 2012 for further discussion).

The effect of the BDNF Val(66)Met polymorphism, priorly implicated in MDD (see above), on dopamine-mediated responses to stress, cognitive regulation, and anticipatory responses in ventral tegmental area (VTA) and nucleus accumbens (NAc) circuitry were researched in a different study (Peciña et al. 2014) via [ $^{11}\text{C}$ ]raclopride PET scans with or without placebo administration of a potentially analgesic substance. A significant effect of BDNF Val(66)Met was demonstrated on baseline D2/3 receptor availability and pain-stress-induced dopamine release in the NAc and responses to anticipation of monetary losses (Met > Val/Val). On the other hand, met carriers didn't react to monetary gains and presented with attenuated dopamine response to analgesic placebo in NAc. This study provides first human evidence on the effect of BDNF Val(66)Met on dopaminergic responses.

### 6.3.2 The Dopamine Transporter

Another frequently studied target is the dopamine transporter (DAT), a member of the monoamine transporter family that transports dopamine back into the presynaptic neuron immediately after its release into the extracellular space. DAT is the main target of therapeutic drugs, for example, methylphenidate, used to treat attention-deficit hyperactivity disorder (ADHD) or the antidepressant bupropion. DAT is also the main site of action of drugs of abuse such as cocaine or amphetamines. Radiolabeled DAT blockers such as [ $^{11}\text{C}$ ]methylphenidate or [ $^{11}\text{C}$ ]cocaine have successfully been used to quantify DAT availability in the human brain.

Several polymorphisms within the DAT gene or in its promoter region have been identified and studied for association with psychiatric disorders, and there is extensive literature on imaging the effects of some of these polymorphisms on DAT binding in the human brain (Willeit and Praschak-Rieder 2010). The most frequently imaged polymorphism is a 40-bp (base pair) VNTR (variable number of tandem

**Table 6.3** Effects of dopamine D2 receptor polymorphisms on D2 receptor binding, dopamine transporter (DAT) binding, and dopamine synthesis

	Polymorphism	Number of subjects <sup>a</sup>	Radioligand	Results	Region	Comment
Pohjalainen et al. (1998)	Taq1 A	54	[ <sup>11</sup> C]raclopride (2 scans)	A1 carriers < A2/A2	Striatum	No effect on D2 affinity
Lamelle et al. (1998)	Taq1 A, Taq1 B	70 (23 SCZ) <sup>b</sup>	[ <sup>123</sup> I]IBZM	No effect	Striatum	
Jönsson et al. (1999)	Taq1 A Taq1 B -141C Ins/Del	56	[ <sup>11</sup> C]raclopride	A1 carriers < A2/A2 B1 carriers < B2/B2 Del carriers > Ins/Ins	Striatum	
Pohjalainen et al. (1999)	-141C Ins/Del	52	[ <sup>11</sup> C]raclopride	No effect	Striatum	Same sample as in Pohjalainen et al. (1998)
Hirvonen et al. (2004)	C957T	45	[ <sup>11</sup> C]raclopride	T/T > C/T > CC	Striatum	Same sample as in Pohjalainen et al. (1998)
Hirvonen et al. (2009)	C957T	45	[ <sup>11</sup> C]raclopride (2 scans)	Effect on affinity: $K_D$ C/C > C/T > TT No effect on $B_{max}$	Striatum	Same sample as in Pohjalainen et al. (1998)
Bertolino et al. (2010)	D2 rs1076560 (G > T)	37	[ <sup>123</sup> I]IBZM [ <sup>123</sup> I]FP-CIT	D2 binding T < G DAT binding T < G	Striatum	Effect on correlation between D2 binding and prefrontal activity
Laakso et al. (2005)	Taq1 A -141C Ins/Del C957T	33	[ <sup>18</sup> F]FDOPA	Higher uptake in A1 carriers; no effect of -141C Ins/Del, C957T	Striatum	
Laine et al. (2001)	Taq1 A	29 AD <sup>c</sup>	[ <sup>123</sup> I] β-CIT	Higher DAT binding in A1 carriers	Striatum	

(continued)

**Table 6.3** (continued)

	Polymorphism	Number of subjects <sup>a,b</sup>	Radioligand	Results	Region	Comment
Sambataro et al. (2011)	D2 rs1076560 (G > T)	28	[ <sup>123</sup> I]FP-CIT	No effect	Striatum	
Shumay et al. (2012a)	PER2 (Period2)	52	[ <sup>11</sup> C]raclopride	Higher binding in 4R and 3R homozygotes	Striatum	Clock-gene family
Joutsa et al. (2014)	Taq1 A	24 (12 <sup>d</sup> )	[ <sup>11</sup> C]raclopride	A1 carriers < A2/A2	Striatum	Especially for high-reward gambling

Adapted from Willeit and Praschak-Rieder (2010)

<sup>a</sup>Healthy subjects if not otherwise indicated

<sup>b</sup>Schizophrenia

<sup>c</sup>Alcohol dependence

<sup>d</sup>Pathologic gamblers



repeats) polymorphism in the 3' untranslated region of exon 15 of the DAT gene SLC6A3 (Vandenberg et al. 1992, 2000). These studies included healthy subjects as well as patients with alcohol dependence, schizophrenia, ADHD, and Parkinson's disease (Heinz et al. 2000a; Jacobsen et al. 2000; Martinez et al. 2001; Lynch et al. 2003; Cheon et al. 2005; van Dyck et al. 2005; Krause et al. 2006; Lafuente et al. 2007; van de Giessen et al. 2009; Kasparbauer et al. 2015; Table 6.4). A recent methodologically sound meta-analysis (Costa et al. 2011) provides no evidence to support the hypothesis that the DAT 40-bp VNTR is significantly associated with interindividual differences in DAT availability in the striatum. A SPECT study investigating a cluster of polymorphisms flanking a recently described splice variant in the DAT gene (Talkowski et al. 2010) did not show significant differences in [<sup>123</sup>I]-β-CIT binding (van de Giessen et al. 2012). Another group (Pak et al. 2018) investigated impacts of 19 different SNPs previously implicated in Parkinson's disease on DAT and SERT availability with <sup>123</sup>I-FP-CIT SPECT. None of the tested SNPs impacted DAT availability but SERT availability in midbrain and pons ( $p = 0.0007$ ) of healthy males was significantly influenced by rs591323 in Fibroblast Growth Factor 20 on chromosome 8. In summary, the results of these studies give no clear picture on the effects of the DAT 40-bp VNTR polymorphism. However, the number of studies is relatively small, and there is significant heterogeneity between the studies.

### 6.3.3 Measuring Dopamine Transporter Function

Radiolabeled dopamine precursors able to cross the blood-brain barrier (e.g., [<sup>18</sup>F]dihydroxyphenylalanine, [<sup>18</sup>F]DOPA) are converted into the DAT substrate [<sup>18</sup>F]dopamine in dopaminergic neurons. The striatal [<sup>18</sup>F]DOPA influx constant  $K_i$  reflects presynaptic dopamine uptake, synthesis, and storage capacity and is regarded as a good proxy for presynaptic dopamine metabolism in the brain (Cumming 2009). Together with so-called “competition” paradigms, a research technique that allows for measuring the inhibition of radioligand binding at  $D_{2/3}$  receptors by endogenous dopamine, [<sup>18</sup>F]DOPA studies have now clearly shown that presynaptic dopamine synthesis capacity (Reith et al. 1994; Hietala et al. 1995, 1999; Lindstrom et al. 1999; Meyer-Lindenberg et al. 2002; McGowan et al. 2004; Kumakura et al. 2007; Bose et al. 2008; Nozaki et al. 2009), baseline synaptic dopamine levels (Abi-Dargham et al. 2000, 2009; Kegeles et al. 2010; Weidenauer et al. 2020), and pharmacologically induced dopamine release (Laruelle et al. 1996; Breier et al. 1997; Abi-Dargham et al. 1998, 2009) are increased in schizophrenia (for review, see Howes et al. 2012).

Imaging measures of the hyper-dopaminergic state in schizophrenia can as well be conceptualized as an endophenotype of the disorder. Indeed, there are a number of studies on the genetic background of in vivo imaging measures of presynaptic

**Table 6.4** Effects of polymorphisms within the dopamine transporter gene SLC6A3 on in vivo dopamine transporter binding in humans

	Polymorphism	Number of subjects <sup>a</sup>	Radioligand	Results	Region	Comment
Heinz et al. (2000a)	DAT 3' VNTR	25 (14 AD) <sup>b</sup>	[ <sup>123</sup> I]-β-CIT	9-repeat carriers <10/10	Putamen	Alcohol not associated with DAT binding genotype
Jacobsen et al. (2000)	DAT 3' VNTR	44 (14 CA) <sup>c</sup>	[ <sup>123</sup> I]-β-CIT	9-repeat carriers >10/10	Striatum	5-HTTLPR not associated with [ <sup>123</sup> I]-β-CIT binding in midbrain
Martinez et al. (2001)	DAT 3' VNTR	43 (22 SCZ) <sup>d</sup>	[ <sup>123</sup> I]-β-CIT	No effect	Striatum	No genotype effect on amphetamine-induced dopamine release
Lynch et al. (2003)	DAT 3' VNTR	166 (100 PD) <sup>e</sup>	[ <sup>99m</sup> Tc]TRODAT-1	No effect	Striatum	Also no effect of COMT and MAO-B polymorphisms
Van Dyck et al. (2005)	DAT 3' VNTR	96	[ <sup>123</sup> I]-β-CIT	9-repeat carriers >10/10	Striatum	
Cheon et al. (2005)	DAT 3' VNTR	11 children with ADHD <sup>f</sup>	[ <sup>123</sup> I] IPT	9-repeat carriers <10/10	Basal ganglia	Response to methylphenidate better in 9-repeat-allele carriers
Krause et al. (2006)	DAT 3' VNTR	22 ADHD <sup>g</sup>	[ <sup>99m</sup> Tc]TRODAT	No effect	Striatum	
Lafuente et al. (2007)	DAT 3' VNTR	15 SCZ <sup>d</sup>	[ <sup>123</sup> I] FP-CIT	No effect	Striatum	
van de Giessen et al. (2009)	DAT 3' VNTR	79	[ <sup>123</sup> I]-β-CIT	9-repeat carriers >10/10	Striatum	Haplotype 5' T-A 3' 9R highest DAT binding
	DAT 5' VNTR					

Drgon et al. (2006)	DAT 5' VNTR (rs2652511, rs2937639)	15 (6 ADHD) <sup>f</sup>	[ <sup>11</sup> C]-cocaine PET	C-G > T-A	Ventral striatum	5' C-G combination confirmed postmortem
Szobot et al. (2010)	DAT 3' VNTR DRD4 48-bp VNTR	17 ADHD <sup>f</sup>	[ <sup>99m</sup> Tc]TRODAT	Reduced occupancy in DAT10R homozygotes/ DRD4-7R carriers	Striatum	Subjects scanned before and after 3 weeks of methylphenidate treatment
Kasparbauer et al. (2015)	DAT 3' VNTR	35	[ <sup>123</sup> I] FP-CIT	9-repeat carriers >10/10	Striatum	Methylphenidate induced increase of blood oxygen for 9R carriers

Adapted from Willeit and Praschak-Rieder (2010)

<sup>a</sup>Healthy subjects if not otherwise indicated

<sup>b</sup>AD Alcohol dependence

<sup>c</sup>CA Cocaine abuse

<sup>d</sup>Schizophrenia

<sup>e</sup>Parkinson's disease

<sup>f</sup>Attention-deficit hyperactivity disorder

dopamine function (Table 6.5). These studies investigated the influence of polymorphisms in the DAT gene (Martinez et al. 2001; Brody et al. 2006), in  $D_2$  and  $D_3$  receptor-related genes (Brody et al. 2006; Peciña et al. 2012; Savitz et al. 2013), in the gene coding for oxytocin (Love et al. 2012), in the  $\mu$ -opioid receptor gene (Domino et al. 2010), leptin gene (Burghardt et al. 2012), and AKT1 gene (Shumay et al. 2017) on DAT function. In these studies, dopamine release was induced either by administration of d-amphetamine, by nicotine from cigarettes or by behavioral stimuli such as stress and pain. All of the abovementioned studies describe an association with reductions in  $D_{2/3}$  radioligand binding (as an indirect measure for dopamine release) and the respective candidate genes. Although there emerges no consistent picture due to the variety of polymorphisms and the different methods used, all of these studies were able to show that results of “competition studies” are in part driven by genetic variation. Some of the information collected in these studies might thus be used for refining future studies in patients with schizophrenia. Perhaps even more important, these studies were able to show that several neurochemical pathways are involved in regulating dopamine release in the living human brain. Regulatory mechanisms include feedback regulation in dopamine terminal fields (see, e.g., Savitz et al. 2013, brain opioid systems (Domino et al. 2010), and the neuropeptide oxytocin (Love et al. 2012)).

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## 6.4 Other Polymorphisms

### 6.4.1 The Catechol-O-methyltransferase Val158Met Polymorphism

Catechol-O-methyltransferase (COMT) is an enzyme that catalyzes degradation of catecholamines (including epinephrine, norepinephrine, and dopamine) in the periphery and the brain. While regulation of extracellular dopamine levels is heavily dependent on DAT function in subcortical brain areas, COMT is believed to be a key mechanism regulating dopamine activity in cortical, especially frontal cortical brain areas (Ciliax et al. 1999; Lewis et al. 2001; Matsumoto et al. 2003).

Several SNPs have been described in the COMT gene, but without doubt, the SNP causing a valine-to-methionine substitution at position 158 (COMT Val158Met) is one of the most studied polymorphisms in psychiatry and a series of other disorders. The COMT 158Val variant degrades dopamine more efficiently than the COMT 158Met variant (Lachman et al. 1996), and the polymorphism has been associated with several psychiatric and other disorders. A finding that seems to hold true is a difference in executive function mediated mainly by prefrontal cortex dopamine between carriers of the gene variants. Thus, several studies have investigated behavioral, cognitive, functional, and anatomical variance associated with this gene as an endophenotype in various disorders (see, e.g., Schosser et al. 2011; Witte and Floel 2011; Ira et al. 2013).

Two studies investigated the effects of COMT Val158Met using [ $^{18}$ F]DOPA and PET. A study (Meyer-Lindenberg et al. 2005) in healthy volunteers found increased

**Table 6.5** Effects of genetic polymorphisms on dopamine-induced reductions in  $D_{25}$  receptor binding

	Polymorphism	Number of subjects <sup>a</sup>	Outcome measure	Results	Region	Comment
Martinez et al. (2001)	DAT 3' VNTR	59 (28 SCZ) <sup>b</sup>	d-Amphetamine-induced decrease in [ <sup>125</sup> I]BZM binding	No effect	Striatum	Greater reductions in patients; trend for greater reductions in 9-repeat carriers
Brody et al. (2006)	DAT 3' VNTR, D2 Taq A1/A, D4 VNTR, COMT Val158Met	45	Nicotine-induced decrease in [ <sup>11</sup> C]raclopride binding	DAT: 9R > 10/10R D2: no effect D4: greater in <7R Val/Val > Met	Striatum	
Peciña et al. (2012)	D2 rs4274224 (A > G)	52	Stress-induced decrease in [ <sup>11</sup> C] raclopride binding	AG > AA/GG	Caudate, putamen, ventral striatum	
Savitz et al. (2013)	D3 Ser9Gly	36 (10 MDD) <sup>c</sup>	Reward-induced decrease in [ <sup>11</sup> C] raclopride binding	Gly > Ser	Caudate, ventral striatum	No genotype effect on baseline $D_{25}$ binding
Love et al. (2012)	Oxytocin Rs4813625	55	Pain-stress-induced decrease in [ <sup>11</sup> C] raclopride binding	GC/CC > GG	Ventromedial caudate	Significant effects in females ( $n = 32$ ) but not males
Domino et al. (2012)	$\mu$ -Opioid 1 receptor A118G	19 males	Nicotine-induced decrease in [ <sup>11</sup> C]carfentanyl binding	AG/GG > AA	Caudate, ventral putamen	Significant effect of the polymorphism also on plasma cortisol levels
Burghardt et al. (2012)	Leptin rs12706832	50	Stress/pain -induced [ <sup>11</sup> C] raclopride binding	GG > AG/AA	Caudate, putamen, ventral striatum, Nucl. accumbens	Plasma leptin levels also correlated to challenged dopamine release.
Shumay et al. (2017)	AKT1VNTR	91; 54	Baseline DRD2 availability; methylphenidate-induced dopamine release	LL > HL > HH	Caudate, Putamen	Only in ventral striatum, highest in HH > HI > LL

<sup>a</sup>Healthy subjects if not otherwise indicated<sup>b</sup>SCZ schizophrenia<sup>c</sup>MDD major depressive disorder

striatal [ $^{18}\text{F}$ ]DOPA uptake in 158Val carriers when compared to 158Met homozygotes. Another study in patients with early Parkinson's disease (Wu et al. 2012) describes higher [ $^{18}\text{F}$ ]DOPA uptake measured between 150 and 210 min after radioligand injection in 158Met homozygous patients. Besides differences between study samples (young healthy volunteers versus patients with Parkinson's disease), the difference might be due to the fact that higher [ $^{18}\text{F}$ ]DOPA in 158VMet homozygous patients with Parkinson's disease was measured from 150 min after tracer injection onward. At this time, the [ $^{18}\text{F}$ ]DOPA signal is believed to reflect rates of dopamine metabolism to 3,4-dihydroxyphenylacetic acid and homovanillic acid (Ruottinen et al. 2001; Ceravolo et al. 2002).

A study in healthy volunteers investigated the effects of COMT Val158Met on dopamine  $D_{2/3}$  receptor binding using PET and the radioligands [ $^{11}\text{C}$ ]raclopride and [ $^{11}\text{C}$ ]FLB457 for assessing striatal and cortical  $D_{2/3}$  binding, respectively. Since genotype had no effect on binding of either radioligand, the authors concluded that baseline in vivo  $D_{2/3}$  receptor availability does not differ between COMT Val158Met genotypic groups. In contrast, another study using the dopamine  $D_1$  receptor radioligand [ $^{11}\text{C}$ ]NNC112 (Slifstein et al. 2008) found significantly higher cortical  $D_1$  receptor binding in Val158 homozygotes, while no difference was found in striatum. This finding is in good agreement with the functional role of COMT Val158Met, as low dopamine levels have been found to cause upregulation of dopamine  $D_1$  receptors in the rodent brain, while dopamine transmission in the basal ganglia is less affected due to the prominent role of DAT in these brain areas (Gogos et al. 1998; Guo et al. 2003). Hernaus et al. investigated the effect of COMT Val158Met on prefrontal [ $^{18}\text{F}$ ]fallypride displacement in PET and subjective stress response to a psychosocial stress challenge in a mixed sample of healthy volunteers and healthy first-degree relatives of patients with psychotic disorders (Hernaus et al. 2013). They showed less [ $^{18}\text{F}$ ]fallypride displacement in Val/Met and Met/Met than Val/Val and Met-allele carriers also experienced larger subjective stress responses than Val-homozygotes. Separated analysis for controls/ first-degree relatives didn't change the direction of these effects. A different group performed [ $^{11}\text{C}$ ]FLB 457 PET in healthy controls to investigate dopamine release in the PFC and were neither able to show a relationship between COMT genotype and dopamine release nor COMT genotype and baseline  $D_{2/3}$  receptor BPND (Narendran et al. 2016). A study investigating the effects of the COMT Val158Met polymorphism on brain 5-HT $_{1A}$  receptor binding found significantly higher [ $^{11}\text{C}$ ]WAY100635 binding in Val158 homozygotes in orbitofrontal and cingulate cortex, insula, and amygdala (Baldinger et al. 2014). Peak differences were found in the posterior cingulate cortex.

## 6.4.2 Monoamine Oxidase A

Monoamine oxidase is an enzyme that catalyzes the oxidative degradation of biogenic amines. There are two isoforms, MAO-A and MAO-B. MAO-A has high affinity for serotonin, norepinephrine, and epinephrine. MAO-B mainly degrades phenethylamine, a so-called trace amine. Dopamine is a substrate of both

isoenzymes (Kalgutkar et al. 2001). The gene for MAO-A is located on the X chromosome (Xp11.23–11.4). A rare loss-of-function mutation of the gene is associated with behavioral features such as impulsive aggression, hypersexuality, and reduced intelligence (Brunner et al. 1993). A polymorphism in the promoter region of the MAO-A gene affecting transcriptional activity (Sabol et al. 1998) has been linked to aggressive or antisocial behavior in subjects exposed to childhood maltreatment (Caspi et al. 2002; Fergusson et al. 2011) or brain injury (Pardini et al. 2011). The polymorphism consists of a 30-bp repeat sequence present in three to five copies and is termed MAO-A *high* and *low* according to the amount of MAO-A produced in cell lines. As the polymorphism is located on the X chromosome, males are hemizygous for either the MAO-A *high* or *low* allele.

Radioligands for quantifying MAO-A activity in the brain are modified MAO-A inhibitors such as [<sup>11</sup>C]clorgyline (MacGregor et al. 1985) or [<sup>11</sup>C]harmine (Bergstrom et al. 1997; Ginovart et al. 2006). Both radioligands have successfully been used to study the role of MAO-A in aggression (Alia-Klein et al. 2008) or affective disorders (Meyer et al. 2006; Sacher et al. 2010).

To our knowledge, there are no studies published on the effects of genetic heterogeneity on [<sup>11</sup>C]harmine binding in the human brain. A study investigating the MAO-A *high/low* polymorphism in 38 healthy subjects using [<sup>11</sup>C]clorgyline (Fowler et al. 2007) did not find any relationship between genotype and the brain [<sup>11</sup>C]clorgyline binding. Males are hemizygous for this polymorphism (the MAO-A gene is located on the X chromosome); the effects of this polymorphism, if functional in vivo, should therefore be clearest in a male-only sample. A possible explanation for the negative finding is epigenetic regulation: methylation of the MAO-A promoter has indeed been shown to have a clear influence on brain [<sup>11</sup>C]clorgyline distribution volumes (Shumay et al. 2012b). Another study investigating the effects of this polymorphism on binding of the 5-HT<sub>1A</sub> receptor radioligand [<sup>11</sup>C]WAY 100635 (Mickey et al. 2008) found higher binding in females carrying the MAO-A *high* allele. No genotype effect was found in males.

### 6.4.3 18-kD Translocator Protein

Immune cells express an 18-kD translocator protein (TSPO) formerly known as the peripheral benzodiazepine receptor (Anholt et al. 1986). Expression of TSPO has been studied as a marker of microglia activation and inflammation in various disorders, including Alzheimer's disease, multiple sclerosis, frontotemporal dementia, and schizophrenia (Debruyne et al. 2002; Cagnin et al. 2004; Edison et al. 2008; Kreisl et al. 2012) using a variety of TSPO radioligands. The TSPO radioligand [<sup>11</sup>C]PBR28 was observed to bind to TSPO with high affinity in some subjects, with low affinity in others and with intermediate affinity in a third group (Fujita et al. 2008; Kreisl et al. 2009; Owen et al. 2010, 2011), an observation that has been related to a polymorphism (rs6971) affecting binding of the radioligand PBR28 to TSPO in human platelets (Owen et al. 2011).

A study using the TSPO radioligand [ $^{18}\text{F}$ ]FEPPA (Wilson et al. 2008) has shown clear effects of the rs6971 polymorphism in the human brain (Mizrahi et al. 2012). Another study using the radioligand [ $^{11}\text{C}$ ]PBR28 confirmed these effects and showed increased [ $^{11}\text{C}$ ]PBR28 binding in patients with schizophrenia when accounting for genotype effects (Kreisl et al. 2012). This study showed that data on TSPO binding need to be stratified by genotype in order to get information on TSPO expression in various disorders.

#### **6.4.4 Translocase of Outer Mitochondrial Membrane 40 (TOMM40)**

The encoded protein of the translocase of outer mitochondrial membrane 40 gene (TOMM40) is a channel-forming subunit of the translocase that is essential for import of protein precursors to mitochondria. Different splice variants of the gene have been identified and linked to Alzheimer's disease because of its linkage disequilibrium with the apolipoprotein E gene and its neurotoxic potential (Roses 2010). Morphology changes of the medial temporal lobe manifest early within the disease progression of Alzheimer's disease. A group (Siddarth et al. 2018) therefore researched the effects of the two genes via regional brain PET with 2-(1-{6-[(2 [F18]fluoroethyl) (methyl) amino]-2-naphthyl} ethylidene)malononitrile (FDDNP) in the medial temporal lobe. FDDNP binding differed significantly between groups and TOMM40 S/S exhibited significantly lower binding compared to TOMM40 S/VL and APOE-4 carriers.

#### **6.4.5 AKT1**

AKT1 is a protein kinase, also referred to as protein kinase B (PKB), that can be found in different cell types of the human body and is involved in various signaling pathways, such as cell apoptosis, proliferation, differentiation, and metabolism. The AKT1 gene is located on chromosome 14 (14q32.33), and somatic mutations have been implemented for different types of cancer. Further, several polymorphisms in the AKT1 gene have been linked to schizophrenia and dopamine neurotransmission, which might be due to the function of AKT1 in G-protein-independent DA receptor signaling, particularly internalization and desensitization of the receptor (Li et al. 2016). The AKT1 activity is further inversely regulated by extracellular DA content, and binding to the DRD2 induces phosphorylation of AKT1 (Zheng et al. 2013; Jönsson et al. 1999).

A fairly recently discovered variant number of tandem repeat (VNTR) polymorphism in AKT1, intron3 [major alleles: L- (eight repeats), H- (nine repeats)], has lately been tied to striatal dopamine D2/3 receptor availability and dopamine release in a healthy cohort. The high CG content of the gene (amenability to DNA methylation) and the location of the considered polymorphism within the nucleosome exclusion region highlight its susceptibility to epigenetic modification.



Shumay et al. used [<sup>11</sup>C]raclopride and PET to determine baseline D2R availability and methylphenidate-induced dopamine release. LL phenotypes showed higher D2R availability than HH in caudate and putamen but not ventral striatum. Dopamine increases further were stronger in the ventral striatum in HH phenotypes, but no increases were observed in caudate or putamen after methylphenidate administration (Shumay et al. 2017). These results suggest a modulating role of AKT1 in striatal dopamine signaling but lack conclusions for schizophrenia or acute (drug-induced) psychoses.

The finding that AK1<sup>-/-</sup> mice present with an amplified reaction to amphetamine and ketamine (Emamian et al. 2004; Featherstone 2013) leads to the belief that a reduced AKT function might cause a supersensitive DA state. Therefore, the HH allele of the gene, for which increased DA release in VS was demonstrated, might serve as a marker for schizophrenia and other psychotic disorders.

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## 6.5 Summary and Outlook

This review shows that combining genetic research with neuroreceptor imaging using PET and SPECT has produced a large array of findings suggesting effects of genetic variants on protein expression in the living brain. Some genetic effects, for example, those of polymorphisms in the TSPO gene, seem strong enough as to produce plausible results with large effect sizes in PET imaging studies. Other results, such as those on the effects of 5-HTTLPR on 5-HTT binding or Taq1 A on dopamine D<sub>2/3</sub> receptor binding in the brain, show less clear-cut effects, and despite having been replicated independently, they are contrasted by studies with differing or negative results. Many studies in the field either are not replicated at all or are replicated with slightly different imaging methods or in differing study collectives, thus creating a “penumbra” of a 0.05 significance level (Kapur et al. 2012) around the original findings without being able to answer the question whether a certain genetic variant does really have an effect on protein binding in the brain.

The way genetic research in psychiatry is conducted has changed dramatically in the last years. Typical studies at the beginning of this research had sample sizes in the low three-digit number range. The picture that emerged from these studies was quite similar to what we see in PET and SPECT imaging genetics today: many significant findings, few replications, and no definite answers on candidate gene loci. Researchers have thus started to combine their efforts and to implement logistics for large international consortia that are able to collect data from tens or hundreds of thousands of subjects (Sullivan 2010). Similar efforts have been made in the field of MRI-based imaging genetics (the IMAGEN consortium started in 2007 is a good example; <http://www.imagen-europe.com>).

The single most important limitation of all studies reviewed here is a lack of power. Sample sizes in PET and SPECT imaging genetic studies are maybe even a bit larger than those in other PET and SPECT studies, most probably because genotyping is sometimes used to extract some additional scientific output from already existing projects sharing a radioligand (the mean sample size in the here reported

studies is 44 subjects, the range 11–166). Still, due to the relatively small effect of most genetic variants and the large number of other factors influencing availability of target receptors, much larger samples would be needed to have adequately powered studies in this field. Neuroreceptor PET is an expensive and logistically difficult research technology available in a limited number of research centers around the world. It is time for the field to follow the example of other research areas and to start combining efforts and cooperate. This would benefit not only imaging genetics but also all other research topics addressed with PET and SPECT imaging techniques.

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**Part II**  
**Systems**



# PET Imaging of Acetylcholinesterase

# 7

Hitoshi Shinotoh, Shigeki Hirano, and Hitoshi Shimada

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## Abstract

Acetylcholinesterase (AChE) is an enzyme that hydrolyzes acetylcholine and thereby regulates the concentration of the transmitter at the cholinergic synapse. AChE is frequently described as a marker of cholinergic neurons in the brain. Several cholinesterase inhibitors have been labeled with carbon-11 as radiotracers for visualizing AChE-binding sites. Among these, carbon-11-labeled methyl-4-piperidyl acetate (MP4A) and methyl-4-piperidyl propionate (MP4P or PMP) are successfully in clinical use for in vivo measurement of AChE activity. Both [<sup>11</sup>C]MP4A and [<sup>11</sup>C]PMP opened a window on cerebral cholinergic function in the human brain and have elucidated the role of cholinergic system in Alzheimer's disease, Parkinson's disease, and related disorders.

## 7.1 Introduction

Acetylcholine-mediated neurotransmission has been implicated in complex functions such as attention, memory, cognition, and consciousness (Bartus et al. 1982; Benarroch 2010). Its gradual loss, as in Alzheimer's disease (AD) and Parkinson's disease (PD), is associated with progressive deterioration of cognitive function (Bartus et al. 1982; Benarroch 2010; Hampel et al. 2018). Choline acetyltransferase (ChAT) is an enzyme that synthesizes acetylcholine, and acetylcholinesterase (AChE) is an enzyme that hydrolyzes acetylcholine and thereby regulates the concentration of the transmitter at the cholinergic synapse. ChAT is cytoplasmic within cholinergic neurons, mainly presynaptic, but also axonal (Mesulam and Geula 1992). In contrast, AChE is localized within the intersynaptic space, membrane bound predominantly on presynaptic cholinergic neurons, and on postsynaptic cholinergic neurons (Mesulam and Geula 1992). AChE and ChAT are frequently described as equivalent markers of cholinergic neurons in the brain.

AChE activity varies grossly across cerebral structures. The relative ratio of cerebral cortex/thalamus/cerebellar hemispheres/striatum is 1/3.5/10/33–45 for AChE activity in postmortem normal human brains (Atack et al. 1986; Arai et al. 1984). It may be challenging to obtain accurate measurement of AChE binding or activity in all regions with a single radiotracer. There are also species differences in brain AChE activity. AChE activity in the cerebellar hemispheres is similar to or smaller than that in the cerebral cortex of mice, rats, and rabbits (Shinotoh et al. 2004; Tavitian et al. 1993a; De Vos et al. 2000), while human cerebellar AChE activity is ten times higher than that in the cerebral cortex.

Acetylcholine is also cleaved by a less specific enzyme, butyrylcholinesterase (BuChE). BuChE can act as a molecular décor for natural anti-AChEs by reacting with these toxins before they reach AChE, but may also play a role in the regulation of cholinergic neurotransmission in specific brain regions (Darvesh et al. 2003). In postmortem AD brains, BuChE activity was found to be increased as the number of

senile plaques increased, while ChAT and AChE activities decreased (Perry et al. 1978). Therefore, radiotracers for AChE are required to have high selectivity for AChE over BuChE.

## 7.2 PET Radiotracers for Acetylcholinesterase

Several positron-emission tomography (PET) radiotracers for AChE binding sites or activity have been developed since the 1990s, while no PET radiotracer has been developed to measure ChAT-binding sites or activity until now. There are two major approaches to measuring AChE-binding sites or activity by radiotracers. One is the use of cholinesterase (ChE) inhibitors that bind to AChE. Another approach is the use of acetylcholine analogues that serve as substrates for AChE.

### 7.2.1 Cholinesterase Inhibitors

Several ChE inhibitors have been labeled with carbon-11 as radiotracers for visualizing AChE-binding sites (Shinotoh et al. 2004). Tacrine (THA) is a potent ChE inhibitor. An analogue of tacrine, 1,2,3,4-tetrahydro-9-methyl-amino-acridine (N-methylTHA, MTHA), was labeled with carbon-11, and the kinetics of [ $^{11}\text{C}$ ]MTHA in baboon brains were investigated by PET (Tavitian et al. 1993a, b). The regional distribution of [ $^{11}\text{C}$ ]MTHA, however, did not parallel that of AChE concentrations in baboon or human brains. This could be attributed to the fact that THA does not have selectivity for AChE over BuChE.

Physostigmine is a competitive ChE inhibitor that reversibly binds to AChE. Radioactivity distribution in the brain following intravenous injection of [ $^{11}\text{C}$ ]physostigmine was found to correspond to the known distribution of AChE in baboons and humans (Tavitian et al. 1993b; Blomqvist et al. 2001). However, the kinetics of [ $^{11}\text{C}$ ]physostigmine are so complex *in vivo* that this may hamper the quantitative assessment of AChE-binding sites (Blomqvist et al. 2001).

Some N-benzylpiperidines are potent and selective AChE inhibitors. Of these compounds, carbon-11-labeled CP126,998 was developed as a PET radiotracer for AChE (Bencherif et al. 2002). The regional brain uptake of [ $^{11}\text{C}$ ]CP126,998 corresponded to the rank order of regional AChE distribution in mouse and human brains. The demethylated form of CP126,998 (CP118,954) is a more potent and selective AChE inhibitor than CP126,998. CP118,954 was labeled with fluorine-18 and evaluated for *in vivo* mapping of AChE in the mouse (Lee et al. 2004). Autoradiography showed high radioactivity accumulation in the striatum, an AChE-rich region.

Donepezil is a noncompetitive, high-affinity, reversible AChE inhibitor, and is widely used for the treatment of AD. Donepezil has over thousandfold more selectivity of AChE over BuChE. Donepezil was initially labeled with carbon-11 at the 6' position (De Vos et al. 2000). Regional distribution studies in rabbit, however, did not show any correlation between the uptake of radioactivity and the amount of AChE. Subsequently, donepezil was labeled with carbon-11 at the 5' position

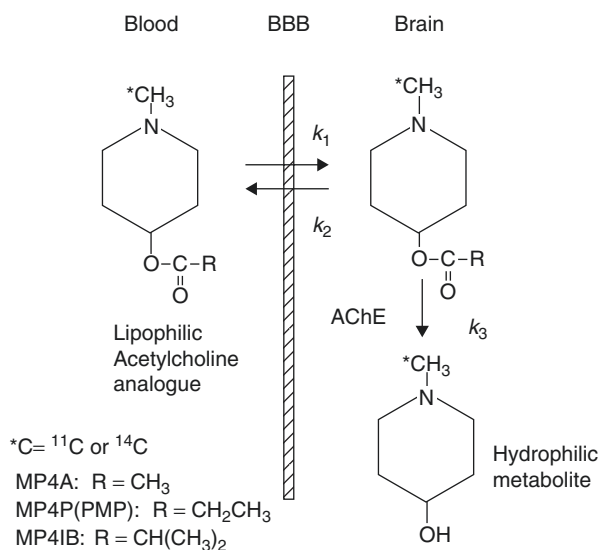
([5-<sup>11</sup>C-methoxy]donepezil), and the regional distribution study demonstrated heterogeneous radioactivity uptake (striatum, brainstem > cerebral cortex > cerebellum) in the rat brain (Funaki et al. 2003). A PET study in humans demonstrated that intravenously injected [5-<sup>11</sup>C methoxy]donepezil rapidly enters the brain and is mainly distributed in the striatum, thalamus, and cerebellum, which contain high densities of AChE compared with the cerebral cortex (Okamura et al. 2008). However, donepezil has moderate binding affinity for  $\sigma$ 1-receptors that are widely distributed throughout the brain including the cerebral cortex, hippocampus, and cerebellum (De Vos et al. 2000; Ishikawa et al. 2009). Therefore, [5-<sup>11</sup>C-methoxy]donepezil may not be a suitable PET ligand for measuring AChE-binding sites in the brain. Recently, [5-<sup>11</sup>C-methoxy]donepezil has been used for the measurement of AChE density in human peripheral organs (Gjerløff et al. 2014).

## 7.2.2 Acetylcholine Analogue Substrates

### 7.2.2.1 Carbon-11-Labeled Compounds

Irie et al. (1994, 1996; Namba et al. 1994) examined several piperidyl esters as acetylcholine analogue substrates as radiotracers for the measurement of AChE activity *in vivo* in the brain (Fig. 7.1). N-methyl-3-piperidyl esters and N-methyl-4-piperidyl esters have a tertiary-amine structure, thereby making them lipophilic, and readily cross the blood-brain barrier. Both compounds are hydrolyzed by AChE in the brain, and yield a hydrophobic and polar metabolite, N-methylpiperidinol, which does not readily cross the blood-brain barrier and is thus trapped locally in the brain. Esters with a different acyl group or its position on the piperidine ring showed a wide range of enzymatic reactivity, about 40-fold, and specificity of the esters to AChE depends

**Fig. 7.1** Two-tissue compartment model with three parameters describing the kinetic behavior of acetylcholine analogue substrates in the brain. AChE: acetylcholinesterase, BBB: blood-brain barrier



on the acyl group. The acetates (methyl-3-piperidyl acetate: MP3A; methyl-4-piperidyl acetate: MP4A) and propionates (methyl-3-piperidyl propionate: MP3P; methyl-4-piperidyl propionate: MP4P) have high specificity of more than 96% for AChE, followed by isobutyrate (methyl-3-piperidyl isobutyrate: MP3IB, 89%; methyl-4-piperidyl isobutyrate: MP4IB, 82%) and butyrate (MP3B, 45%). All of the esters (after labeling with carbon-14) showed rapid and high uptakes in the mouse brain followed by a slow decrease of radioactivity. Rank orders of brain uptake of these esters at 30 min postinjection were in accord with that of AChE activity in the mouse brain: striatum > cerebral cortex > cerebellar cortex. Chemical forms of  $^{14}\text{C}$  radioactivity in the mouse brain after intravenous injections of MP3A and MP4A were analyzed by a thin-layer chromatography method. More than 95% of radioactivity was found to exist as hydrolytic metabolites ( $^{14}\text{C}$ MP3OH or  $^{14}\text{C}$ MP4OH). When  $^{14}\text{C}$ MP3OH or  $^{14}\text{C}$ MP4OH was intravenously injected in the mouse, only low brain uptakes were observed, indicating that these N-methylpiperidinols are so hydrophilic that they do not readily cross the blood-brain barrier. The 4-piperidyl series do not have asymmetrical carbon, while the 3-piperidyl series are asymmetrical and thus have optical isomers, and their kinetics have to be taken as averaged values of two optical isomers, a confounding factor in the interpretation of the kinetics of these radiotracers. Taken together, MP4A and MP4P are thought to be the most promising radiotracers for the measurement of AChE activity in the brain.

Furthermore, the brain kinetics of  $^{14}\text{C}$ MP4A and  $^{14}\text{C}$ MP4P were investigated in rats with a unilateral lesion in the nucleus basalis magnocellularis (nBM) (Irie et al. 1996). The nBM lesion reduced cortical AChE activity by 30–50% compared with that in the contralateral side of the cerebral cortex. Radioactivity uptakes 30 min postinjection of  $^{14}\text{C}$ MP4A and  $^{14}\text{C}$ MP4P were reduced by 21% and 27%, respectively, in the lesioned side of the cortex, suggesting that both radiotracers have sufficient sensitivity to capture less than 50% of AChE activity changes in the cerebral cortex (Irie et al. 1996).

Kilbourn et al. (1996) examined the kinetics of  $^{11}\text{C}$ MP4A and  $^{11}\text{C}$ MP4P (or  $^{11}\text{C}$ PMP; they preferred to call  $^{11}\text{C}$ PMP) in mice and monkeys. They found that regional brain distributions after intravenous injection of  $^{11}\text{C}$ PMP showed better discrimination between regions of high, intermediate, and low AChE activities than  $^{11}\text{C}$ MP4A because the hydrolysis rate of acetylcholine by MP4P is about one-fourth that by MP4A (Table 7.1) (Shinotoh et al. 2004; Irie et al. 1996; Kilbourn et al. 1996).

Currently, both  $^{11}\text{C}$ MP4A and  $^{11}\text{C}$ PMP are in clinical use for studies of various neurodegenerative diseases by PET (Tables 7.2 and 7.3).  $^{11}\text{C}$ MP4A has higher specificity for AChE over BuChE than  $^{11}\text{C}$ PMP in the human cerebral cortex (Table 7.1) (Shinotoh et al. 2004; Irie et al. 1996).  $^{11}\text{C}$ MP4A-PET gives precise estimates of  $k_3$ , an index of AChE activity, in the cerebral cortex, while  $^{11}\text{C}$ PMP provides precise measurement of  $k_3$  in the cerebral cortex and thalamus. In the striatum, both radiotracers give poor estimates of  $k_3$  values because of the delivery limitation effect.

**Table 7.1** Characteristics of two radiotracers for measurement of acetylcholinesterase activity

Radiotracer	Selectivity for AChE				PET scan time	Optimum brain regions for measurement in humans
	Mouse brain (Irie et al. 1994)	Human cerebral cortex (Irie et al. 1996)	Relative hydrolysis rate (Irie et al. 1994)			
[ <sup>11</sup> C]MP4A	99%	94%	4.43		40–60 min	Cerebral cortex (thalamus)
[ <sup>11</sup> C]PMP	97%	86%	0.97		60–80 min	Cerebral cortex, thalamus, (cerebellar cortex) (striatum)

AChE: acetylcholinesterase, [<sup>11</sup>C]MP4A: [<sup>11</sup>C]N-methylpiperidine-4-yl acetate, [<sup>11</sup>C]PMP: [<sup>11</sup>C]N-methylpiperidine-4-yl propionate

**Table 7.2** PET studies in Alzheimer's disease and related disorders

Radiotracer	Author	Year	Subjects	Main findings on AChE activity
[ <sup>11</sup> C]MP4A	Iyo et al. (1997)	1997	5 AD, 8 HC	Reduction in neocortex (19–38%) in AD patients
[ <sup>11</sup> C]PMP	Kuhl et al. (1999)	1999	14 AD, 26 HC	Reduction in neocortex and hippocampus (25–33%) in AD patients Inhibition (50%) in brain AChE activity after 1.5 mg dose of physostigmine infusion
[ <sup>11</sup> C]MP4A	Shinotoh et al. (2000)	2000	30 AD, 14 HC	Reduction in neocortex (23%), hippocampus (21%), and amygdala (27%) in EOAD Reduction in temporoparietal cortex (14%, 18%) and amygdala (26%) in LOAD Progressive loss of cortical AChE activity in 7 repeat PET scans in AD Correlation between cortical AChE activity and MMSE scores in 37 AD patients
[ <sup>11</sup> C]PMP	Kuhl et al. (2000)	2000	9 AD	Inhibition (27%) of cortical AChE activity after 5–10 mg/day donepezil treatment
[ <sup>11</sup> C]MP4A	Shinotoh et al. (2001)	2001	3 AD	Inhibition (39 ± 5%) of cortical AChE activity after 3–5 mg/day donepezil treatment
[ <sup>11</sup> C]MP4A	Kaasinen et al. (2002)	2002	11 AD	Inhibition (28–39%) of cortical AChE activity after 10 mg/day donepezil treatment Inhibition (28–37%) of cortical AChE activity after 9 mg/day rivastigmine treatment
[ <sup>11</sup> C]MP4A	Rinne et al. (2003)	2003	13 AD, 12 MCI, 12 HC	Reduction of AChE activity in hippocampus in MCI patients (17%) and in AD patients (27%)
[ <sup>11</sup> C]MP4A	Herholz et al. (2004)	2004	9 AD, 13 HC	Reduction of cortical AChE activity (18–34%) in AD, [ <sup>11</sup> C]MP4A uptake preserved in nbM
[ <sup>11</sup> C]PMP	Bohnen et al. (2005a)	2005	15 AD, 12 HC	Cortical AChE activity associated with performance on a test of attention and working memory

**Table 7.2** (continued)

Radiotracer	Author	Year	Subjects	Main findings on AChE activity
[ <sup>11</sup> C]PMP	Bohnen et al. (2005b)	2005	14 AD	Inhibition (19.1%) of cortical AChE after 10 mg/day donepezil treatment Degree of cortical AChE inhibition correlates with executive and attentional functional changes
[ <sup>11</sup> C]MP4A	Eggers et al. (2006)	2006	19 AD	Preserved cortical AChE activity in ApoE4-positive patients compared with negative patients
[ <sup>11</sup> C]donepezil	Okamura et al. (2008)	2007	10 AD, 6 HC	Reduction (18–30%) of cortical [ <sup>11</sup> C]donepezil binding in AD Inhibition (62–63%) of cortical [ <sup>11</sup> C]donepezil binding after 5 mg/day donepezil treatment
[ <sup>11</sup> C]PMP	Kadir et al. (2008)	2008	18 AD	Inhibition (30–40%) of cortical AChE activity after galantamine (16–24 mg/day) treatment
[ <sup>11</sup> C]PMP	Bohnen et al. (2009a)	2009	28 elderly subjects	Severity of periventricular WMH was related to cortical AChE activity
[ <sup>11</sup> C]MP4A	Marcone et al. (2012)	2012	10 MCI, 7 AD, 4 DLB, 9 HC	Reduction of cortical and hippocampal AChE activity in MCI patients and in AD patients
[ <sup>11</sup> C]MP4A	Haense et al. (2012)	2012	17 MCI, 21 HC	Decline of cortical AChE activity, most pronounced in temporal regions in MCI
[ <sup>11</sup> C]MP4A	Garibotto et al. (2013)	2013	9 MCI, 7 AD, 9 HC	Correlation between education and hippocampal AChE activity Correlation between occupation and AChE activity in right posterior cingulate cortex
[ <sup>11</sup> C]MP4A	Richter et al. (2014)	2014	14 elderly subjects	AChE activity in right medial temporal lobe correlated with episodic memory function
[ <sup>11</sup> C]MP4A	Shimada et al. (2015)	2015	14 DLB, 25 AD, 18 HC	Reduction of cortical AChE activity was consistently prominent in DLB than AD patients
[ <sup>11</sup> C]MP4A	Richter et al. (2017)	2017	17 MCI, 18 HC	Periventricular WML negatively correlated with brain AChE activity in MCI and HC
[ <sup>11</sup> C]MP4A	Richter et al. (2018)	2018	14 MCI, 16HC	Reduction of cortical AChE, especially in lateral temporal lobe Association of change in neural activation in response to rivastigmine with local AChE activity
[ <sup>11</sup> C]MP4A	Hirano et al. (2018)	2018	22 EOAD, 26 LOAD, 16 HC	Reduction of wider cortical AChE activity in EOAD than LOAD

AD: Alzheimer's disease, HC: healthy controls, EOAD: early-onset AD, LOAD: late-onset AD, MCI: mild cognitive impairment, nbM: nucleus basalis of Meynert, WMH(L): white matter hyperintensities (load), DLB: dementia with Lewy bodies

**Table 7.3** PET studies in Parkinson's disease and related disorders

Radiotracer	Author	Year	Subjects	Main findings on AChE activity
[ <sup>11</sup> C]MP4A	Shinotoh et al. (1999)	1999	16 PD, 12 PSP, 13 HC	Reduction of cortical (17%) and thalamic (13%) AChE activity in PD Prominent reduction (38%) of thalamic but not cortical AChE activity in PSP
[ <sup>11</sup> C]PMP	Bohnen et al. (2003)	2003	11 PD, 14 PDD, 12 AD, 10 HC	More severe reduction of cortical AChE activity in PD with dementia than AD
[ <sup>11</sup> C]MP4A	Hilker et al. (2005)	2005	17 PD, 1 PDD, 31 HC	Reduction (10.7%, 29.7%) of cortical AChE activity in PD and PDD
[ <sup>11</sup> C]MP4A	Hirano et al. (2006)	2006	2 FTDP-17	Prominent reduction of cortical (13% to 44%) and thalamic (57%) AChE activity
[ <sup>11</sup> C]PMP	Bohnen et al. (2007)	2006	12 PD, 6 PDD, 10 HC	Inverse correlation between cortical AChE activity and depression scale
[ <sup>11</sup> C]PMP	Hirano et al. (2008)	2008	8 MSA-C, 7 SCA-3, 3 SCA-6, 13 HC	Reduction of AChE activity in thalamus (27%) and cerebellar cortex (36%) in MSA-C Reduction of AChE activity in thalamus (23%) in SCA-3
[ <sup>11</sup> C]MP4A	Shimada et al. (2009)	2009	18 PD, 21 PDD/DLB, 26HC	Reduction (12%) of cortical activity, especially in medial occipital cortex in PD Severe reduction (27%) of cortical AChE activity in PDD/DLB
[ <sup>11</sup> C]PMP	Bohnen et al. (2009b)	2009	44 PD, 15 HC	Reduction of cortical (12.3%) and thalamic (11.8%) AChE activity in PD fallers PD non-fallers showed reduction (6.6%) of AChE activity only in cortex
[ <sup>11</sup> C]MP4A	Hirano et al. (2010)	2010	7 CBS, 12 PSP, 8 FTD, 16 HC	Reduction of cortical (17.5%) but not thalamic AChE activity in CBS Reduction of cortical (9.4%) and thalamic (24%) AChE activity in PSP No reduction of brain AChE activity in FTD
[ <sup>11</sup> C]PMP	Gilman et al. (2010)	2010	12 PD, 13 MSA-P, 4 PSP, 22 HC	Cortical AChE reductions were 15.3%, 14.6%, 13.2% in PD, MSA-P, PSP Subcortical AChE activities were more decreased in MSA-P and PSP than in PD

**Table 7.3** (continued)

Radiotracer	Author	Year	Subjects	Main findings on AChE activity
[ <sup>11</sup> C]MP4A	Miyoshi et al. (2010)	2010	3 Presymptomatic FTDP-17	One presymptomatic subject showed reduction of cortical AChE activity
[ <sup>11</sup> C]PMP	Bohnen et al. (2012)	2012	101 PD, 29 HC	Low neocortical AChE activity PD subgroup had lower cognitive performance More than half of low thalamic AChE activity PD subgroup had history of falls
[ <sup>11</sup> C]PMP	Kotagal et al. (2012)	2012	80 PD	Reduction of cortical and thalamic AChE activity in PD with RBD ( <i>n</i> = 27) in comparison to those without RBD ( <i>n</i> = 53)
[ <sup>11</sup> C]PMP	Bohnen et al. (2013)	2013	125 PD, 32 HC	PD group with low cortical AChE activity had slower gait speed
[ <sup>11</sup> C]PMP	Bohnen et al. (2014)	2014	143 PD	90% of PD with freezing of gait had cortical AChE activity reduction or amyloid deposition
[ <sup>11</sup> C]PMP	Weather et al. (2014)	2014	30 PD	Less thalamic ACh activity in risky drivers compared to safe drivers
[ <sup>11</sup> C]PMP	Bohnen et al. (2015)	2015	143 PD	Cortical cholinergic denervation frequency increased with increasing cognitive impairment
[ <sup>11</sup> C]PMP	Müller et al. (2015)	2015	137 PD	Combination of RBD and fall history is a predictor for thalamic and cortical deficits
[ <sup>11</sup> C]PMP	Chou et al. (2016)	2016	133 PD	Cholinergic uptake was not a predictor of fatigue in PD
[ <sup>11</sup> C]donepezil	Gjerløff et al. (2015)	2015	12 PD, 12 HC	Reduction of [ <sup>11</sup> C]donepezil binding in small intestine (35%), and pancreas (22%)
[ <sup>11</sup> C]donepezil	Fedorova et al. (2017)	2017	19 early PD, 16 HC	Reduction of [ <sup>11</sup> C]donepezil binding in small intestine (14%), colon (22%), and kidney (14%)
[ <sup>11</sup> C]PMP	Liu et al. (2018)	2018	14 LRRK2 PD, 16 LRRKs mutation carriers, 8 PD, 11 HC	Increased cortical AChE activity in LRRK2 mutation carriers compared with HC Increased cortical and thalamic AChE activity in LRRK2 PD compared with PD

PD: Parkinson's disease, PSP: progressive supranuclear palsy, HC: healthy controls, PDD: PD with dementia, FTDP-17: frontotemporal dementia with parkinsonism linked to chromosome 17, MSA-C: cerebellar syndrome of multiple system atrophy, SCA-3: spinocerebellar atrophy type 3, DLB: dementia with Lewy bodies, CBS: corticobasal syndrome, FTD: frontotemporal dementia, MSA-P: parkinsonian syndromes of multiple system atrophy, RBD: REM sleep behavior disorder



### 7.2.2.2 Kinetics Analysis of [<sup>11</sup>C]MP4A and [<sup>11</sup>C]PMP

[<sup>11</sup>C]MP4A and [<sup>11</sup>C]PMP are irreversible PET radiotracers. Irreversible tracers are generally analyzed using a two-tissue compartment model—one representing an authentic radiotracer, and a second representing a metabolically trapped radiotracer (Fig. 7.1). The two-tissue compartment configuration has three rate parameters,  $K_1$  to  $k_3$ .  $K_1$  is defined as the transfer rate of an authentic tracer from plasma to tissue,  $k_2$  is the efflux rate of authentic tracer from tissue to plasma, and  $k_3$  is the hydrolysis rate of authentic tracer by AChE. The model does not require the parameter  $k_4$  because hydrolysis by AChE is completely irreversible. The three parameters are estimated by fitting regional brain time-radioactivity curves with plasma input function to the following theoretical function in a nonlinear least-squares method:

$$C_t = C_b + C_m, \quad (7.1)$$

$$C_b = \int_0^T K_1 C e^{-(k_2+k_3)(T-t)} dt, \quad (7.2)$$

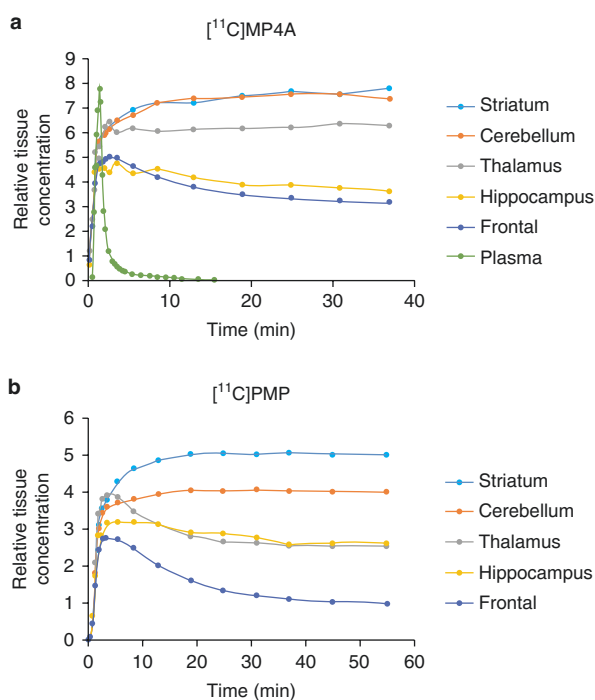
$$C_m = \int_0^T k_3 C_b dt, \quad (7.3)$$

where  $C_b$  and  $C_m$  represent authentic and metabolized tracers, respectively.

An irreversible radiotracer is required to be matched well between the rate of trapping ( $k_3$ ) and the rate of blood-brain barrier transport ( $K_1$  and  $k_2$ ) (Koeppel et al. 1999). The ratio of the metabolic rate ( $k_3$ ) and the efflux rate ( $k_2$ ), i.e.,  $\alpha \equiv k_3/k_2$ , can affect the reliability of data from compartment model analysis (Koeppel et al. 1999). A simulation study showed that the optimal  $\alpha$ -value with the maximal precision of  $k_3$  parameter is around 0.2 in the two-tissue compartment model with the three parameters (Ohya et al. 2011). The good precision of  $k_3$  estimates was maintained when the tracer  $\alpha$ -value was within a range of approximately 0.05–1.0 (optimal  $\alpha$  range). With  $k_2$  values of about 0.13 min<sup>-1</sup> (Namba et al. 1999), the  $\alpha$ -value was calculated to be 0.48–0.73 in the cerebral cortex in healthy volunteers in a [<sup>11</sup>C]MP4A PET study, although  $\alpha$ -values in the thalamus, cerebellum, and striatum fell outside of the optimal range. In a [<sup>11</sup>C]PMP PET study, the  $\alpha$ -value was about 0.15–3.3 in the brain, and  $\alpha$ -values in the striatum fell outside of the optimal range (Koeppel et al. 1999).

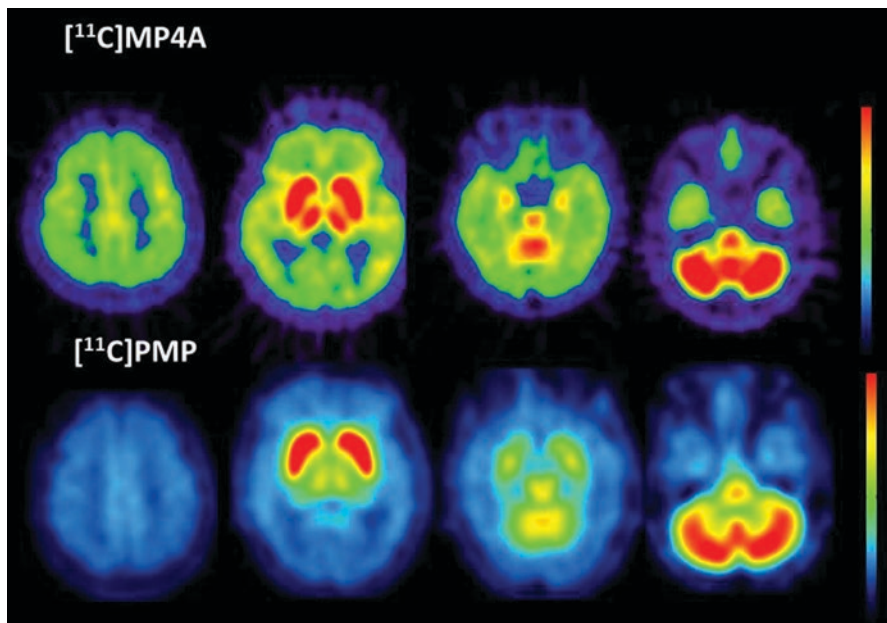
Radioactivity in the cerebral cortex gradually decreased, reaching a plateau after an initial high uptake following intravenous injection of [<sup>11</sup>C]MP4A in 20 healthy volunteers, while that in the cerebellum and striatum gradually increased to reach a plateau at about 20 min following an initial rapid uptake (Fig. 7.2) (Namba et al. 1999). The radioactivity curve of the thalamus is between those of the cerebellum and the cerebral cortex. The plateau level of radioactivity in those regions suggests that washout of the radioactive metabolite from the brain is negligible during PET measurement. Each of the plateau levels was in accord with the regional distribution of AChE activity in the brain (Figs. 7.2 and 7.3). The fraction of [<sup>11</sup>C]MP4A in the total plasma radioactivity rapidly decreases along bi-exponential curves to reach a

**Fig. 7.2** Time-activity curves in the brain and plasma in a 66-year-old healthy man following intravenous injection of [ $^{11}\text{C}$ ]MP4A (a), and time-activity curves in the brain in a 63-year-old healthy man following intravenous injection of [ $^{11}\text{C}$ ]PMP (b)



low level of less than 5% within 15 min (Fig. 7.2a) (Namba et al. 1999). [ $^{11}\text{C}$ ]MP4A PET scans are usually performed for 40–60 min. In the two-tissue compartment model with the three parameters, the  $k_3$  values in the cerebral cortices were estimated with coefficients of variation of less than 15% in 20 healthy subjects by a nonlinear least-squares method (Namba et al. 1999). In the regions with higher AChE activity (thalamus, cerebellum, and striatum), an obvious washout phase of time-activity curves is not observed, and thus  $k_2$  is estimated to be almost zero, leading to underestimation of  $k_3$  values. Reasonable  $k_3$  values in the regions with high AChE activity would be obtained by constraining  $K_1/k_2$  values. We constrained the  $K_1/k_2$  values in the thalamus, cerebellum, and striatum to be the mean  $\pm$  2SD of fitted  $K_1/k_2$  values in the cerebral cortices for each subject (Namba et al. 1999). This constraint of the  $K_1/k_2$  values was based on the assumption that the tissue distribution volume of authentic tracers is approximately uniform across the brain. In this analysis, the  $k_3$  values in the thalamus were estimated with coefficients of variation of 20% (Namba et al. 1999). The coefficients of variation of  $k_3$  values in the cerebellum and striatum were 46% and 161% in this analysis, suggesting that  $k_3$  estimates in those regions are not reliable.

Based on the characteristic kinetics of [ $^{11}\text{C}$ ]MP4A, a unique kinetic analysis was developed (Nagatsuka et al. 2001; Herholz et al. 2001). Since metabolism of [ $^{11}\text{C}$ ]MP4A by AChE in the striatum and cerebellum is very high, it is assumed that all [ $^{11}\text{C}$ ]MP4A molecules entering these regions would be trapped there (metabolic



**Fig. 7.3** PET images of a 66-year-old healthy man 20–40 min following intravenous injection of [ $^{11}\text{C}$ ]MP4A (upper row), and PET images of a 63-year-old healthy man 40–60 min following i.v. injection of [ $^{11}\text{C}$ ]PMP. These images were scanned at the National Institutes for Quantum and Radiological Science and Technology, Japan

trapping). This assumption allows the use of the striatum or cerebellum as reference regions to obtain input function data in the form of a single integral with respect to PET scan time:

$$C_{\text{R}}(t) = K_1^{\text{REF}} \int_0^t C_{\text{a}}(\theta) d\theta$$

where  $C_{\text{R}}(t)$  is radioactivity in the reference region, and  $K_1^{\text{REF}}$  is the value of  $K_1$  in this region.

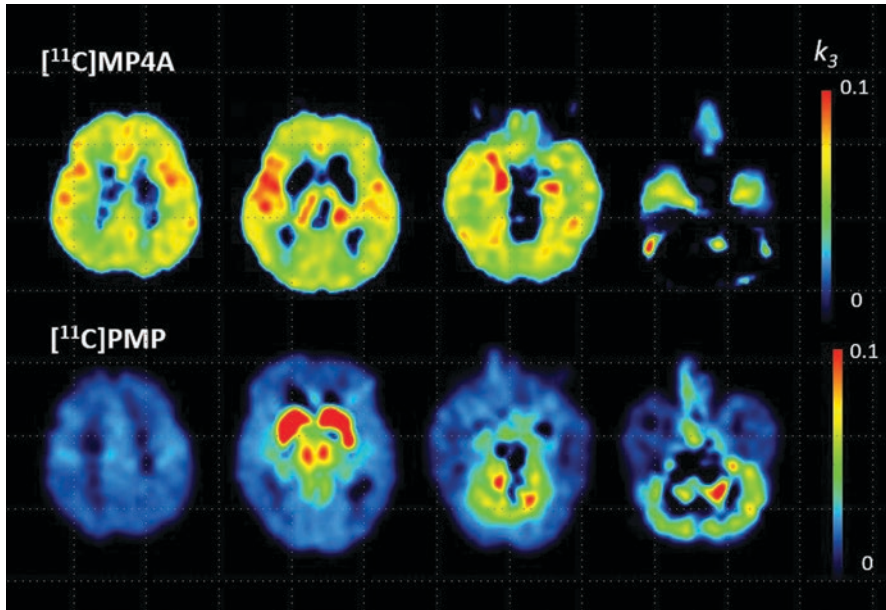
When such input function data is applied to Blomqvist's linear approach (Blomqvist 1984), the kinetics parameter  $k_3$  can be estimated from PET data alone without arterial blood sampling: (Nagatsuka et al. 2001; Herholz et al. 2001)

$$C_{\text{T}}(t) = p_1 C_{\text{R}}(t) + p_2 \int_0^t C_{\text{R}}(\theta) d\theta + p_3 \int_0^t C_{\text{T}}(\theta) d\theta \quad (7.4)$$

$$p_1 = \frac{K_{1\text{Target}}}{K_{1\text{REF}}} \equiv K_{\text{R}} \quad (7.5)$$

$$p_2 = K_{\text{R}} \cdot k_3 \quad (7.6)$$

$$p_3 = -(k_2 + k_3) \quad (7.7)$$



**Fig. 7.4** Parametric ( $k_3$ ) images of a 66-year-old healthy man in a [ $^{11}\text{C}$ ]MP4A PET study (upper row), and parametric ( $k_3$ ) images of a 63-year-old healthy man in a [ $^{11}\text{C}$ ]PMP PET study. Both parametric images of [ $^{11}\text{C}$ ]MP4A were obtained by Nagatsuka's reference tissue-based linear least-squares (RLS) analysis implemented in PMOD 4.0 (PMOD Technologies LLC, Zürich, Switzerland). Note that  $k_3$  could not be estimated in the striatum and cerebellum by [ $^{11}\text{C}$ ]MP4A PET and a part of the cerebellum and brainstem by [ $^{11}\text{C}$ ]PMP PET

In this approach, the rate of tracer influx into the target region is obtained as a relative value  $K_R$ .

This method is applicable to [ $^{11}\text{C}$ ]PMP PET. In the case of [ $^{11}\text{C}$ ]PMP, the striatum is preferable to the cerebellum because of its slower kinetics. This kinetic analysis without blood sampling is known as reference tissue-based linear-least squares (RLS) analysis, and is often employed in clinical [ $^{11}\text{C}$ ]MP4A or [ $^{11}\text{C}$ ]PMP PET studies (Fig. 7.4).

The time-activity curves of [ $^{11}\text{C}$ ]PMP in the brain and blood are similar but slower than those of [ $^{11}\text{C}$ ]MP4A. Koeppe et al. (1999) examined the kinetics of [ $^{11}\text{C}$ ]PMP in 12 healthy volunteers. Sequential [ $^{11}\text{C}$ ]PMP PET scans were performed for 80 min. They found that [ $^{11}\text{C}$ ]PMP can be used with dynamic PET to provide an in vivo index of local AChE activity ( $k_3$ ) with excellent precision in the cerebral cortex (coefficient of variation less than 10% in healthy volunteers) and other regions of low enzyme activity by unconstrained nonlinear least-squares fit to the theoretical model. The excellent precision was thought to be due to its bolus-like input function shape, complete irreversibility, and near-optimal kinetic properties. Estimates of  $k_3$  were poorer in regions of high enzyme activity such as the striatum and cerebellum because of excessive bias and variability caused by

delivery-limitation effects. The constraint on  $K_1/k_2$  was required to yield meaningful estimates. Koeppe et al. constrained  $K_1/k_2$  values by two methods. In one method, the  $K_1/k_2$  ratio was fixed at 4.0. This estimate of the AChE activity index was referred to as the constrained least-squares estimate of  $k_3c$ . In the other method,  $*k_3$  (modified  $k_3$ ) was defined as unconstrained  $k_3$  multiplied by unconstrained  $K_1/k_2$ . This was based on the following facts. In regions with high AChE activity, the estimates of  $k_2$  and  $k_3$  become extremely variable, but they remain highly correlated. If  $k_3$  is underestimated,  $k_2$  is underestimated, while  $K_1$  continues to be estimated precisely. Modified  $k_3$  value was defined as follows:

$$*k_3 = k_3 \left[ (K_1 / k_2) / 4.0 \right] (\text{min}^{-1}) \quad (7.8)$$

This estimate of the AChE activity index was referred to as the constrained least-squares estimate of  $*k_3$ . These constrained least-square estimates of  $k_3c$  and  $*k_3$  yielded meaningful estimates of AChE activity in the thalamus, cerebellum, and striatum in the [ $^{11}\text{C}$ ]PMP PET study.

Koeppe et al. (1999) proposed another method to estimate  $k_3$  entirely from the shape of the tissue time-activity curve without arterial blood sampling or plasma metabolite analysis (shape analysis) (Koeppe et al. 1999). The following conditions are required for this approach to be valid: the concentration of authentic tracer in arterial plasma must reach zero during the study period, and hydrolysis of the substrate PMP is completely irreversible. Under these assumptions, two further conditions hold: (Bartus et al. 1982) all radiotracer in brain is in the free compartment at the start of the study, and all radiotracer in the brain is in the metabolite compartment by the end of the study. Shape analysis is applicable only to brain areas of low activity such as the cerebral cortex because estimates of  $k_3$  values in brain regions with high enzyme activity become progressively more inaccurate. This shape analysis is also applicable to [ $^{11}\text{C}$ ]MP4A, although the sensitivity of shape analysis for detecting neocortical regions with abnormally low  $k_3$  in AD patients is less than the conventional compartment analysis with arterial sampling (Tanaka et al. 2001).

### 7.2.2.3 Fluorine-18-Labeled Compounds

There have been several attempts to develop acetylcholine analogues labeled with fluorine-18 (Kikuchi et al. 2005, 2010, 2013; Shao et al. 2005; Zhang et al. 2003). Fluorine-18-labeled derivatives of MP4A or PMP may allow for the delivery of radiotracers to other PET centers since fluorine-18 has a longer half-life (109.8 min) than carbon-11 (20.4 min). *N*-[ $^{18}\text{F}$ ]fluoroethylpiperidin-4-yl acetate ([ $^{18}\text{F}$ ]FETP4A) was synthesized as an [ $^{18}\text{F}$ ]fluoroethylated MP4A derivative.<sup>33</sup> Disappointingly, an in vitro study of [ $^{18}\text{F}$ ]FETP4A showed that this compound has a much lower specificity for AChE compared with [ $^{11}\text{C}$ ]MP4A and a much slower (approximately one-twentieth) AChE-mediated hydrolysis rate compared with MP4A in rat cerebral cortical homogenate or test tube with pure enzyme.

A series of *N*-[ $^{14}\text{C}$ ]ethylpiperidin-3-yl or -4-ylmethanol esters (acetyl and propionyl esters) were newly designed and evaluated in an in vitro screening study (Kikuchi et al. 2005, 2010, 2013). *N*-[ $^{18}\text{F}$ ]fluoroethylpiperidin-4-ylmethyl acetate ([ $^{18}\text{F}$ ]FEP-4MA) was identified as a potential  $^{18}\text{F}$ -labeled compound for measuring

cerebral AChE activity. In rat experiments, [ $^{18}\text{F}$ ]FEP-4MA showed high brain uptake of the authentic ester, high AChE specificity, a moderate hydrolysis rate, and low membrane permeability of the metabolite, although the metabolite was slowly eliminated from the brain, probably by efflux transporters (Kikuchi et al. 2010, 2013). A simulation study and a monkey study of [ $^{18}\text{F}$ ]FEP-4MA were performed. In the simulation study, it was investigated how the tracer  $\alpha$ -value affected  $k_3$  precision under a different efflux rate constant ( $k_{e1}$ ) (Ohya et al. 2011). A nonlinear least-squares analysis of the [ $^{18}\text{F}$ ]FEP-4MA PET data showed that the precision of the  $k_3$  estimate was as high as that for [ $^{11}\text{C}$ ]MP4A, as the ratio of  $k_{e1}$  to the efflux rate ( $k_2$ ) was relatively small ( $<0.1$ ), suggesting that [ $^{18}\text{F}$ ]FEP-4MA may be useful as a radiotracer for the measurement of AChE activity in humans (Ohya et al. 2011).

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## 7.3 Clinical Applications of Acetylcholinesterase Imaging

### 7.3.1 Healthy Older Adults

Aging is a major risk factor for AD, and it is therefore of interest to study the effect of age on brain AChE activity. There was no alteration of cortical AChE activity in 23 healthy volunteers ranging in age from 24 to 89 years old in a [ $^{11}\text{C}$ ]MP4A PET study (Namba et al. 1998, 1999). Likewise, Kuhl et al. (1999) did not find any effect of age on cortical AChE activity measured by [ $^{11}\text{C}$ ]PMP PET in 26 healthy volunteers ranging in age from 23 to 80 years. There was also no gender effect on cortical AChE activity (Kuhl et al. 1999).

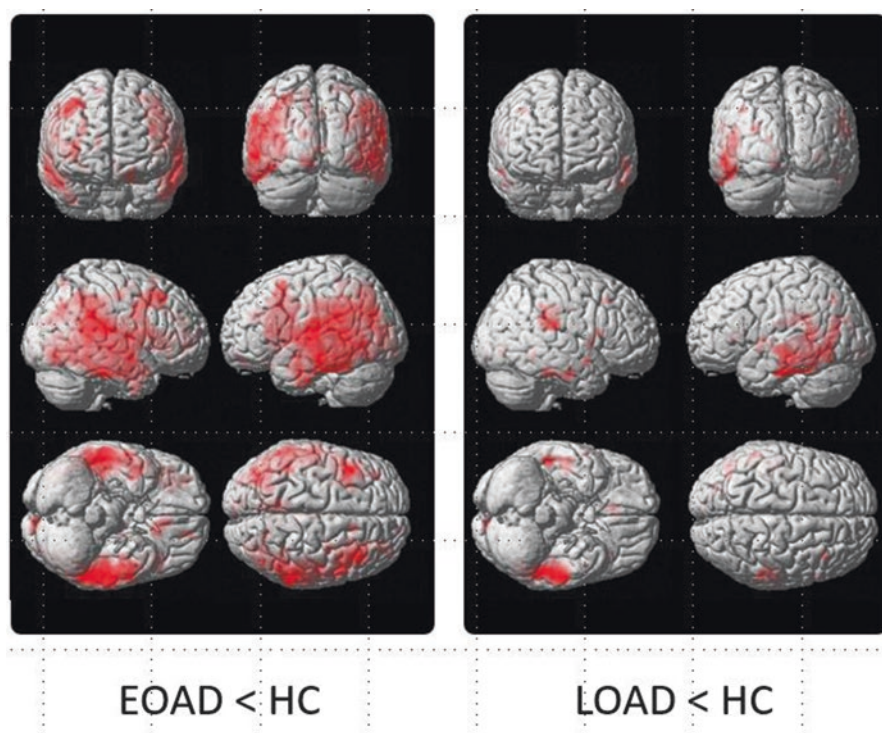
Richter et al. (2014) investigated the relationship between brain AChE activity measured by [ $^{11}\text{C}$ ]MP4A PET and memory performance in 14 healthy older adults ranging in age from 53 to 77 years. They found a significant correlation between AChE activity in the right medial temporal lobe and number of words recalled after a delay, suggesting that the integrity of the cholinergic system underlies inter-individual variability in memory function.

### 7.3.2 Alzheimer's Disease and Related Disorders

Impairment of memory and learning by cholinergic antagonists was demonstrated by pharmacological studies of healthy volunteers in the 1970s (Bartus et al. 1982). In the mid-1970s, depletions of presynaptic cholinergic markers, ChAT and AChE activity, were found in postmortem brains with AD (Perry et al. 1978; Davies and Maloney 1976). In 1982, pronounced loss of cholinergic neurons in the nucleus basalis of Meynert in postmortem brains with AD was demonstrated (Whitehouse et al. 1982). These discoveries led to the cholinergic hypothesis of AD (Bartus et al. 1982).

In 1997, we showed an average reduction of 31% of cortical AChE activity in 5 mild AD patients compared with 12 healthy controls in a [ $^{11}\text{C}$ ]MP4A PET study with arterial input function (Iyo et al. 1997). Subsequently, we showed different reduced distributions of brain AChE activity between 14 patients with early-onset

AD (EOAD) and 14 patients with late-onset AD (LOAD) by [ $^{11}\text{C}$ ]MP4A PET (Shinotoh et al. 2000). AChE activity was reduced in the entire neocortex (23% on average), hippocampus (21%), and amygdala (27%) in EOAD patients, while AChE activity was reduced only in the temporoparietal cortex (14, 18%) and amygdala (26%) in LOAD patients. Seven AD patients had repeat PET scans with a mean interval of 2 years, and cortical AChE activity had reduced further in the second PET scans, suggesting a progressive loss of the ascending cholinergic system from the nucleus basalis of Meynert in AD. There was a highly significant correlation between cortical AChE activity and Mini-Mental State Examination (MMSE) scores, supporting the cholinergic hypothesis in AD. Recently, we conducted voxel-based AChE activity estimation of [ $^{11}\text{C}$ ]MP4A PET data in an unconstrained nonlinear least-squares method with arterial input in 22 EOAD patients, 26 LOAD patients, and 16 healthy controls (Hirano et al. 2018). Both the EOAD and LOAD groups showed cortical AChE activity decrement in parietal, temporal, and occipital cortices, with wider and stringent cortical involvement in the EOAD group, most prominently in the temporal region, compared with healthy controls (Fig. 7.5). Herholz



**Fig. 7.5** Acetylcholinesterase (AChE) activity reduction in early-onset AD (EOAD) ( $n = 22$ ) and late-onset AD (LOAD) ( $n = 26$ ) in comparison with 16 healthy controls (HC). Statistical parametric T-score maps, with red-yellow color representing AChE activity reduction, are superimposed on a single subject T1, thresholded at voxel-level  $p < 0.05$ ; family-wise error (FEW) corrected with cluster above 30 voxels

et al. (2004) found reduction of AChE activity in the cerebral cortex and amygdala but not in the nucleus basalis of Meynert in mild-to-moderate AD in a [ $^{11}\text{C}$ ]MP4A PET study, suggesting that neocortical and amygdaloid function changes of the cholinergic system are an early and leading event in AD, rather than the consequence of neurodegeneration of basal nuclei.

Kuhl et al. (1999) found a reduction of neocortical (25–33%) and hippocampal AChE activity in 14 moderate-severe AD patients with a mean MMSE score of 14 by [ $^{11}\text{C}$ ]PMP PET. There was no reduction of AChE activity in the caudate, putamen, thalamus, pons, and cerebellum.

Bohnen et al. (2005a) found that cortical AChE activity is more robustly associated with functions of attention and working memory (WAIS-III Digit Span,  $R = 0.46$ ,  $P = 0.01$ ) compared to performance on primary memory tests (California Verbal Learning Test).

The E4 allele of apolipoprotein  $\epsilon$  (ApoE4) is the major genetic risk factor for AD, and increases the formation of amyloid plaques and neurofibrillary tangles. Eggers et al. (2006) examined the relation between ApoE genotype and cortical AChE activity in 19 patients with mild-to-moderate AD by [ $^{11}\text{C}$ ]MP4A PET. The ApoE4-positive patients ( $n = 11$ ) had significantly higher AChE levels than the ApoE4-negative ( $n = 8$ ) patients, while the two groups were comparable with regard to age and dementia severity. Higher AChE levels in ApoE4-positive patients could be attributed to AChE activity in amyloid plaque in the brain.

Rinne et al. (2003) measured brain AChE activity in 12 patients with mild cognitive impairment (MCI), 13 AD patients, and 12 healthy elderly volunteers by [ $^{11}\text{C}$ ]MP4A PET. There was a significant difference in AChE activity only in the hippocampus between the groups, and hippocampal AChE activity was only slightly reduced in MCI (17%) and early AD (27%). They concluded that the value of in vivo AChE activity measurements in detecting early AD is limited.

Marcone et al. (2012) assessed AChE activity with [ $^{11}\text{C}$ ]MP4A PET in 10 patients with multi-domain amnesic MCI (aMCI), 7 AD patients, and 9 healthy controls. They found widespread AChE reductions in several cortical regions and in hippocampus in all AD and aMCI patients who progressed to AD with a follow-up of 2 years. There was also a significant correlation between hippocampal AChE activity and long-term verbal and nonverbal memory. In addition, Haense et al. (2012) found a significant decline of cortical AChE activity in 17 MCI patients.

Richter et al. (2018) measured brain AChE activity by [ $^{11}\text{C}$ ]MP4A PET in 14 patients with MCI due to AD who had positive cerebrospinal fluid biomarkers indicative of AD pathology, and 16 age-matched healthy controls. AChE activity was significantly decreased in the cerebral cortex (11%), and this decrease was most pronounced in the lateral temporal lobes of MCI patients relative to healthy controls. The MCI patients showed less memory-related neural activation in the fusiform gyrus and impaired deactivation in the posterior cingulate cortex, relative to healthy controls, by functional magnetic resonance imaging (fMRI). These differences were attenuated under cholinergic stimulation with rivastigmine. The change in neural activation in response to rivastigmine was negatively associated with local AChE activity.



Garibotto et al. (2013) investigated reserve capacity by [ $^{11}\text{C}$ ]MP4A PET in nine patients with MCI, seven patients with early AD, and nine healthy controls. They found a positive correlation between education and AChE activity in the hippocampus bilaterally, and between occupation and AChE activity in the right posterior cingulate gyrus.

White matter lesions (WML) are observed in cognitively normal older persons and may be associated with cognitive dysfunction. One possible mechanism of cognitive decline in individuals with WML is disruption of cholinergic fibers by strategically located WMLs. Bohnen et al. (2009a) investigated the relationship between WML and cortical cholinergic AChE activity in 18 subjects without dementia ranging in age from 55 to 84 years. The severity of periventricular WML but not non-periventricular WMH was inversely related to global cortical AChE activity. There was an inverse correlation between cortical AChE activity and reaction time in Conners Continuous Performance Test. On the way from the nucleus basalis of Meynert, cholinergic fibers are initially tightly packed in bundles, fanning out toward the cortex after passing the basal ganglia and ventricles (Selden et al. 1998). In proximity to the ventricles, their density is much higher than in deep white matter, which could explain why only periventricular white matter is related to cortical AChE activity.

Richter et al. (2017) investigated the relationship between periventricular WML, cortical AChE activity, and memory function in 18 patients with MCI due to AD and 18 cognitively normal older participants. Only periventricular, not non-periventricular, WML load negatively correlated with cortical AChE activity in both groups. AChE activity predicted memory function better than WML load, suggesting that the effects of WML load on memory were fully mediated by AChE activity.

AChE PET allows a direct measurement of the degree of AChE inhibition by therapeutic AChE inhibitors. Kuhl et al. (2000) found that mean AChE inhibition in the cerebral cortex of AD patients was  $26 \pm 12\%$  and  $27 \pm 11\%$  after donepezil doses of 5 or 10 mg/day for at least 5 weeks. Since 80–90% red blood cell AChE inhibition in humans by these doses of donepezil had been reported, they concluded that AChE inhibition in vivo in living Alzheimer cerebral cortex by these doses of donepezil was limited. We found that 3–5 mg/day donepezil treatment reduced cortical AChE activity by 29–39% in three AD patients (Shinotoh et al. 2001). Bohnen et al. (2005b) observed a correlation between the degree of cortical AChE inhibition and changes in executive and attentional function by donepezil treatment in AD patients in a [ $^{11}\text{C}$ ]PMP study. Kaasinen et al. (2002) reported a similar inhibition of cortical AChE activity between 10 mg/day donepezil and 9 mg/day rivastigmine (28–39% vs. 28–37%). They also found that the pooled effects of donepezil and rivastigmine are greater in the frontal cortex compared to the temporal cortex in AD, which could be attributed to the prominent temporoparietal reduction of AChE in AD. Kadir et al. (2008) observed 30–40% inhibition of cortical AChE activity after 16–24 mg/day of galantamine treatment.

Dementia with Lewy bodies (DLB) is the second most frequent neurodegenerative dementia after AD. We measured brain AChE activity in 14 DLB patients, 25 late-onset AD patients, and 18 age-matched healthy controls (Shimada et al. 2015).

Cortical AChE activity was more prominently reduced in DLB patients (27.8%) than late-onset AD patients (8.2%). Regional brain AChE activity was well able to discriminate DLB from late-onset AD.

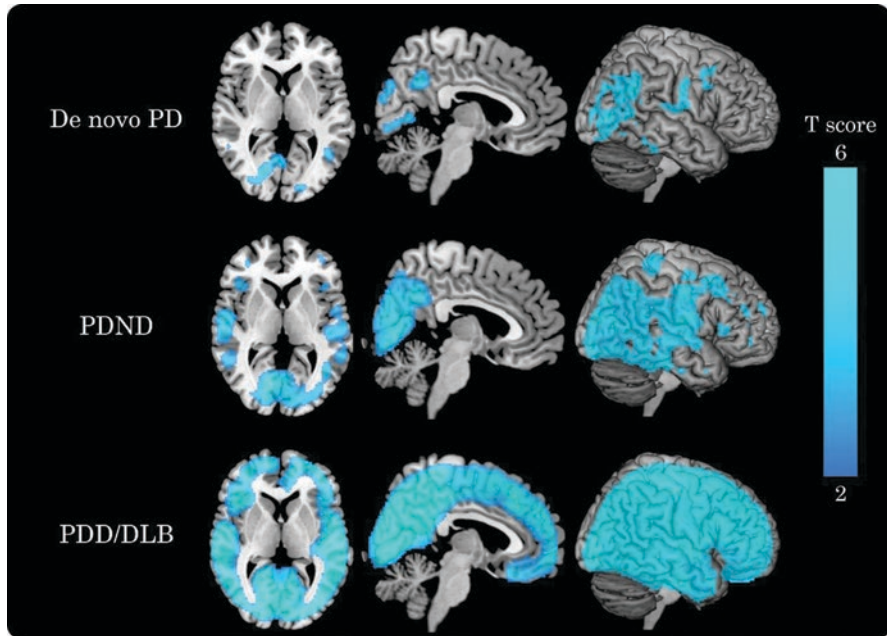
Perry et al. (1978) found that BuChE activity is increased by 40% in the cerebral cortex of postmortem AD brains. It has not been settled, however, whether cholinesterase inhibitor therapy should target BuChE in AD, not AChE alone, because both enzymes hydrolyze acetylcholine. Snyder et al. (2001) investigated a series of 1-methyl-4-piperidinyl esters, and identified 1-[<sup>11</sup>C]methyl-4-piperidinyl n-butyrate ([<sup>11</sup>C]BMP) as a most promising ligand for the in vivo measurement of cerebral BuChE activity. Kuhl et al. (2006) performed [<sup>11</sup>C]BMP PET scan in 15 AD patients and 12 healthy controls. Contrary to prediction, cerebral BuChE activity was decreased by 18%. BuChE in the neuritic plaques is considered to account for the incremental BuChE observed in postmortem AD cerebral cortex tissue homogenates. The PET method would underestimate BuChE activity in plaques, because delivery of [<sup>11</sup>C]BMP tracer to the plaque enzyme would be limited.

### 7.3.3 Parkinson's Disease and Related Disorders

Neurochemical analysis of postmortem brains with PD in the 1980s revealed a decrement of AChE activity in association with ChAT activity (Ruberg et al. 1986). Since then, however, cholinergic deficits in PD had not drawn much attention. PET imaging of AChE has elucidated the role of cholinergic dysfunction in PD and related disorders in the last 20 years.

We measured brain AChE activity in 16 patients with PD, 12 patients with progressive supranuclear palsy (PSP), and 13 age-matched healthy controls by [<sup>11</sup>C]MP4A PET (Shinotoh et al. 1999). We found a significant reduction (17%) of cortical AChE activity associated with a nonsignificant reduction (13%) of thalamic AChE activity in PD patients compared with healthy controls. PD patients with dementia and hallucination showed more prominent reduction of cortical AChE activity than PD without dementia. PSP patients showed a nonsignificant reduction (10%) of cortical AChE activity but a prominent reduction (38%) of thalamic AChE activity. There is a second major set of cholinergic nuclei in the brainstem, the pedunculopontine nucleus, and the laterodorsal tegmental nucleus, which gives rise to a major ascending projection to the thalamus (Mesulam et al. 1984; Tubert et al. 2019). The pedunculopontine nucleus plays a major role in the control of gait (Tubert et al. 2019). The prominent reduction of thalamic AChE activity suggests that the ascending cholinergic system from the brainstem to the thalamus is severely impaired in PSP, probably leading to gait instability.

By voxel-by-voxel analysis of AChE activity, we found a significant reduction (12%) in the cerebral cortex, and especially in the medial occipital cortex in PD patients ( $n = 18$ ) in comparison with healthy controls ( $n = 26$ ) (Fig. 7.6) (Shimada et al. 2009). AChE activity reduction was seen in the medial occipital cortex even in de novo PD patients ( $n = 5$ ), suggesting that cholinergic denervation occurs early in



**Fig. 7.6** Cortical AChE activity reduction in de novo Parkinson's disease (PD) ( $n = 5$ ), PD without dementia (PDND) ( $n = 18$ ), PD with dementia (PDD), and dementia with Lewy bodies (DLB) ( $n = 21$ ), compared with 26 healthy controls. Statistical parametric T-score maps with blue color representing AChE activity reduction are superimposed on SPM T1 MRI template, thresholded at voxel-level  $p < 0.01$ ; false discovery rate (FDR) corrected with cluster above 26 voxels, age as covariate

PD. There was more profound (27%) and widespread reduction of cortical AChE activity in both PDD and DLB patients.

Bohnen et al. (2003) found that cortical AChE activity is more severely affected in PD patients with dementia (20%,  $n = 14$ ) than in patients with mild AD (9.1%,  $n = 12$ ) relative to healthy controls ( $n = 10$ ) in a [ $^{11}\text{C}$ ]PMP PET study with shape analysis. PD patients without dementia showed intermediate cortical AChE activity reduction (12.9%,  $n = 11$ ).

Hilker et al. (2005) found severe reduction (29.7%) of cortical AChE activity in PDD ( $n = 10$ ) and moderate reduction of cortical AChE activity in PD ( $n = 17$ ) in comparison with healthy controls ( $n = 31$ ) in their [ $^{11}\text{C}$ ]MP4A-PET study. They also found a significant covariance of striatal reduction of [ $^{18}\text{F}$ ]fluorodopa uptake and decreased cortical AChE activity in PDD, suggesting a common disease process leading to a transmitter deficiency syndrome in PDD.

Bohnen et al. (2009b) compared brain ACh activity between 17 PD patients with a history of falls and 27 PD patients without such a history. Cortical AChE activity was reduced in the PD fallers group (12.3%), followed by PD non-fallers (6.6%), compared with healthy controls ( $n = 15$ ). Thalamic AChE activity was lower only in the PD fallers (11.8%), suggesting that impairment of the pedunculopontine nucleus

contributes to postural control and gait dysfunction in PD, like in PSP. In another study, Bohnen et al. (2013) also found that low cortical AChE activity was associated with slow gait speed. Subsequently, Chung et al. (2010) did clinical trials of donepezil, a central cholinesterase inhibitor, in 23 PD patients who reported falling or nearly falling more than twice per week and found that subjects with PD fell approximately half as often during the 6 weeks on donepezil than on placebo. A randomized, double-blind, placebo-controlled, phase 2 trial of rivastigmine in 130 PD patients showed that PD patients who were assigned to rivastigmine had improved gait stability and possible reduction in the frequency of falls (Henderson et al. 2016).

Bohnen et al. (2010) investigated olfactory dysfunction, brain AChE activity, and cognitive impairment in 58 PD patients, and found that odor identification scores correlated positively with AChE activity in hippocampal, amygdala, and cortical AChE activity, and scores on cognitive measures of episodic verbal learning. The results suggest that greater deficits in odor identification may identify patients with PD at risk for cognitive impairment. Bohnen et al. found heterogeneity of cholinergic denervation in PD. (Bohnen et al. 2012) They found that cortical cholinergic denervation frequency increased monotonically with increasing cognitive impairment in 143 non-demented PD patients (Bohnen et al. 2015). They investigated the relationship of brain AChE activity with depressive symptoms, rapid eye movement sleep behavior disorder, freezing of gait, risky driving, and fatigue (Bohnen et al. 2007, 2014, 2015; Kotagal et al. 2012; Müller et al. 2015; Weathers et al. 2014; Chou et al. 2016).

We investigated brain AChE activity in corticobasal syndrome and found a significant reduction of cortical (17.5%) but not thalamic AChE activity by [<sup>11</sup>C]MP4A PET (Hirano et al. 2010). AChE activity was reduced in the thalamus and the posterior lobe of the cerebellar cortex but not in the cerebral cortex in seven patients with cerebellar syndrome of multiple system atrophy (MSA-C) (Hirano et al. 2008). Gilman et al. (2010) found prominent reduction of AChE activity in the thalamus, mesencephalon, pons, cerebellum, and caudate putamen and mild reduction in the cerebral cortex in patients with parkinsonian syndrome of multiple system atrophy (MSA-P) in a [<sup>11</sup>C]PMP PET study.

Two symptomatic patients with frontotemporal dementia with parkinsonism linked to chromosome 17 (FTDP-17) carrying N279K mutation showed prominent reduction of cortical and thalamic AChE activity (Hirano et al. 2006). One of the three asymptomatic gene carriers with N279K mutation showed a decrease in temporal and parietal AChE activity (Miyoshi et al. 2010).

[<sup>11</sup>C]donepezil is used for assessment of the autonomic nervous system in the abdomen in PD. Gjerløff et al. (2015) found reduction of [<sup>11</sup>C]donepezil binding in the small intestine (35%) and pancreas (22%) in 12 PD patients. Reduction of [<sup>11</sup>C]donepezil binding is also seen in early PD patients (Fedorova et al. 2017).

Autosomal dominant inherited PD caused by LRRK2 mutations is one of the commonest forms of genetic parkinsonism. Liu et al. (2018) found that LRRK2 mutation carriers without PD ( $n = 16$ ) had increased cortical AChE activity compared with healthy controls ( $n = 11$ ). PD patients with LRRK2 mutations had

significantly higher AChE activity in some cortical regions and thalamus compared with idiopathic PD patients. Changes in cholinergic activity might represent early and sustained attempts to compensate for LRRRK2-related dysfunction. Another possible cause for increased AChE activity might represent alteration of AChE activity in non-neuronal cells.

### 7.3.4 Other Disorders

Östberg et al. (2018) investigated cholinergic function in 17 subjects with traumatic brain injury (TBI). They found significant reduction of neocortical AChE activity in TBI subjects compared to healthy controls ( $n = 12$ ).

Patients with multiple sclerosis (MS) often present with cognitive impairment. Virta et al. (2011) investigated the brain cholinergic system in ten MS patients with marked cognitive impairment by [ $^{11}\text{C}$ ]MP4A PET. There were no differences in cortical AChE activity between the MS patients and ten healthy controls.

AChE is expressed in the inflamed vessel wall of patients with giant-cell arteritis. Two patients with large-vessel giant-cell arteritis were scanned with [ $^{11}\text{C}$ ]donepezil and [ $^{18}\text{F}$ ]fluorodeoxy glucose. PET scan revealed extensive vascular FDG uptake but no [ $^{11}\text{C}$ ]donepezil (Therkildsen et al. 2019).

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## 7.4 Conclusions

1. *N*-methylpiperidyl esters have been extensively characterized as synthetic substrates for AChE.
2. Carbon-11-labeled analogues of two of these esters, [ $^{11}\text{C}$ ]MP4A and [ $^{11}\text{C}$ ]PMP, are in successful clinical use for in vivo measurement of AChE activity. These radiotracers give precise estimates of AChE activity in brain regions with low enzyme activity because of its bolus-like input function shape, complete irreversibility, and near-optimal kinetic properties. It is challenging to estimate AChE activity in regions with high enzyme activity such as the striatum.
3. [ $^{11}\text{C}$ ]MP4A and [ $^{11}\text{C}$ ]PMP opened a window on cerebral cholinergic function (Reed and Jagust 1999) and have elucidated the role of the cholinergic system in AD, PD, and related disorders.

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# Imaging of Adenosine Receptors

# 8

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## 8.1 Introduction

Adenosine is a fundamental molecule of life. It is a part of the DNA and the main degradation product of the central currency of energy metabolism in humans and animals—adenosine triphosphate (ATP). The nucleoside adenosine is composed of the nucleobase adenine and the sugar ribose. It has a short biological half-life of about 1–2 s and the endogenous extracellular level of adenosine is in the range of several hundred nano mol in heart and brain of humans (Zeitzer et al. 2006). Adenosine is formed by breakdown of ATP to ADP to AMP and finally to adenosine by the enzyme 5'-nucleotidase. Adenosine is metabolized by adenosine deaminase to inosine and mainly by

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phosphorylation by adenosine kinase to AMP. Extra- and intracellular concentrations of adenosine can equilibrate quickly by equilibrative nucleoside transporters. Under pathological conditions like hypoxia the adenosine concentration can rise severalfold—up to micromolar concentrations. Adenosine is a part of the heterogeneous group of neuromodulators. In contrast to classical neurotransmitters it is not stored and released from vesicles. The net effect of adenosine on excitable tissue is inhibitory affecting the release of classical neurotransmitters like glutamate, GABA (gamma-aminobutyric acid), and dopamine. Adenosine, for example, reduces the contraction frequency of a tachycardic heart or reduces epileptic activity in the brain. The most prominent and most widespread used substance that interacts with adenosine in daily life is caffeine. The widely used neurostimulant caffeine exerts its effects as an antagonist at adenosine receptors. Four different types of adenosine receptors have been described in mammals:  $A_1$ ,  $A_{2A}$ ,  $A_{2B}$ , and  $A_3$  which are all G-protein-coupled receptors (Fredholm et al. 2001; Hess 2001; Palmer and Stiles 1995). The coupling of  $A_1$  and  $A_3$  to a  $G_i$ -protein leads to inhibition of adenylate cyclase whereas the coupling of  $A_{2A}$  and  $A_{2B}$  to  $G_s$ - or  $G_o$ -proteins leads to stimulation of adenylate cyclase. In humans the  $A_1$  adenosine receptor ( $A_1AR$ ) shows the most abundant distribution and highest concentrations (in descending order) in brain, heart, adipose tissue, stomach, testis, spleen, kidney, aorta, liver, eye, and bladder. The  $A_{2A}$  adenosine receptor ( $A_{2A}AR$ ) can be found in striatum, nucleus accumbens, olfactory tubercle, heart, lung, and blood vessels.  $A_{2B}$  adenosine receptors ( $A_{2B}AR$ ) are expressed in low levels in all tissues and  $A_3$  adenosine receptors ( $A_3ARs$ ) show low levels in brain and high levels in lung and liver (Jacobson and Gao 2006). The affinities of endogenous adenosine for the  $A_1AR$  and the  $A_{2A}AR$  are in the nanomolar range whereas the affinities for the  $A_{2B}AR$  and the  $A_3AR$  are in the micromolar range (Dunwiddie and Masino 2001). This discrepancy in affinity in combination with normal adenosine levels in the nanomolar range suggests that  $A_{2B}AR$  and  $A_3AR$  are primarily activated under pathological conditions and massively increased levels of intracellular adenosine. An important aspect is that  $A_{2A}ARs$  form heterodimers with  $A_1ARs$  or dopamine  $D_2$  receptors and  $A_1AR$  with dopamine  $D_1$  receptors which in turn influence the affinity of the adenosine receptors (Ciruela et al. 2006; Franco et al. 2007; Cristovao-Ferreira et al. 2011).

Over the last 30 years adenosine receptor ligands, agonists, as well as antagonists have emerged as a class of useful therapeutics (for a comprehensive review see van Waarde et al. 2018; Jacobson 2009; Jacobson and Gao 2006). The requirements for compounds suitable for noninvasive *in vivo* imaging of adenosine receptors (radiopharmaceuticals, radiotracers) with positron-emission tomography (PET) are in some aspects different from those for therapeutic drugs. This difference is even more pronounced in the case of radiotracers involved in neurotransmission research. One possible process for adenosine receptor ligand development in combination with adequate evaluation methods is depicted in Table 8.1.

In contrast to therapeutically dosed drugs the amounts of intravenously (*i.v.*) administered radiopharmaceuticals are extremely small. Thus, biochemical equilibria are not or only marginally disturbed allowing quantitative receptor scintigraphy (determination of concentration of binding sites,  $B_{max}$ ).

Depending on the amount of applied radioactivity and on the molar activity of the respective radiotracer the applied mass of a PET-radiopharmaceutical typically lies in the range of nano- to micrograms. Thus, even toxic drugs, which are not suitable as therapeutics, can be used as radiotracers for PET.

**Table 8.1** Requirements for a suitable radiotracer

Criterion	Requirements	Evaluation system
Affinity and reversibility	$K_D$ or $K_I$ in nM range	Assays with brain tissue or cell homogenates, autoradiography
Specificity	Selectivity compared to receptors of the same subfamily and to receptors mainly present in the region of interest	
Nonspecific binding potential	Low	
Agonism	Antagonists (usually for receptor quantification; for determination of endogenous ligand concentrations agonistic or partially agonistic properties are useful)	Functional assays, for instance cAMP
Penetration of BBB	>1% ID/g @ $t_{max}$	Organ distribution in animals, animal PET or SPECT
Metabolism	Preferably no metabolism in the target organ (brain) or biologically inactive metabolites	
Bioavailability	Half-life of radionuclide compatible with physiological kinetics Fraction of “free” ligand should be as high as possible (>1%)	
Mutagenicity	Very low	Specific assays for instance chromosome aberration, Ames test
Toxicity	Low << effects thousandfold of applied mass	Theoretically: <i>Quantitative Structure-Activity Relationship</i> Animal experiments LD <sub>50</sub>

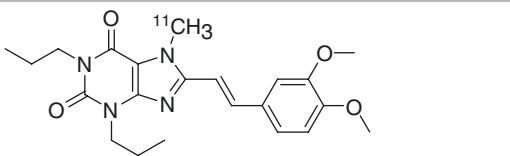
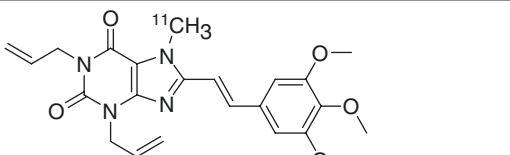
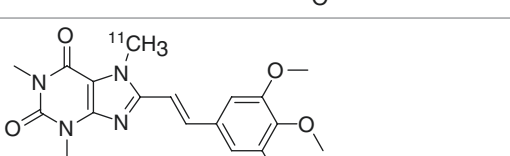
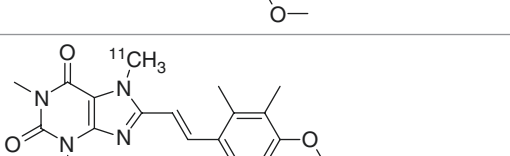
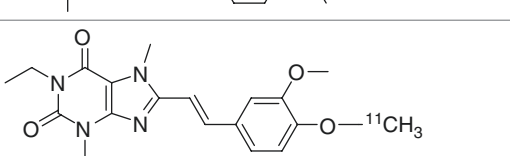
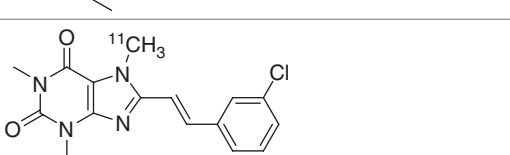
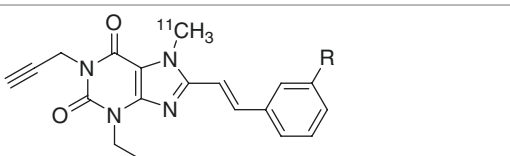
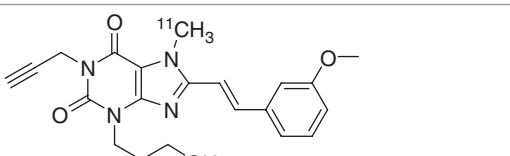
A brain radiotracer, in order to be suitable for imaging, has to fulfill a number of criteria with regard to its physical and physiological properties (Ding and Fowler 2005; Halldin et al. 2001; Mason and Mathis 2003). The ligand should have nanomolar to subnanomolar affinity ( $K_D$ ) and sufficient selectivity for the target receptor. Ideally, binding to the receptor should be reversible over the experimental time range because this allows the use of biomathematical equilibrium approaches for the quantification of ligand-receptor interaction. Lipophilicity ( $\log P$ ) of the tracer should be appropriate for crossing the blood-brain barrier (BBB). For neuroradioligands there is an optimal, rather narrow “window” of lipophilicity in the  $\log P$  range between 1 and 4 (Dishino et al. 1983; Moerlein et al. 1985). As a rule of thumb molecules with a  $\log P$  value <1 will not cross the BBB, while those with a  $\log P > 4$  generally tend to exhibit high nonspecific binding. Other criteria for a successful radiotracer are sufficient in vivo stability in plasma and, in respect to tracer availability, low affinity for plasma proteins. With regard to image analysis, there should be no radiolabeled metabolites which are capable of crossing the BBB. Further, a suitable radiotracer should show a kinetic profile that allows adequate quantification of the obtained data.

Radiolabeled subtype-specific adenosine receptor ligands currently under development are depicted in Table 8.2.

**Table 8.2** Selection of PET ligands for adenosine receptors

A <sub>1</sub>	R=CH <sub>3</sub> CH <sub>2</sub> <sup>11</sup> CH <sub>2</sub> , [ <sup>11</sup> C]KF15372 R=CH <sub>3</sub> <sup>11</sup> CH <sub>2</sub> , [ <sup>11</sup> C]EDPX R= <sup>11</sup> CH <sub>3</sub> , [ <sup>11</sup> C]MDPX (Ishiwata et al. 1995; Noguchi et al. 1997)	
	[ <sup>18</sup> F]CPFPX (Holschbach et al. 2002)	
	R=H, [ <sup>18</sup> F]CBX R=CH <sub>3</sub> , [ <sup>18</sup> F]1-MCBX (Kreft et al. 2017)	
	R=H, [ <sup>18</sup> F]CBX R=CH <sub>3</sub> , [ <sup>18</sup> F]1-MCBX (Schneider et al. 2019a, b)	
	[ <sup>11</sup> C]FR194921 (Matsuya et al. 2005)	
	[ <sup>131</sup> I]CPIPX (Sihver et al. 2003)	
	[ <sup>11</sup> C]22b A <sub>1</sub> AR partial agonist radioligand (Guo et al. 2018)	

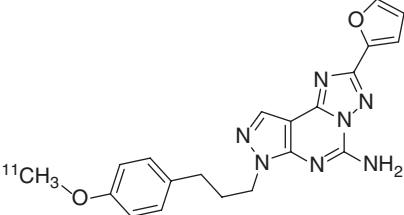
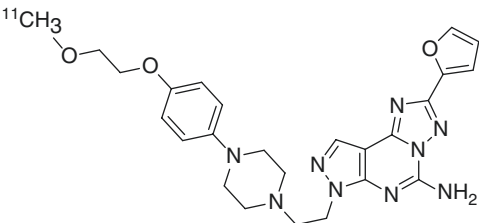
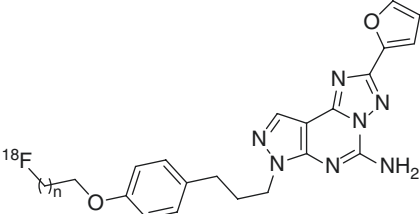
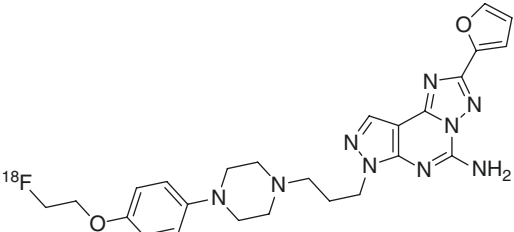
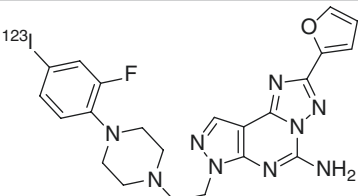
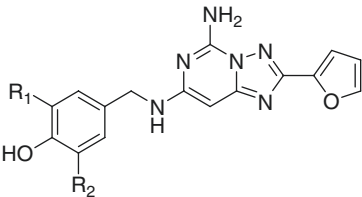
**Table 8.2** (continued)

A <sub>2A</sub>	[ <sup>11</sup> C]KF117837 (Ishiwata et al. 1996)	
	[ <sup>11</sup> C]KF19631 (Ishiwata et al. 2000a, b)	
	[ <sup>11</sup> C]KF18446, [ <sup>11</sup> C]TMSX (Ishiwata et al. 2000a, b, c, d)	
	[ <sup>11</sup> C]KF21213 (Wang et al. 2000)	
	[ <sup>11</sup> C]KW6002 (Hirani et al. 2001)	
	[ <sup>11</sup> C]CSC (Ishiwata et al. 2000a, b, c, d; Marian et al. 1999)	
	R=Br, [ <sup>11</sup> C]BS-DMPX R=I, [ <sup>11</sup> C]IS-DMPX (Ishiwata et al. 2000a, b, c, d)	
	[ <sup>11</sup> C]MSX-2 (Holschbach et al., 2000)	

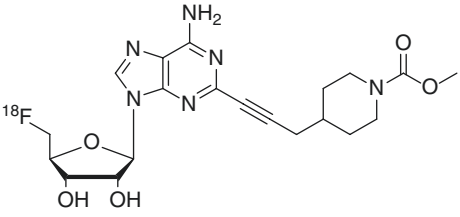
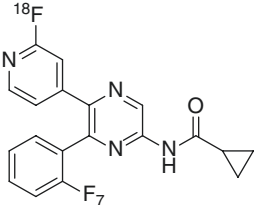
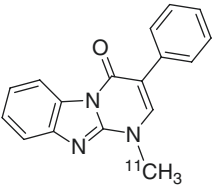
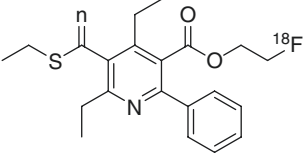
(continued)



**Table 8.2** (continued)

<p>[<sup>11</sup>C]SCH442416 (Moresco et al. 2005; Todde et al. 2000)</p>	
<p>[<sup>11</sup>C]SCH420814, [<sup>11</sup>C]Preladenant (Zhou et al. 2014)</p>	
<p><math>n = 1</math>, [<sup>18</sup>F]fluoroethyl SCH420814  <math>n = 2</math>, [<sup>18</sup>F]fluoropropyl SCH420814 (Khanapur et al. 2014a, b)</p>	
<p>[<sup>18</sup>F]MNI-444 (Barret et al. 2015; Vala et al. 2016)</p>	
<p>[<sup>123</sup>I]MI-420 (Vala et al. 2016; Tavares et al. 2013)</p>	
<p><math>R_1 = H, R_2 = ^{131}I</math>: [<sup>131</sup>I]iodo-ZM241385  <math>R_1 = I, R_2 = ^{131}I</math>: [<sup>131</sup>I]diiodo-ZM241385 (Poucher et al. 1995; Sihver et al. 2004)</p>	

**Table 8.2** (continued)

	[ <sup>18</sup> F]FDA-PP1-labeled A <sub>2A</sub> AR agonist (enzymatic fluorination) (Lowe et al. 2017)	
A <sub>2B</sub>	[ <sup>18</sup> F]7a-pyrazine based radioligand (Lindemann et al. 2018)	
	[ <sup>11</sup> C]-labeled triazinobenzimidazole radioligand (Petroni et al. 2016)	
A <sub>3</sub>	[ <sup>18</sup> F]FE@SUPPY (Wadsak et al. 2008)	

A<sub>1</sub>AR and A<sub>2A</sub>AR antagonists are either xanthines or non-xanthine monocyclic or fused bi- or tricyclic substituted heteroaromatic compounds, some of them containing hetero ring atoms such as nitrogen, oxygen, and sulfur.

Three characteristics identify xanthines that are potent and selective antagonists at the A<sub>1</sub>AR. They possess long alkyl chains in positions 1 and 3 as well as a bulky (bi)cycloalkyl substituent at C-8, with the optimum substituents at N-1- and N-3 being propyl groups. Additionally, N-7 must be unsubstituted in order to act as a hydrogen bond donor, a feature which is essential for binding to the A<sub>1</sub>AR. Whereas some modifications of the 1- and 3-substituents (fluorination, iodination, substitution of an aralkyl group at N-3) preserve or at least do not seriously affect activity, the criteria based on N-7 and C-8 are required for antagonist activity at the A<sub>1</sub>AR.

Xanthine-type A<sub>2A</sub>AR antagonists with high A<sub>2A</sub>AR affinity contain a (substituted) (*E*)-styryl substituent at C-8 in combination with alkyl/alkynyl moieties at N-1 and N-3; a methyl group at N-7 reduces affinity to the A<sub>1</sub>AR by suppressing the formation of hydrogen bonds and can sterically interact with the styryl group by

forcing the vinylic double bond into a favored conformation for binding to the  $A_{2A}$ AR.

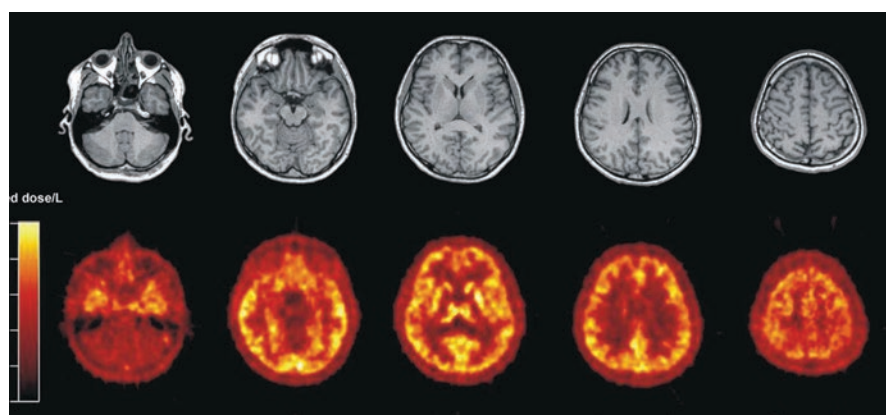
Non-xanthine  $A_1$ AR antagonists are often structurally based on adenine, having a 6-amino group substituted with bulky cycloaliphatic moieties and variable nitrogen atoms in the heterocyclic core.

Nitrogen polyheterocyclic  $A_{2A}$ AR ligands, especially those based on a triazolopyrimidine structure, display good affinity and selectivity when they bear a furan moiety of the core aromatic scaffold, a prerequisite which is crucial for binding. Moreover, bi- and tricyclic fused nitrogen heterocyclic antagonists, in order to be active, often contain a long-chain aralkyl moiety. PET ligands for adenosine receptors at different states of development have been reviewed recently (Vuorimaa et al. 2017; Li et al. 2019; Mishina and Ishiwata 2014; Khanapur et al. 2014a, b; Bauer and Ishiwata 2009; Ishiwata et al. 2007a, b).

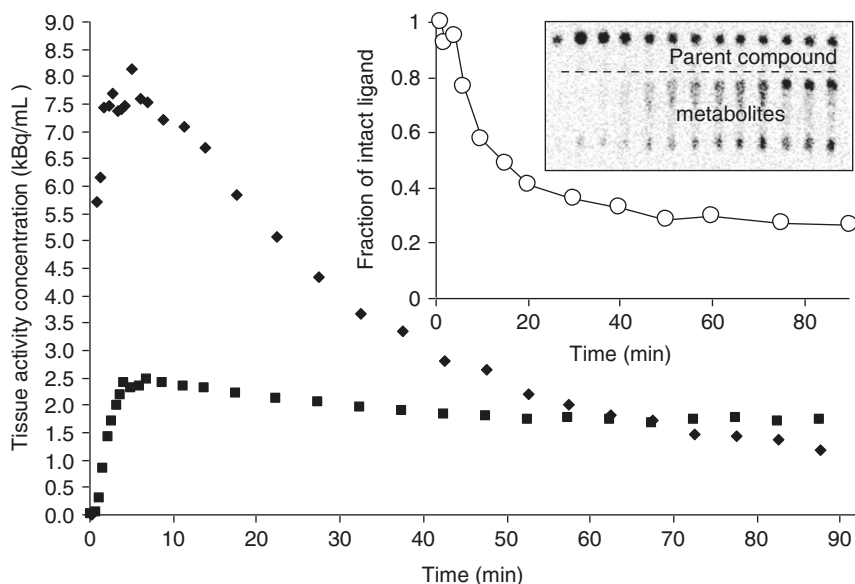
## 8.2 $A_1$ Adenosine Receptor Ligands

So far two PET tracers have been applied and evaluated to image  $A_1$ ARs in humans: [ $^{11}\text{C}$ ]MPDX (Noguchi et al. 1997) and [ $^{18}\text{F}$ ]CPFPX (Holschbach et al. 2002). Both are xanthine derivates. Potential scientific applications and findings for  $A_1$ AR imaging have been reviewed (Paul et al. 2011a, b; Mishina and Ishiwata 2014; Li et al. 2019).

The antagonist [ $^{18}\text{F}$ ]CPFPX ( $\text{Log } P = 2.1$ ) shows a high affinity ( $K_D$  1.26 nM in cells constitutively expressing the human  $A_1$ AR) and high selectivity ( $A_{2A}$ AR affinity 940 nM) for the  $A_1$ AR (Bauer et al. 2003; Holschbach et al. 2002). After i.v. injection it rapidly (within 5 min) reaches a maximum brain uptake of approximately 3% of injected dose (Bauer et al. 2003; Meyer et al. 2004) (Fig. 8.1). Free



**Fig. 8.1** In vivo distribution of [ $^{18}\text{F}$ ]CPFPX: individual MRI (top) and summed (5–60 min postinjection) PET (bottom) of a representative healthy volunteer. Note the high accumulation of [ $^{18}\text{F}$ ]CPFPX in cerebral cortex, basal ganglia, and thalamus



**Fig. 8.2** Examples of representative bolus (diamonds) and bolus plus constant infusion (boxes) experiments in humans with [ $^{18}\text{F}$ ]CPFPX. Depicted is the tissue time-activity curve of the frontal lobe. Steady state is reached after approximately 50 min. Inserted figure shows [ $^{18}\text{F}$ ]CPFPX metabolism, fraction of parent compound of the bolus/infusion experiment, and thin-layer chromatography radioactivity distribution

fraction of parent compound in plasma is about 1–2%. [ $^{18}\text{F}$ ]CPFPX is metabolized fast in the liver by the cytochrome isoenzyme CYP1A2 so that the fraction of parent compound is reduced to approximately 50% after 10 min and to less than 10% after 60 min (Matusch et al. 2006; Meyer et al. 2004). Formed metabolites are hydrophilic and do not cross the blood-brain barrier (>98% unchanged ligand in mouse brain tissue after 60 min) (Holschbach et al. 2002). The chemical structure of seven main metabolites (Fig. 8.2) and their pathways have been described (Holschbach et al. 2009, 2017; Bier et al. 2006). The regional cortical and subcortical distribution of in vivo binding is in agreement with the human *postmortem* distribution in autoradiographic measurements.

The suitability of [ $^{18}\text{F}$ ]CPFPX for clinical applications has been shown in a dosimetry study (Herzog et al. 2008) which revealed that an injected dose of 300 MBq posed a radiation burden of 5.3 mSv on a patient. This dose is comparable to other  $^{18}\text{F}$ -labeled PET tracers in clinical use.

[ $^{18}\text{F}$ ]CPFPX has been administered i.v. as bolus and as bolus plus subsequent infusion (Meyer et al. 2004, 2005a, b). Pharmacokinetic modeling of an injection of a bolus can be described significantly better with a two-compartment model than with a one-compartment model. Nevertheless the individual rate constants were highly correlated leading to high standard errors of the estimates and the derived outcome parameter  $BP_p$  denoting the specific binding (Meyer et al. 2004). The

parameter total distribution volume ( $V_T$ ) is a more valid measure as it combines the specific and nonspecific binding.  $V_T$  can also be determined with Logan's graphical analysis which allows voxel-wise determination and statistical parametric mapping. The noise-dependent bias in  $V_T$  parametric images can be reduced by spatial smoothing before applying the graphical analysis. A relatively short scan duration of 60 min is sufficient (Meyer et al. 2006a). Including the free fraction into the model increases variation and is therefore often omitted. Because of the high fraction of metabolites at late time points a blood volume correction should be applied. It had been shown that not only arterial but also venous input function for the graphical analysis allows reliable quantification of  $V_T$  (Meyer et al. 2005a, b).

Displacement studies with unlabeled CPFPPX revealed that about one-third of cerebellar binding and at least two-thirds of cortical binding (depending on the cortical region) are specific ( $V_S$ ) (Meyer et al. 2006b). [ $^{18}\text{F}$ ]CPFPPX is displaceable by caffeine in rats (Meyer et al. 2003) and humans (Elmenhorst et al. 2012). Therefore it is compulsory that subjects refrain from caffeine intake at least 36 h before participating in a PET study. Due to the selective liver metabolism special caution has to be taken in patients with liver dysfunction where the elimination half-life of caffeine is extremely prolonged (Matusch et al. 2006).

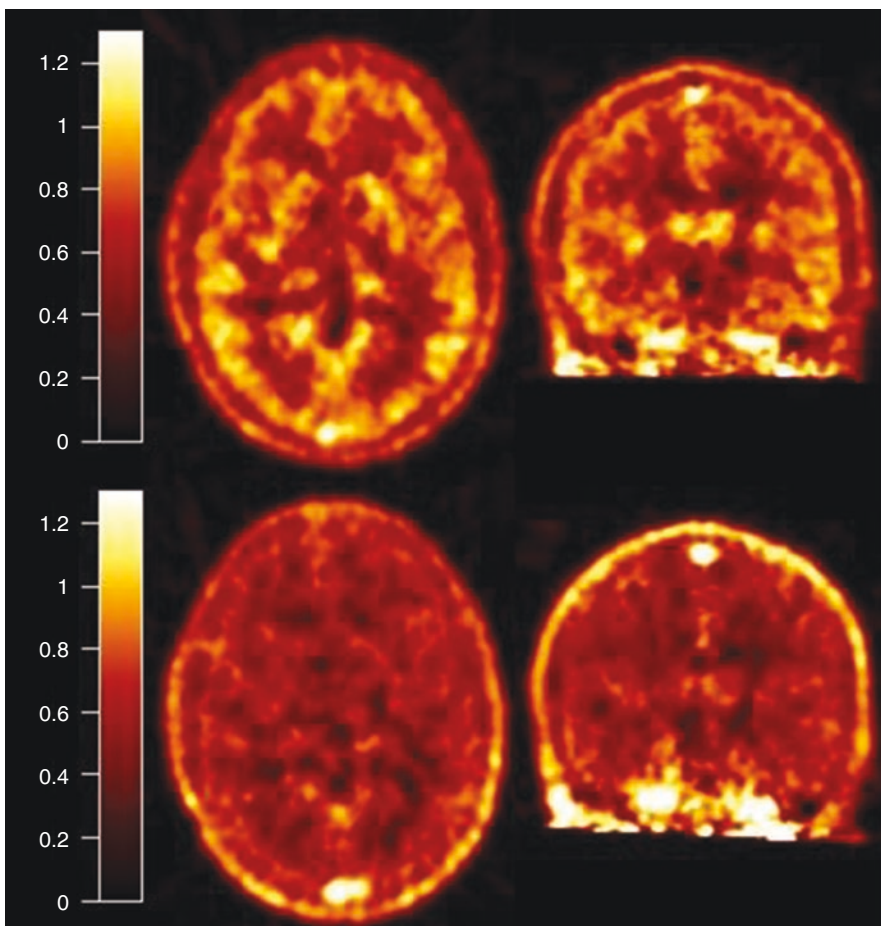
The in vitro inhibitory concentration of caffeine that displaces 50% of the binding ( $IC_{50}$ ) of [ $^3\text{H}$ ]CPFPPX at human brain *postmortem*  $A_1\text{ARs}$  is in the range of 113–170  $\mu\text{M}$  (Elmenhorst et al. 2011). Therefore the dose of caffeine after common consumption of coffee or tea has probably a direct impact on the in vivo quantification of  $A_1\text{ARs}$  with the PET ligand [ $^{18}\text{F}$ ]CPFPPX. Despite the acute effects of caffeine and various contradicting in vitro findings of the long-term effects of caffeine, an effect on  $A_1\text{AR}$  availability was not observed in a rat model of chronic caffeine exposure (30 mg/kg body weight/day, corresponding to 4–5 cups of coffee per day in humans for 12 weeks) (Nabbi-Schroeter et al. 2018).

The bolus plus subsequent infusion method allows quantifying modulations of binding (e.g., displacement) within a single scan. After establishing a steady state of ligand concentration in plasma and brain tissue, the total distribution volume  $V_T$  can be determined as a ratio of the concentrations in brain tissue and plasma. Due to the fast kinetics of [ $^{18}\text{F}$ ]CPFPPX equilibrium can be achieved within approximately 60 min (Meyer et al. 2005a, b).

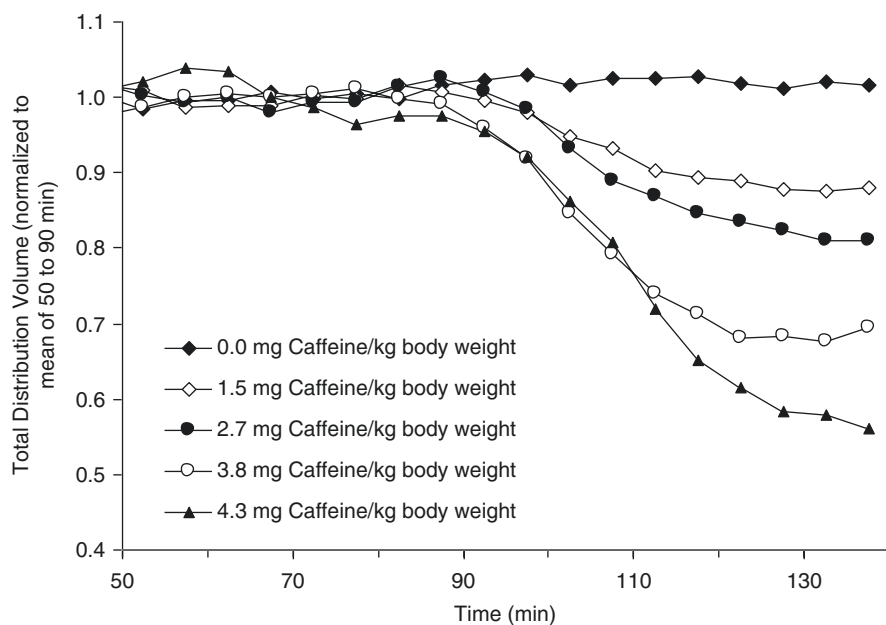
In the case of [ $^{18}\text{F}$ ]CPFPPX the relatively low fraction of specific binding (about 2/3 of  $V_T$ ) reduces the sensitivity for possible changes in  $V_T$ . Another shortcoming is the dependence on error-prone individual plasma input functions, which increases variability. These drawbacks can be avoided by using a reference region input for  $BP_{\text{ND}}$  (specific vs. non-displaceable input) which is independent on blood sampling (Meyer et al. 2006b; Elmenhorst et al. 2007a, b). The cerebellar cortex is used as a reference region. Although the cerebellar cortex is the region with the lowest  $A_1\text{AR}$  density in the human brain, one-third of the  $V_T$  is displaceable with unlabeled CPFPPX (Meyer et al. 2006b). Therefore changes in  $BP_{\text{ND}}$  deserve careful consideration with regard to their dependency on  $V_S$  changes in the reference region.  $BP_{\text{ND}}$  can be determined by Logan's noninvasive graphical analysis. Test-retest evaluation of the variability and reproducibility revealed that reference-based measures were

superior to those based on plasma input function (Elmenhorst et al. 2007a, b). Exemplary values for  $V_T$  and  $BP_{ND}$  in the frontal cortex are 0.80 mL/mL and 0.92, respectively.

In an approach to determine the in vivo  $IC_{50}$  value of caffeine at the  $A_1AR$  in humans 18 subjects underwent bolus plus constant-infusion PET experiments after caffeine abstinence. Caffeine in varying concentrations (0.5–4.3 mg/kg body weight) was administered during the steady-state phase of ligand delivery (Fig. 8.3). [ $^{18}F$ ]CPFPX binding was displaced in a concentration-dependent manner ranging from 5 to 44% (Fig. 8.4). No displacement (0.3%) was found after placebo administration.  $IC_{50}$  was estimated to be 65  $\mu M$  in plasma which corresponds to 460 mg in a 70 kg subject (approximately 4.5 cups of coffee). Given both the biological



**Fig. 8.3** Equilibrium imaging of caffeine action: parametric total distribution volume PET images before (50–90 min, top) and after caffeine infusion (100–140 min, bottom) of a subject receiving 4.3 mg caffeine per kg body weight

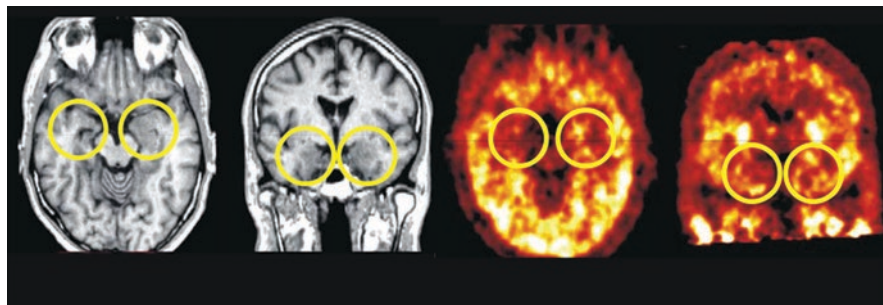


**Fig. 8.4** Displacement of [ $^{18}\text{F}$ ]CPFPX  $A_1\text{AR}$  binding by caffeine under equilibrium conditions (bolus/infusion PET scanning): Representative [ $^{18}\text{F}$ ]CPFPX displacement studies employing increasing doses of caffeine. Intravenous infusion of 0.0–4.3 mg caffeine per kg body weight (equivalent to approximately 0–4 cups of coffee) between 90 and 100 min. Note the rapid and dose-dependent displacement of [ $^{18}\text{F}$ ]CPFPX (x-axis: time after start of bolus/infusion study, y-axis: total volume of distribution normalized to the mean of 50–90 min)

half-life of caffeine of about 5 h and the repeated consumption of caffeinated beverages during a day, occupancies of 50% of the cerebral  $A_1\text{AR}$  are probably common conditions in caffeine consumers (Elmenhorst et al. 2012).

The  $A_1\text{AR}$  is known to be involved in numerous physiological and pathophysiological processes. Age-related changes were investigated over a time span of approximately 50 years showing a range of decline of  $BP_{\text{ND}}$  between  $-17\%$  (putamen) and  $-34\%$  (postcentral gyrus) (Meyer et al. 2006c). Further, it is hypothesized that adenosine is involved in the induction of sleep after prolonged wakefulness. 24 and 52 h of sleep deprivation lead to increased  $V_T$  values in a region-specific pattern in the brain with a maximum increase in the orbitofrontal cortex (15.3%). The upregulation in cortical and subcortical brain regions after prolonged wakefulness is likely to indicate that  $A_1\text{AR}$  expression is contributing to the homeostatic sleep regulation. Furthermore, correlations with psychomotor vigilance underscore the importance of  $A_1\text{AR}$  for cognitive performance (Elmenhorst et al. 2007a, b, 2017, 2018).

Interestingly ethanol exposure leads to an increase in  $A_1\text{AR}$  availability in humans measured with [ $^{18}\text{F}$ ]CPFPX (Elmenhorst et al. 2018) which confirms previous reports of in vitro and in vivo experiments in rodents. In rats a 35–55% increase in [ $^{11}\text{C}$ ]MPDX distribution volume was measured shortly after ethanol exposure along with an adenosine kinase inhibitor (Paul et al. 2011a, b). Recent results from



**Fig. 8.5** Corresponding MRI (left) and [ $^{18}\text{F}$ ]CPFPX PET (right) planes of a patient suffering from mesial temporal lobe epilepsy (focus in the right hemisphere). Note the considerably reduced  $A_1\text{AR}$  binding in the left mesial temporal lobe

mice suggest that sleep deprivation attenuates the sensitivity to ethanol via reduction of extracellular adenosine accumulation. The magnitude of the 35% increase in cerebral [ $^{18}\text{F}$ ]CPFPX  $V_T$  after ethanol is comparable to a respective decrease observed after four cups of espresso.

In a pilot study on a patient with recurrent glioblastoma multiforme increases in  $A_1\text{AR}$  density in the direct vicinity of the tumor confirmed corresponding results in a F98 glioma-bearing rat model (Bauer et al. 2005). In a case of temporal lobe epilepsy there was evidence that  $A_1\text{AR}$  binding is reduced in the epileptic focus (Fig. 8.5).

The inverse antagonist [ $^{11}\text{C}$ ]MPDX ( $\log P < 2.2$ ) (Ishiwata et al. 1995) shows a slightly lower affinity ( $K_i$  4.2 nM) (Noguchi et al. 1997) for the  $A_1\text{AR}$  compared to [ $^{18}\text{F}$ ]CPFPX. After i.v. injection it rapidly (within 2.5 min) reaches a maximum brain uptake of approximately 6% of injected dose (Kimura et al. 2004). [ $^{11}\text{C}$ ]MPDX is metabolized slowly, so that the fraction of parent compound is reduced to approximately 75% after 60 min. The relative density distribution pattern of [ $^{11}\text{C}$ ]MPDX is comparable to [ $^{18}\text{F}$ ]CPFPX. Pharmacokinetic modeling of  $V_T$  was evaluated and performed with both a two-tissue-compartment model (with  $K_1/k_2$  parameter estimates based on cerebellum as a reference region) and a graphical approach in the time span between 10 and 60 min (Logan plot) (Fukumitsu et al. 2003). Using an arterial input function with and without metabolite correction produced only a small negative bias of 4%. Parametric  $BP_{ND}$  images based on  $V_T$  determined by graphical analysis and cerebellum as reference region have been published (Fukumitsu et al. 2003). A blocking experiment in cats with DPCPX revealed that about 53% of  $V_T$  was specific binding. Exemplary values for  $V_T$  and  $BP_{ND}$  in the frontal cortex are 0.82 mL/mL and 0.23, respectively (Kimura et al. 2004). Interestingly, the radioactivity concentration of [ $^{11}\text{C}$ ]MPDX in whole blood was slightly higher than in plasma suggesting an uptake in or binding to red blood cells (Fukumitsu et al. 2005). A noninvasive method to determine the outcome parameters based on independent component analysis has been proposed. Plasma time-activity curves could be estimated from dynamic PET images because the slow metabolism of [ $^{11}\text{C}$ ]MPDX allowed to disregard metabolite correction. P-gp regulates the ability of endogenous and exogenous compounds to cross the BBB. Pretreatment of mice with the P-gp



blocker cyclosporine A before [ $^{11}\text{C}$ ]TMSX and [ $^{11}\text{C}$ ]MPDX PET increased both the blood-to-brain ratio and the cerebral uptake (Ishiwata et al. 2007a, b). [ $^{11}\text{C}$ ]TMSX and [ $^{11}\text{C}$ ]MPDX binding are therefore dependent on P-gp function (at least in mice).

The suitability of [ $^{18}\text{F}$ ]CPFPX and [ $^{11}\text{C}$ ]MPDX for investigating cerebral  $\text{A}_1\text{ARs}$  in longitudinal study protocols has been reported repeatedly for  $\text{A}_1\text{AR}$  availability and occupancy in a reliable (test-retest) and quantitative (pharmacokinetic modeling) manner in rats (Elmenhorst et al. 2013; Kroll et al. 2014; Paul et al. 2014a, b; Sijbesma et al. 2016).

PET imaging of  $\text{A}_1\text{ARs}$  has been evaluated as a tool to predict the severity of ischemic cerebral insults. In 13 cats the right middle cerebral artery was occluded transiently. [ $^{11}\text{C}$ ]MPDX uptake was significantly lower in those animals with a severe ischemic damage and the authors concluded that the binding after reperfusion was a sensitive predictor of severity (Nariai et al. 2003). Stimulation of cerebral  $\text{A}_1\text{AR}$  dampens microglial activation after tissue trauma and has neuroprotective effects in animal models. Neuroinflammatory processes (HSV-1 infection) increased [ $^{11}\text{C}$ ]MPDX uptake in infected regions in concordance with immunohistochemical analyses (Paul et al. 2014).

The previously mentioned decline of adenosine receptor densities in human brains with increasing age measured with [ $^{18}\text{F}$ ]CPFPX was replicated for the  $\text{A}_1$  subtype with [ $^{11}\text{C}$ ]MPDX (Mishina et al. 2012, 2017a, b). Interestingly, in the same study no effect of aging on the  $\text{A}_{2\text{A}}\text{AR}$  was found. In an explorative study though, a putative interaction of  $\text{A}_{2\text{A}}\text{ARs}$  and  $\text{A}_1\text{ARs}$  in an inverse manner was observed between carriers of specific single-nucleotide polymorphisms in the  $\text{A}_{2\text{A}}\text{AR}$  gene who showed a regionally higher  $\text{A}_1\text{AR}$  density (Hohoff et al. 2014).

$\text{A}_1\text{ARs}$  have been investigated so far in various neurodegenerative diseases: In patients suffering from Alzheimer's disease a decreased binding of [ $^{11}\text{C}$ ]MPDX was detected in temporal cortex and thalamus (Fukumitsu et al. 2008). As the pattern of binding differs from the widely used [ $^{18}\text{F}$ ]FDG, adenosine receptor imaging might provide an additional diagnostic value. A cross sectional investigation of Huntington's disease patients indicated a switch of  $\text{A}_1\text{AR}$  availability from supra- to subnormal levels during phenoconversion of the disease (Matusch et al. 2014). [ $^{11}\text{C}$ ]MPDX PET in 10 drug-naïve patients with early Parkinson's disease did not show altered  $\text{A}_1\text{AR}$  in comparison to controls (Mishina et al. 2017). In patients in chronic stages of traumatic brain injury (diffuse axonal injury,  $n = 10$ ) an increased [ $^{11}\text{C}$ ]MPDX  $\text{BP}_{\text{ND}}$  was measured in lower frontal and posterior cingulate cortex and the Rolandic area (Hayashi et al. 2018). The recent preclinical evaluation of [ $^{11}\text{C}$ ]22b, a  $\text{A}_1\text{AR}$  partial agonist PET radiotracer, showed besides suitable uptake, region-specific distribution, selective blocking, and test-retest stability the promise of detecting endogenous adenosine fluctuations (Guo et al. 2018). This feature might open up a new field of in vivo adenosine imaging.

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### 8.3 $\text{A}_{2\text{A}}$ Adenosine Receptor Ligands

The  $\text{A}_{2\text{A}}\text{AR}$  antagonist [ $^{11}\text{C}$ ]TMSX (calculated  $\log P = 1.1$ ) (Ishiwata et al. 2000a, b, c, d) was the first radioligand to visualize  $\text{A}_{2\text{A}}\text{ARs}$  in humans (Ishiwata et al. 2005). The compound presents a high affinity to  $\text{A}_{2\text{A}}\text{ARs}$  ( $K_1 = 5.9$  nM) and a high

selectivity compared to the  $A_1AR$  ( $K_i = 1600$  nM) (Ishiwata et al. 2003). It was confirmed in vitro that TMSX has a very low affinity for 13 different neuroreceptors (Ishiwata et al. 2000a, b, c, d). After i.v. injection it rapidly (within 1.5–2.5 min) reaches a maximum brain uptake of approximately 1–2% of injected dose (estimated from the data provided by Naganawa et al. 2007). In humans [ $^{11}C$ ]TMSX is very slowly metabolized (over 90% parent compound after 60 min) (Naganawa et al. 2007). Exemplary values for  $V_T$  and  $BP_{ND}$  in the putamen are 1.75 mL/mL and 1.26, respectively. In a single subject Ishiwata et al. (2005) showed that [ $^{11}C$ ]TMSX binding is blocked by a simultaneous and constant infusion of theophylline.  $V_T$  was reduced 8% in caudate nucleus and 4.5% in putamen but not in other brain areas. Theophylline infusion had an inhibiting effect on the metabolism of [ $^{11}C$ ]TMSX. Quantification of  $A_{2A}ARs$  based on arterial blood samples and metabolite analyses in five healthy volunteers were described by a two-compartment model and showed highest distribution volumes in putamen and thalamus (Mishina et al. 2007). Simplified methods for the determination of  $BP_{ND}$  by Logan's graphical analysis and distribution volume ratio to the reference region (centrum semiovale) have been reported for [ $^{11}C$ ]TMSX. One method is based on the generation of an estimated plasma time-activity curve by independent component analysis (EPICA), which is applicable to [ $^{11}C$ ]TMSX modeling because of the negligible metabolism of the compound (Naganawa et al. 2007). The second method using an intersectional searching algorithm and clustering (called EPISA) is independent of the metabolism of a ligand. In contrast to Logan, noninvasive EPISA-based  $V_T$  calculations to obtain  $BP_{ND}$  with a reference region circumvent  $k_2$  parameter determinations which necessitate arterial blood sampling and metabolite correction (Naganawa et al. 2008). This is achieved by obtaining an unscaled integral of the metabolite-corrected plasma input function which is determined by clusters of distinct tissue time-activity curves. An alternative approach extracted reference region data from the PET images (cortical gray matter by using supervised clustering) and linked those to a population-based arterial input function which correlated well with standard plasma input analyses (Rissanen et al. 2015). Test-retest stability of [ $^{11}C$ ]TMSX  $V_T$  and  $BP_{ND}$  was confirmed in five subjects (Naganawa et al. 2014).

Analogs of SCH442416 called [ $^{18}F$ ]-FESCH and [ $^{18}F$ ]-FPSCHE were preclinically evaluated in rats (Khanapur et al. 2017). Blocking studies and pharmacokinetic modeling revealed that a 60-min dynamic scan of [ $^{18}F$ ]-FESCH is suitable for blood- or reference region (cerebellum)-based  $A_{2A}R$  quantification. The same group developed a radiolabeled version of the drug preladenant. The in vivo assessment of [ $^{11}C$ ]preladenant in rats showed a larger striatum-to-cerebellum ratio compared to previously presented  $A_{2A}R$  radioligands in rats. Pharmacokinetic quantification of  $A_{2A}R$  (striatal  $BP_{ND} \sim 6$ ,  $V_T \sim 10$ ) in a rat model was possible and reliable, and formation of metabolites in plasma was low (17% after 60 min) (Zhou et al. 2017a, b, c). Transferring organ residence times of the rat model to a human radiation dosimetry model estimated common  $^{11}C$  radiotracer doses of 6  $\mu Sv/MBq$  for [ $^{11}C$ ]preladenant (Zhou et al. 2017a, b, c). Finally, first human data ( $n = 8$ ) confirmed the low effective dose range, slow metabolization (23% after 60 min), and high target-to-background ratio (putamen  $BP_{ND} \sim 5$ ,  $V_T \sim 8$ ) (Sakata et al. 2017).

Another promising, although from a PET imaging perspective less favorable, derivate of preladenant called [ $^{18}\text{F}$ ]-MNI-444 was recently tested in monkeys (Barret et al. 2014) and human subjects (Barret et al. 2015). Kinetics in humans were relatively slower (90-min scanning overestimated 210-min striatal  $\text{BP}_{\text{ND}}$  by  $\sim 10\%$ ). Binding in putamen was high ( $\text{BP}_{\text{ND}} \sim 5$ ,  $V_{\text{T}} \sim 3$ ) and the effective dose comparable to other [ $^{18}\text{F}$ ] radioligands.

Due to the heterodimerization between dopamine  $\text{D}_2$  and  $\text{A}_{2\text{A}}\text{ARs}$ , adenosine antagonists were explored as therapeutics for Parkinson's disease. Using PET it was investigated whether the disease affects basal cerebral  $\text{A}_{2\text{A}}\text{AR}$  density and whether therapeutic interventions with standard antiparkinsonian medication had an impact on  $\text{A}_{2\text{A}}\text{AR}$  binding. It was found that the binding potential in the putamen was higher in patients with additional dyskinetic symptoms compared to controls (Mishina et al. 2011). In another study with the  $\text{A}_{2\text{A}}\text{AR}$  ligand [ $^{11}\text{C}$ ]SCH442416 this finding could be reproduced as patients with dyskinesia had higher binding potentials in contrast to patients without dyskinesia who showed binding potentials comparable to the control subjects (Ramlackhansingh et al. 2011). In a preliminary analysis the EPICA method has been used to determine the  $\text{BP}_{\text{ND}}$  of patients with Parkinson's disease.  $\text{BP}_{\text{ND}}$  was significantly lower on the more affected side of the posterior putamen in Parkinson's disease while [ $^{11}\text{C}$ ]raclopride binding to dopamine receptors was increased. A recent study with the new  $\text{A}_{2\text{A}}\text{AR}$  ligand [ $^{11}\text{C}$ ]preladenant showed no significant differences in the basal ganglia between Parkinson's disease and control groups in  $\text{BP}_{\text{ND}}$  values (Ishibashi et al. 2018). In the 6-hydroxydopamine rat model [ $^{11}\text{C}$ ]preladenant binding was reduced after induction and increased after L-DOPA treatment (Zhou et al. 2017a, b, c).

[ $^{11}\text{C}$ ]TMSX was used to investigate the changes of  $\text{A}_{2\text{A}}\text{AR}$  availability in patients ( $n = 8$ ) with normal-appearing white matter of secondary progressive multiple sclerosis. Patients showed an  $\sim 20\%$  higher  $V_{\text{T}}$  compared to healthy controls (Rissanen et al. 2015).

Besides the aforementioned application of [ $^{11}\text{C}$ ]TMSX in brain imaging studies it has additionally been used to study the human  $\text{A}_{2\text{A}}\text{AR}$  in skeletal muscle, heart, and brown adipose tissue. An infusion of theophylline resulted in a reduction of binding suggesting specific binding of [ $^{11}\text{C}$ ]TMSX in human muscles (Ishiwata et al. 2004). Increased adenosine  $\text{A}_{2\text{A}}\text{AR}$  densities were found in the muscles of endurance-trained vs. untrained men (Mizuno et al. 2005). This finding could not be confirmed in a subsequent study (Heinonen et al. 2008). Exposure to cold ambient temperature ( $16\text{ }^\circ\text{C}$ ) leads to a reduction of [ $^{11}\text{C}$ ]TMSX binding in brown adipose tissue (neck region) which was interpreted as an occupancy by endogenous adenosine (Lahesmaa et al. 2019).

Regarding human occupancy studies of drug candidates, [ $^{11}\text{C}$ ]SCH442416 has been used to assess adenosine  $\text{A}_{2\text{A}}\text{AR}$  occupancy of vipadenant (BIIB014) in healthy subjects (Brooks et al. 2010) and [ $^{11}\text{C}$ ]KW-6002 (istradefylline) to find optimal oral doses of the same unlabeled compound (Brooks et al. 2008). The human in vivo  $\text{A}_{2\text{A}}\text{AR}$  occupancy of this Parkinson medication was measured with the new radioligand [ $^{11}\text{C}$ ]preladenant confirming the approved dose range (Ishibashi et al. 2018).

## 8.4 A<sub>2B</sub> Adenosine Receptor

Although several A<sub>2B</sub>AR agonists and antagonists have been evaluated for preclinical applications (Kalla et al. 2009), none of these compounds has so far been introduced as a preclinical or clinical PET ligand.

[<sup>11</sup>C]4a triazinobenzimidazole-based radioligand has been introduced and tested *in vivo* showing very high chemical and blood stability but low brain uptake in rats (Petroni et al. 2016). The [<sup>18</sup>F]7a pyrazine-based radioligand ( $K_D$  for A<sub>2B</sub>AR = 4.2 nM) (Lindemann et al. 2018) showed brain uptake in mice (SUV = 1) but formation of a blood-brain barrier-penetrating metabolite. Another drawback of this compound is its low A<sub>1</sub>AR selectivity ( $K_D$  19 nM).

## 8.5 A<sub>3</sub> Adenosine Receptor

There are no reports on A<sub>3</sub>AR *in vivo* imaging studies so far although some radioligands were evaluated in biodistribution studies in rodents. Distribution and metabolism experiments have been performed in rodents with the fluoroethyl ester [<sup>18</sup>F]FE@SUPPY and its fluoroethyl thioester analog [<sup>18</sup>F]FE@SUPPY:2 (Haeusler et al. 2015, 2010; Wadsak et al. 2008). Unfortunately, both tracers are metabolized to BBB-permeating metabolites which complicate pharmacokinetic evaluation of the binding. Fifteen minutes after injection of [<sup>18</sup>F]FE@SUPPY:2 the metabolites accounted for ~40% of brain tissue radioactivity and this fraction increased to ~80% after 60 min. [<sup>18</sup>F]FE@SUPPY was more favorable in this respect as no metabolites were detected in the brain tissue up to 15 min after injection. However, the concentration increased to ~80% after 60 min (Haeusler et al. 2010). In another biodistribution study the [<sup>76</sup>Br]-labeled antagonist MRS5147 was evaluated in rats showing an increased uptake in A<sub>3</sub>AR-rich testes (Kiesewetter et al. 2009). However, in a dipping study with whole brain and testes no specific binding of [<sup>76</sup>Br]MRS5147 was detected. In colorectal cancer A<sub>3</sub>ARs are described to be highly expressed. In a xenograft mouse model *in vivo* detection of A<sub>3</sub>AR with [<sup>18</sup>F]FE@SUPPY was tested but showed insufficient tumor uptake (Balber et al. 2018).

## 8.6 Conclusion

Adenosine and its receptors play an important role in the physiology and pathophysiology of the brain as well as in other organs, particularly kidney, bowel, and vascular system, and in tumors. At present, only the A<sub>1</sub>AR and the A<sub>2A</sub>AR are accessible with molecular *in vivo* imaging. So far, these tools revealed important insights into neurologic and psychiatric disorders, sleep physiology, and cancer. The constantly increasing amount of data on adenosine and its important role in multiple organ systems as well as in the pathophysiology of frequent diseases underscore the need for further research in the field of ligand development for adenosine receptors.

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# Imaging of Central Benzodiazepine Receptors in Chronic Cerebral Ischemia

# 9

Hiroshi Yamauchi

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## Abstract

In patients with atherosclerotic internal carotid artery (ICA) or middle cerebral artery (MCA) occlusive disease, fluctuation of cerebral perfusion under conditions of chronic hemodynamic compromise may cause not only cerebral infarction but also selective neuronal damage in the cerebral cortex that is not detectable as infarction on CT or MRI. Imaging of the central-type benzodiazepine receptors (cBZR), which are expressed by most cortical neurons, has enabled visualization of neuronal alterations induced by ischemia in vivo, in humans. Using PET and  $^{11}\text{C}$ -flumazenil, we showed that selective neuronal damage demon-

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strated as a decrease in cBZR in the normal-appearing cerebral cortex was associated with increased oxygen extraction fraction (OEF), the so-called misery perfusion, in patients with atherosclerotic ICA or MCA occlusive disease in the chronic stage. Follow-up examinations of the patients without ischemic episode showed that a decrease in cBZR was associated with an increase in OEF (hemodynamic deterioration). A decrease in cBZR was also associated with executive dysfunction. Chronic hemodynamic ischemia causes selective neuronal damage observed as a decrease in cBZR that has an impact on the functional outcomes. Therapies for preventing neuronal damage are needed, and imaging of cBZR may be useful to monitor the effects of such therapies.

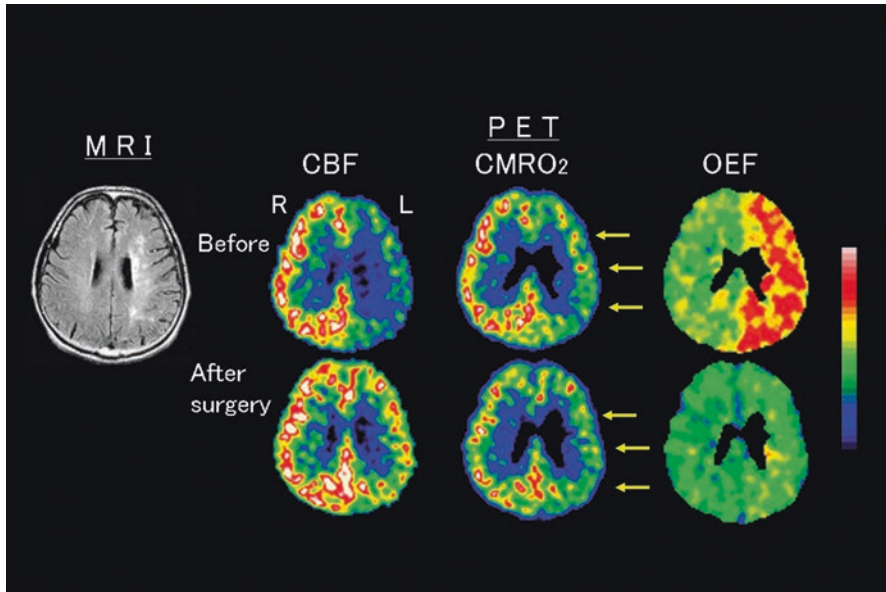
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## 9.1 Introduction

### 9.1.1 Chronic Hemodynamic Compromise and Risk for Stroke

In patients with atherosclerotic internal carotid artery (ICA) or middle cerebral artery (MCA) occlusive disease, chronic reduction in cerebral perfusion pressure (chronic hemodynamic compromise) increases the risk of cerebral ischemic damage (Baron et al. 1981; Klijn and Kappelle 2010; Powers 1991; Yamauchi et al. 1992). Previous studies have shown that chronic hemodynamic compromise, as indicated by increased oxygen extraction fraction (OEF; misery perfusion) (Baron et al. 1981) on positron-emission tomography (PET) or severely decreased vasodilatory capacity, is a risk factor for subsequent ischemic stroke in atherosclerotic ICA or MCA occlusive disease (Derdeyn et al. 1999; Grubb et al. 1998; Kuroda et al. 2001; Markus and Cullinane 2001; Ogasawara et al. 2002; Vernieri et al. 1999; Yamauchi et al. 1996b, 1999, 2012). Furthermore, follow-up examinations of patients without ischemic episodes showed that an increase in OEF (hemodynamic deterioration) was associated with an increase in the risk for subsequent stroke (Yamauchi et al. 2000b). Thus, understanding the pathophysiology of chronic hemodynamic cerebral ischemia is essential for the management of patients with atherosclerotic ICA or MCA occlusive disease.

Vascular reconstruction surgery can improve chronic hemodynamic compromise (Yamauchi et al. 1994), which may prevent subsequent ischemic stroke of hemodynamic origin (Fig. 9.1). Extracranial-to-intracranial (EC-IC) bypass surgery—which has no benefit in patients with ICA or MCA occlusive disease in general (The EC/IC Bypass Study Group 1985)—may prevent recurrent strokes in select patients with hemodynamic compromise. Two randomized clinical trials used hemodynamic criteria for patient selection (Ogawa 2012; Powers et al. 2011), to test this hypothesis. One trial in Japan (Japanese EC-IC bypass trial; JET study) demonstrated a benefit from bypass surgery for preventing disabling stroke (modified Rankin scale >2) (2-year ipsilateral ischemic stroke rate including perioperative stroke: 2.9% in



**Fig. 9.1** Reduced cerebral cortical oxygen metabolism without morphological changes. A patient with occlusion of the left internal carotid artery showed decreased cerebral blood flow (CBF) and increased oxygen extraction fraction (OEF, misery perfusion) in the hemisphere with occlusion. Surgical reperfusion with bypass improved misery perfusion. However, we should note that cerebral metabolic rate of oxygen, which was decreased in the normal-appearing cerebral cortex before surgery, was unchanged after surgical reperfusion. Some irreversible damage may occur in the cerebral cortex

the surgical group vs. 10.7% in the medical group) (Ogawa 2012); in contrast, another study (Carotid Occlusion Surgery Study; COSS) did not demonstrate a benefit (any stroke, 21% vs. 23%) (Powers et al. 2011). The perioperative stroke rate in the two studies differed (disabling stroke, 0% vs. any stroke, 15%), which may have led to their different conclusions. Because of the high prevalence of Moyamoya disease, EC-IC bypass with postoperative management has been an essential skill for neurosurgeons in Japan (Kuroda and Houkin 2008), which may have resulted in the 0% perioperative disabling stroke rate of the Japanese EC/IC Bypass Trial (Ogawa 2012). Achieving low operative risk could lead to the benefit of bypass surgery.

Although correct evaluation of hemodynamic status is essential to determine prognosis, therapeutic strategies to prevent recurrent strokes, which should differ between patients with and those without hemodynamic compromise, are not clearly established. Continuing efforts are warranted to determine the strategies for selecting treatments based on hemodynamic measurements.

### 9.1.2 Selective Neuronal Damage/Loss or Incomplete Infarction

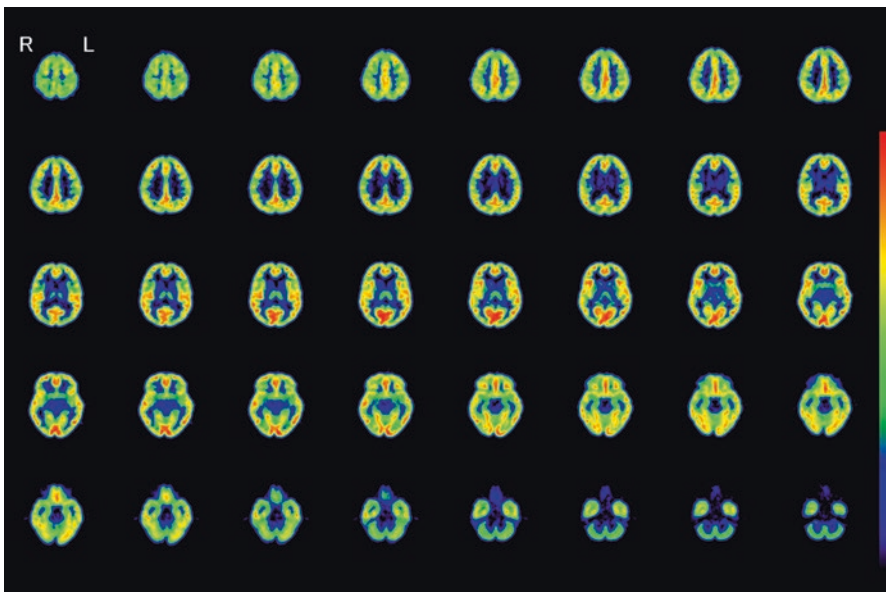
In atherosclerotic ICA or MCA occlusive disease, chronic hemodynamic compromise increases the risk for cerebral ischemic damage regardless of the development of symptoms. Hemodynamic cerebral ischemia due to ICA or MCA occlusive disease may cause not only cerebral infarction but also minor tissue damage in the cerebral cortex that is not detectable as infarction on CT or magnetic resonance imaging (MRI). Neuropathological studies have demonstrated cortical granular atrophy or watershed territory microemboli in the cerebral cortex distal to a large cerebral arterial occlusion (Graham 1992). Observations in both humans and animals also suggest that cortical selective neuronal loss results from acute occlusion of the large cerebral arteries accompanied by ischemia of moderate severity (Garcia et al. 1996; Lassen et al. 1983). Fifteen-minute distal MCA occlusion in spontaneously hypertensive rats, mimicking true cortical TIA, resulted in normal MRI findings, but consistent cortical selective neuronal loss was observed (Ejaz et al. 2015a, b). However, selective neuronal loss in patients with atherosclerotic ICA or MCA disease has not been confirmed pathologically in the human brain. It is unclear whether selective neuronal ischemic change develops in patients with atherosclerotic occlusion of the large cerebral arteries because a direct demonstration of the cortical neuronal loss in the living brain is not possible.

Indirect markers for neuronal damage, including decreased metabolism and brain atrophy, have been investigated to address the issue of neuronal loss without infarction in ICA or MCA occlusive diseases. Decrease in glucose or oxygen metabolism in the normal-appearing cerebral cortex is one of the sequelae of ischemic damage to the tissue; however, it can also be caused by a decrease in neural input from distant regions with fiber connections (Feeney and Baron 1986). CT studies in ICA occlusive disease have revealed only hemiatrophy in some patients (Radu and Moseley 1978). In patients with ICA or MCA occlusive disease, including Moyamoya disease, MRI studies have shown decreases in cortical thickness in the hemispheres with arterial disease (Fierstra et al. 2010). Atrophy of the corpus callosum may be an indirect but sensitive indicator of ischemic cortical neuronal loss. The largest fraction of neurons projecting into the corpus callosum comprises the large pyramidal cells in layer 3, which constitute one of the groups of cortical neurons most vulnerable to ischemia (Graham 1992). Therefore, cortical ischemic loss may lead to callosal atrophy. Reduced cerebral cortical oxygen metabolism in the normal-appearing cerebral cortex, which surgical reperfusion cannot improve, has been demonstrated by using PET in ICA or MCA occlusive disease, suggesting that selective neuronal damage may occur (Powers et al. 1984; Yamauchi et al. 1990). Our studies showed that a decrease in bilateral cerebral cortical oxygen metabolism occurs in association with callosal atrophy in patients with ICA occlusive disease, suggesting that ischemic neuronal loss may contribute to the reduced cortical oxygen metabolism (Yamauchi et al. 1993, 1995, 1996a, 2000b). However, to explore this aspect further, a more direct marker for neuronal damage is needed.

## 9.2 Imaging of Central-Type Benzodiazepine Receptors

Development of imaging techniques makes it possible to visualize the distribution of central-type benzodiazepine receptors (cBZR) in humans by using  $^{11}\text{C}$ -labeled flumazenil (FMZ) for PET or  $^{123}\text{I}$ -labeled iomazenil (IMZ) for single-photon emission computed tomography (SPECT) (Beer et al. 1990; Persson et al. 1985; Samson et al. 1985). cBZR is coupled with  $\gamma$ -aminobutyric acid receptors (GABAA/BZ receptor complex), and cBZR exists in the membrane of neurons (Olsen 1981). Most cortical neurons express cBZR. Binding of FMZ is high in the cerebral cortex and low in the cerebellum, thalamus, and basal ganglia, while it is lacking in the white matter (Fig. 9.2). As shown in the case of crossed cerebellar diaschisis, a decrease in neural input may not decrease cBZR (Dong et al. 1997; Hatazawa et al. 1995a). Thus, imaging of the cBZR may detect cortical neuronal damage, which could not be evaluated directly by other neuroimaging modalities (Baron et al. 2014; Garcia et al. 1996; Hatazawa and Shimosegawa 1998; Sette et al. 1993). Cortical neuronal damage should lead to a decrease in cBZR.

Histopathological correlations between decreased cBZR and selective neuronal loss have recently been confirmed (Ejaz et al. 2013). A rat ischemia-reperfusion model with 45-min occlusion of the distal MCA was examined with FMZ PET and immunohistochemistry 28 days after ischemia. The study showed that a decrease in



**Fig. 9.2** BZR imaging with  $^{11}\text{C}$ -flumazenil PET. Images of a male normal volunteer of 49 years old show high binding in the cerebral cortex



FMZ binding in the normal-appearing cerebral cortex on T2-weighted MRI was associated with selective neuronal loss on stains with an antibody for neurons. Imaging of cBZR by using  $^{11}\text{C}$ -FMZ and PET can detect selective neuronal damage or loss.

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### **9.3 Pathophysiology of Selective Neuronal Damage Demonstrated as Decreased BZR**

#### **9.3.1 Selective Neuronal Damage and Chronic Hemodynamic Cerebral Ischemia**

Several small studies in patients with atherosclerotic ICA or MCA disease in the chronic stage demonstrated a reduction in cBZR in the normal-appearing cortical areas on CT or MRI, suggesting selective neuronal damage (Dong et al. 1997; Kuroda et al. 2004, 2006; Moriwaki et al. 1998; Sasaki et al. 1997; Yamauchi et al. 2000a). The IMZ uptake on SPECT was significantly but weakly correlated with the cerebral metabolic rate of oxygen, while no significant correlation was found of the IMZ uptake with the blood flow, the cerebrovascular reactivity, or the cerebral metabolic rate of glucose, which suggested that the reduction in IMZ uptake may be due to selective neuronal loss that may not be revealed by other imaging tracers (Dong et al. 1997; Moriwaki et al. 1998; Sasaki et al. 1997). These findings were also shown by using FMZ and PET (Kuroda et al. 2004, 2006). A decrease in cortical BZR binding was correlated with corpus callosum atrophy, which supported that decreased cortical BZR binding may reflect cortical neuronal damage (Yamauchi et al. 2000a).

However, the incidence, extent, and severity of selective neuronal damage demonstrated as decreased cBZR are unclear. Furthermore, the factors responsible for decreased cBZR are not well understood. Specifically, although a chronic hemodynamic mechanism may be responsible for the decreased cBZR, the relationship between decreased BZR and chronic hemodynamic compromise is unknown. The clarification of this relationship is essential because vascular reconstruction surgery can improve chronic hemodynamic compromise, which may prevent the development of selective neuronal damage as well as infarction.

#### **9.3.2 Selective Neuronal Damage and Low-Flow Infarction**

Border zone infarction has been demonstrated to be associated with ICA occlusive disease and hemodynamic compromise (Momjian-Mayoy and Baron 2005; Yamauchi et al. 1990, 1991). Hemodynamic ischemia due to ICA occlusive disease may cause not only border-zone infarction but also selective neuronal damage beyond the regions of infarcts that may be detected by a decrease in cBZR in the normal-appearing cerebral cortex. To determine whether selective neuronal damage is associated with border-zone infarction in ICA occlusive disease, we measured

cBZR using PET and  $^{11}\text{C}$ -FMZ in 62 nondisabled patients with ICA steno-occlusive lesions in the chronic stage (Yamauchi et al. 2005). FMZ-binding potential (BP) was calculated using the dynamic data and Logan graphical analysis with the reference tissue, with the pons as a reference region (Logan et al. 1996; Okazawa et al. 2004). The infarcts on MRI, which were categorized as territorial, border zone (external or internal), striatocapsular, lacunar, and other white matter infarcts, were correlated with mean cerebral/cerebellar cortical BP ratio in the hemisphere with ICA occlusive disease. Patients with border-zone infarction ( $n = 18$ ) had significantly decreased FMZ-BP ratio in the hemisphere with ICA disease, compared with patients without border-zone infarction ( $n = 44$ ) and normal control subjects ( $n = 10$ ). Both external and internal border-zone infarctions were associated with a decreased FMZ-BP ratio. Multivariate analysis showed that external border-zone infarction was an independent predictor of the decreased FMZ-BP ratio. In ICA occlusive disease, selective neuronal damage demonstrated as decreased cBZR is associated with border-zone infarction. Microemboli can cause border-zone infarction, particularly in cases of ICA stenosis (Moustafa et al. 2010). However, it is less likely that border-zone infarction due to microemboli causes a significant decrease in BZR beyond the regions of the infarct because ischemia due to microemboli is restricted to the border zone region. Therefore, hemodynamic ischemia leading to border-zone infarction may cause selective neuronal damage beyond the regions of infarcts in the chronic stage.

Decreased cBZR was strongly associated with decreased oxygen metabolism. However, the correlation of decreased cBZR with increased OEF (hemodynamic compromise) was weak. Selective neuronal damage caused by hemodynamic ischemia, consequently, decreases the degree of hemodynamic compromise due to ICA occlusive disease by causing reduced metabolic demand of the tissue (Kuroda et al. 2004; Yamauchi et al. 2004), which may lead to normalization of OEF after the development of selective neuronal damage. The association of decrease in cBZR with border-zone infarction suggests that severe hemodynamic compromise, which was present in the acute stage of infarction, disappeared according to the decrease in the metabolic needs of the tissue within the surviving penumbra between the time of infarction and PET examination (Morris et al. 2018). Careful medical management in the acute stage for patients with border-zone infarction should prevent selective neuronal damage, even when infarct size is small, and symptom is mild. Subsequently, the preserved metabolic needs of the cortical tissue in the chronic stage may permit the persistence of the hemodynamic compromise that can be improved by vascular reconstruction surgery. This strategy results in preservation of cortical metabolism that may lead to good functional outcomes. Recognition and prevention of selective neuronal damage are important for the management of patients with ICA occlusive disease.

The association between border-zone infarction (internal border-zone infarction) and a decrease in cBZR in the overlying cerebral cortex was also found in 62 nondisabled patients with atherosclerotic occlusive disease of the MCA and no cortical infarction (Yamauchi et al. 2009).

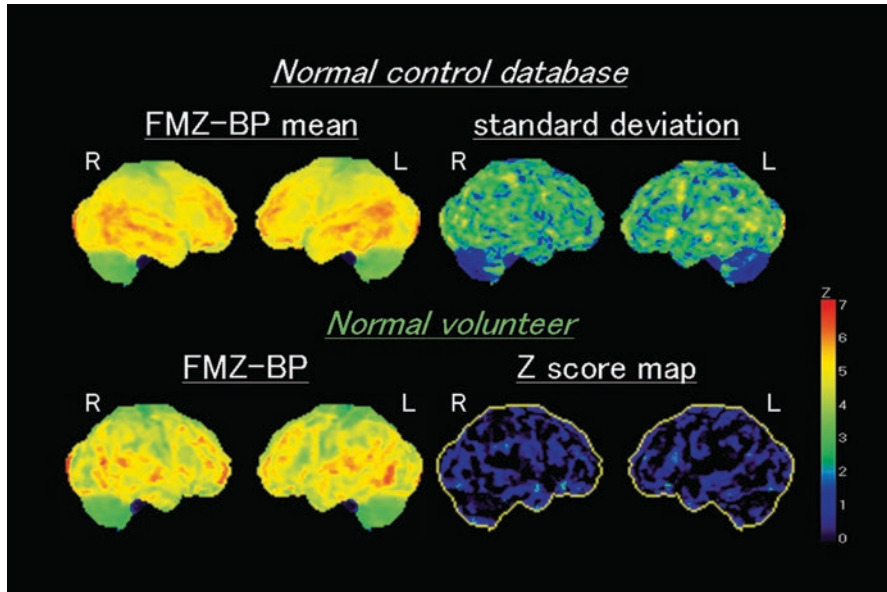
### 9.3.3 Selective Neuronal Damage and Misery Perfusion

#### 9.3.3.1 A Cross-Sectional Study

The notion that normalization of OEF may occur after the development of selective neuronal damage suggests that it may not be easy to demonstrate an association of decreased cBZR with chronic hemodynamic compromise (increased OEF) in the chronic stage. To overcome this issue, we studied 105 nondisabled patients with atherosclerotic ICA or MCA occlusive disease and *no cortical infarction* on routine MRI images (T1-, T2-weighted, or fluid-attenuated inversion recovery images) and measured cBZR and OEF using PET. We investigated the association of selective neuronal damage demonstrated as a decrease in cBZR in the normal-appearing cerebral cortex with increased OEF (misery perfusion) (Yamauchi et al. 2007).

A three-dimensional stereotactic surface projection (3D-SSP) technique was used to analyze FMZ BP (Minoshima et al. 1995). After anatomical standardization, a method of data extraction is performed in which the cortical activity is projected onto the brain surface. The algorithm searches the highest pixel value in an inward direction along the vector to a six-pixel depth into the cortex on an individual's anatomically standardized PET image set and assigns the maximum value to the surface pixel. The surface projection technique minimizes the effect of cortical atrophy. Then, a comparison of the resultant cortical projections between a patient and controls is performed. Pixel-by-pixel  $Z$  scores were used.  $Z$  scores ( $[(\text{mean normalized pixel value of controls}) - (\text{normalized pixel value of each patient})] / (\text{SD of controls})$ ) were calculated for each surface pixel ( $Z$  score map) (Fig. 9.3) to quantify a decrease in FMZ BP. A positive  $Z$  score represents a reduced FMZ BP in patients relative to controls. To assess the degree of abnormal FMZ BP reduction in each patient quantitatively, the BZR index [(the extent of the pixels with  $Z$  score more than 2 compared with controls)  $\cdot$  (average  $Z$  score in those pixels)] of the MCA distribution with arterial disease in the cerebral cortex was calculated by the stereotactic extraction estimation (SEE) method (Mizumura et al. 2003).

All patients had pixels with abnormally decreased cBZR in the cerebral cortex of the MCA distribution, with the extent varying from 0.04% to 60.91% (Fig. 9.4). The BZR index was significantly correlated with the mean hemispheric value of OEF or CBF in the ipsilateral hemisphere. Multivariate analysis showed that the BZR index was positively correlated with the value of OEF and the history of stroke, which suggested that misery perfusion might cause selective neuronal damage. On the other hand, the BZR index was negatively correlated with the presence of statin treatment. Several studies have shown that statins have beneficial effects on the cerebral circulation and brain parenchyma during ischemic stroke and reperfusion (Vaughan and Delanty 1999). The antioxidative properties of statins might be involved in their beneficial effects against neuronal damage in cerebral ischemia (Nagotani et al. 2005). Statins might also reduce microemboli by plaque stabilization in patients with ICA or MCA stenosis. Therefore, statins may reduce the occurrence of selective neuronal damage due to ischemia in atherosclerotic ICA or MCA occlusive disease.

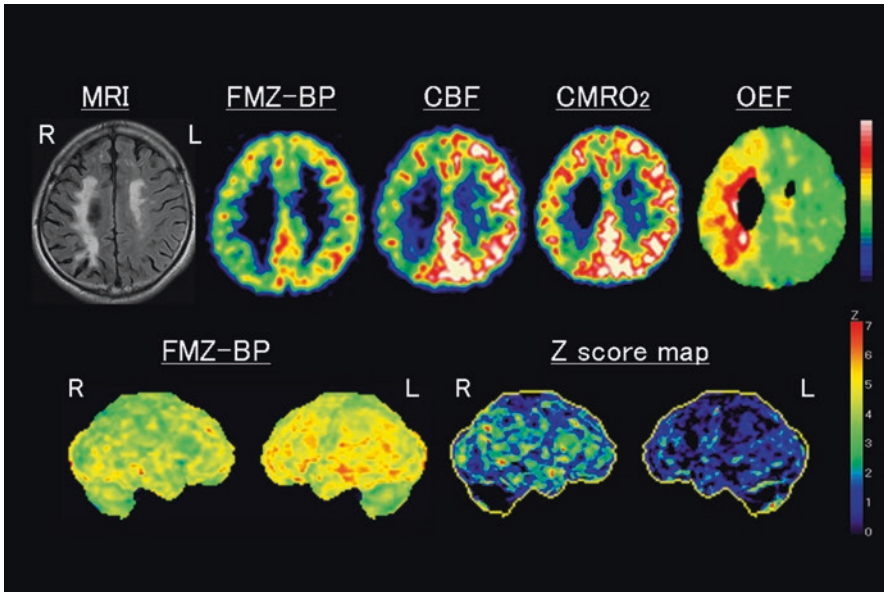


**Fig. 9.3** Three-dimensional stereotactic surface projection (3D-SSP) analysis. Upper row: 3D-SSP images showing the value of mean and standard deviation for normal control database ( $N = 10$ , mean age 57 years). Lower row: 3D-SSP images and Z score maps of a male normal volunteer of 49 years old shown in Fig. 9.2

### 9.3.3.2 Follow-Up Study

It is essential to show a correlation between an increase in OEF and progression of selective neuronal damage using a longitudinal study to demonstrate a causal relationship between misery perfusion and selective neuronal damage more directly. Therefore, we evaluated the distribution of cBZR twice using PET and  $^{11}\text{C}$ -flumazenil over time in 80 medically treated patients, aged  $63 \pm 8$  years, with atherosclerotic ICA or MCA occlusive disease who had no ischemic episodes during follow-up (Yamauchi et al. 2016). The total change of the BZR index or the OEF value in the cerebral cortex of the MCA distribution with arterial disease was calculated by subtracting the value obtained at the second scanning from that obtained at the first scanning.

Follow-up examinations in seven normal subjects, aged  $56 \pm 8$  years, demonstrated no significant change in Z index during follow-up (mean  $41 \pm 3$  months). The mean  $\pm$  SD value of changes in the index in the controls was  $0.94 \pm 1.38$ . In patients, an increase of the index beyond the upper 95% limit (the mean plus  $6t_{0.05}\text{SD}$ ) defined in normal subjects (above 4.32) was considered to be an increased BZR index (progression of neuronal damage) during follow-up.



**Fig. 9.4** Selective cortical neuronal damage and misery perfusion. Upper row: Examples of PET images showing a decrease of flumazenil-binding potential (FMZ-BP), cerebral blood flow (CBF), and cerebral metabolic rate of oxygen (CMRO<sub>2</sub>) with increased oxygen extraction fraction (OEF) in a patient with right (R) internal carotid artery occlusion who showed internal border-zone infarction on the corresponding MRI images. This patient presented with left hemiparesis and spatial neglect. Lower row: 3D-SSP images and Z score maps showing a decrease of FMZ-BP in the right hemisphere. The extent of abnormally decreased benzodiazepine receptors (BZR) was 47.6% in the right middle cerebral artery distribution; the mean z score in those pixels was 2.92, and the BZR index was 139.2

In the hemisphere affected by arterial disease, the BZR index in 40 patients (50%) was increased during follow-up (mean  $26 \pm 20$  months). In multivariable logistic regression analyses, increases in the BZR index were associated with the decreased CBF at baseline and an increased OEF during follow-up. Increases in the OEF during follow-up were associated with a lack of statin use. In patients with atherosclerotic ICA or MCA disease, the progression of cortical neuronal damage was associated with hemodynamic impairment at baseline and hemodynamic deterioration during follow-up. Statin use may be beneficial against hemodynamic deterioration and therefore be neuroprotective.

Follow-up data show that hemodynamic deterioration causes selective neuronal damage demonstrated as decreased BZR without an overt episode of stroke. This study confirmed the relationship between selective neuronal damage and chronic hemodynamic compromise for the first time. In patients with chronic hemodynamic compromise, the perfusion may fall below the penumbra threshold for a matter of minutes, causing selective neuronal damage.

## 9.4 Silent Cortical Neuronal Damage in Asymptomatic Patients

The finding that selective neuronal damage manifested as loss of cBZR occurs in association with hemodynamic deterioration without an overt episode of stroke suggests that selective neuronal damage may occur in association with hemodynamic compromise in patients whose atherosclerotic ICA or MCA disease is asymptomatic. To test this hypothesis, we measured cBZR using PET and  $^{11}\text{C}$ -FMZ in 79 patients with asymptomatic atherosclerotic ICA or MCA disease and no cortical infarction (Yamauchi et al. 2011b). 3D-SSP was used to calculate the BZR index, a measure of abnormally decreased cBZR in the cerebral cortex within the MCA distribution. The BZR index was abnormal ( $>5.84$ ) in 71% of the asymptomatic patients compared to that in healthy control subjects. This abnormality was mild compared with that previously reported for symptomatic patients (the BZR index,  $16.0 \pm 14.5$  vs.  $48.2 \pm 54.1$ ). Multiple regression analysis showed the BZR index to be positively correlated with the value of OEF, presence of silent subcortical infarcts, and presence of ischemic heart disease, which suggested that hemodynamic compromise is associated with selective neuronal damage in asymptomatic patients. Hemodynamic compromise may cause silent cortical neuronal damage in asymptomatic patients as well. Patients at elevated risk of stroke may have already suffered ischemic cerebral damage by the time their asymptomatic ICA or MCA disease is discovered. Therapeutic strategies to prevent neuronal damage, including vascular reconstruction surgery, may be called for in asymptomatic patients with chronic hemodynamic compromise.

On the other hand, the BZR index was negatively correlated with the treatment of hypertension with angiotensin receptor blockers (ARBs). ARBs were associated with preservation of cortical cBZR. Several experimental studies have shown that ARBs can attenuate ischemic neuronal injury through various mechanisms, including decreased oxidative stress and improved arterial compliance (Mogi and Horiuchi 2009). ARBs might have beneficial effects against neuronal damage in atherosclerotic ICA or MCA disease.

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## 9.5 Clinical Impact of Selective Neuronal Damage: Cognitive Impairment

In atherosclerotic ICA or MCA disease, chronic hemodynamic compromise causes selective neuronal damage, which can be detected as a decrease in cBZR in the normal-appearing cerebral cortex. However, the impact of selective neuronal damage on clinical outcomes is unclear.

One to three months after acute stroke, selective neuronal loss—as documented by a decrease in cBZR—within the rescued penumbra was shown to impede neural activation on functional MRI, suggesting the presence of functional impairments

(Carrera et al. 2013). In cases of subcortical hemorrhage, patients with aphasia or anosmia showed an ipsilateral/contralateral IMZ uptake ratio of  $\leq 0.78$  in the cortical area (Hatazawa et al. 1995b). Another SPECT study also showed that cerebral hyperperfusion after carotid endarterectomy resulted in a postoperative decrease in IMZ uptake that correlated with postoperative cognitive impairment (Chida et al. 2009). Therefore, selective neuronal damage may contribute to the development of subtle poststroke cognitive impairment, depending on the degree of neuronal damage.

In patients with atherosclerotic ICA or MCA disease, a few subjects with reduced cBZR in the cerebral cortex overlying subcortical infarcts were reported to show aphasia (Hatazawa et al. 1995a; Moriwaki et al. 1998), and a decrease in cortical cBZR was shown to be correlated with the atrophy of corpus callosum that is associated with global cognitive impairment (Yamauchi et al. 1996a, 2000a). However, the relationship between a decrease in cBZR and neurological symptoms has not been systematically investigated. Clinical correlates are essential to establish the clinical significance of cBZR imaging.

Executive dysfunction may be an early sign of vascular cognitive impairment and may have a considerable impact on the functional outcomes of patients with stroke (Roman and Royall 1999). The Wisconsin Card Sorting Test (WCST) is a widely used clinical test for assessing executive functions (Nagahama et al. 1996; Nelson 1976). To determine whether a decrease in the cBZRs in the non-infarcted cerebral cortex is associated with poor performance on the WCST, we measured cBZR using PET and  $^{11}\text{C}$ -FMZ in 60 nondisabled patients with unilateral atherosclerotic ICA or MCA disease and no cortical infarction (Yamauchi et al. 2011a). Using 3D-SSP, the BZR indices in the cerebral cortex of the anterior cerebral artery (ACA) and the MCA territories were calculated and found to be correlated with the patient's score on the WCST. We used the Keio-Fukuoka-Shimane version of the WCST that can be performed on a personal computer (Kobayashi 2002). Because the WCST method used in this study needs verbal mediation during the performance, the language-related left hemisphere may contribute more than the right hemisphere to the performance of the WCST. The BZR index was associated with an increase in OEF in the ACA and MCA territories of the hemisphere with ICA or MCA disease. Based on the WCST results, 39 patients were considered abnormal (low categories achieved) for their age. The BZR index of the ACA territory in the hemisphere affected by arterial disease was significantly higher in abnormal patients than in normal patients. The BZR index of the MCA territory differed significantly between the two groups when patients with left arterial disease ( $n = 28$ ) were analyzed separately. Among the BZR indices for the subregions of the ACA territory, those of the anterior cingulate gyrus and the medial frontal gyrus were positively correlated with the total number of errors after controlling for the effect of age through multiple regression analysis. In patients with left arterial diseases, among the BZR indices for the subregions of the MCA territory, the BZR index of the middle frontal gyrus was also positively correlated with the total number of errors. These subregions have been reported to be essential for the performance on WCST (Buckley et al. 2009; Nagahama et al. 1996).

In atherosclerotic ICA or MCA disease, selective neuronal damage that manifests as a decrease in cBZR in the non-infarcted cerebral cortex may contribute to the development of executive dysfunction. PET imaging of cBZR may be useful as an objective measure of cognitive impairments in atherosclerotic occlusive disease of the major cerebral artery. In this study, a decrease in cBZR was also associated with hemodynamic compromise. Interestingly, a recent IMZ SPECT study showed that improvement of hemodynamic compromise after carotid endarterectomy resulted in a postoperative increase in IMZ uptake that correlated with postoperative improvement of cognitive impairment (Chida et al. 2010). In patients with severe hemodynamic compromise, a part of the cBZR decrease may be reversible, and improvement of hemodynamic compromise may lead to the improvement of cognitive impairment with cBZR reduction.

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## 9.6 Selective Neuronal Damage as Outcome Measures

In patients with atherosclerotic ICA or MCA disease, the progression of selective cortical neuronal damage occurs in the absence of an overt stroke episode and is associated with both decreased CBF at baseline and an increased OEF during follow-up. Decreased CBF at baseline may reflect misery perfusion with cortical neuronal damage, while an increased OEF during follow-up may reflect hemodynamic deterioration. Thus, to prevent the progression of selective cortical neuronal damage, vascular reconstruction surgery to improve hemodynamic impairment and appropriate control of vascular risk factors (e.g., the use of statin or antihypertensive agents) may be useful.  $^{11}\text{C}$ -flumazenil PET would be useful to monitor the effects of the therapies.

### 9.6.1 Vascular Reconstruction Surgery

Bypass surgery can improve misery perfusion and may prevent neuronal damage. A recent IMZ SPECT study showed that improvement of hemodynamic compromise after bypass surgery resulted in a postoperative improvement of BZR that was associated with cognitive improvement (Chida et al. 2011). An MRI study also showed an improvement in cortical thinness after bypass surgery in patients with ICA or MCA occlusive diseases, including Moyamoya disease (Fierstra et al. 2011). However, surgery often involves a potential risk for neuronal damage.

To determine whether the progression of cortical neuronal damage in surgically treated patients with misery perfusion is more significant than in those without, or in medically treated patients with misery perfusion, we evaluated the distribution of cBZR using  $^{11}\text{C}$ -flumazenil PET in patients with atherosclerotic ICA or MCA disease, including 18 surgically treated patients with no history of perioperative stroke, among which 9 presented with misery perfusion, and 8 medically treated patients with misery perfusion but no intervening ischemic event (Yamauchi et al. 2017).



We quantified abnormal decreases in the cBZRs distributed within the MCA of the cerebral cortex, and compared the changes in the BZR index among the three groups. The change in the BZR index in surgically treated patients with misery perfusion ( $27.5 \pm 15.6$ ) during  $7 \pm 5$  months was significantly greater than that in surgically treated patients without misery perfusion ( $-5.2 \pm 9.4$ ) during  $6 \pm 4$  months ( $p < 0.001$ ), and in medically treated patients with misery perfusion ( $3.2 \pm 15.4$ ) during  $16 \pm 6$  months ( $p < 0.01$ ).

Therefore, the results suggest that the progression of cortical neuronal damage in surgically treated patients with misery perfusion and no perioperative stroke may be more extensive than in medically treated patients with misery perfusion and no intervening ischemic event. The progression of cortical neuronal damage should be considered while using bypass surgery for treating patients with misery perfusion.

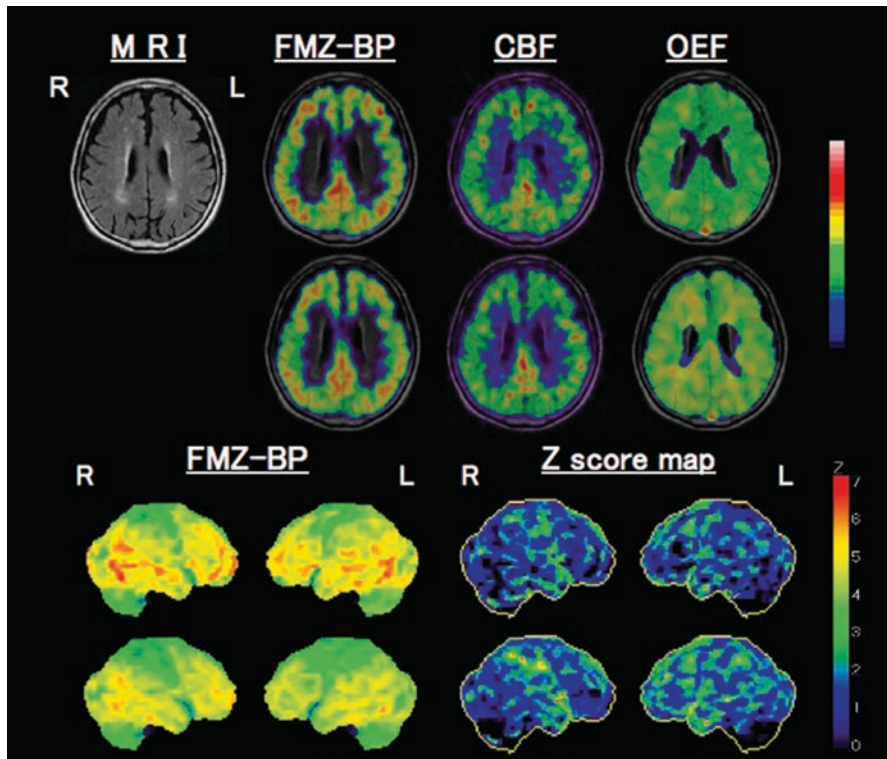
### 9.6.2 Blood Pressure Control

The treatment of hypertension is generally beneficial to patients with atherosclerotic major cerebral artery diseases (Rashid et al. 2003); however, the optimal blood pressure (BP) level to achieve maximal benefits among patients with this condition remains unknown. In patients with atherosclerotic major cerebral artery disease, low BP may impair cerebral perfusion, thereby worsening the risk of ischemic cerebral damage. It is thus debatable whether BP regulated to lower levels may be associated with the reduced risks of a recurrent stroke and hemodynamic impairment in these patients (Amin-Hanjani et al. 2017; Powers et al. 2014; Yamauchi et al. 2013, 2015).

Our previous studies in patients with ICA or MCA disease showed that the incidence of ischemic stroke in the territory of the diseased artery was significantly higher in patients with low systolic BP (SBP) ( $<130$  mmHg) (Yamauchi et al. 2013, 2015). This was particularly true in patients with impaired perfusion, characterized by a decreased CBF/cerebral blood volume ratio. Furthermore, low SBP during follow-up may be associated with hemodynamic deterioration (Yamauchi et al. 2019a) and consequently with a risk of selective neuronal damage.

To determine whether low BP at follow-up is associated with increased selective neuronal damage, we retrospectively analyzed data from 76 medically treated patients with atherosclerotic ICA or MCA disease with no ischemic episodes reported on a follow-up of 6 months or more (Yamauchi et al. 2019b). In all patients the distribution of cBZR was measured using  $^{11}\text{C}$ -flumazenil PET. The changes in the BZR index during follow-up were negatively correlated with SBP at follow-up. More significant increases in the BZR index indicating neuronal damage were observed in patients with lower SBP levels and decreased CBF than in patients without such decreases (Fig. 9.5). Therefore, low SBP may contribute to the risk of cortical neuronal damage in hemispheres with decreased CBF.

Further studies are needed to determine the optimal BP in patients with atherosclerotic major cerebral artery disease for favorable management of the condition.



**Fig. 9.5** Progressive cortical neuronal damage and low blood pressure in a patient with a right internal carotid artery occlusion and a left internal carotid artery stenosis (mild). Collateral pathways included the anterior communicating artery and leptomeningeal collaterals from the posterior cerebral artery. The first PET study (first row) revealed a mild decrease in flumazenil-binding potential (FMZ-BP) in the right (R) hemisphere with internal carotid artery occlusion and subcortical ischemic lesions (MRI), while cerebral blood flow (CBF) was decreased and oxygen extraction fraction (OEF) was increased slightly. A follow-up 48 months later (second row) revealed decreased FMZ-BP and CBF with increased OEF in the right hemisphere. Mean hemispheric CBF decreased from 28.4 to 27.5 (mL/100 g/min) while OEF increased from 51.5 to 57.0 (%). 3D-SSP images and Z score maps from the first (third row) and second (fourth row) examinations demonstrated a decrease in FMZ-BP in the right hemisphere, especially in the frontoparietal region. The BZR index increased from 35.9 to 103.1 between baseline and follow-up. The patient's systolic blood pressure was 119 (mmHg) at baseline and 107 at follow-up. Antihypertensive therapy was discontinued proximally to the follow-up examinations. The patient showed a mild attention deficit (a Mini Mental State Examination score, 26) (Yamauchi et al. 2019b)

## 9.7 Conclusions

1. In patients with atherosclerotic ICA or MCA occlusive disease, misery perfusion contributes to selective neuronal damage characterized by a decreased cBZR distribution in the apparently unremarkable cerebral cortex.
2. Hemodynamic deterioration may contribute to selective neuronal damage in the absence of an overt episode of stroke.
3. Selective neuronal damage may affect the functional outcomes.
4. Careful medical management and successful vascular reconstruction surgery in patients with misery perfusion may prevent selective neuronal damage.
5. Statins and angiotensin receptor blockers may reduce the occurrence of selective neuronal damage.
6. Imaging of cBZR is a potential technique to quantitatively evaluate the success of future therapeutic interventions to protect neurons from ischemic damage.

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# PET Imaging of Cyclooxygenases in Neuroinflammation

# 10

Atul Bhardwaj and Frank Wuest

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## Abstract

Neuroinflammation has detrimental effects on brain functions and plays critical roles in the pathogenesis and progression of several neurodegenerative diseases, including Alzheimer's and Parkinson's disease, and dementia. Cyclooxygenase isoenzymes COX-1 and COX-2 are responsible for the conversion of arachidonic acid to prostanoids such as prostaglandins, thromboxanes, and prostacyclins. Aberrant expression and function of COX-1 and COX-2 isoforms in the brain are linked to various inflammatory conditions leading to neural death and manifestation of neurodegenerative disorders. Consequently, COX-1 and COX-2 have emerged as important biomarkers for functional imaging of neuroinflammation and associated neurodegenerative diseases with positron-emission tomography (PET). This chapter reviews recent advances of PET imaging agents for the non-invasive visualization and characterization of COX-1 and COX-2 enzymes in neuroinflammation.

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## 10.1 Introduction

### 10.1.1 Neuroinflammation

Inflammation is an essential and vital part of the body's defense mechanism to respond to injuries and infections. The inflammatory response is a crucial signal to the body's immune system to trigger the healing and repair of injured tissue, as well as to mobilize the defense system against invasive microbes such as viruses and bacteria (Medzhitov 2008; DiSabato et al. 2016; Schmid et al. 2009; Schain and Kreisl 2017a; Glass et al. 2010). If the inflammatory response is not successful and fails to counteract the effects of tissue injury and infection, conditions can lead to chronic inflammation and start of vicious processes resulting in further damage of injured tissue and surrounding healthy tissue. Aberrant inflammatory responses have detrimental effects on organ and tissue function and contribute to the development and manifestation of many chronic diseases. Molecular imaging of inflammatory processes and crucial biomarkers of inflammation represents an important tool for early and accurate diagnosis and staging of inflammatory disease as well as for monitoring the response to therapeutic interventions (Narayanaswami et al. 2018; Winkeler et al. 2010).

This becomes particularly important when the inflammatory response is linked to the central nervous system (CNS) as damaged neural tissues can remain unnoticed for several years before the development of symptoms. Inflammation of the CNS (neuroinflammation) is a primary driver and contributor to the development and progression of several neurodegenerative diseases such as Alzheimer's disease (Graeber 2010; Wyss-Coray and Rogers 2012; Glass et al. 2010; Ransohoff 2016; Schain and Kreisl 2017b; Wyss-Coray and Mucke 2012; Craft et al. 2005), Parkinson's disease (Chen et al. 2005; Hirsch and Hunot 2009; Phani et al. 2012),

amyotrophic lateral sclerosis, multiple sclerosis, and traumatic brain injury (Bartels and Leenders 2010; Calsolaro and Edison 2016; Stephenson et al. 2018; Amor et al. 2014; Lymana et al. 2014; Tansey et al. 2007; Ransohoff 2016).

### 10.1.2 Cyclooxygenases, Neuroinflammation, and Neurodegenerative Diseases

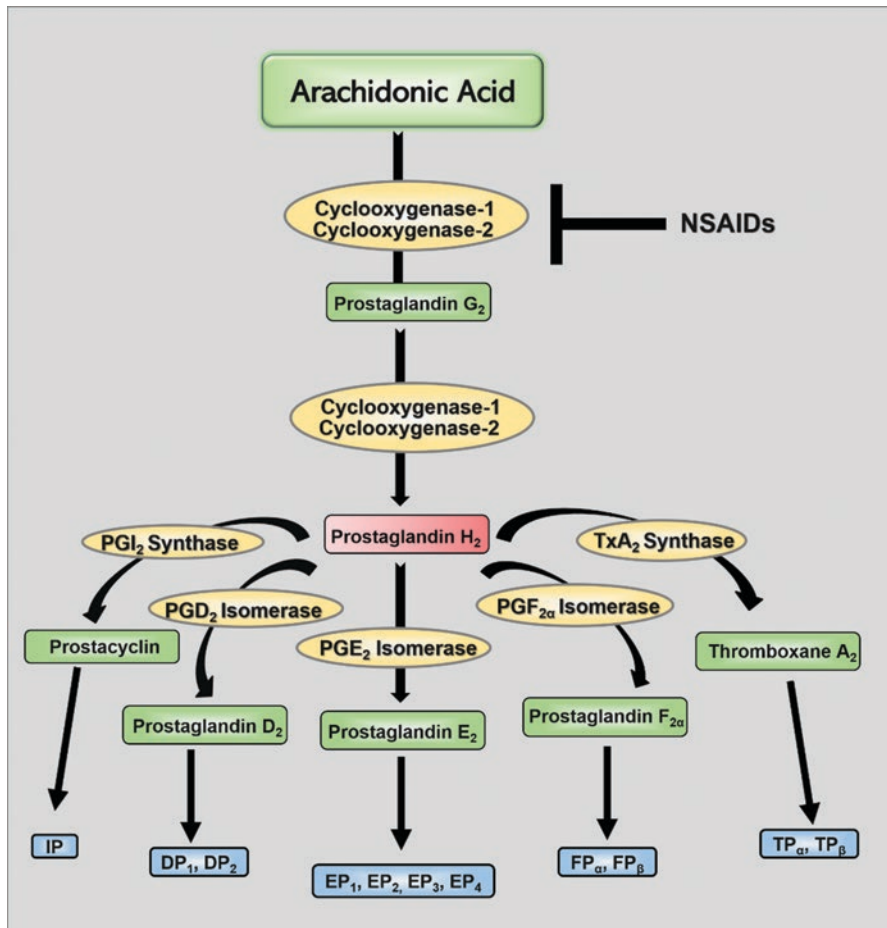
Cyclooxygenases (COXs) play a central role in the arachidonic acid metabolism cascade. COXs are involved in the activation of various inflammatory pathways that generate prostaglandins (PGs), chemokines, and cytokines as well as various reactive oxygen species (Smith et al. 2011; van der Donk et al. 2002; Blobaum and Marnett 2007; Marnett 2009). The excessive and sustained release of these substances can lead to chronic inflammation and oxidative stress which is harmful to any organ, including the brain. The COX-mediated release of prostaglandins as inflammatory mediators is a significant contributor to neuroinflammation and neurodegenerative diseases (Choi et al. 2009a; Ricciotti and FitzGerald 2011; Aid and Bosetti 2011; Dargahi et al. 2011; Phillis et al. 2006). Therefore, COXs continue as a primary drug target for the treatment and as a biomarker for the diagnosis of chronic inflammatory conditions, including pain, fever, rheumatoid arthritis, cancer, and neurodegenerative diseases. Several classes of COX inhibitors have been developed, also referred to as nonsteroidal anti-inflammatory drugs (NSAIDs). Many COX inhibitors were tested as therapeutic candidates for various CNS disorders (Imbimbo et al. 2010; Vlad et al. 2008; Rogers et al. 1993; McGeer and McGeer 2007; Cudaback et al. 2014).

COXs are rate-limiting enzymes in the arachidonic acid pathway (Fig. 10.1).

COXs first oxidize arachidonic acid to prostaglandin G<sub>2</sub> (PGG<sub>2</sub>) which is further reduced into prostaglandin H<sub>2</sub> (PGH<sub>2</sub>) (Smith et al. 1996; Kurumbail et al. 1996a; Smith et al. 2000; Simmons et al. 2004; Chandrasekharan et al. 2002a; Mitchell and Warner 1999). Various tissue-specific enzymes (isomerases) metabolize PGH<sub>2</sub> into prostanoids, including prostaglandins (PGs: PGE<sub>2</sub>, PGD<sub>2</sub>, PGF<sub>2</sub>α), prostacyclin (PGI<sub>2</sub>), and thromboxanes (TXs). Prostanoids bind to G-protein-coupled receptors (e.g., EP<sub>1-4</sub>) and mediate a variety of physiological and pathological processes, including inflammation.

COXs exist in three isoforms: a constitutive isoform, COX-1; a mitogen-inducible isoform, COX-2; and a splice variant of COX-1, COX-3 (Smith et al. 1996; Marnett et al. 1999; Kurumbail et al. 1996b; Chandrasekharan et al. 2002b; Choi et al. 2009b). The human COX-1 enzyme (599 amino acids, 68.686 kDa) is constitutively expressed in most tissues and responsible for the physiological production of PGs involved in the regulation of processes such as thrombogenesis and homeostasis of the gastrointestinal tract and kidneys.

In addition to its homeostatic functions in various organs, the COX-1 isoform is also considered as a major player in neuroinflammation. Several studies indicated that COX-1 is expressed in the midbrain, pons, and medulla. Moreover, COX-1 plays a critical role in brain injuries induced by pro-inflammatory stimuli including



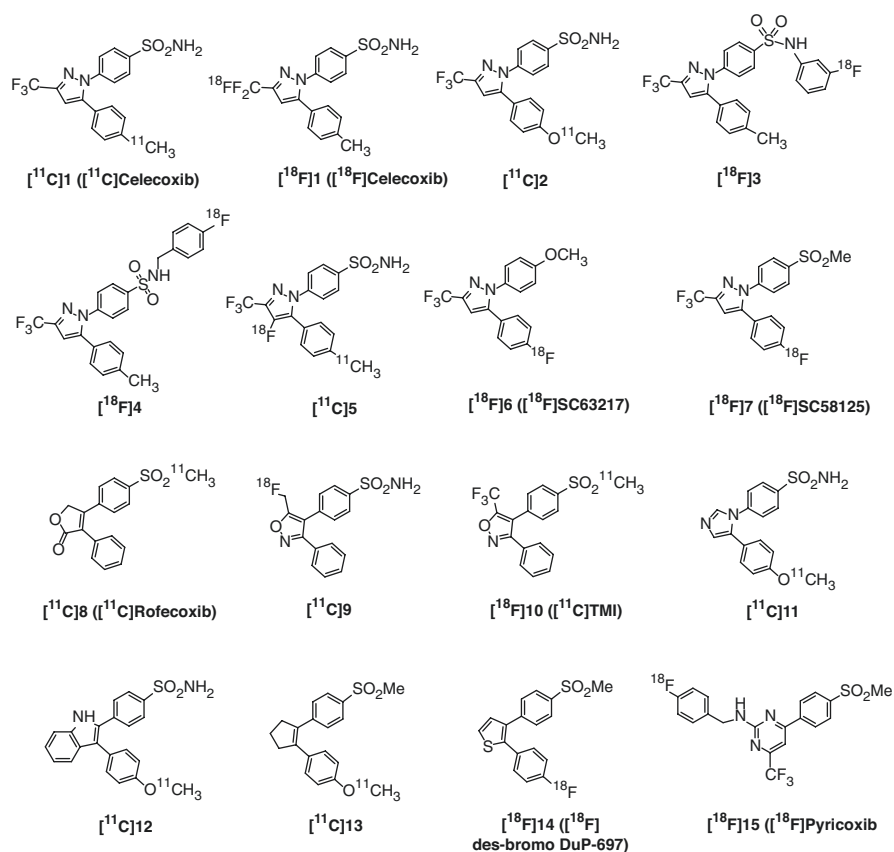
**Fig. 10.1** Cyclooxygenase-mediated metabolic pathway of arachidonic acid

prostaglandin E2 (PGE2) or cytokines such as interleukin (IL)-1 $\beta$ , lipopolysaccharide, IL-1 $\beta$ , and TNF $\alpha$  (Hoozemans et al. 2001; García-Bueno et al. 2009; Yermakova et al. 1999; Bosetti 2007). COX-1 is also involved in microglial activation during neuroinflammation (Aid et al. 2010; Toscano et al. 2007). The pharmacological inhibition or genetic deletion of COX-1 isoform resulted in the reduction of oxidative stress and neuronal damage (Choi et al. 2008; Choi and Bosetti 2009; Matousek et al. 2010).

The human COX-2 isoform (604 amino acids, 68.996 kDa) is only scarcely expressed in most tissues and organs under normal conditions and is mainly induced by several inflammatory stimuli, hormones, and growth factors. COX-2 is an established therapeutic target for various chronic inflammatory diseases, including cancer (Wang and DuBois 2010, 2013; Meric et al. 2006). Although COX-2 is mostly referred to as the “inducible” COX isoform, COX-2 is constitutively expressed throughout the forebrain, including hippocampal and cortical glutamatergic

neurons. The COX-2 isoform plays a central role in synaptic activity and long-term synaptic plasticity (Wang et al. 2005; Stefanovic et al. 2006; Hewett et al. 2006; Slanina and Schweitzer 2005). Possible involvement of COX-2 isoform in neurotoxicity is still subject to controversial debates but several studies demonstrated that overexpression of COX-2 drives neurological disorders such as Alzheimer's disease, Parkinson's disease, multiple sclerosis, long-term potentiation (LTP), stroke, epilepsy, schizophrenia, and bipolar disorder (Yamagata et al. 1993; Yang and Chen 2008; Teismann et al. 2003; Blais et al. 2005).

The discovery of novel COX-1 and COX-2 selective inhibitors also referred to as Coxibs has stimulated the development of COX-imaging agents for imaging biomarker research (Bhardwaj et al. 2013, 2017; Marnett 2012; Uddin et al. 2010; Kaur et al. 2015; Laube et al. 2013; Morgenroth et al. 2017; Tietz et al. 2016). During the last two decades, several COX-1 and COX-2 selective fluorescent and radiolabeled imaging agents have been developed and tested. A selection of  $^{11}\text{C}$ - and  $^{18}\text{F}$ -labeled PET imaging agents is depicted in Fig. 10.2 (Tietz et al. 2013; Pacelli et al. 2014; de Vries 2006).



**Fig. 10.2** COX-1- and COX-2-targeted PET imaging probes labeled with  $^{11}\text{C}$  and  $^{18}\text{F}$

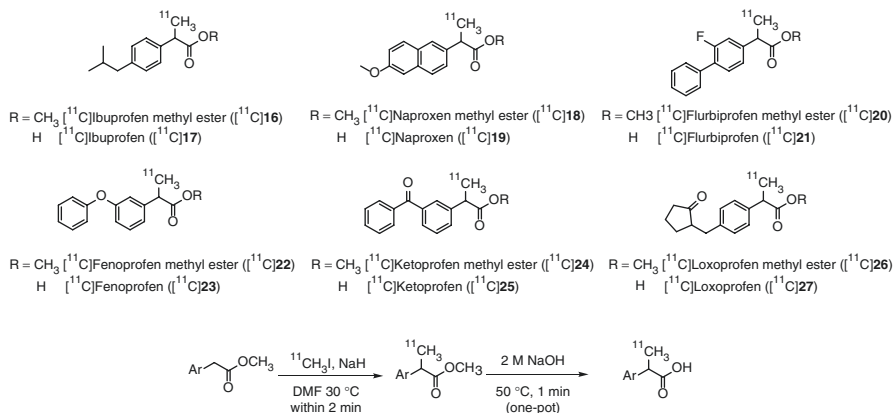
However, to date, no COX PET imaging agent has received approval for clinical use. Significant challenges and limitations of PET imaging agents for COXs include low in vivo affinity, high blood pool retention, poor metabolic stability, low uptake in target organs and tissues, and nonspecific binding. The following sections discuss the current status of radiotracer development for PET imaging of COX-1 and COX-2 in the brain.

## 10.2 PET Imaging Agents for COX-1

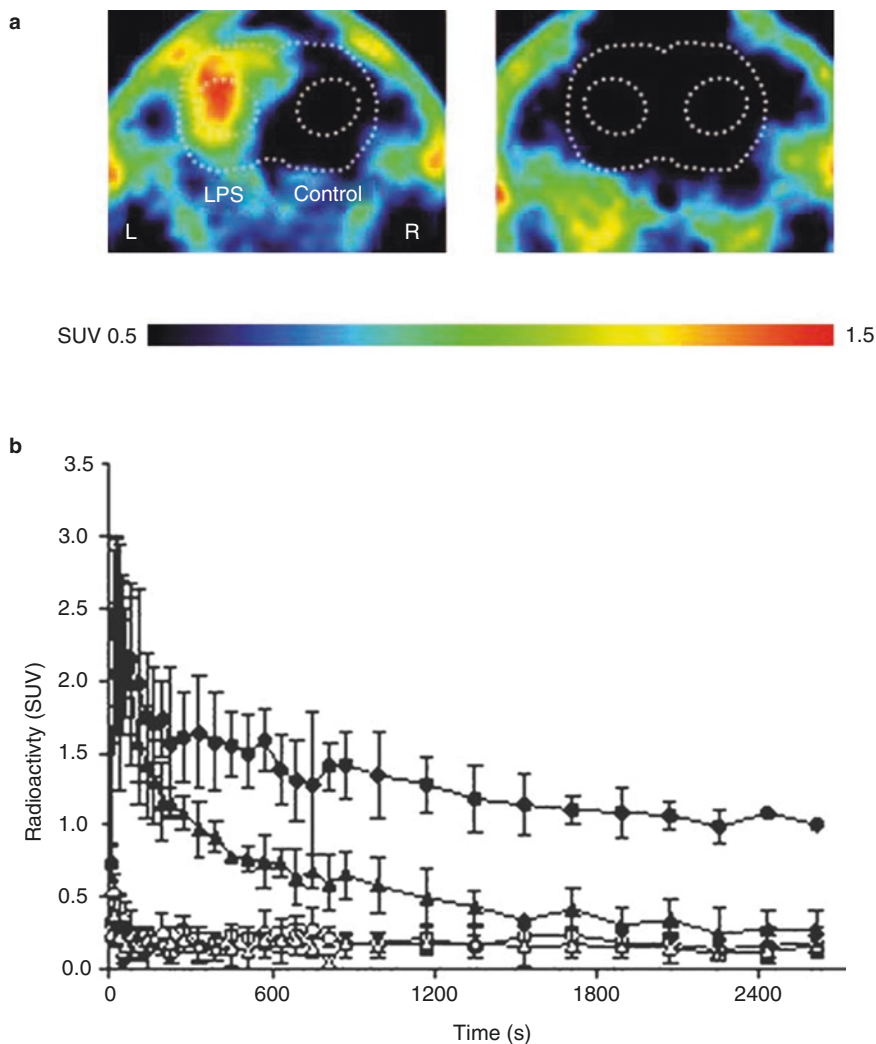
### 10.2.1 [ $^{11}\text{C}$ ]Ketoprofen Methyl Ester

PET imaging of COX-1 was advanced to clinical testing using  $^{11}\text{C}$ -labeled ketoprofen methyl ester ( $^{11}\text{C}$ -KTP-Me, [ $^{11}\text{C}$ ]**24**) as a radiotracer. The first PET imaging of COX-1 in activated microglia was successfully achieved with racemic (*R,S*)- $^{11}\text{C}$ -KTP-Me (Takashima-Hirano et al. 2010). The radiosynthesis of racemic (*R,S*)- $^{11}\text{C}$ -KTP-Me was reported along with the preparation of a series of  $^{11}\text{C}$ -labeled arylpropionic acids of six NSAIDs (ibuprofen, naproxen, flurbiprofen, fenoprofen, ketoprofen, and loxoprofen) and their respective methyl esters (Fig. 10.3).

COX-1 selective inhibitor ketoprofen is widely used as an analgesic and anti-pyretic drug. The use of the respective methyl ester (*R,S*)- $^{11}\text{C}$ -KTP-Me was motivated to overcome the poor blood-brain barrier (BBB) permeability of ketoprofen. Racemic (*R,S*)- $^{11}\text{C}$ -KTP-Me was synthesized by a [ $^{11}\text{C}$ ]methylation reaction of (3-benzophenyl)acetic acid methyl ester with [ $^{11}\text{C}$ ]CH $_3$ I in the presence of NaH at 30 °C, and the obtained chemical and radiochemical purity was >99% (Takashima-Hirano et al. 2010). Corresponding (*R,S*)- $^{11}\text{C}$ ketoprofen was prepared by treating methyl ester (*R,S*)- $^{11}\text{C}$ -KTP-Me with 2 M NaOH. Extension of the [ $^{11}\text{C}$ ]methylation reaction time beyond 2 min and use of reaction temperatures of 50 °C resulted in lower yield and decomposition of (*R,S*)- $^{11}\text{C}$ -KTP-Me into unknown side products.



**Fig. 10.3**  $^{11}\text{C}$ -labeled arylpropionic acid derivatives (Takashima-Hirano et al. 2010)



**Fig. 10.4** (a) Summarized PET images (from 5 to 45 min after radiotracer injection) of  $[^{11}\text{C}]\mathbf{24}$  (left panel) and  $[^{11}\text{C}]\mathbf{25}$  (right panel) in rat brain inflammation induced by LPS (50 mg) injection into the left striatum. (b) Time-activity curves of  $[^{11}\text{C}]\mathbf{24}$  ( $n = 2$ ) and  $[^{11}\text{C}]\mathbf{25}$  ( $n = 3$ ) in LPS-injected inflammatory area (LPS) and contralateral region (control). Data are expressed as mean  $\pm$  SD ( $\bullet$ : LPS  $[^{11}\text{C}]\mathbf{24}$ ,  $\blacktriangle$ : control  $[^{11}\text{C}]\mathbf{25}$ ;  $\circ$ : LPS  $[^{11}\text{C}]\mathbf{24}$ ,  $\Delta$ : control  $[^{11}\text{C}]\mathbf{25}$ ) (Takashima-Hirano et al. 2010)

PET imaging with  $(R,S)\text{-}^{11}\text{C}\text{-KTP-Me}$  in rat brain demonstrated specific accumulation of radioactivity in the inflamed region (Fig. 10.4) (Takashima-Hirano et al. 2010). The obtained SUV of  $(R,S)\text{-}^{11}\text{C}\text{-KTP-Me}$  was comparable to that of  $[^{11}\text{C}]\text{PK11195}$ , a commonly used PET radiotracer for imaging peripheral benzodiazepine receptors in activated microglia during neuroinflammation. Radiotracer

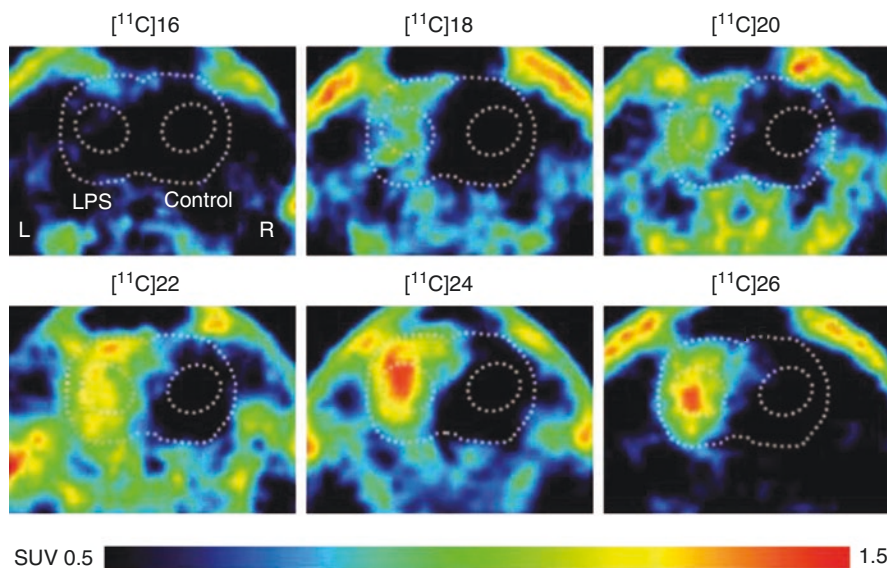
(*R,S*)- $^{11}\text{C}$ -ketoprofen ( $^{11}\text{C}$ 24) was not accumulated in the brain due to its poor blood-brain barrier (BBB) permeability (Fig. 10.4) (Takashima-Hirano et al. 2010).

Metabolite analysis of (*R,S*)- $^{11}\text{C}$ -KTP-Me in the rat brain and blood showed that >90% of the radiotracer hydrolyzed to its pharmacologically active form (*R,S*)- $^{11}\text{C}$ -ketoprofen within 5 min postinjection. In the rat model of neuroinflammation model, all 2-aryl  $^{11}\text{C}$ propionic acid methyl esters showed high accumulation in the area of lipopolysaccharide (LPS)-induced inflammation, except for  $^{11}\text{C}$ ibuprofen methyl ester  $^{11}\text{C}$ 24 (Fig. 10.5) (Takashima-Hirano et al. 2010).

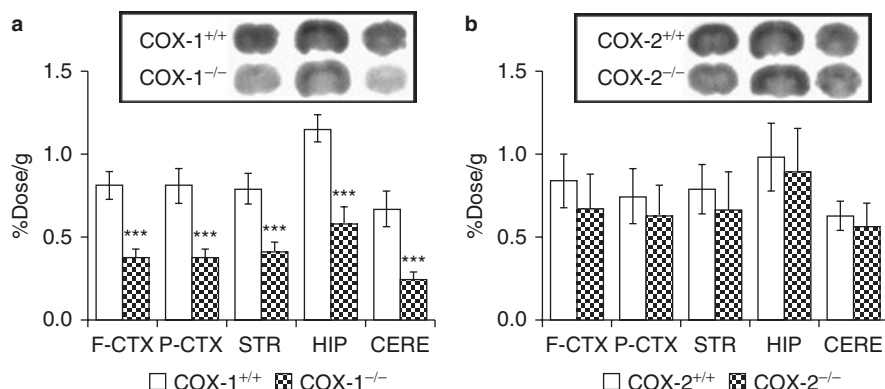
(*R,S*)- $^{11}\text{C}$ -KTP-Me displayed the highest radioactivity accumulation in inflammatory regions compared to all other tested radiotracers. The observed significant blocking effect with pharmacological doses of ketoprofen methyl ester on the radioactivity accumulation of (*R,S*)- $^{11}\text{C}$ -KTP-Me in brain regions of inflammation was indicative of high binding specificity of the radiotracer to COX.

Additional research involved ex vivo and in vivo imaging studies with  $^{11}\text{C}$ -KTP-Me to study the COX-1 and COX-2 selectivity and specificity profile of the radiotracer (Shukuri et al. 2011). High accumulation of radiotracer of  $^{11}\text{C}$ -KTP-Me was observed in the hippocampus, cerebral cortex, and cerebellum of wild-type (COX-1<sup>+/+</sup>, COX-2<sup>+/+</sup>) mice, whereas uptake of  $^{11}\text{C}$ -KTP-Me was significantly lower in COX-1<sup>-/-</sup> mice (Fig. 10.6a). No differences were detected between COX-2<sup>-/-</sup> and wild-type COX-2<sup>+/+</sup> mice (Fig. 10.6b).

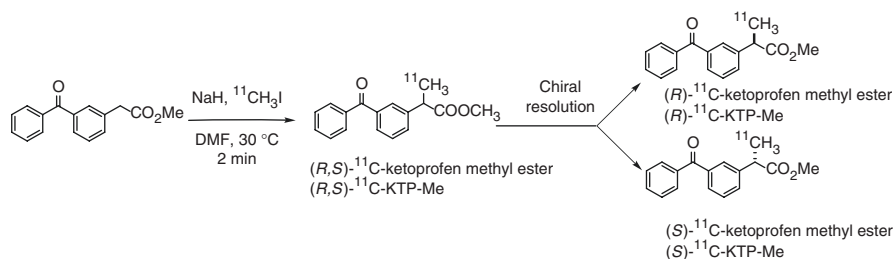
This experiment provided important evidence that radiotracer  $^{11}\text{C}$ -KTP-Me selectively binds to the COX-1 isoform in the brain (Fig. 10.6) (Shukuri et al. 2011).



**Fig. 10.5** PET images of 2-aryl $^{11}\text{C}$ propionic acid methyl esters in rat brain inflammation induced by LPS (50 mg) injection into the left striatum (Takashima-Hirano et al. 2010)



**Fig. 10.6** Accumulation of  $^{11}\text{C}$ -ketoprofen methyl ester  $^{11}\text{C}$ -KTP-Me in the brain of  $\text{COX-1}^{-/-}$  and  $\text{COX-2}^{-/-}$  mice. Ex vivo autoradiography and regional brain tissue uptake of  $^{11}\text{C}$ -KTP-Me in  $\text{COX-1}^{+/+}$  and  $\text{COX-1}^{-/-}$  mice (a) and  $\text{COX-2}^{+/+}$  and  $\text{COX-2}^{-/-}$  mice (b) were obtained 15 min after injection of the radiotracer. F-CTX 5 frontal cortex; P-CTX 5 parietal cortex; STR 5 striatum; HIP 5 hippocampus; CERE 5 cerebellum. Data are mean  $\pm$  SD ( $n = 6$ ). \*\*\* $P$ , 0.001 versus wild-type mice (Shukuri et al. 2011)



**Fig. 10.7** Synthesis of (*R*)- and (*S*)- $^{11}\text{C}$ -KTP-Me via rapid  $\text{C-}^{11}\text{C}$ -methylation and chiral resolution (Shukuri et al. 2016)

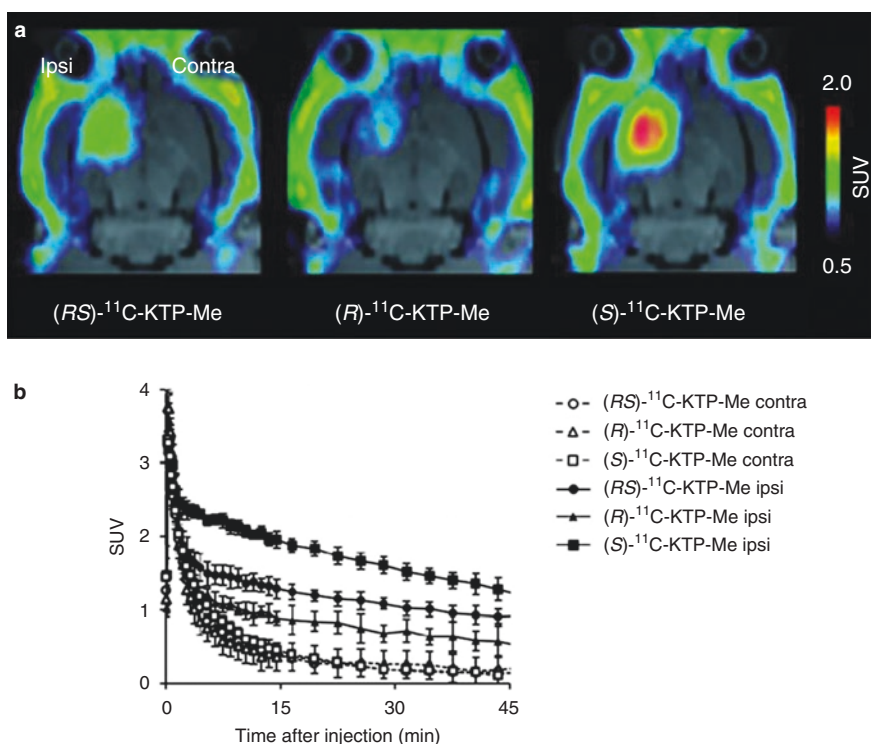
To this point, racemic (*R,S*)- $^{11}\text{C}$ -KTP-Me was used for all COX-targeted PET imaging studies in the brain, although the (*S*)-enantiomer of ketoprofen is known to be pharmacologically more active than the (*R*)-enantiomer. Thus, Onoe et al. separated both isomers using a solid chiral resolution method with semi-preparative high-performance liquid chromatography (HPLC) (Fig. 10.7) (Shukuri et al. 2016).

The (*S*)- $^{11}\text{C}$ -KTP-Me enantiomer was identified as an improved PET imaging agent with high COX-1 specificity. Radiotracer (*S*)- $^{11}\text{C}$ -KTP-Me was further investigated for COX-1 imaging during the progression of Alzheimer's disease in a transgenic mouse model. In contrast to COX-1 inhibitory activity of (*R*)- $^{11}\text{C}$ -KTP-Me ( $\text{IC}_{50} = 126.7 \mu\text{M}$ ), both racemic (*R,S*)- $^{11}\text{C}$ -KTP-Me ( $\text{IC}_{50} = 0.031 \mu\text{M}$ ) and (*S*)- $^{11}\text{C}$ -KTP-Me ( $\text{IC}_{50} = 0.011 \mu\text{M}$ ) showed higher inhibitory potency for COX-1 (Shukuri et al. 2016). Moreover, PET imaging of neuroinflammation in rat brain also showed 1.5-fold higher radioactivity accumulation in LPS-induced inflamed region after

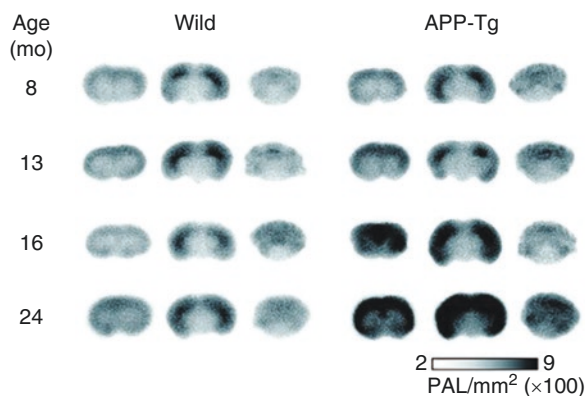


injection of (*S*)- $^{11}\text{C}$ -KTP-Me compared to racemic (*R,S*)- $^{11}\text{C}$ -KTP-Me. Enantiomer (*R*)- $^{11}\text{C}$ -KTP-Me displayed only low and nonspecific radioactivity accumulation in the brain (Shukuri et al. 2016). Time-activity curve analyses of all three radiotracers revealed rapid washout from the brain within 15 min under normal conditions, whereas radioactivity remained in the striatum for 45 min in the case of (*R,S*)- $^{11}\text{C}$ -KTP-Me and (*S*)- $^{11}\text{C}$ -KTP-Me after the LPS-induced neuroinflammation. Moreover, retention of radioactivity was improved significantly for (*S*)- $^{11}\text{C}$ -KTP-Me compared to racemic (*R,S*)- $^{11}\text{C}$ -KTP-Me (Fig. 10.8) (Shukuri et al. 2016).

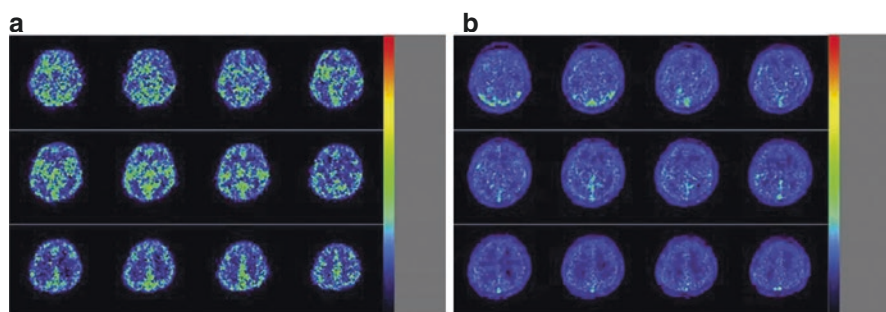
Ex vivo autoradiography analyses showed an age-dependent increase in (*S*)- $^{11}\text{C}$ -KTP-Me accumulation in all brain regions of amyloid precursor protein transgenic (APP-Tg) mice in direct comparison with radiotracer accumulation in wild-type mice (Fig. 10.9) (Shukuri et al. 2016).



**Fig. 10.8** Representative PET images and time-radioactivity curves of each  $^{11}\text{C}$ -KTP-Me enantiomer in rat brain after LPS injection. (a) Transaxial rat brain views of SUV-summed PET images from 5 to 45 min after tracer injection were coregistered with individual MR images at baseline. (*S*)- $^{11}\text{C}$ -KTP-Me showed higher accumulation and superior specificity in rat brain hemisphere than (*R,S*)- and (*R*)- $^{11}\text{C}$ -KTP-Me. (b) Quantitative time-radioactivity curves of each  $^{11}\text{C}$ -KTP-Me enantiomer in contralateral (Contra) and ipsilateral (Ipsi) striatum. Data are expressed as SUV and are mean  $\pm$  SD ((*R*)- $^{11}\text{C}$ ]KTP-Me,  $n = 3$ ; others,  $n = 4$ ) (Shukuri et al. 2016)

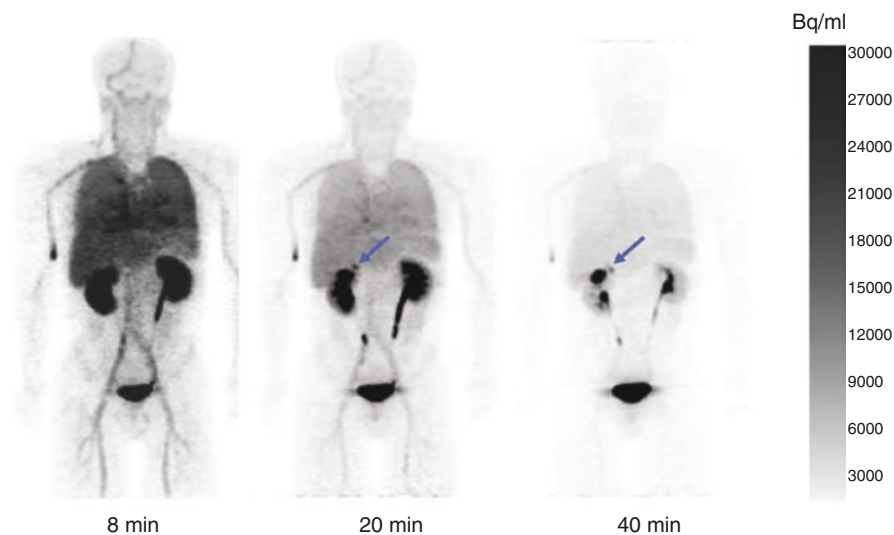


**Fig. 10.9** Age-dependent changes in (*S*)- $^{11}\text{C}$ -KTP-Me accumulation in the brain of APP-Tg 2576 and wild-type mice. Representative ex vivo autoradiographs of coronal sections of mice at 15 min after PET tracer injection. Progressive and remarkable increases in (*S*)- $^{11}\text{C}$ -KTP-Me in the frontal cortex, parietal occipital cortex, and hippocampus of APP-Tg mice of 16 months old. The range bar represents the intensity of photostimulated luminescence (PSL/ $\text{mm}^2$ )



**Fig. 10.10** Brain summation image during 2 min after injection of (*R,S*)- $^{11}\text{C}$ KTP-Me in a young normal volunteer (a). Late-phase brain summation image created from three whole-body scans (8, 20, and 40 min postinjection) (b) (Ohnishi et al. 2014)

The encouraging PET imaging results in animals stimulated efforts to explore the clinical utility of racemic (*R,S*)- $^{11}\text{C}$ KTP-Me for PET imaging of COX-1 in neuroinflammation, and a first-in-human study was performed in healthy volunteers (Ohnishi et al. 2014). In this study, the safety, radiation dosimetry, pharmacokinetics, biodistribution, and brain uptake of (*R,S*)- $^{11}\text{C}$ -KTP-Me were investigated. Optimization of the radiosynthesis in an automated synthesis unit based on previously reported methods provided the radiotracer in high radiochemical purity >99%. No adverse effects were observed after radiotracer administration. The radiotracer was rapidly metabolized to more hydrophilic  $^{11}\text{C}$ -KTP ( $^{11}\text{C}$ ]17) and cleared from the blood. The whole-brain and cerebral cortex time-activity curves indicated that radioactivity readily entered the brain, with an average peak cortical SUV of 1.5 in the cortex at 2 min postinjection. Then, radioactivity was gradually washed out presumably due to the lack of neuroinflammation (Fig. 10.10) (Ohnishi et al. 2014).



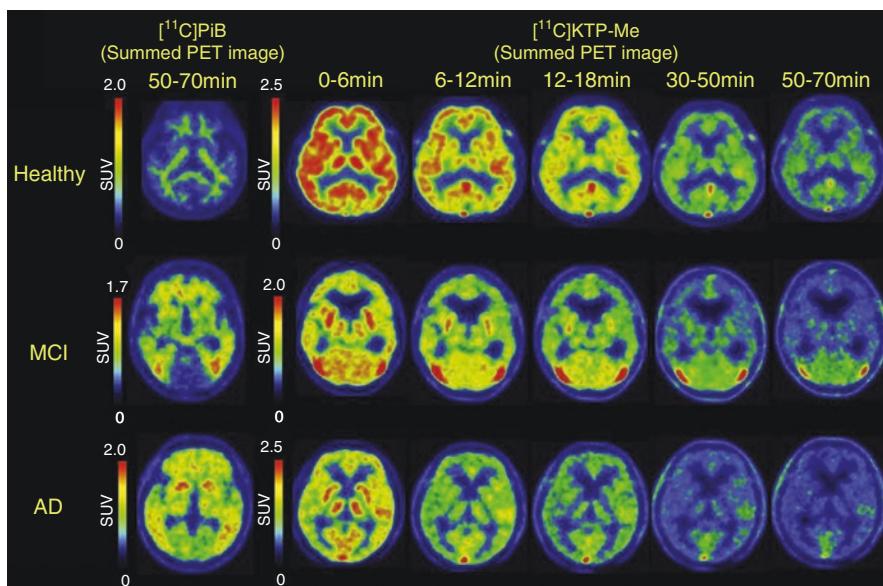
**Fig. 10.11** Biodistribution of radioactivity in a healthy volunteer obtained with whole-body scans starting 8, 20, and 40 min after injection of  $(R,S)$ - $^{11}\text{C}$ -KTP-Me. Images are decay corrected. Arrows ( $\leftarrow$ ) indicate biliary excretion (Ohnishi et al. 2014)

The biodistribution analyses revealed that urinary bladder was the major excretory pathway, and the organs with the highest absorbed radioactivity dose were the urinary bladder ( $41 \mu\text{Gy}/\text{MBq}$ ), the kidneys ( $23 \mu\text{Gy}/\text{MBq}$ ), and the lungs ( $13 \mu\text{Gy}/\text{MBq}$ ) (Fig. 10.11). Similar to other  $^{11}\text{C}$ -labeled PET radiotracers the calculated effective dose of  $(R,S)$ - $^{11}\text{C}$ -KTP-Me was  $4.7 \mu\text{Sv}/\text{MBq}$ .

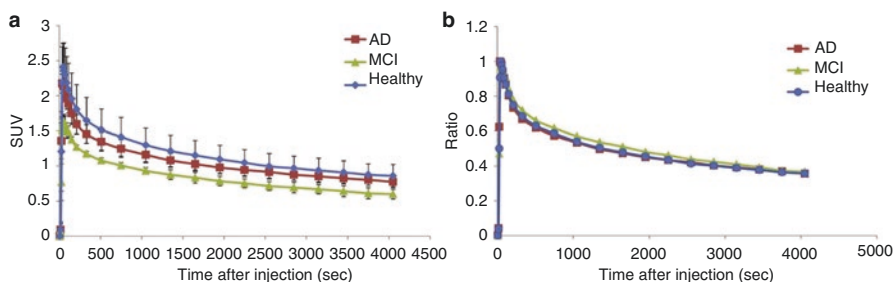
Radiotracer  $(R,S)$ - $^{11}\text{C}$ -KTP-Me was further investigated as a diagnostic biomarker for neuroinflammatory processes in Alzheimer's disease (AD) (Ohnishi et al. 2016).  $(R,S)$ - $^{11}\text{C}$ -KTP-Me was intravenously administered to healthy subjects, Pittsburgh compound-B (PiB)-positive patients with mild cognitive impairment (MCI), and PiB-positive AD patients. PET scans were recorded for 70 min (Fig. 10.12) (Ohnishi et al. 2016).

In all subjects, no abnormal symptoms, physical examination, vital signs, electrocardiogram, urine tests, or blood tests after injection of  $(R,S)$ - $^{11}\text{C}$ -KTP-Me were observed. Unfortunately, the analyses of average peak cortical SUV revealed no statistically significant difference between healthy and MCI/AD patients (Fig. 10.13) (Ohnishi et al. 2016).

The evaluation of regional uptake in healthy subjects and patients showed no connection between the distribution profile of  $(R,S)$ - $^{11}\text{C}$ -KTP-Me and  $^{11}\text{C}$ -PiB in the patient population. Overall, the results of this study indicated that  $^{11}\text{C}$ -KTP-Me enters the human brain but it was found not useful as a diagnostic biomarker for neuroinflammation in MCI/AD. Because of better COX-1 affinity of  $(S)$ - $^{11}\text{C}$ -KTP-Me over racemic  $^{11}\text{C}$ -KTP-Me, the authors suggested the evaluations with  $S$ -enantiomer in AD, MCI, and other neuroinflammatory conditions.



**Fig. 10.12** Brain summation images 50–70 min after injection of  $[^{11}\text{C}]\text{PiB}$  and during 0–6 min, 6–12 min, 12–18 min, 30–50 min, and 50–70 min after injection of  $(R,S)\text{-}[^{11}\text{C}]\text{KTP-Me}$  in a representative healthy subject, MCI patient, and AD patient. No obvious differences were observed in cortical or regional retention among the subjects (Ohnishi et al. 2016)



**Fig. 10.13** (a) Mean time-activity curves and SD of the cerebral cortex after injection of  $(R,S)\text{-}[^{11}\text{C}]\text{KTP-Me}$  in healthy subjects, MCI patients, and AD patients. (b) Time course of the ratio to the peak of each curve. No apparent differences were observed in the washout curve among the three groups (Ohnishi et al. 2016)

### 10.2.1.1 $[^{11}\text{C}](1,5\text{-bis}(4\text{-methoxyphenyl})\text{-}3\text{-}(2,2,2\text{-trifluoroethoxy})\text{-}1\text{H}\text{-}1,2,4\text{ triazoles}$ ( $[^{11}\text{C}]\text{PS13}$ , $[^{11}\text{C}]\text{PS1}$ , $[^2\text{H}_2, ^{18}\text{F}]\text{PS2}$ , and $[^{11}\text{C}]\text{PS13}$ )

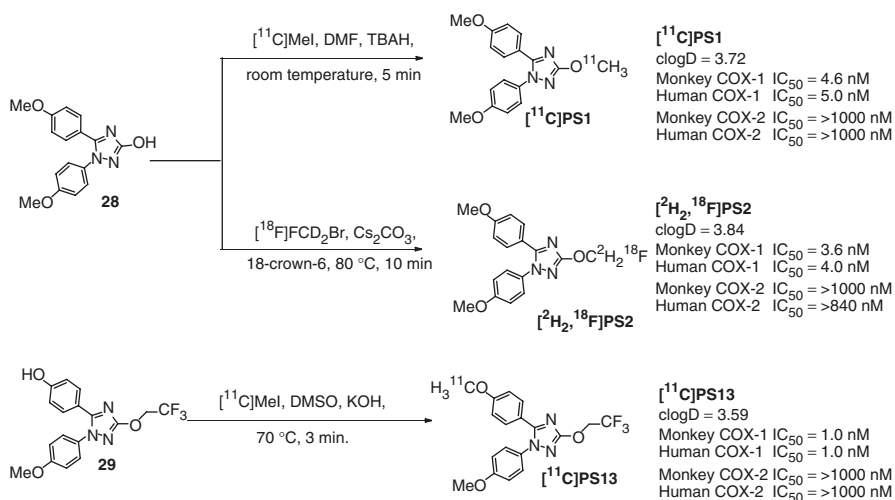
The limitations of using radiotracer  $[^{11}\text{C}]\text{KTP-Me}$  for PET imaging of COX-1 prompted the development of other  $^{11}\text{C}$ -labeled imaging agents such as  $[^{11}\text{C}]\text{PS13}$  (1,5-bis(4-methoxyphenyl)-3-(2,2,2-trifluoroethoxy)-1H-1,2,4 triazole) for

studying neuroinflammation in monkeys and humans. Seventeen analogs of potent and selective COX-1 inhibitor FK881 (PS1) ( $IC_{50} = 4.9$  nM (COX-1);  $IC_{50} = 3.2$   $\mu$ M (COX-2);  $clog D = 3.72$ ) were prepared and tested (Imanishi et al. 2011; Singh et al. 2018).

The high inhibitory potency and COX-1 selectivity profile and calculated moderate lipophilicities drove the development of  $^{11}C$ - and  $^{18}F$ -labeled radiotracer candidates for PET imaging in monkeys: 1,5-(4-methoxyphenyl)-1*H*-1,2,4-triazole [ $^{11}C$ ]PS1; 3- [ $^{18}F$ ]fluoromethoxy-substituted and deuterated compound [ $^{18}F$ ]PS2; and 3-(1,1,1-trifluoroethoxy) compound [ $^{11}C$ ]PS13 (Fig. 10.14).

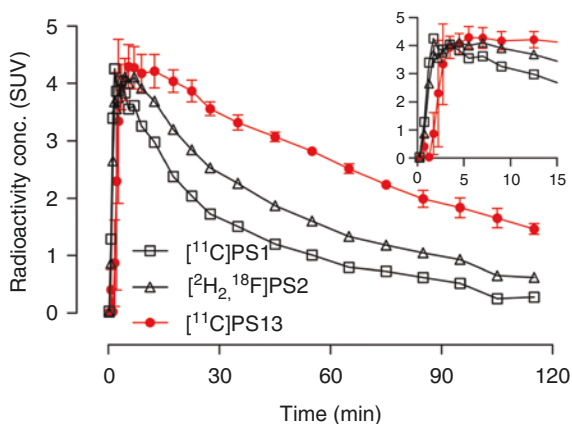
Radiotracer [ $^{11}C$ ]PS1 was synthesized by *O*-methylation of the 1,5-bis-(4-methoxy-phenyl)-1*H*-[1,2,4]triazol-3-ol with [ $^{11}C$ ]MeI in  $20 \pm 3\%$  radiochemical (from [ $^{11}C$ ]CO<sub>2</sub>) in  $>99\%$  radiochemical purity and with a molar activity of  $125 \pm 62$  GBq/ $\mu$ mol, radiotracer [ $^{18}F$ ]PS2 was obtained in  $14 \pm 4\%$  radiochemical yield in  $>99\%$  radiochemical purity and with a molar activity of  $53 \pm 17$  GBq/ $\mu$ mol by treating 1,5-bis-(4-methoxy-phenyl)-1*H*-[1,2,4]triazol-3-ol with deuterated  $d^2$ - [ $^{18}F$ ]fluorobromomethane, and compound [ $^{11}C$ ]PS13 was prepared in  $9.8 \pm 3.5\%$  radiochemical yield (from [ $^{11}C$ ]CO<sub>2</sub>) with  $>99\%$  radiochemical purity and with a molar activity of  $176 \pm 73$  GBq/ $\mu$ mol by the reaction of desmethoxy precursor 4-[2-(4-methoxy-phenyl)-5-(2,2,2-trifluoro-ethoxy)-2*H*-[1,2,4]triazol-3-yl]-phenol with [ $^{11}C$ ]MeI (Singh et al. 2018; Shrestha et al. 2018).

The PS1 displayed a high COX1 affinity ( $IC_{50} = 5$  nM) and selectivity over COX-2. In the monkey brain, radioactivity concentration for radioligand [ $^{11}C$ ]PS1 peaked at an SUV of 4.26 at 1.75 min, followed by a washout reaching an SUV of



**Fig. 10.14** Radiosynthesis of [ $^{11}C$ ]PS1, [ $^{18}F$ ]PS2, and [ $^{11}C$ ]PS13 (Singh et al. 2018; Shrestha et al. 2018)

**Fig. 10.15** PET time-activity curves from rhesus monkey whole brain (gray matter) following intravenous injection of [ $^{11}\text{C}$ ]PS1 ( $n = 1$ ), [ $^2\text{H}_2, ^{18}\text{F}$ ]PS2 ( $n = 1$ ), and [ $^{11}\text{C}$ ]PS13 (mean  $\pm$  SD,  $n = 3$ ). The inset expands the data presentation over the early stage of scanning (Shrestha et al. 2018)



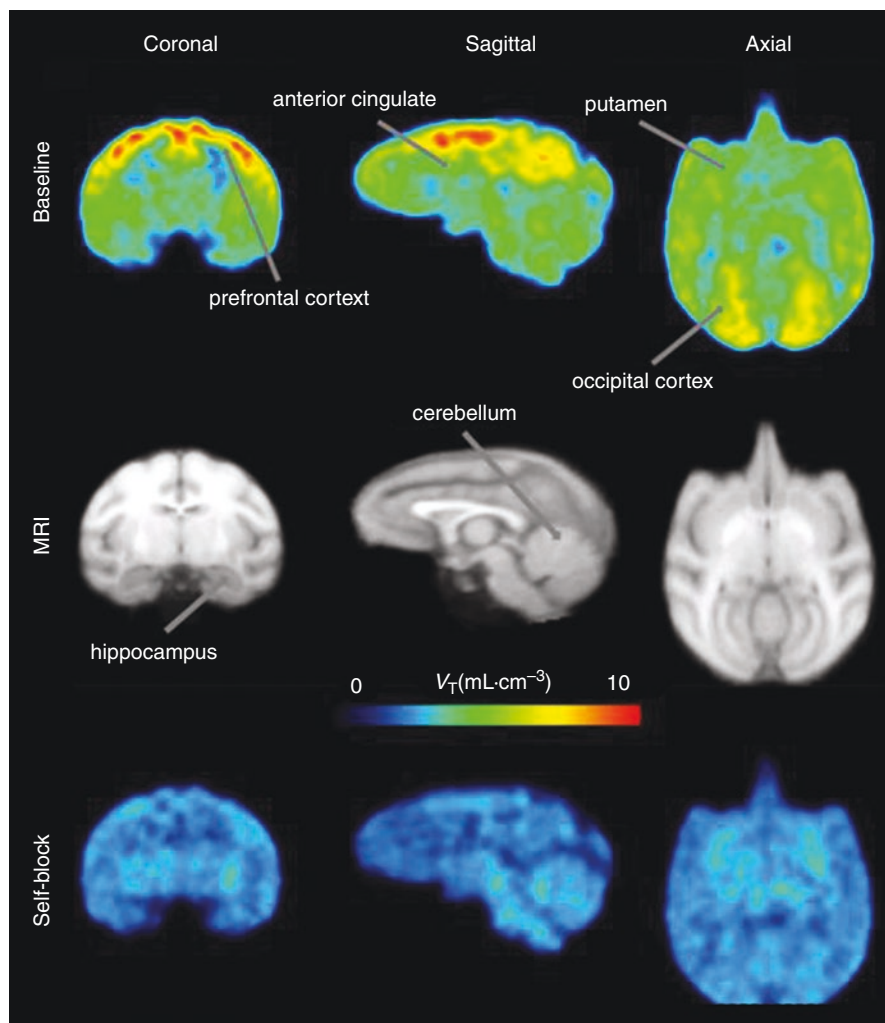
0.62 at 85 min. Compound PS2 also showed a high affinity for human COX1 (4 nM) with  $>200$  selectivity over COX-2. In the monkey brain, radiotracer [ $^2\text{H}_2, ^{18}\text{F}$ ]PS2 showed an SUV of 4.11 at 7 min postinjection which declined by 74% reaching an SUV = 1.05 at 85 min (Fig. 10.15) (Singh et al. 2018; Shrestha et al. 2018).

Despite the use of two deuterium atoms in the [ $^2\text{H}_2, ^{18}\text{F}$ ]fluoromethoxy group to reduce in vivo radiodefluorination, brain images with radiotracer [ $^2\text{H}_2, ^{18}\text{F}$ ]PS2 confirmed high radioactivity uptake in the skull which is indicative of radiodefluorination. Radiodefluorination and accumulation of radioactivity in the skull challenged the quantification of radiotracer [ $^2\text{H}_2, ^{18}\text{F}$ ]PS2 in the monkey brain, and radiotracer [ $^2\text{H}_2, ^{18}\text{F}$ ]PS2 was not used for further studies. COX-1 inhibitor PS13 had a high affinity for both monkey and human COX-1 ( $\text{IC}_{50} \sim 1$  nM) and a thousandfold selectivity over COX-2.

Radiotracer [ $^{11}\text{C}$ ]PS13 showed high radioactivity concentration in the monkey brain (SUV = 4.3, at 9 min) which was washed out by 54% reaching an SUV of 1.99 at 85 min (Fig. 10.15) (Shrestha et al. 2018). Among all tested 1,2,4-triazole radioligands (PS1, PS2, PS3), radiotracer [ $^{11}\text{C}$ ]PS13 displayed the most favorable characteristics for COX-1 imaging and quantification in the brain.

The COX-1 specificity of [ $^{11}\text{C}$ ]PS13 was further confirmed with blocking experiments, using pharmacological doses of PS13 and KTP-Me. From brain volume distribution ( $V_T$ ) estimations, high uptake of [ $^{11}\text{C}$ ]PS13 in the monkey brain was noticed in the neocortex, temporal cortex, striatum, thalamus, mid-brain, and hippocampus, in descending order (Fig. 10.16, top row). Blocking with PS13 or KTP-Me resulted in a reduction in  $V_T$  (Figs. 10.16 and 10.17) (Shrestha et al. 2018).

The excellent in vivo selectivity and PET imaging profile of radiotracer [ $^{11}\text{C}$ ]PS13 encouraged biodistribution studies in healthy human subjects (Anaya et al. 2019).

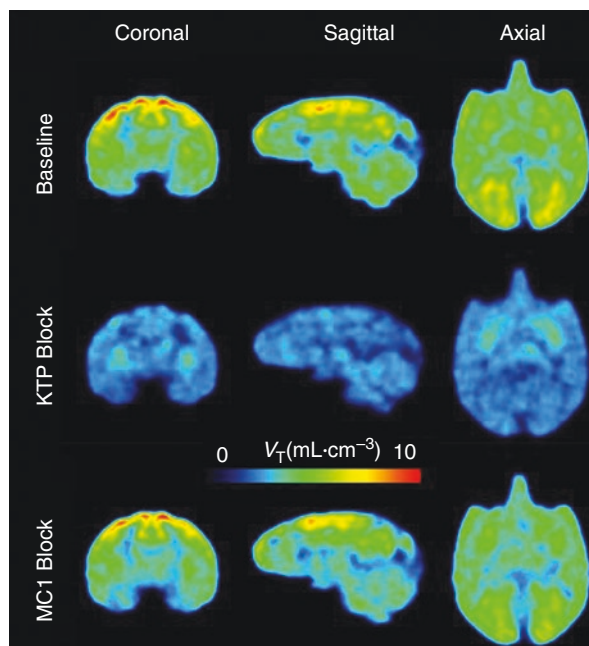


**Fig. 10.16** Parametric images of  $V_T$  for [ $^{11}\text{C}$ ]PS13 in the brain of a normal rhesus monkey at baseline (top row) and after blocking with PS13 (1 mg/kg, i.v., bottom row). Kinetic analysis was performed at the voxel level using a Logan plot so that the resulting  $V_T$  values are shown as PET images. The middle row shows template MRI scans (Shrestha et al. 2018)

### 10.3 PET Imaging Agents for COX-2

Despite the remarkable success in the development of potent and selective COX-2 inhibitors, no COX-2 targeted radioligands are currently available for clinical use. Research progress for the development of radiotracers for molecular imaging of

**Fig. 10.17** [ $^{11}\text{C}$ ]PS13 binds selectively to COX-1. Parametric images of  $V_T$  in the brain of a normal rhesus monkey at baseline (top row) and after administration of either KTP-Me (3 mg/kg, i.v.; as prodrug of COX-1-specific KTP; middle row) or MC1 (0.3 mg/kg, i.v.; COX-2-specific; bottom row). Only KTP-Me reduced whole-brain uptake of [ $^{11}\text{C}$ ]PS13 (Shrestha et al. 2018)



COX-2 has been summarized in several comprehensive review articles (Tietz et al. 2013; Pacelli et al. 2014; de Vries 2006). Selected examples of COX-2-targeted radiotracers are presented in Fig. 10.1. Among available COX-2 radiotracers, two compounds ([ $^{11}\text{C}$ ]MC1 and [ $^{11}\text{C}$ ]VA426) have recently been reported for PET imaging of COX-2 in the brain.

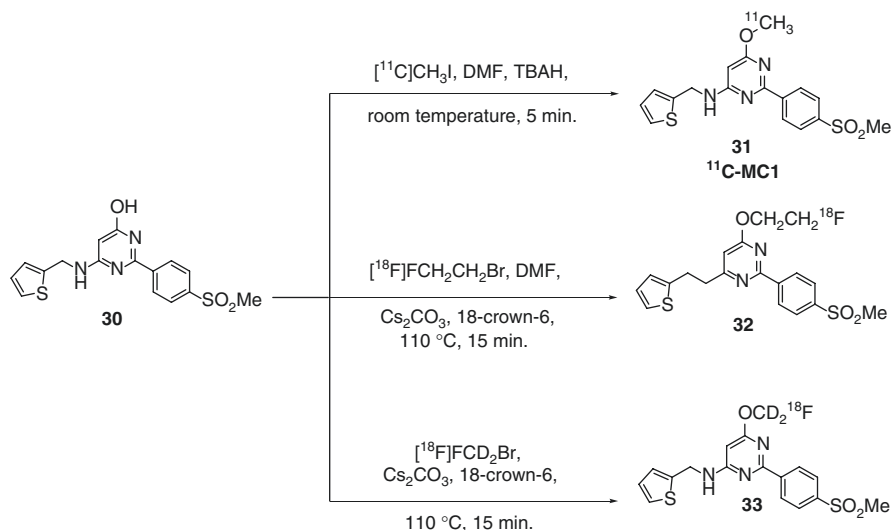
### 10.3.1 (6-[ $^{11}\text{C}$ ]Methoxy-2-(4-(methylsulfonyl)phenyl)-*N*-(thiophen-2-ylmethyl)-pyrimidin-4-amine ([ $^{11}\text{C}$ ]MC1)

In parallel to studies with COX-1-specific radiotracer [ $^{11}\text{C}$ ]PS13, compound 6-methoxy-2-(4-(methylsulfonyl)phenyl)-*N*-(thiophen-2-ylmethyl)pyrimidin-4-amine (MC1) was tested and also evaluated as a COX-2 selective inhibitor (Singh et al. 2018). The radiosynthesis [ $^{11}\text{C}$ ]MC1 and related  $^{18}\text{F}$ -labeled imaging agents **32** and **33** are displayed in Fig. 10.18.

Compound MC1 displayed high affinity and selectivity for monkey ( $\text{IC}_{50} = 2.58 \text{ nM}$ ) and human ( $\text{IC}_{50} = 2.89 \text{ nM}$ ) COX-2 and a favorable  $\log D$  (3.74) (Cortes-Salva et al. 2018).

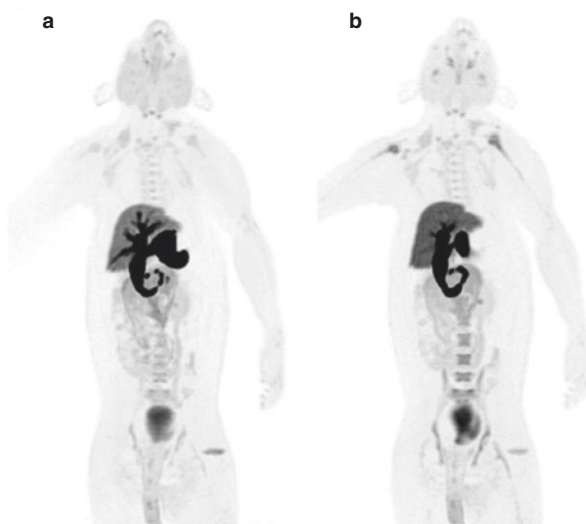
Radiotracer [ $^{11}\text{C}$ ]MC1 was used for whole-body biodistribution and specific binding studies (Kim et al. 2018). The highest uptake of [ $^{11}\text{C}$ ]MC1 in rhesus monkeys was noticed in excretory organs liver, gallbladder, and urinary bladder





**Fig. 10.18** Radiosynthesis of  $[^{11}\text{C}]\text{MC1}$  and radiotracers **32** and **33** (Cortes-Salva et al. 2018)

**Fig. 10.19** Maximum-intensity projections after injection of  $[^{11}\text{C}]\text{MC1}$  under baseline conditions (a) and after injection of MC1 (0.3 mg/kg) (b). High uptake was observed only in excretory organs at baseline and was not blocked by pharmacologic doses of MC1 (Kim et al. 2018)



(Fig. 10.19). No blocking effect was noticed with MC1 in excretory organs, although a significant blocking by MC1 and celecoxib was observed in the ovaries (Table 10.1 and Fig. 10.20).

In a follow-up study, the brain uptake of  $[^{11}\text{C}]\text{MC1}$  in LPS-neuroinflammation monkey model was investigated, and percentage of specific uptake was 41% and 35% on postinjection days 3 and 8, respectively (Cortes et al. 2017).

**Table 10.1** Uptake of [<sup>11</sup>C]MC1 in monkeys and percentage blockade with pharmacologic doses of COX-1 and COX-2 inhibitors (Kim et al. 2018)

Organ	SUV <sub>10-90</sub> (n = 8)	Blocking effect (%) <sup>a</sup>			
		COX-1		COX-2	
		PS13 (n = 1)	Aspirin (n = 1)	MC1 (n = 1)	Celecoxib (n = 1)
Brain	1.22 ± 0.25	17	0	21	0
Lungs	0.22 ± 0.05	7	0	0	0
Heart	0.70 ± 0.11	7	0	0	0
Liver	9.10 ± 2.42	21	0	0	14
Spleen	0.63 ± 0.10	12	0	0	5
Gastrointestinal tract	1.10 ± 0.26	19	0	14	0
Kidney cortex	1.73 ± 0.36	16	5	16	22
Kidney medulla	1.91 ± 0.27	22	12	35	30
Ovaries	0.82 ± 0.23	12	12	38	46

<sup>a</sup>Corrected by plasma parent radioactivity at four time points

### 10.3.1.1 [1-(4-Fluorophenyl)-3-(2-[<sup>11</sup>C]methoxyethyl)-2-methyl-5-(4-(methylsulfonyl)phenyl)-1H-pyrrole] (<sup>11</sup>C-VA426)

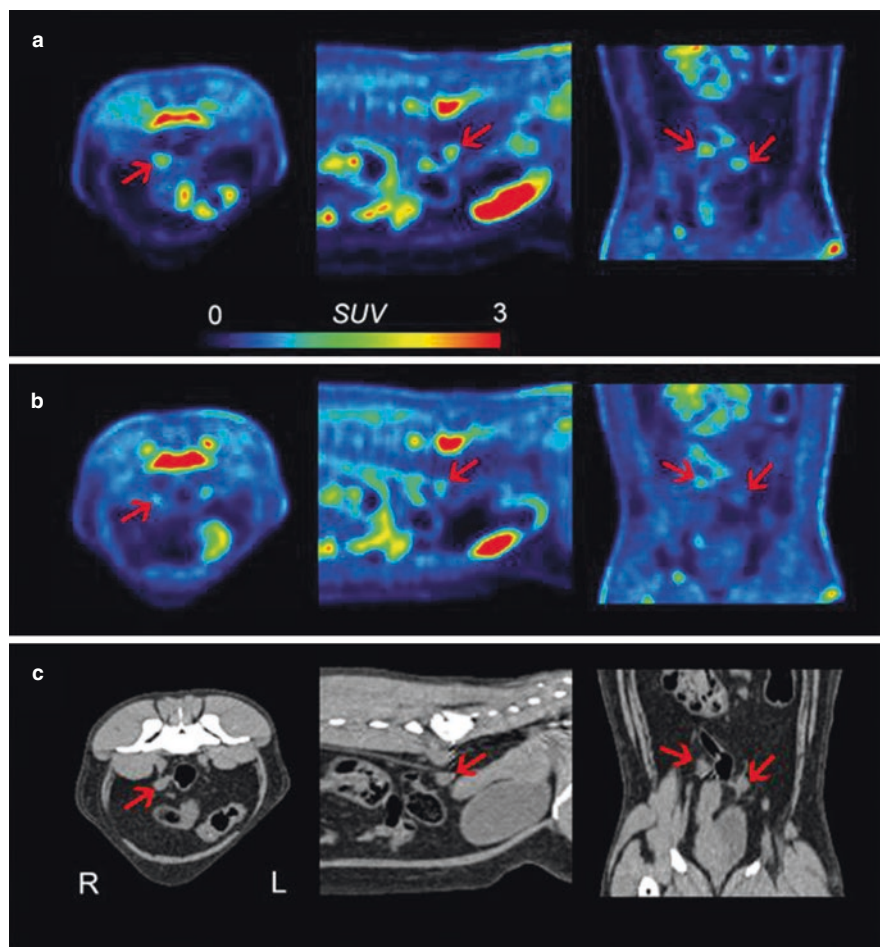
The favorable COX-2 inhibitory activity and selectivity profile of 1-(4-fluorophenyl)-3-(2-methoxyethyl)-2-methyl-5-(4-(methylsulfonyl)phenyl)-1H-pyrrole (VA426) (Anzini et al. 2008) prompted Carpinelli et al. to optimize and validate an automated radiosynthesis of [<sup>11</sup>C]VA426 (Fig. 10.21) (Carpinelli et al. 2019).

The radiotracer was synthesized in 15 ± 2% radiochemical yield (based on [<sup>11</sup>C]MeOTf) and high radiochemical purity >95%. The molar activity ranged between 37 and 148 GBq/μmol. Radiotracer [<sup>11</sup>C]VA426 was studied in models of LPS-induced brain neuroinflammation (Carpinelli et al. 2019). Ex vivo biodistribution experiments with radiotracer [<sup>11</sup>C]VA426 confirmed fast radioactivity uptake in the peripheral organs (kidney, lung, and heart). Highest uptake values were observed in the liver and kidneys at 10 min postinjection, and the kidney radioactivity was washed out thereafter whereas liver radioactivity slightly increased over time (Fig. 10.22).

Radioactivity peaked at 10 min in the analyzed brain regions followed by slow washout. Ex vivo study in the LPS-induced neuroinflammation model revealed no significant difference between radioactivity levels in the LPS-treated and healthy striatal regions (Carpinelli et al. 2019).

PET imaging studies with radiotracer [<sup>11</sup>C]VA426 in healthy mice showed rapid radioactivity accumulation mainly in the liver and kidneys which slightly decreased at 30 min (Fig. 10.23).

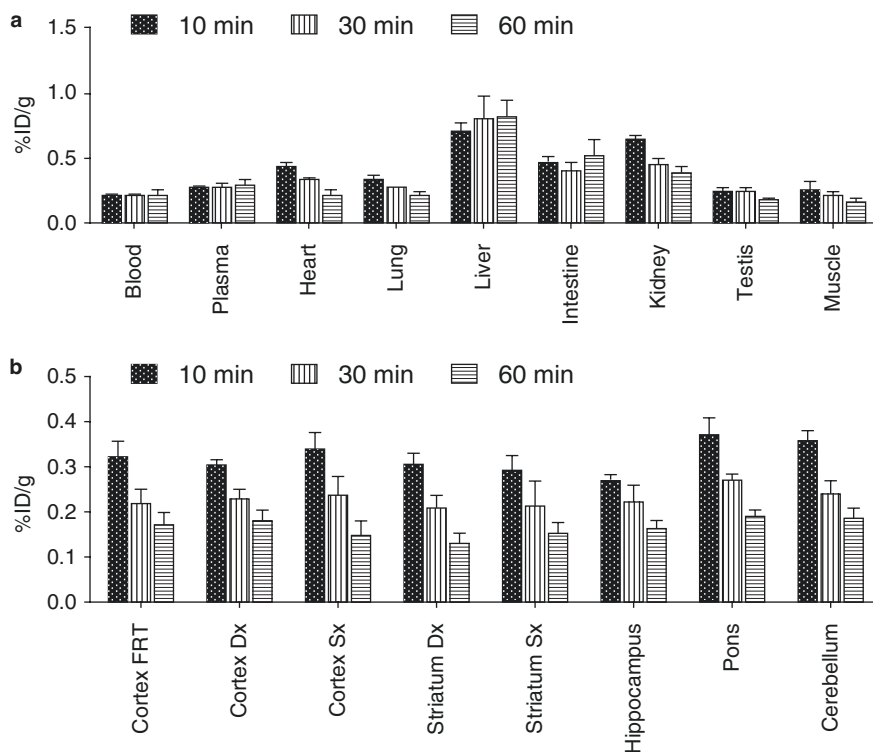
Pre-dosing with pharmacological doses of celecoxib resulted in some reduction of radioactivity uptake in the intestine regions (Carpinelli et al. 2019). Moreover, PET imaging of [<sup>11</sup>C]VA426 under blocking conditions with celecoxib in a model of peripheral inflammation revealed a reduced radioactivity uptake in all organs,



**Fig. 10.20** Whole-body PET images after the injection of [ $^{11}\text{C}$ ]MC1 under baseline conditions (a) and after injection of MC1 (0.3 mg/kg) (b), as well as corresponding CT images (c). High uptake was observed in both ovaries (arrows) and was blocked by a pharmacologic dose of MC1, consistent with specific binding to COX-2. PET images are shown as SUV<sup>91</sup>



**Fig. 10.21** Radiosynthesis of [ $^{11}\text{C}$ ]VA426



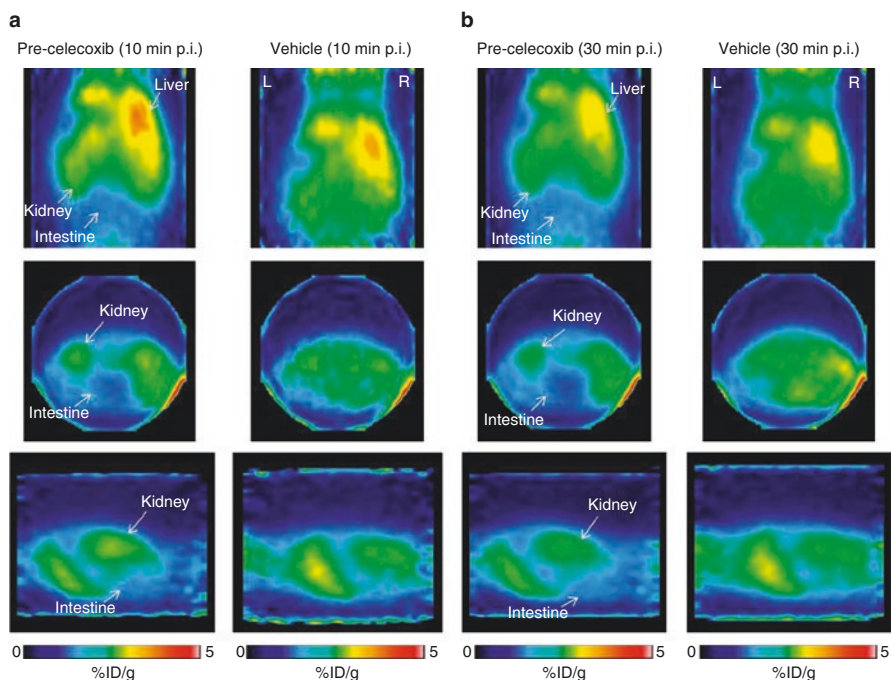
**Fig. 10.22** Ex vivo biodistribution of [ $^{11}\text{C}$ ]VA426 (a) in the periphery and (b) in the brain of healthy rats ( $n = 9$ , three per time point). Uptake value areas are expressed as %ID/g<sup>94</sup>

including blood and plasma indicating that the blocking effect was not linked to the inhibition of COX-2-binding sites (Fig. 10.24).

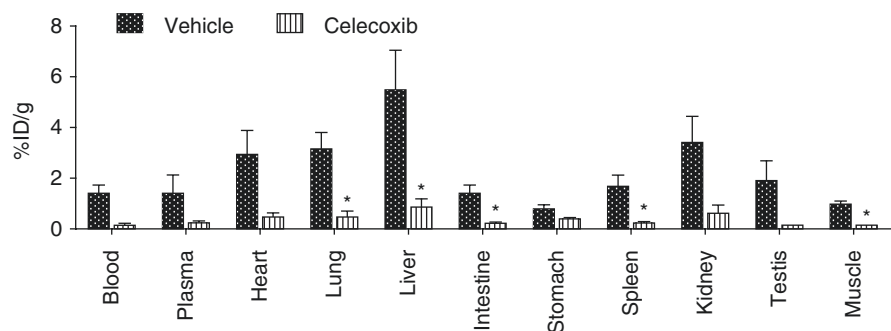
In vivo stability analyses showed that [ $^{11}\text{C}$ ]VA426 was not metabolically stable, and more than 50% of the detected radioactivity was related to radiometabolites of [ $^{11}\text{C}$ ]VA426. Taken together, despite the favorable COX-2 inhibitory and selectivity profile of VA426, radiotracer [ $^{11}\text{C}$ ]VA426 was not suitable for further development as a clinical COX-2 PET imaging agent.

### 10.3.1.2 4-(5-([ $^{11}\text{C}$ ]Methoxymethyl)-3-phenylisoxazol-4-yl) benzenesulfonamide ([ $^{11}\text{C}$ ]MOV)

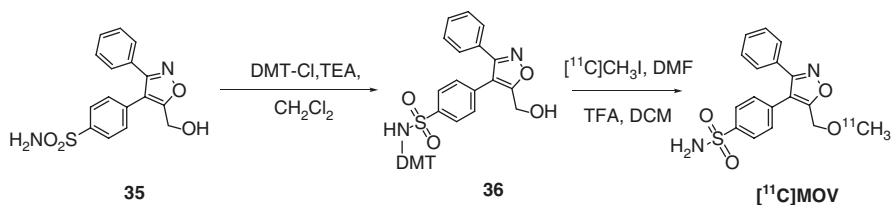
Prabhakaran et al. used hydroxymethyl valdecoxib **35**, an in vivo metabolite of valdecoxib, for the development and evaluation of novel COX-2 imaging agent [ $^{11}\text{C}$ ]MOV (Prabhakaran et al. 2018). The first step of synthesis involved the protection of the sulfonamide group of hydroxymethyl valdecoxib **35** to form dimethoxytrityl (DMT) intermediate **36**. Compound **36** was reacted with [ $^{11}\text{C}$ ]CH<sub>3</sub>I in DMF with KO<sup>t</sup>Bu as a base. Upon the removal of DMT-protecting group, radiotracer [ $^{11}\text{C}$ ]MOV was obtained in 40 ± 10% radiochemical yield (based on [ $^{11}\text{C}$ ]CH<sub>3</sub>I) (Fig. 10.25) (Prabhakaran et al. 2018).



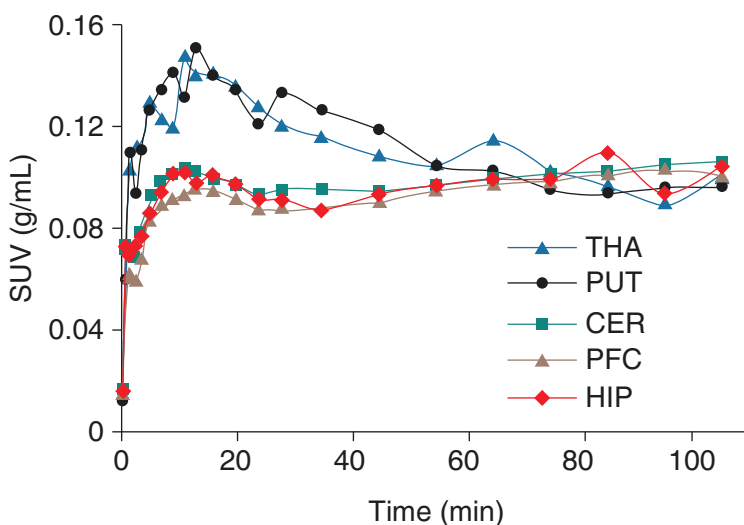
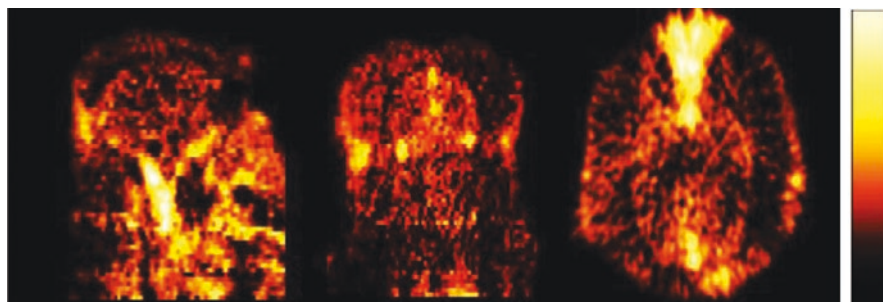
**Fig. 10.23** PET images of  $[^{11}\text{C}]\text{VA426}$  distribution in mouse (a) at 10 and (b) 30 min after injection. Each acquisition was shown in coronal, transaxial, and sagittal sections, respectively. Mice were anesthetized with a mixture of 2% isoflurane in air, injected intravenously with 1.85 MBq (mouse 1) and 3.7 MBq (mouse 2) of  $[^{11}\text{C}]\text{VA426}$  and, respectively, pretreated (5 min before) with celecoxib or vehicle (DMSO). Dynamic PET data were acquired for 30 min (four scans of 2.5 min followed by four of 5 min). White arrow indicates the main anatomical regions of  $[^{11}\text{C}]\text{VA426}$  distribution. In coronal views, R (right) and L (left) indicate the spatial orientation of mouse (Carpinelli et al. 2019)



**Fig. 10.24** Inhibition study in the peripheral inflammation model; inflammation was induced by intraperitoneal injection of LPS 6 h before the study. Mice were treated with celecoxib ( $n = 3$ ) or vehicle ( $n = 3$ ) prior to  $^{11}\text{C}\text{-VA426}$  injection and sacrificed after 10 min. Uptake values in peripheral areas are expressed as a percentage of injected dose per gram of tissue (%ID/g) ( $*p < 0.05$  vs. vehicle) (Carpinelli et al. 2019)



**Fig. 10.25** Radiosynthesis of  $[^{11}\text{C}]\text{MOV}$



**Fig. 10.26** Top: Sum of 0–120 min PET images of  $[^{11}\text{C}]\text{MOV}$  in a representative anesthetized baboon (left: sagittal, middle: coronal, right: transaxial). Bottom: Time-activity curves of  $[^{11}\text{C}]\text{MOV}$  in baboon brain (*CER* cerebellum, *HIP* hippocampus, *PFC* prefrontal cortex, *PUT* putamen, *THA* thalamus) (Prabhakaran et al. 2018)

PET imaging with  $[^{11}\text{C}]\text{MOV}$  in anesthetized baboons confirmed blood-brain barrier penetration of the radiotracer. However, the retention of radioactivity in the brain was very low (Fig. 10.26) (Prabhakaran et al. 2018).

Radioactivity accumulation was observed in the heart and duodenum. Metabolite analysis of [ $^{11}\text{C}$ ]MOV confirmed a rapid metabolism of the radiotracer. Both the very low retention of radioactivity in the brain and the rapid metabolic degradation make radiotracer [ $^{11}\text{C}$ ]MOV not suitable for COX-2 imaging in neuroinflammation.

## 10.4 Summary and Conclusion

Despite their structural similarities, both cyclooxygenase isozymes (COX-1 and COX-2) display quite different distribution and regulation patterns in various tissues and organs. In the central nervous system (CNS), the COX-1 isoform is particularly upregulated in the midbrain, pons, and medulla. Moreover, COX-1 upregulation and activation in the CNS are part of the defense mechanism in response to acute and chronic neuroinflammatory conditions. Overall, recent data demonstrated that COX-1 plays a crucial role in the process of neuroinflammation and neurodegeneration. The role of inducible COX-2 isoform in neuroinflammation is more controversial. COX-2 may mediate a neurotoxic or anti-inflammatory role depending on the stimulus and cell type. As COX-1 and COX-2 have become important drug targets in neuroinflammatory and neurodegenerative diseases, imaging techniques like positron-emission tomography (PET) have emerged as valuable tools to detect and visualize the functional expression of COXs *in vivo*.

The development of PET imaging agents is most advanced for COX-1-targeted radiotracers with a clear dominance for compounds labeled with the short-lived positron emitter carbon-11 ( $^{11}\text{C}$ ,  $t_{1/2} = 20.4$  min). The COX-1 radiotracer (*S*)-[ $^{11}\text{C}$ ]KTP-Me has progressed to first-in-human studies in healthy volunteers and patients with mild cognitive impairment and Alzheimer's disease. However, the radiotracer (*S*)-[ $^{11}\text{C}$ ]KTP-Me was not successful as a COX-1-imaging agent in neuroinflammation, and radiotracer [ $^{11}\text{C}$ ]PS3 was developed as a more potent and COX-1 selective PET imaging agent. The obtained excellent PET images with radiotracer [ $^{11}\text{C}$ ]PS3 in monkeys encouraged first human biodistribution and dosimetry studies in healthy volunteers, but detailed clinical studies in patients with neurodegenerative diseases are still pending.

To date, no COX-2 radiotracers were tested in humans for PET imaging of neuroinflammatory conditions. Despite favorable COX-2 inhibitory potencies and selectivities, most radiotracers lack sufficient COX-2 binding capacity in the brain *in vivo*.

As a result, all tested COX-2 radiotracers displayed rapid washout from the brain without retention of radioactivity related to COX-2-binding sites as demonstrated in rat brains with neuroinflammation after intracerebral LPS injection.

Overall, the most promising radiotracers [ $^{11}\text{C}$ ]PS3 and [ $^{11}\text{C}$ ]MC1 represent interesting lead compounds for further optimization towards the development of suitable PET imaging agents for detecting and visualizing COX-1 and COX-2 in neuroinflammation. Optimization should involve the use of the positron emitter fluorine-18 ( $^{18}\text{F}$ ,  $t_{1/2} = 109.8$  min) which would allow the preparation of COX1/2 PET imaging

agents at higher molar activities. High molar activity is an essential requirement to achieve favorable PET imaging contrast *in vivo* for saturable binding sites and intracellular targets with low abundance like COX-1 and COX-2.

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# PET and SPECT Imaging of the Central Dopamine System in Humans

# 11

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## Abstract

The neurotransmitter dopamine plays a role in many different functions of the human brain, ranging from psychomotor planning to cognition. This short review addresses which parts of the dopamine system can be imaged quantitatively in the living human brain using positron-emission tomography (PET) or single-photon emission computed tomography (SPECT).

Nowadays, imaging of the nigrostriatal dopaminergic pathway in humans can be performed quantitatively using radiotracers like the aromatic amino acid decarboxylase (AADC) substrate [ $^{18}\text{F}$ ]FDOPA, vesicular monoamine transporter 2 (VMAT-2) radioligands derived from tetrabenazine or PET/SPECT radioligands that bind to the dopamine transporter (DAT). Using PET, also several other dopaminergic projection pathways (e.g. mesocortical projections) can be assessed in humans. Several antagonist PET radioligands for the dopamine  $\text{D}_1$  receptor have been developed successfully. In addition, well-validated antagonist PET and SPECT radioligands are available for imaging of dopamine  $\text{D}_2$  and dopamine  $\text{D}_3$  ( $\text{D}_{2/3}$ ) receptors in the living human brain. Also agonist PET radioligands for the dopamine  $\text{D}_{2/3}$  receptors have become available, which afford the opportunity to evaluate the existence of the high-affinity state of these receptors *in vivo*. These agonist radiopharmaceuticals may also prove more sensitive to changes in dopamine concentrations (e.g. induced by the dopamine releaser amphetamine). Finally, selective antagonist PET radioligands for the dopamine  $\text{D}_4$  receptor have recently been synthesized and evaluated successfully in small laboratory animals, although these radioligands have not yet been reported as applied in human subjects.

In conclusion, after almost four decades of research, several relevant parts of the central dopamine system can be assessed quantitatively in the living human brain using PET or SPECT. Future studies may include application of agonist radioligands and more dopamine receptor subtype selective radioligands.

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## 11.1 Introduction

Numerous cognitive, psychomotor and emotional functions are regulated, at least in part, by dopaminergic circuits in the brain. The neurotransmitter dopamine plays, for example, an important role in movement and the experience of pleasure (hedonia). Dopaminergic neurons in the midbrain area are relevant in coding the value of neuronal signals, and indeed, these neurons are activated by rewarding stimuli (such as amphetamines, which induce dopamine release), but may also play a role in reward learning, prediction of error signalling and interpretation of the salience of events (Fiorillo et al. 2003; Nakahara et al. 2004; Morris et al. 2006; Zijlstra et al. 2008; Enomoto et al. 2011; Schultz, 2016; Takahashi et al. 2017; Wang et al. 2019). In addition, neurodegeneration of the dopaminergic neurons in the midbrain can lead to severe locomotor dysfunction such as in Parkinson's disease. Given the broad spectrum of functions of dopamine in the brain, it is reasonable that imaging

of the central dopamine system has been used frequently as a tool to assess different aspects of dopaminergic functioning in health and disease, particularly in a variety of neuropsychiatric disorders.

Here we review shortly which parts of the central dopamine system can be imaged quantitatively in the living human brain using positron-emission tomography (PET) or single-photon emission computed tomography (SPECT). The findings of SPECT and PET studies of the dopamine system in different neuropsychiatric disorders will be discussed only shortly, since they will be discussed in depth in other chapters of this book.

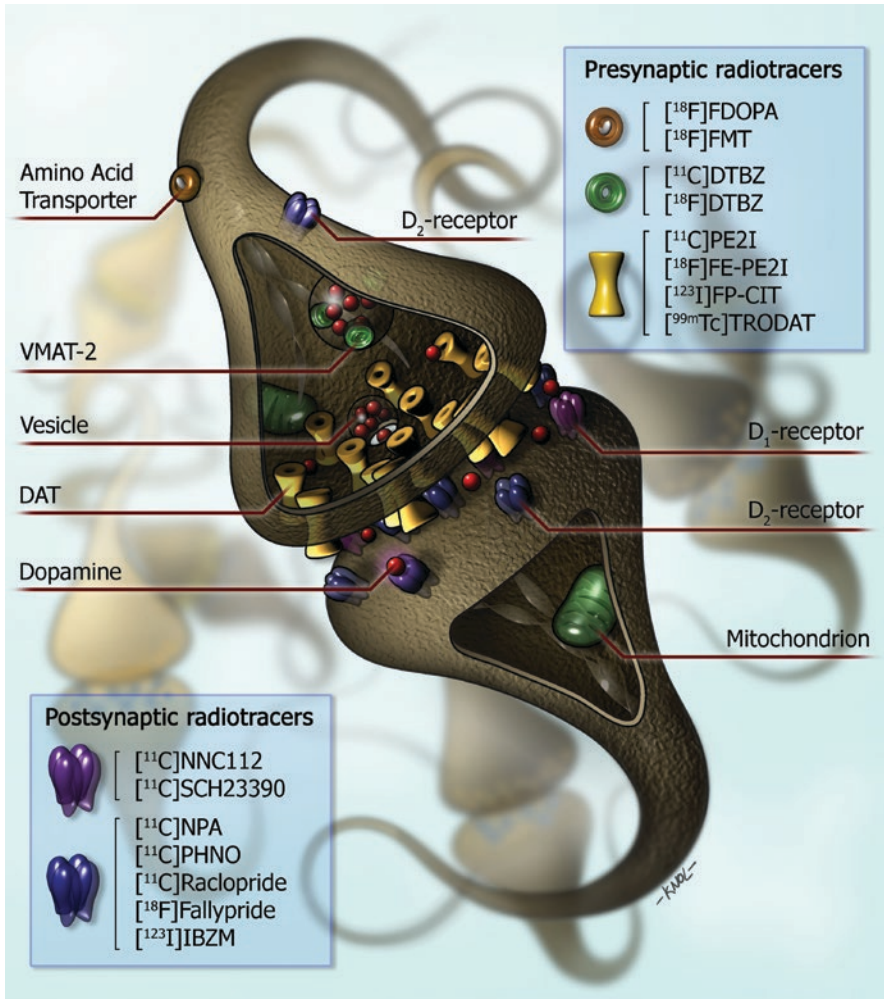
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## 11.2 Imaging of the Presynaptic Dopamine System

### 11.2.1 [ $^{18}\text{F}$ ]FDOPA and [ $^{18}\text{F}$ ]FMT

The majority of the cell bodies of dopaminergic neurons are located in the midbrain area, with axons predominantly projecting to the striatum (the nigrostriatal dopaminergic pathway). The scintigraphic techniques PET and SPECT offer unique means to assess the *in vivo* integrity of the nigrostriatal pathway. The biochemical integrity of presynaptic dopamine neurons can be assessed with the PET radioligands 6- $^{18}\text{F}$ -fluoro-*L*-DOPA ([ $^{18}\text{F}$ ]FDOPA) or 6- $^{18}\text{F}$ -fluoro-meta-tyrosine ([ $^{18}\text{F}$ ]FMT) (for reviews, see Booij et al. 1999; Kumakura and Cumming 2009; Sarikaya 2015).

[ $^{18}\text{F}$ ]FDOPA is a widely applied PET tracer commonly radiolabelled with fluorine-18, although in some clinical studies DOPA has been applied as radiolabelled with carbon-11. After injection, the radiotracer will be taken up into dopaminergic neurons by the amino acid transporter (Fig. 11.1), decarboxylated (by aromatic amino acid decarboxylase; AADC) to [ $^{18}\text{F}$ ]fluorodopamine (or [ $^{11}\text{C}$ ]dopamine when [ $^{11}\text{C}$ ]DOPA is administered) and then temporarily stored in vesicles within the nerve terminals. Therefore, [ $^{18}\text{F}$ ]FDOPA accumulation in dopamine neurons reflects a regulated aspect of the synthesis of dopamine in presynaptic neurons. The trapping of [ $^{18}\text{F}$ ]fluorodopamine in nigrostriatal dopamine terminals has, however, been demonstrated not to be completely irreversible (Holden et al. 1997; Cumming et al. 2001). Indeed, steady-state kinetic analyses of [ $^{18}\text{F}$ ]FDOPA PET studies have shown that besides the synthesis, also the turnover of [ $^{18}\text{F}$ ]FDOPA to *O*-methyl-[ $^{18}\text{F}$ ]FDOPA can be assessed (Kumakura et al. 2005). Importantly, in most clinical studies, static striatal [ $^{18}\text{F}$ ]FDOPA uptake was assessed relative to uptake in a reference tissue (reflecting the nonspecific radioactivity concentration) and using linear graphical analysis. This approach is convenient for clinical studies in which radioactivity in arterial blood is commonly not measured (Kumakura and Cumming 2009; Jauhar et al. 2018; Majuri et al. 2018). However, the kinetic properties of [ $^{18}\text{F}$ ]FDOPA are complex and include not only the washout of radiotracer but also the entry of a radiolabelled metabolite of [ $^{18}\text{F}$ ]FDOPA into the brain. Consequently, relatively long acquisitions of dynamic PET images, as well as the analysis of the metabolite-corrected arterial [ $^{18}\text{F}$ ]FDOPA input function, are required to yield a more physiological index of [ $^{18}\text{F}$ ]FDOPA utilization, including [ $^{18}\text{F}$ ]FDOPA turnover (for a review, see Kumakura and Cumming 2009).



**Fig. 11.1** Simplified diagram of a striatal dopaminergic synapse. On the presynaptic side, potential markers for imaging of the integrity of dopaminergic neurons in humans are shown.  $[^{18}\text{F}]\text{FDOPA}$  and  $[^{18}\text{F}]\text{FMT}$  PET provide measures of the structural and biochemical integrity of the dopaminergic neurons.  $[^{11}\text{C}]\text{DTBZ}$  and  $[^{18}\text{F}]\text{DTBZ}$  are commonly used radiopharmaceuticals for the vesicular monoaminergic transporter. Substituted (nor)phenyltropanes ( $[^{11}\text{C}]\text{PE2I}$ ,  $[^{18}\text{F}]\text{FE-PE2I}$ ,  $[^{123}\text{I}]\text{FP-CIT}$  and  $[^{99\text{m}}\text{Tc}]\text{TRODAT}$ ) are frequently used PET and SPECT radioligands for imaging of the DAT in humans. On the postsynaptic side,  $[^{11}\text{C}]\text{NNC112}$  and  $[^{11}\text{C}]\text{SCH23390}$  are commonly used antagonist radiopharmaceuticals for the dopamine D<sub>1</sub> receptor. Dopamine D<sub>2</sub> receptors are predominantly expressed on the postsynaptic side as compared to the presynaptic side of the dopaminergic synapse.  $[^{11}\text{C}]\text{NPA}$  and  $[^{11}\text{C}]\text{PHNO}$  are agonist radioligands for dopamine D<sub>2/3</sub> receptors. Commonly used antagonist radioligands for D<sub>2/3</sub> receptors are substituted benzamides ( $[^{11}\text{C}]\text{raclopride}$ ,  $[^{11}\text{C}]\text{FLB 457}$ ,  $[^{18}\text{F}]\text{fallypride}$  and  $[^{123}\text{I}]\text{IBZM}$ ). *DAT* dopamine transporter, *VMAT-2* vesicular monoaminergic transporter 2

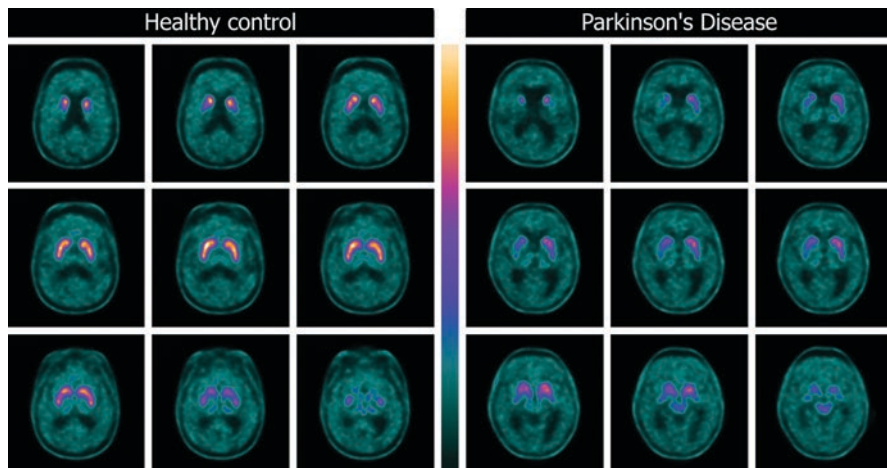


As an alternative to [ $^{18}\text{F}$ ]FDOPA, the tyrosine derivative [ $^{18}\text{F}$ ]FMT has been developed. This radiotracer is similar to [ $^{18}\text{F}$ ]FDOPA in that both radiotracers are substrates of AADC. However, unlike [ $^{18}\text{F}$ ]FDOPA, [ $^{18}\text{F}$ ]FMT is not a substrate for catechol-*O*-methyl-transferase (COMT), an enzyme essential in the breakdown of dopamine. Therefore, there are no radioactive *O*-methyl-metabolites which contribute to the nonspecific radioactivity in the brain, which enables the use of simplified kinetic modelling approaches (Dejesus et al. 2001).

The enzyme AADC plays an important role in the synthesis of dopamine, in that it converts *L*-DOPA to dopamine. It is important to consider that AADC also plays a role in the synthesis of other monoamine transmitters, e.g. in the conversion of 5-hydroxytryptophan to serotonin (5-HT). Since both [ $^{18}\text{F}$ ]FDOPA and [ $^{18}\text{F}$ ]FMT are substrates of AADC, these tracers are converted not only in dopaminergic neurons but also in serotonergic and noradrenergic neurons (Brown et al. 1999; Moore et al. 2003). The striatal accumulation of these tracers does however predominantly reflect conversion in nigrostriatal dopaminergic neurons (Pavese et al. 2012).

Initial PET studies applying [ $^{18}\text{F}$ ]FDOPA or [ $^{18}\text{F}$ ]FMT focused on the evaluation of the nigrostriatal pathway, by studying the distribution of these tracers in the striatum (Antonini et al. 1995). Due to the better performance characteristics of the PET systems, as well as improved PET data analysis methods, also extrastriatal radioactivity concentrations can nowadays be quantified (e.g. cortical areas). However, and as mentioned before, due to the expression of AADC in all monoaminergic neurons, accumulation in the raphe nuclei and locus coeruleus likely reflects predominantly serotonergic and noradrenergic activity, respectively, instead of accumulation in dopaminergic neurons (Ito et al. 2008; Lewis et al. 2012; Pavese et al. 2012), while accumulation in the thalamus may represent a combination of serotonergic and noradrenergic activity (Stokholm et al. 2018). An important consideration when studying extrastriatal brain regions is that the reliability of measurements in extrastriatal regions is lower than in striatum (Egerton et al. 2010).

Neuropsychiatric diseases like Parkinson's disease and dementia with Lewy bodies are characterized by severe loss of dopaminergic nigrostriatal neurons (Fig. 11.2; for more details on imaging of the dopaminergic system in Parkinsonian syndromes, see Chap. 27 in "PET and SPECT in Neurology"). Consequently, many PET studies have utilized [ $^{18}\text{F}$ ]FDOPA and [ $^{18}\text{F}$ ]FMT to demonstrate the ability to detect loss of dopaminergic neurons in cohorts of these patients (Brooks et al. 1990; Antonini et al. 1995; Hu et al. 2000; Pavese et al. 2012). Also, [ $^{18}\text{F}$ ]DOPA may already detect the loss of dopaminergic neurons in idiopathic rapid eye movement (REM) sleep behaviour disorder, a prodromal phenotype of Parkinson's disease (Stokholm et al. 2017; Knudsen et al. 2018). In addition, several [ $^{18}\text{F}$ ]FDOPA PET studies have been performed in patients suffering from psychosis, with the majority of them demonstrating an elevated striatal [ $^{18}\text{F}$ ]FDOPA accumulation, which suggests an increased synthesis rate as well as dopamine turnover in these patients (Kumakura et al. 2007; Howes and Kapur 2009; for systematic reviews see Fusar-Poli and Meyer-Lindenberg 2013; McCutcheon et al. 2018). Interestingly, the results of recent studies suggested that the increased [ $^{18}\text{F}$ ]FDOPA uptake may be



**Fig. 11.2** Transversal planes of a [ $^{18}\text{F}$ ]DOPA PET image obtained in a subject without nigrostriatal cell loss (left panel), and in a patient suffering from Parkinson's disease (PD; right panel). Please note the asymmetric striatal uptake as well as the severe loss of [ $^{18}\text{F}$ ]DOPA uptake particularly in the putamen of the PD case

only observed in patients that respond to classic antipsychotics, but not in patients with treatment-resistant schizophrenia (Demjaha et al. 2012; Kim et al. 2017).

[ $^{18}\text{F}$ ]FDOPA and [ $^{18}\text{F}$ ]FMT PET have also been applied to evaluate the role of dopamine in the human reward system (Dreher et al. 2008; Schlegelhauf et al. 2013; Deserno et al. 2015; Smith et al. 2016). Finally, in healthy controls [ $^{18}\text{F}$ ]FDOPA and [ $^{18}\text{F}$ ]FMT PET have been used to assess ageing effects and cognitive functions (Braskie et al. 2008; Dreher et al. 2008; Kumakura et al. 2010).

### 11.2.2 Imaging of the VMAT-2

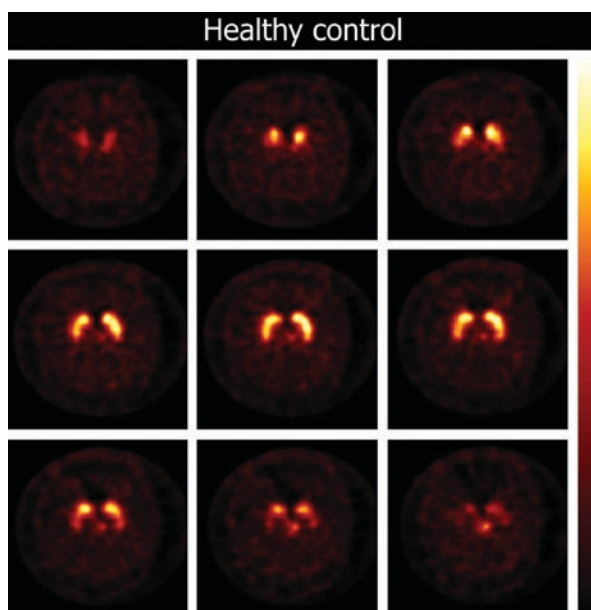
The vesicular monoamine transporter type 2 (VMAT-2; Fig. 11.1) is located in pre-synaptic dopaminergic neurons. Initially, carbon-11-labelled tetrabenazine derivatives were developed for PET imaging to visualize and quantify the VMAT-2 in humans (for a review, see Kilbourn 1997). More recently, also fluorine-18-labelled radioligands for the VMAT-2 have been developed successfully (Lin et al. 2010; for a recent review see Kilbourn and Koeppe, 2019), which provides the opportunity to use these radioligands also in hospitals not equipped with a cyclotron.

VMAT-2 is not exclusively present in dopaminergic neurons, but also in other monoaminergic neurons, including serotonergic (Guillot and Miller 2009), although the vast majority of the striatal radioactivity concentration represents binding in dopaminergic nerve terminals (Wang et al. 2010). In this regard, it is of interest that it has been suggested that VMAT-2 radioligands can also be used to visualize loss of pancreatic beta-cell mass in type 1 and type 2 diabetes (Goland et al. 2009; Cline et al. 2018), since beta-cells express VMAT-2. However, the specific binding of VMAT-2 PET radioligands binding to beta-cell mass has been disputed (Virostko

et al. 2011). Like [ $^{18}\text{F}$ ]FDOPA and [ $^{18}\text{F}$ ]FMT use in PET studies, VMAT-2 studies in humans have demonstrated that VMAT-2 imaging is a sensitive technique to detect reductions of nigrostriatal dopaminergic terminals in patients suffering from Parkinson's disease or dementia with Lewy bodies (Frey et al. 1996; Okamura et al. 2010; Burke et al. 2011a; Villemagne et al. 2011; Shah et al. 2016). In addition, VMAT-2 binding has been evaluated in relation to natural ageing effects (Bohnen et al. 2006), although findings are not consistent (Lin et al. 2013). Until now, SPECT radioligands have not been developed successfully for imaging of the VMAT-2.

### 11.2.3 Imaging of the Dopamine Transporter

The dopamine transporter (DAT) is mainly located in the membrane of terminals of presynaptic dopaminergic neurons (Fig. 11.1), but is also expressed at much lower levels in cell bodies in the substantia nigra (Ma et al. 1999; Fazio et al. 2018). Radioligands for the DAT have been developed successfully for both PET and SPECT. Regarding SPECT radioligands, radiopharmaceuticals derived from cocaine, e.g. [ $^{99\text{m}}\text{Tc}$ ]TRODAT-1, [ $^{123}\text{I}$ ] $\beta$ -CIT and [ $^{123}\text{I}$ ]FP-CIT, are nonselective DAT radioligands, although binding in the striatum predominantly reflects binding to DAT (Laruelle et al. 1993; Booij et al. 1997a, 2007; Dresel et al. 1998; Ziebell et al. 2010; Koopman et al. 2012; Fig. 11.3). Indeed, [ $^{123}\text{I}$ ]FP-CIT SPECT studies have

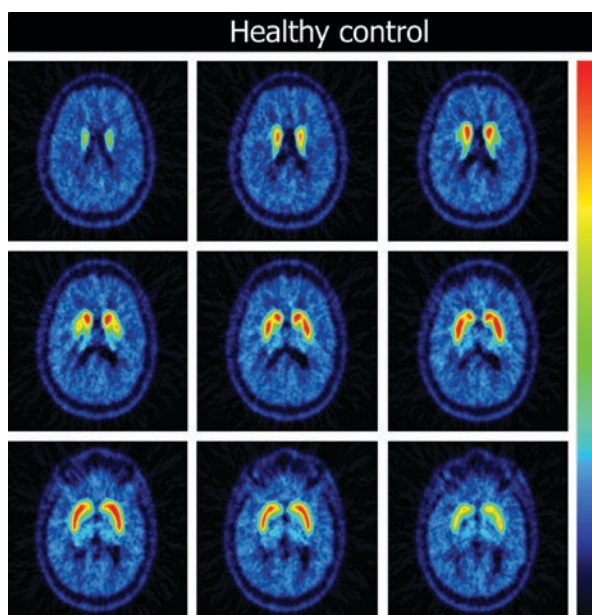


**Fig. 11.3** Transversal SPECT images obtained 3 h after i.v. injection of [ $^{123}\text{I}$ ]FP-CIT (radiotracer for the dopamine transporter) into a healthy female control subject. Intense symmetric and specific binding is visible in the caudate nucleus and putamen. The used colour scale represents a maximum intensity scale. This study has been acquired on a brain-dedicated SPECT system (InSPira)

shown that extrastriatal binding (in the diencephalon and midbrain) can be blocked by a selective serotonin reuptake inhibitor, indicating predominant binding to serotonin transporters in these brain areas (Booij et al. 2007; Ziebell et al. 2010). In addition, also promising and DAT-selective SPECT ligands, like [ $^{123}\text{I}$ ]PE2I, have been developed and used in human subjects (Kuikka et al. 1998; Ziebell et al. 2010; Thomsen et al. 2013).

PET radioligands for imaging of the DAT have been developed numerously over the last two decades (see for reviews Varrone and Halldin 2012a, b). Initially, non-selective DAT radioligands (e.g. [ $^{11}\text{C}/^{18}\text{F}$ ]CFT, Rinne et al. 2002; Nurmi et al. 2003; [ $^{18}\text{F}$ ]FP-CIT; Oh et al. 2012) were applied in human studies, but more recently reported radioligands have an improved DAT selectivity (e.g. [ $^{11}\text{C}$ ]PE2I (Hirvonen et al. 2008) (Fig. 11.4) or [ $^{18}\text{F}$ ]FE-PE2I (Varrone et al. 2009; Sasaki et al. 2012; Sonni et al. 2016)). The developed PET radioligands also afford the opportunity to assess extrastriatal DAT binding, i.e. binding in the midbrain area and the orbito-frontal area (Jucaite et al. 2005; Hirvonen et al. 2008; Yagi et al. 2010; Fazio et al. 2018), while with SPECT only DAT binding in the striatum can be assessed accurately.

The DAT plays an important role in regulating dopaminergic neurotransmission. In this regard, appropriate regulation of the DAT expression is critical. Indeed, the



**Fig. 11.4** Transversal planes of a mean PET summation image (6–93 min) obtained after i.v. injection of [ $^{11}\text{C}$ ]PE2I (selective radiotracer for the dopamine transporter) into a healthy male control subject. Intense specific binding is visible in the caudate nucleus and putamen and lower specific binding in extrastriatal brain areas (thalamus/midbrain). This study has been acquired on a brain-dedicated PET system (HRRT) (images courtesy of Dr. Andrea Varrone and Professor Christer Halldin, Karolinska Institutet)

DAT expression undergoes dynamic control by cellular protein kinases and phosphatases (Ramamoorthy et al. 2011; Won et al. 2018). So, the DAT is not only expressed on the plasma membrane of dopaminergic neurons, as internalization of the DAT has been reported (Eriksen et al. 2009; Chen et al. 2010). Until now, it is not clear whether the abovementioned DAT radioligands label only plasma membrane-bound transporters or also internalized transporters when applied in PET/SPECT studies evaluating in vivo conditions (Earley et al. 2011), although a recent PET study provided indirect evidence that DAT PET tracers may bind preferentially to DAT located on cell membrane (Kim et al. 2016). More specifically, in mice it was shown that environmental enrichment induced internalization of the DAT (confirmed by surface biotinylation studies) and reduced striatal [ $^{18}\text{F}$ ]FP-CIT binding. It might thus be that such tracers do not pass the cell membrane or, more likely, the radioligand binding characteristics to internalized transporters may differ from those transporters that are not internalized. For example, Afonso-Orama and co-workers showed that downregulation of the DAT consisted of a decrease of glycosylated (mature) DAT on the plasma membrane and accumulation of non-glycosylated (immature) DAT intracellularly (Afonso-Oramas et al. 2010). So, it might be that tracers like [ $^{18}\text{F}$ ]FP-CIT bind with higher affinity to the mature than immature DAT. Interestingly, Guo and co-workers have demonstrated that radioligands that bind to the dopamine  $D_{2/3}$  receptors have a lower affinity to internalized than to non-internalized receptors (Guo et al. 2010). Future studies are necessary to evaluate whether DAT internalization is a relevant consideration for in vivo imaging of DAT.

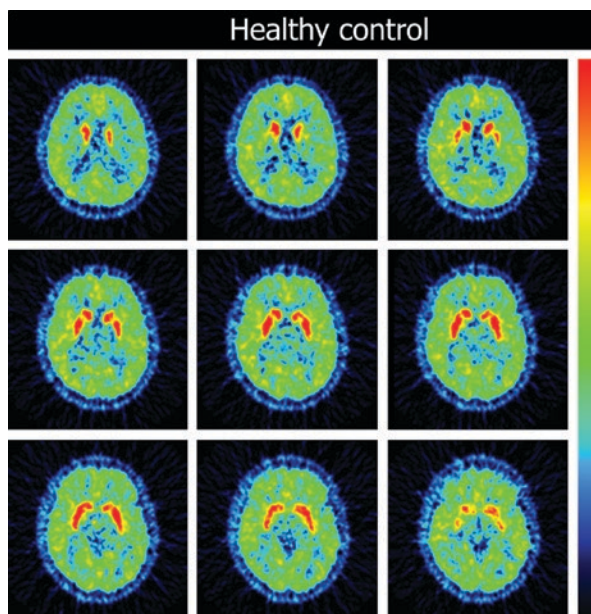
Numerous PET and SPECT studies evaluating DAT binding have shown the possibility to detect in vivo degeneration of nigrostriatal cells in disorders like Parkinson's disease and dementia with Lewy bodies (Booij et al. 1997b; Varrone et al. 2001; McKeith et al. 2007; Yagi et al. 2010; Oh et al. 2012; Fazio et al. 2018). Also, DAT binding has been examined in relation to ageing and gender effects in healthy controls, as well as the effects of polymorphism of the DAT gene (Lavalaye et al. 2000; van Dyck et al. 2005; van de Giessen et al. 2009; Troiano et al. 2010; Burke et al. 2011b; Varrone et al. 2013). Generally speaking, brain SPECT imaging of neuroreceptors/transporters is more accessible and cheaper than PET imaging, and consequently DAT imaging with SPECT is more commonly used in routine clinical studies to exclude or detect loss of nigrostriatal neurons in individual patients (Booij et al. 2001; Løkkegaard et al. 2002; Catafau et al. 2004; Ziebell et al. 2012; for reviews see Ba and Martin 2015; Suwijn et al. 2015).

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## 11.3 Imaging of the Postsynaptic Dopaminergic System

### 11.3.1 Imaging of Dopamine $D_1$ Receptors

Dopamine exerts its pharmacological action through G-protein-coupled receptors. These transmembrane receptors can be divided into two subfamilies based on their pharmacological properties: the  $D_1$ - ( $D_1$ ,  $D_5$ ) and  $D_2$ -like receptor subfamily ( $D_2$ ,  $D_3$ ,  $D_4$ ; Stoof and Kebabian 1981; Strange 1993; Beaulieu and Gainetdinov 2011).



**Fig. 11.5** Transversal plane of a mean PET summation image (12–63 min) obtained after i.v. injection of [ $^{11}\text{C}$ ]SCH23390 (radiotracer for the dopamine  $\text{D}_1$  receptor) into a healthy male control subject. Intense specific binding is visible in the caudate and putamen, and lower specific binding in the extrastriatal brain areas (cortex). This study has been acquired on a brain-dedicated PET system (ECAT EXACT HR) (images are courtesy of Per Stenkrona and Professor Christer Halldin, Karolinska Institutet)

The dopamine  $\text{D}_1$  receptor is a highly prevalent dopamine receptor in the striatum and neocortex and is located postsynaptically (Cortés et al. 1989; Volkow et al. 1996). The radiopharmaceuticals [ $^{11}\text{C}$ ]NNC 112 and [ $^{11}\text{C}$ ]SCH23390 have successfully been developed to image dopamine  $\text{D}_1$  receptors in the living human brain using PET (Fig. 11.5; for a recent review see Cervenka, 2019). Both radioligands are high-affinity dopamine  $\text{D}_1$  receptor antagonists, although they also have been reported to bind to serotonin  $2\text{A}$  ( $5\text{-HT}_{2\text{A}}$ ) receptors, estimated as approximately 25% of cortical binding being due to binding to  $5\text{-HT}_{2\text{A}}$  receptors (Ekelund et al. 2007). To circumvent this limitation, a study proposed the use of the  $5\text{-HT}_{2\text{A}}$  receptor antagonist ketanserin. After ketanserin administration, the  $5\text{-HT}_{2\text{A}}$  receptor binding is inhibited, and cortical  $\text{D}_1$  receptor binding can be accurately assessed using [ $^{11}\text{C}$ ]NNC 112 in humans (Catafau et al. 2010).

Although [ $^{11}\text{C}$ ]NNC 112 and [ $^{11}\text{C}$ ]SCH23390 bind to both dopamine  $\text{D}_1$  and  $5\text{-HT}_{2\text{A}}$  receptors, their affinity to dopamine  $\text{D}_2$ -like receptors is substantially lower (Andersen et al. 1992) although binding to dopamine  $\text{D}_5$  receptors cannot be excluded (Sunahara et al. 1991; Chou et al. 2006). While dopamine  $\text{D}_1$  receptors are expressed at high levels in the striatum and cortical areas, the expression of dopamine  $\text{D}_5$  receptors in the brain is low (Beaulieu and Gainetdinov 2011). Consequently,

in vivo binding to dopamine D<sub>5</sub> receptors, as compared to D<sub>1</sub> receptors, with PET radioligands like [<sup>11</sup>C]NNC 112 is likely negligible.

Dopamine receptors have been demonstrated to exist in two affinity states in vitro. The two receptor states are convertible and consist of a state of high and low affinity for the endogenous agonist dopamine (or exogenous agonists) (Sibley et al. 1982; Chio et al. 1994). The high-affinity state represents the active form of the receptors that are coupled to G-proteins (Zahniser and Molinoff 1978). Regarding dopamine D<sub>1</sub> receptors, results of in vitro experiments suggest that 20–40% of these receptors are in the high-affinity state (Richfield et al. 1989; Mamelak et al. 1993; McCauley et al. 1995). Importantly, while antagonist radioligands label both the high- and low-affinity state of the receptors, agonist radioligands may label receptors only in its high-affinity state (for a recent review see Shalgunov et al. 2019). So far, the available radioligands [<sup>11</sup>C]NNC 112 and [<sup>11</sup>C]SCH23390 are both antagonists for the dopamine D<sub>1</sub> receptor. Developments of agonist radioligands for the dopamine D<sub>1</sub> receptor are ongoing and will enable to examine in future studies whether the high-affinity state of the dopamine D<sub>1</sub> receptor exists in living humans (Palner et al. 2010; Finnema et al. 2013), although human studies with such novel tracers have not been performed yet. Also, dopamine agonist radioligands may be more sensitive to detect changes in dopamine concentrations in the synapse, e.g. changes induced by dopamine releasers like amphetamines (see the following section).

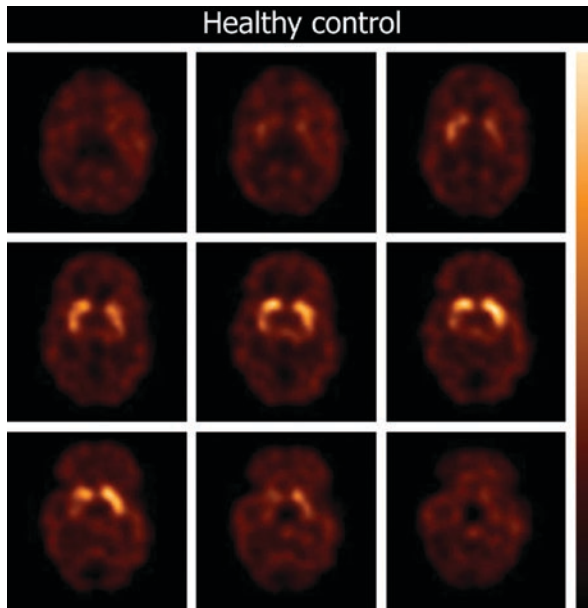
Since dopamine D<sub>1</sub> receptors have been implicated in cognitive performance, dopamine D<sub>1</sub> PET studies in healthy volunteers have been used to examine changes in working memory (McNab et al. 2009; Roffman et al. 2016), reward learning (de Boer et al. 2019), amygdala response to affective stimuli (Takahashi et al. 2010) and natural ageing (Jucaite et al. 2010) and to study the effects of genes involved in the dopaminergic tone (Slifstein et al. 2008). Dopamine D<sub>1</sub> receptor imaging in neuropsychiatric disorders has so far focused on schizophrenia (and the mechanism of action of antipsychotics), cocaine addiction and movement disorders (Nordström et al. 1995; Karlsson et al. 2002; Hirvonen et al. 2006; Martinez et al. 2009; Karimi et al. 2013; Thompson et al. 2014). Until now, the development of SPECT radioligands for the dopamine D<sub>1</sub> receptor has not been successful.

### 11.3.2 Imaging of Dopamine D<sub>2</sub>-Like Receptors

#### 11.3.2.1 Dopamine D<sub>2/3</sub> Receptor Imaging

In the past three decades, PET as well as SPECT radioligands have been developed successfully to image and quantify dopamine D<sub>2/3</sub> receptors in the human brain.

Regarding SPECT imaging, the commonly used benzamide [<sup>123</sup>I]IBZM is an antagonist radioligand which binds with high affinity to dopamine D<sub>2/3</sub> receptors (i.e. in the low nanomolar range) (Verhoeff et al. 1993; Videbaek et al. 2000; de Haan et al. 2003; Boot et al. 2008; Visser et al. 2008; Figuee et al. 2014). With this radioligand striatal dopamine D<sub>2/3</sub> receptors can be assessed accurately, and extrastriatal distribution may be visualized, particularly when using brain-dedicated SPECT systems (Fig. 11.6). The radioligand [<sup>123</sup>I]epidepride is a dopamine D<sub>2/3</sub>



**Fig. 11.6** Transversal [ $^{123}\text{I}$ ]IBZM slices obtained in a healthy control, 2 h after start of bolus/constant infusion. Intense specific binding is visible in the striatum, and much lower specific binding in the extrastriatal brain areas (thalamus/midbrain). The used colour scale represents a maximum intensity scale. This study has been acquired on a brain-dedicated SPECT system (NeuroFocus)

receptor antagonist with an enhanced affinity (picomolar range) for the dopamine  $D_{2/3}$  receptors (Kessler et al. 1991). This ultrahigh-affinity SPECT radioligand allows assessing dopamine  $D_{2/3}$  receptor availability in extrastriatal brain areas, such as the midbrain, diencephalon and cortical areas (Kornhuber et al. 1995; Varrone et al. 2000; Glenthøj et al. 2006; Tuppurainen et al. 2010; Nørbak-Emig et al. 2016).

Regarding PET imaging, the benzamide [ $^{11}\text{C}$ ]raclopride is an antagonist radioligand that binds with high affinity to dopamine  $D_{2/3}$  receptors (in the low nanomolar range; Farde et al. 1986) and is commonly used in human studies (Ito et al. 2011; Martinez et al. 2011; Wang et al. 2019). The radioligand [ $^{11}\text{C}$ ]N-methylspiperone binds also with high affinity to dopamine  $D_{2/3}$  receptors, although binding has been shown to be partly related to 5-HT $_{2A}$  receptors (Nyberg et al. 1999). Also antagonist PET radioligands with ultrahigh affinity for the dopamine  $D_{2/3}$  receptors have been developed successfully. For example, [ $^{11}\text{C}$ ]FLB 457 (Halldin et al. 1995) and [ $^{18}\text{F}$ ]fallypride (Mukherjee et al. 1995; Kasanova et al. 2018) are radioligands with a very high dopamine  $D_{2/3}$  receptor affinity allowing for evaluation of extrastriatal dopamine  $D_{2/3}$  receptors (picomolar range; Mukherjee et al. 1995; for a review see Högberg 1993). Finally, the antagonist [ $^{18}\text{F}$ ]N-methylbenperidol binds selectively to dopamine  $D_2$  receptors (Karimi et al. 2011).



More recently, many agonist PET radioligands for dopamine  $D_{2/3}$  receptors have been synthesized and evaluated (for a recent review see Shalgunov et al. 2019), and at least three were tested successfully in humans, that is, [ $^{11}\text{C}$ ]PHNO, [ $^{11}\text{C}$ ]NPA and [ $^{11}\text{C}$ ]MNPA (Willeit et al. 2006; Laymon et al. 2009; Otsuka et al. 2009). The apomorphine derivatives NPA and MNPA bind with high affinity to both dopamine  $D_2$  and dopamine  $D_3$  receptors (Seeman et al. 1985; Neumeyer et al. 1990). In contrast, *in vitro* studies have shown that the affinity of PHNO is 30- to 50-fold lower for the dopamine  $D_2$  than for the dopamine  $D_3$  receptor (Freedman et al. 1994; van Vliet et al. 2000). Indeed, a recent study in humans demonstrated that binding of [ $^{11}\text{C}$ ]PHNO in several brain regions, e.g. hypothalamus, substantia nigra, ventral pallidum and globus pallidus, is predominantly to dopamine  $D_3$  receptors. On the other hand, in striatal subregions, *in vivo* binding is predominant to dopamine  $D_2$  receptors (Tziortzi et al. 2011). Although the agonist [ $^{11}\text{C}$ ]PHNO binds predominantly to dopamine  $D_3$  receptors in certain brain areas, selective agonist radioligands for dopamine  $D_2$  or dopamine  $D_3$  receptors are not available yet.

Regarding dopamine  $D_2$  receptors, *in vitro* experiments on homogenized striata from rats and mice show that roughly 15–30% of the receptors are in the high-affinity state with a range up to 80% depending on the detection method used. In more intact tissue (brain slices), 90% of these receptors may even be in the high-affinity state (for review see van Wieringen et al. 2013). Initial results of *in vivo* PET measurements using amphetamine in nonhuman primates suggested that 60–80% of these receptors are in the high-affinity state (Narendran et al. 2004). However, although several attempts have been made, formally the existence of the high- and low-affinity state has until now not been confirmed or disputed *in vivo* (Finnema et al. 2009; see for reviews Finnema et al. 2010; Skinbjerg et al. 2012; Shalgunov et al. 2019). The results of the recent application of agonist radioligands in human PET studies suggest that 60–65% of dopamine  $D_2$  receptors are in the high-affinity state *in vivo* (Narendran et al. 2010; Shotbolt et al. 2012).

As discussed earlier, dopamine agonist radioligands may be more sensitive to detect changes in dopamine concentrations in the synapse, e.g. changes induced by dopamine releasers like amphetamines (Laruelle et al. 1995) or depletion induced by AMPT (Laruelle et al. 1997; Boot et al. 2008). Indeed, recent studies have indicated that the agonist radioligands ([ $^{11}\text{C}$ ]NPA and [ $^{11}\text{C}$ ]PHNO) are more vulnerable to endogenous striatal dopamine competition than the antagonist radioligand [ $^{11}\text{C}$ ]raclopride in humans (Narendran et al. 2010; Shotbolt et al. 2012; Gallezot et al. 2014). Also, it may be possible to use [ $^{11}\text{C}$ ]PHNO PET as a tool to assess not only striatal dopamine release but also release in the substantia nigra (Tseng et al. 2018). It can be concluded that agonist dopamine  $D_{2/3}$  receptor radioligands hold promise for studying changes in dopamine concentrations although further confirmation in human subjects is required.

The dopamine  $D_{2/3}$  receptors are expressed not only on the plasma membrane of postsynaptic neurons, but also presynaptically (Fig. 11.1). There are two splice variants of dopamine  $D_2$  receptors, the dopamine  $D_2$  long ( $D_2\text{L}$ ) and the short variant ( $D_2\text{S}$ ) (Chio et al. 1990). These isoforms have different localizations, with  $D_2\text{S}$

primarily located presynaptically (as an autoreceptor) (Khan et al. 1998). From a pharmacological point of view, these splice variants cannot be discriminated, and consequently, radioligands for the dopamine  $D_{2/3}$  receptors may bind to both  $D_2L$  and  $D_2S$ . However, since the striatal density of  $D_2L$  is much higher than that of  $D_2S$  (Rani and Kanungo 2006), imaging of dopamine  $D_{2/3}$  receptors is frequently referred to as imaging of postsynaptic dopamine  $D_{2/3}$  receptors. Also, imaging of the mid-brain dopamine  $D_{2/3}$  receptors may predominantly visualize dopamine  $D_{2/3}$  autoreceptors (Buckholtz et al. 2010).

Dopamine  $D_2$  receptors can internalize into the intracellular compartment after agonist stimulation, similar as for other G-protein-coupled receptors. Interestingly, Guo and co-workers have reported that dopamine  $D_2$  receptor radioligands have a lower affinity to internalized than to non-internalized receptors (Guo et al. 2010). A frequently used approach to assess dopamine release in vivo is the use of dexamphetamine (see for review Laruelle 2000). Administration of dexamphetamine intravenously or orally in doses of 0.2–0.3 mg/kg body weight induces a substantial release of endogenous dopamine, which will result in a displacement of radioligand binding to dopamine  $D_{2/3}$  receptors (i.e. decrease of, e.g., striatal [ $^{123}$ I]IBZM or [ $^{11}$ C]raclopride binding; Laruelle et al. 1995; Breier et al. 1997). Typically, administration of dexamphetamine leads to a rapid release of extracellular dopamine which is not long lasting (Breier et al. 1997). However, in imaging studies a prolonged displacement of radioligand binding has been observed, which may be caused (partly) by receptor internalization (Laruelle 2000; Cárdenas et al. 2004; Scott et al. 2007). Indeed, studies in knockout mice, which are incapable of internalizing dopamine  $D_2$  receptors, suggest that the prolonged displacement is mainly due to internalization of  $D_2$  receptors (Skinbjerg et al. 2010). Interestingly, a recent human [ $^{11}$ C]raclopride PET study examined whether the prolonged displacement (or return of the [ $^{11}$ C]raclopride binding to baseline), as a measure of dopamine  $D_{2/3}$  receptor internalization, was impaired in schizophrenia (Weinstein et al. 2018). Their findings suggested that this was not the case. Additional studies are needed to demonstrate whether molecular imaging studies are indeed able to assess internalization of dopamine  $D_2$  receptors in humans, preferentially using simpler paradigms than that presented by Weinstein and co-workers (Weinstein et al. 2018), which may be relevant to study the role of a possible abnormal internalization of these receptors in neuropsychiatric disorders.

### 11.3.2.2 Dopamine $D_4$ Receptor Imaging

The dopamine  $D_4$  receptor was cloned for the first time in 1991 (Van Tol et al. 1991) and is thought to play a role in a variety of neuropsychiatric disorders. This receptor is predominantly expressed postsynaptically (Rivera et al. 2002). In contrast to dopamine  $D_{2/3}$  receptors, dopamine  $D_4$  receptors do not internalize after agonist stimulation (Spooren et al. 2010). Interestingly, the antipsychotic clozapine, for instance, demonstrates a substantially higher affinity for the dopamine  $D_4$  than for the dopamine  $D_2$  receptor subtype (Van Tol et al. 1991; Eisenegger et al. 2010; Smith 2010). Recently, progress has been reported on the development of dopamine  $D_4$  receptor radioligands. This has been a challenging task as the density of

dopamine D<sub>4</sub> receptors in the human brain is much lower than that of other dopamine D<sub>2</sub>-like receptors (Marazziti et al. 2009; Lacivita et al. 2010; Kügler et al. 2011). Although Lacivita and co-workers reported that their radioligand mainly demonstrated high binding in the retina of non-primates (Lacivita et al. 2010), it is promising that a relatively recent study demonstrated clear brain uptake of a dopamine D<sub>4</sub> receptor antagonist radioligand in mice (Kügler et al. 2011), although it remains challenging to identify an optimized radiotracer (Leopoldo et al. 2014).

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## 11.4 Conclusion

Imaging of the nigrostriatal pathway in humans can be performed quantitatively with PET and SPECT techniques, using the PET radiotracer [<sup>18</sup>F]FDOPA or [<sup>18</sup>F]FMT, PET radioligands for the VMAT-2 and PET or SPECT radioligands for the DAT. With PET, also several other dopaminergic projections (e.g. mesocortical projections) can be assessed adequately in the human brain. Several antagonist PET radioligands for the dopamine D<sub>1</sub> receptor have been developed successfully. In addition, well-validated antagonist PET and SPECT radioligands are available to image dopamine D<sub>2/3</sub> receptors in the living human brain. Relatively recently, also agonist PET radioligands for the dopamine D<sub>2/3</sub> receptors have been developed, which affords the opportunity to evaluate the existence of the high-affinity state of these receptors *in vivo*, and these radiopharmaceuticals may be more sensitive to detect changes in dopamine concentrations. Finally, although selective antagonist PET radioligands for the dopamine D<sub>4</sub> receptor have been synthesized and evaluated successfully in small laboratory animals, these radioligands have not yet been applied in human research.

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# PET Imaging of the Endocannabinoid System

# 12

Garth E. Terry, Vanessa Raymont, and Andrew G. Horti

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## Abstract

The endocannabinoid system consists of cannabinoid receptors (which mediate the actions of cannabis), their endogenous ligands (endocannabinoids), and the enzymes and proteins associated with their regulation. In brain, the endocannabinoid system, in part, functions to modulate the release of other neurotransmitters via the subtype 1 (CB<sub>1</sub>) receptor. Abnormalities of CB<sub>1</sub> receptor expression or endocannabinoid transmission have been associated with several neuropsychiatric diseases. Subtype 2 (CB<sub>2</sub>) receptors are found primarily on immune and neuroimmune cells, and are overexpressed in activated microglia and neuroinflammation. Both receptors are of particular interest for biomarker development and therapeutic targets. Fatty acid amide hydrolase (FAAH) and monoacylglycerol lipase (MAGL) are responsible for the breakdown of endocannabinoids. Growing evidence supports the regulation of endocannabinoids as involved in neuropsychiatric and neuroinflammatory diseases. However, the function of endocannabinoid system components *in vivo* remains difficult to assess. Therefore, the use of functional imaging techniques, such as positron-emission tomography (PET), may be particularly useful in the assessment and quantification of the endocannabinoid system. This review covers the current understanding of PET imaging that directly targets the endocannabinoid system.

## Abbreviations

2-AG	2-Arachidonoyl glycerol
AAI	Aminoalkylindoles
AChE	Acetylcholinesterase
AD	Alzheimer's disease
AEA	N-arachidonylethanolamine, aka anandamide
AIDS	Acquired immunodeficiency syndrome
ALS	Amyotrophic lateral sclerosis

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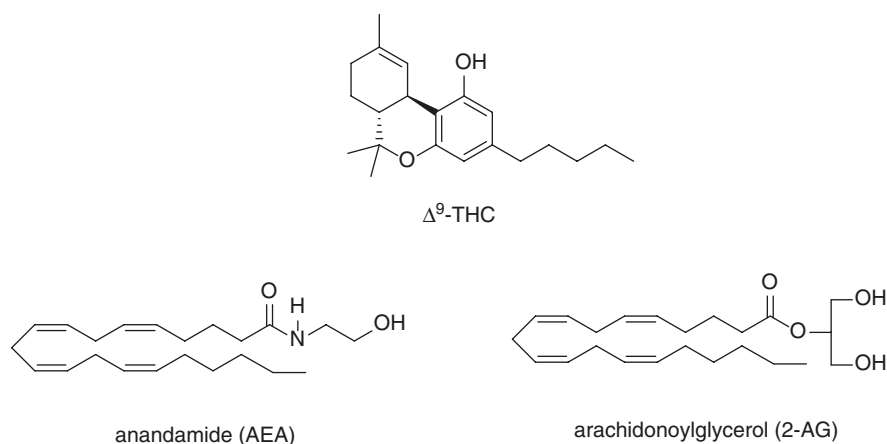
AM404	An endocannabinoid transporter inhibitor
BAT	Brown adipose tissue
BBB	Blood-brain barrier
$B_{\max}$	Binding maximum (i.e., total receptor density)
BMI	Body mass index
BP	Binding potential
cAMP	3'-5'-Cyclic adenosine monophosphate
CB <sub>1</sub>	Cannabinoid subtype 1
CB <sub>2</sub>	Cannabinoid subtype 2
<i>CNR1</i>	Gene encoding cannabinoid subtype 1 receptor
<i>CNR2</i>	Gene encoding cannabinoid subtype 2 receptor
CNS	Central nervous system
COX-2	Cyclooxygenase type 2
CP-55,940	A nonclassical cannabinoid agonist
D <sub>2</sub>	Dopamine subtype 2
DAGL	Diacylglycerol lipase
EAE	Experimental autoimmune encephalomyelitis
ECS	Endocannabinoid system
FAAH	Fatty acid amide hydrolase
FUR	Fractional uptake ratio
GABA	Gamma-aminobutyric acid
GPCR	G-protein-coupled receptor
GPR55	G-protein-coupled receptor 55
GTP	Guanosine-5'-triphosphate
HD	Huntington's disease
HRRT	High-resolution research tomograph
ICC	Intraclass correlation coefficient
$K_D$	Dissociation constant
$K_i$	Inhibition constant
LTD	Long-term depression
LTP	Long-term potentiation
MAGL	Monoacylglycerol lipase
<i>MDR1</i>	Multidrug resistance gene 1 (encodes P-gp)
mRNA	Messenger ribonucleic acid
MS	Multiple sclerosis
MS	Multiple sclerosis
mSUV	Modified standardized uptake value
MW	Molecular weight
NAPE-PLD	N-acyl-phosphatidylethanolamine-selective phospholipase D
PANSS	Positive and negative syndrome scale
PD	Parkinson's disease
PET	Positron-emission tomography
P-gp	P-glycoprotein (an efflux transporter)
PSA	Polar surface area
SNP	Single-nucleotide polymorphism

SPECT	Single-photon emission computed tomography
SUV	Standardized uptake value
TCM	Tissue compartment model
TGN	Trigeminal nerve
THC, $\Delta^9$ -THC	$\Delta^9$ -Tetrahydrocannabinol
TRPV1	Transient receptor potential vanilloid 1 (a cation channel)
TSPO	Translocator protein (18 kDa)
URB597	A FAAH inhibitor
$V_T$	Total distribution volume
WIN 55,212-2	An aminoalkylindole cannabinoid agonist

## 12.1 Introduction

### 12.1.1 Endocannabinoid System

Cannabis has been known to have medicinal and narcotic effects for thousands of years and is the most commonly used illegal drug in the world today (Anthony et al. 2017; Copeland and Swift 2009). However, it was not until 1964 that  $\Delta^9$ -tetrahydrocannabinol ( $\Delta^9$ -THC) was described by Gaoni and Mechoulam as the principal psychoactive component of cannabis (Gaoni and Mechoulam 1964) (Fig. 12.1). While over 100 plant-derived cannabinoids (phytocannabinoids) have been discovered, to date most remain insufficiently characterized (Turner et al. 2017). For approximately the last 20 years,  $\Delta^9$ -THC and cannabidiol (CBD) have received increasingly focused attention for their potential therapeutic effects and their greater concentration in cultivated cannabis. In particular,  $\Delta^9$ -THC



**Fig. 12.1** Structure of  $\Delta^9$ -THC and the endocannabinoids, anandamide (AEA), and 2-arachidonoylglycerol (2-AG)



concentration in cannabis has increased from less than 1.5% in 1980 to over 17% in 2017, and nearly 30% in regions with legalized cannabis<sup>1</sup> (ElSohly et al. 2000; Chandra et al. 2019; Smart et al. 2017). This has prompted the need for additional research of cannabis and cannabinoids, their impact on health, and the biology and pharmacology they have uncovered.

The effects of  $\Delta^9$ -THC are primarily mediated by G-protein-coupled receptors (GPCR) discovered in the 1990s and termed cannabinoid receptors. The first cannabinoid receptor subtype, CB<sub>1</sub>, is highly concentrated in brain and was found to be responsible for the psychotropic effects of cannabinoids (Matsuda et al. 1990; Devane et al. 1988). CB<sub>1</sub> receptors are abundantly expressed in hippocampus, striatum, cerebral cortex, and molecular layer of the cerebellum, and with the lowest density in the brainstem, thalamus, and white matter (Howlett et al. 2002; Lynn and Herkenham 1994; Herkenham et al. 1990, 1991; Glass et al. 1997; Iversen 2003). CB<sub>1</sub> is also widely expressed in the gut, liver, brown adipose tissue, and bone marrow (Massa et al. 2005).

The second cannabinoid receptor subtype, CB<sub>2</sub>, was discovered shortly thereafter and found primarily on cells associated with the immune system (Howlett et al. 2002; Munro et al. 1993). Broadly CB<sub>2</sub> receptors participate in the regulation of cytokine release and function (Klein 2005). They are typically in low density in the brain (Van Sickle et al. 2005; Onaivi et al. 2006), but are overexpressed during states of neuroinflammation on activated microglia (Howlett et al. 2002). In the periphery, they have high density in spleen and leukocytes, particularly B-lymphocytes (Munro et al. 1993).

The discovery of cannabinoid receptors prompted the search for their endogenous ligands, termed endocannabinoids (*endogenous cannabinoids*), which could activate them. The first identified endocannabinoid, anandamide (N-arachidonylethanolamine; AEA), was named for the Sanskrit word *ananda*, meaning “bliss,” and has been characterized as a partial agonist for cannabinoid receptors (Fig. 12.1) (Devane et al. 1992). Subsequently, 2-arachidonoylglycerol (2-AG) was identified as a full agonist and at substantially higher concentrations in brain than anandamide, though with a lower binding affinity (Stella et al. 1997). Both anandamide and 2-AG are able to mimic several pharmacological effects of  $\Delta^9$ -THC (Paradisi et al. 2006).

In contrast to monoamine neurotransmitters which are synthesized, stored in vesicles, and released upon signaling by presynaptic neurons, endocannabinoids are synthesized on demand following Ca<sup>2+</sup> elevation in the postsynaptic neuron. Anandamide is cleaved from N-acyl-phosphatidylethanolamine by phospholipase D (NAPE-PLD), and 2-AG is generated from diacylglycerol hydrolyzed by

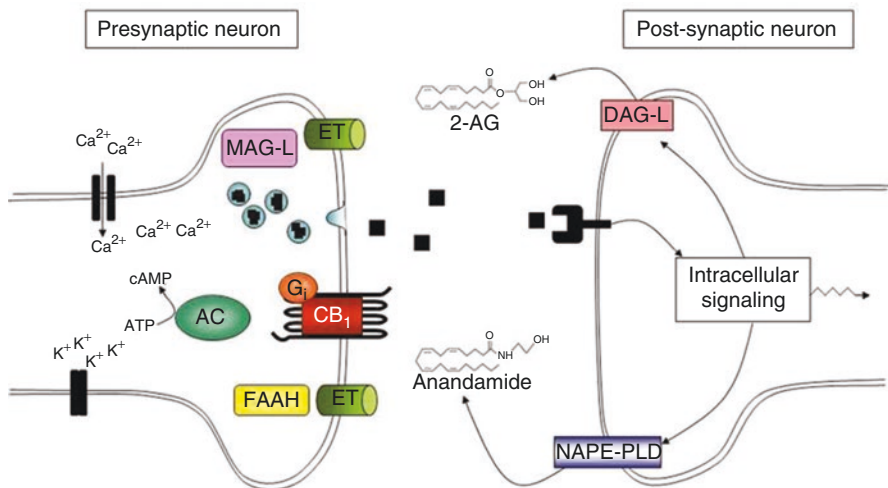
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<sup>1</sup>As of this writing (mid-2019) cannabis remains largely illegal throughout the world. Uruguay and Canada have legalized cannabis for recreational use, while South Africa and Georgia have legalized the possession and consumption, but not the sale, of cannabis for recreational use. Many other countries have legalized cannabis for medicinal uses only. Cannabis remains illegal in the United States on a federal level, while laws are mixed across states and territories ranging from legal for recreational and medical use, legal for medical use only, and illegal but decriminalized to fully illegal.

diacylglycerol lipase (DAG-L) (Di Marzo et al. 2011; Marsicano et al. 2003). They subsequently diffuse across the synapse in a retrograde manner where they act on presynaptic CB<sub>1</sub> receptors, which in turn propagate second messenger signaling through G-stimulatory and G-inhibitory proteins, among other mechanisms. The resulting cascade of events leads to decreased cAMP production, closing of Ca<sup>2+</sup> channels, and reduced mobilization of neurotransmitter-containing vesicles, and therefore an inhibition of neurotransmitter release (e.g., GABA, glutamate, dopamine).

Endocannabinoids undergo a reuptake and degradation process. Fatty acid amide hydrolase (FAAH) is primarily responsible for the breakdown of anandamide, whereas the monoacylglycerol lipase (MAGL) plays a key role in the enzymatic hydrolysis of about 85% of 2-AG (Dinh et al. 2002), with  $\alpha/\beta$  hydrolase domain-6 and -12 (ABHD6 and ABHD12) additionally contributing to 2-AG catabolism. Cyclooxygenase 2 (COX-2), in addition to its production of prostaglandins from arachidonic acid, metabolizes endocannabinoids to prostacannabinoids, which in turn can participate in cell signaling (Kozak et al. 2002). Indirect evidence supports the presence of an endocannabinoid transporter on postsynaptic membranes; however no protein structure has been elucidated to confirm its existence (Felder et al. 2006; Fowler 2012). Acetaminophen (paracetamol), a widely used pain reliever which has long lacked an understood mechanism, is metabolized *in vivo* to N-(4-hydroxyphenyl)arachidonylethanolamide (AM404), an inhibitor of the putative endocannabinoid transporter and a compound known to reduce pain (Hogestatt et al. 2005; La Rana et al. 2006; Bertolini et al. 2006).

Cannabinoid receptors, endocannabinoids, and enzymes for endocannabinoid synthesis, uptake, and degradation constitute the endocannabinoid system (ECS) (Fig. 12.2) (Joshi and Onaivi 2019). Other receptors identified as targets of



**Fig. 12.2** Endocannabinoid system signaling (reprinted from Terry (2009))

phyto- and endocannabinoids (e.g., transient receptor potential cation channel subfamily V member 1 [TRPV1], orphan receptors such as GPR18 and GPR55) (Di Marzo and De Petrocellis 2010; Irving et al. 2017), putative endocannabinoids (e.g., 2-arachidonyl glyceryl ether [noladin ether]), endocannabinoid-like compounds (e.g., N-palmitoylethanolamide [PEA]), and cannabinoid receptor-binding endogenous compounds (e.g., hemopressin-like peptides) are considered to be closely involved with the ECS, but not definitively classified as part of it (Pertwee 2015; Pertwee et al. 2010). In particular, TRPV1 is a receptor at which anandamide has been identified as a full agonist, and may be partly responsible for the analgesic properties of the phytocannabinoid cannabidiol (CBD), which has low binding affinity to CB<sub>1</sub> and CB<sub>2</sub> (Muller et al. 2018). The ECS is fundamental to physiology, and has widespread influences on the human brain, immune system, and other organs. It is found in fish, reptiles, and all mammals, suggesting that the ECS is evolutionarily an ancient system (Elphick and Egertova 2005). Abnormalities of this system have been implicated in many cognitive, neuropsychiatric, and metabolic disorders (Pacher et al. 2006).

### 12.1.2 Physiology of the Endocannabinoid System

The ECS plays an important modulating role in homeostasis, metabolism, and immunological reactions as well as motor functions, learning, and reward processing. In the brain, endocannabinoids mostly act on presynaptic CB<sub>1</sub> receptors as retrograde messengers and modulate the neurotransmitters GABA-A, dopamine, glutamate, and opioids, as described above (Schlicker and Kathmann 2001; Wilson and Nicoll 2002). While this represents a rather straightforward model, cannabinoid receptors possess a richness of pharmacology and present challenges for research. For example, CB<sub>1</sub> and CB<sub>2</sub> receptors are associated with a variety of GPCR-associated second messenger systems other than G-inhibitory and G-stimulatory proteins, such as mitogen-activated protein kinase (MAPK), phosphatidylinositol-3-kinases (PI3K), and G-protein inward-rectifying potassium (GIRK) channels. In addition, as GPCRs they have the ability to be in high-affinity states (G-protein coupled and agonist preferring), in low-affinity states (G-protein uncoupled), and constitutively active (*i.e.*, engaging in signal transduction without the binding of a ligand). However, due to the lipophilicity of endocannabinoids and their ability to travel within the plasma membrane, they may laterally translocate to the binding site; thus, the description of true inverse agonist activity of these receptors has been called into question in lieu of antagonist and/or partial agonist inhibition of baseline endocannabinoid activity. A substantial portion of CB<sub>1</sub> receptors can be internalized in endosomes, or located on intracellular lysosomes and mitochondria, increasing its receptor reserve (Zou and Kumar 2018). Finally, cannabinoid receptors are able to undergo homo- and heteromerization with other GPCRs with cross-cell transduction signaling.

Generally, increased or decreased endocannabinoid neurotransmission results in decreases or increases in neurotransmitter release, respectively (Iversen 2003),

and additionally, changes in neurotransmitter function can modulate endocannabinoid signaling (Degroot and Nomikos 2007). By inhibiting excitatory or inhibitory signals, endocannabinoids facilitate long-term potentiation (LTP) or depression (LTD) between neurons both directly and indirectly, which enables the process of learning and memory formation (Di Marzo et al. 2004; Riedel and Davies 2005). The process of learning and memory formation likely depends on the timing and location of endocannabinoid release, which permits the extinction of previous memories (Chhatwal et al. 2005). Understandably then, the ECS plays an important role in neural development and synaptic formations given that repeated stimulation between excitatory and inhibitory connections is part of synaptic pruning and neuronal maturation (Dow-Edwards and Silva 2017). Therefore, global stimulation of CB<sub>1</sub> receptors, as from cannabis, interrupts the local formation of LTP and LTD and delays learning and memory, and can impact neuronal development. As well, aberrant expression or function of the ECS may represent change of the system itself, or compensatory changes of other neurotransmission systems, and thus has the potential as a biomarker or therapeutic target in numerous neuropsychiatric disorders.

CB<sub>1</sub> receptors are associated with a wide range of behaviors, including cognition, pain perception, movement, drug addiction, and memory consolidation. Stimulation of these receptors also has anti-inflammatory, antioxidant, and neuroprotective effects (Drysdale and Platt 2003), as well as a role in neuronal gene expression (Piomelli 2003) and endocrinological modulation (Cota et al. 2007). CB<sub>1</sub> receptor expression is affected by the estrous cycle in females (Rodriguez de Fonseca et al. 1994), suggesting its sensitivity to estrogen and potential for sex-related differences in expression or function. As such, when examining the ECS, differences between males, females, and phases of the estrous cycle may need to be taken into consideration.

Less is known about the physiological role of CB<sub>2</sub> receptors. In the absence of inflammatory CNS disease, CB<sub>2</sub> receptors are not expressed in any significant amount in the human brain with the exception of the brainstem, where low concentrations can be found (Van Sickle et al. 2005). However, in inflammatory conditions such as cancer, atherosclerosis, peripheral inflammation, and several brain disorders there is a marked upregulation of CB<sub>2</sub> receptors, primarily in activated microglia, which can be ten times that of basal levels (Maresz et al. 2005; Cabral and Marciano-Cabral 2005). In neurodegenerative brain disorders, CB<sub>2</sub> receptor upregulation was noted postmortem in microglia surrounding and coincident with the beta-amyloid plaques of patients with Alzheimer's disease (Benito et al. 2003), and related to the extent and progression of neuronal loss in Huntington's disease (Sapp et al. 2001). CB<sub>2</sub> agonists have been found to stimulate human macrophages to remove beta-amyloid in *in vitro* studies, thereby representing a potential therapeutic approach in the treatment of neurodegenerative disorders (Tolon et al. 2009). CB<sub>2</sub> receptors have been implicated in mediating pain transmission (*i.e.*, nociception) and in the function and communication of immune cells (*e.g.*, mediating cytokine release). CB<sub>2</sub> receptor agonists have been shown to have analgesic effects and decrease lymphocyte function (Hsieh et al. 2011; Cencioni et al. 2010). In brain cancers, CB<sub>2</sub>

receptor expression has been found to correlate with the stage of tumor malignancy (Ellert-Miklaszewska et al. 2007; Calatuzzolo et al. 2007).

As FAAH is highly involved in controlling the concentration of endocannabinoids, it has a key role in determining endocannabinoid tone. In brain, FAAH expression differs from that of CB<sub>1</sub> receptors with regard to distribution and density. The gene encoding FAAH, *FAAH*, contains an estrogen-binding site on the promoter region (Waleh et al. 2002), again suggesting sex differences in the ECS. Genetic knockout or pharmacologic blockade of FAAH has been shown to reduce anxiety and depression in animal models (Moreira et al. 2008), and an identified single-nucleotide polymorphism on *FAAH* and microdeletion on a pseudogene leading to FAAH loss of function have been associated with reduced anxiety and pain sensitivity (Habib et al. 2019).

### 12.1.3 Pharmacology of the Endocannabinoid System

Cannabis has been consumed in its natural form since 3000 years, and has been utilized as an analgesic, appetite stimulant, anti-inflammatory, and decongestant, among other uses (Booth 2005). The ECS is vital to learning, memory, motor behavior, pain and weight control, as well as inflammation, gut motility, and intracellular apoptosis. Thus, cannabinoid drugs are being used or have the potential for use in treating multiple pathologies, including anorexia (in AIDS and cachexia secondary to cancer), emesis, obesity, pain, inflammation, schizophrenia, multiple sclerosis, neurodegenerative disorders, epilepsy, stroke, glaucoma, osteoporosis, cardiovascular disorders, and cancer (Di Marzo et al. 2004; Kogan and Mechoulam 2007; Howlett 1995; Pertwee 1999; Timpone et al. 1997).

Despite the potential therapeutic value of the cannabinoid partial agonist  $\Delta^9$ -THC and its analogues, side effects have been a problem and the potential for abuse has been a concern (Howlett et al. 2002; Howlett 1995, 2002; Murray et al. 2007), although it has become apparent that adverse effects may be reduced by developing subtype-selective drugs (Hertzog 2004). Two cannabinoid agonists on the market in the United States, dronabinol (Marinol,  $\Delta^9$ -THC) and nabilone (Cesamet, a THC synthetic analogue), are approved for treatment of nausea, vomiting, and appetite stimulation in cancer and AIDS. Nabiximols (Sativex<sup>®</sup>), a  $\Delta^9$ -THC and cannabidiol mixture, is available in some EU countries and Canada, as the treatment for symptoms of multiple sclerosis. Cannabidiol (Epidiolex) has been approved in the United States for seizures associated with the severe pediatric disorders Lennox-Gastaut and Dravet syndromes.

The first clinically useful selective CB<sub>1</sub> inverse agonist, rimonabant, has been used as an anti-obesity medication, and was investigated for use in alcohol and tobacco addiction (Van Gaal et al. 2005; Pi-Sunyer et al. 2006; Rosenstock et al. 2008; Despres et al. 2009; Nissen et al. 2008; Soyka et al. 2008; Cahill and Ussher 2007). In addition to its central mechanism by decreasing appetite, rimonabant mediated improvements in peripheral comorbidities of obesity, including increased HDL, decreased triglyceride levels, and decreased hemoglobin A<sub>1c</sub> (Van Gaal et al.

2005). Later, rimonabant was withdrawn from the market due to its central side effects of depression and anxiety. CB<sub>1</sub> inverse agonists that act predominantly peripherally (thus avoiding psychiatric side effects (Cluny et al. 2010)) are under preclinical investigation (Chorvat 2013). A number of FAAH inhibitors (notably PF-04457845 and URB597) are also under preclinical investigation to target anxiety disorders (Piomelli et al. 2006) and cannabis withdrawal (D'Souza et al. 2019).

### 12.1.4 Special Considerations for Radioligand Development and PET Imaging of the Endocannabinoid System

Until recently, only *in vitro*, animal model, electrophysiological, behavioral, and postmortem data existed on the ECS. However, there is a need for direct functional measurement in human subjects given the differences between species (Howlett et al. 2002; Pertwee et al. 2010). There are several important applications of functional imaging of the central CB<sub>1</sub> receptor using PET and SPECT (Horti and Van Laere 2008; Van Laere 2007). They have a role in planning drug therapy using inverse agonists or antagonists. Quantitative information about the receptor occupancy can be obtained after a titration of individual doses and existing radioligands. They also allow for the investigation of CNS side effects, which are frequent problems, particularly with CB<sub>1</sub> receptor blockade. PET or SPECT imaging allows information from preclinical models to be tested in humans, and may also lead to the development of novel diagnostic biomarkers where marked regional up- or downregulation is present. These imaging techniques can be used to examine the interactions of the ECS with other neurotransmitter systems (*e.g.*, GABA-A, dopamine, and opioid systems). Functional imaging could provide quantitative measurements of endocannabinoid release and allow study of its effects in pathological and cognitive states. However, more data is needed on the sensitivity of existing cannabinoid ligands to displacement by endocannabinoids.

Multiple efforts have been undertaken to probe the ECS through imaging. Methods such as [<sup>15</sup>O]H<sub>2</sub>O or [<sup>18</sup>F]FDG PET, or fMRI, do not directly probe the ECS, though they have provided information regarding the effects of cannabinoids on the brain and behavior. Several studies have evaluated the effect of cannabis and Δ<sup>9</sup>-THC on the dopaminergic system with mixed results; while animal studies have demonstrated dopamine release in the nucleus accumbens after exposure to Δ<sup>9</sup>-THC, *in vivo* human studies using PET have demonstrated little to no dopamine release, suggesting alternative neuromolecular mechanisms for reinforcement and addiction (Zehra et al. 2018; Bloomfield et al. 2016; Urban and Martinez 2012). While these results are intriguing, they are outside the scope of this review, and have been described elsewhere (Martin-Santos et al. 2010; Bloomfield et al. 2019).

#### 12.1.4.1 Radioligand Requirements

Generally, PET radioligands for quantification of cerebral receptors must possess a variety of imaging properties which have been previously described (Eckelman et al. 1979; Waterhouse 2003; Pike 2009; Zhang and Villalobos 2017). Briefly, the

most important characteristics of successful radioligands include high binding affinity with high selectivity for the intended target and low nonspecific binding, and characteristics generally predictive of blood-brain barrier penetration including moderate lipophilicity, low-to-moderate molecular weight, and appropriate polar surface area (PSA). Rapid clearance from the blood, appropriate brain kinetics, and absence of active radiometabolites are also among the crucial imaging properties of cerebral PET radioligands. As well, radioligands must be safe for human use, and possess a low radiation burden and low toxicity. There is certain serendipity in the development of novel PET radioligands because relationships of some physical-chemical properties with imaging properties are still not fully understood.

The typical range of essential parameters for successfully developed PET radioligands is summarized in Table 12.1. It is conventionally agreed that most good CNS PET radioligands with reasonable BBB permeability and minimal nonspecific binding should exhibit lipophilicity of  $\log D_{7.4} = 0-4$ . Radioligands with a low value of  $\log D_{7.4}$  are likely to exhibit a low BBB permeability, whereas very lipophilic compounds bind to plasma proteins and produce a low concentration of free radioligand in the blood and, correspondingly, show poor permeability through the BBB, and can produce high nonspecific binding. The high lipophilicity of most cannabinoids and cannabinomimetic ligands ( $\log D_{7.4} \gg 4$ ) remains a main concern of medicinal radiochemists who are involved in the development of CB<sub>1</sub> and CB<sub>2</sub> PET radioligands. A number of high-affinity-labeled cannabinoids demonstrated poor imaging properties due to their high lipophilicity. Development of high-affinity cannabinoids with reduced lipophilicities represents a difficult undertaking (Katoch-Rouse et al. 2003). Most successful CB<sub>1</sub> and CB<sub>2</sub> radiotracers reviewed and shown below that were developed in the last few years exhibit relatively high lipophilicity values ( $\text{clog}D_{7.4} = 3.3-6.0$ ).

The classic Lipinski “rule of 5” that proposes an upper limit of molecular weight (MW) of 500 as acceptable for orally absorbed drugs seems to be valid for the majority of successful PET radiotracers for brain imaging that are administered intravenously in humans. PSA is used for characterization of the transport and partitioning properties of cerebral drugs and radiotracers. PSA embodies an optimal combination of H-bonding, molecular polarity, and solubility. A common measure of brain/blood distribution ( $\log\text{BB}$ ) can be calculated as a simple function of lipophilicity and PSA. The great majority of CNS drugs exhibit a PSA value of  $<70 \text{ \AA}^2$  (Ertl 2008). Calculation of PSA for all currently available PET radioligands for CNS receptor imaging in humans suggests a window of opportunity PSA  $<84 \text{ \AA}^2$  (Table 12.1) (Horti 2011). It is likely that successful CB<sub>1</sub> and CB<sub>2</sub> radioligands for

**Table 12.1** Molecular descriptors of successful PET radioligands ( $n = 23$ ) for imaging of brain-binding sites in humans (Horti 2011)

Molecular property	Range	Mean $\pm$ SD
Molecular weight (Dalton)	157–490	319 $\pm$ 87
Lipophilicity, $\text{clog}D_{7.4}$	–1.65 to 6.00	2.1 $\pm$ 1.8
Polar surface area, PSA ( $\text{\AA}^2$ )	23–84	47 $\pm$ 19

PET imaging should display the molecular property range that is shown in Table 12.1.

An additional component to the BBB is the presence of efflux transporters, which prevent entry and/or eject substrates from the brain (Pike 2009). In particular, permeability glycoprotein (P-gp)-mediated efflux has proven to be a burden to CNS drug and radiotracer development.  $\Delta^9$ -THC and several other cannabinoid ligands are partial substrates for this particular transporter (Bonhomme-Faivre et al. 2008a). P-gp is normally expressed at the BBB and blood-testes barrier to protect these sensitive tissues from potentially harmful compounds, and is sometimes overexpressed on cancer cells, causing multidrug resistance (hence the gene name, *MDR1*) (Ambudkar et al. 2003). As a result, compounds that act as substrates for P-gp are prevented from entering and accumulating in the brain; many CNS drug and radiotracer candidates have been abandoned due to this limitation.

#### 12.1.4.2 Binding Site Density and the Radioligand Binding Affinity

An advantage of PET is its quantitative ability, and the main quantitative measures are typically standardized uptake value (SUV), distribution volume ( $V_T$ ), and binding potential ( $BP$ ). Organ uptake of radioactivity is generally measured as SUV (or %SUV), which corrects only for injected radioactivity and total body weight, but does not adjust for nonspecific binding, free and unbound radioligand, and peripheral metabolism of the radioligand. Therefore, SUV and its derivative measures can reflect relative biodistribution of the imaging target, but might not accurately reflect target density. At equilibrium, both  $BP$  and  $V_T$  are proportional to the density of available receptor-binding sites ( $B_{max}$ ) and the binding affinity of the radioligand; a ligand's binding affinity is inversely proportional to its  $K_D$  (binding affinity =  $1/K_D$ ;  $BP = B_{max}/K_D$ ). A full description, derivation, and definition of related parameters can be reviewed elsewhere (Innis et al. 2007). The cerebral receptors with a greater value of  $B_{max}$  exhibit greater  $BP$  values than those of the receptors with low  $B_{max}$ . Obviously, the binding affinity of a cerebral radioligand has to be optimally high in order to gather sufficiently quantitative data. However, it is also essential to keep in mind that CNS radioligands with binding affinities that are too high can demonstrate slow or even seemingly irreversible brain kinetics that may complicate quantification of the receptor. Additionally, radioligands must demonstrate sufficient specific signal, and thus have low affinity for other nontarget receptors or proteins, and low nonspecific binding to proteins, lipids, or extracellular matrix.

CB<sub>1</sub> receptor radioligands have been successfully developed in spite of their moderately high lipophilicity due in part to the high  $B_{max}$  of CB<sub>1</sub> receptors in the CNS (Howlett et al. 2002). The low density of CB<sub>2</sub> receptors then likely creates an additional need for CNS radioligands to have a lower or more moderate lipophilicity.

Estimation of required binding affinities using a conventional approach (Eckelman et al. 1979) suggests that a successful CB<sub>1</sub> receptor PET radioligand should exhibit a moderately high binding affinity with a low nanomolar value of  $K_i$  or even lower.  $B_{max}$  data for the CB<sub>2</sub> subtype in normal mammalian brain tissue and during neuroinflammation are not available yet, but expression of *CNRI* gene



transcripts in the normal mouse brain stem is 100 times the *CNR2* gene expression (Onaivi et al. 2006). These results make us to believe that a good CB<sub>2</sub> receptor radioligand for PET should exhibit very high binding affinities with  $K_i$  values in the picomolar range.

## 12.2 Imaging of CB<sub>1</sub> Receptors

Radioligands that have been developed for imaging CB<sub>1</sub> receptors may be broadly categorized as first- and second-generation compounds. As discussed below, first-generation compounds were limited by their low specific binding and poor brain uptake. Second-generation compounds achieved success largely by retaining a high binding affinity with more moderate lipophilicity. Pharmacokinetic modeling of these radioligands is challenging; however these compounds have been used in numerous clinical investigations.

### 12.2.1 Initial CB<sub>1</sub> Radioligands and Imaging Studies

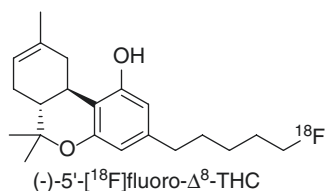
#### 12.2.1.1 (–)-5′-[<sup>18</sup>F]-Δ<sup>8</sup>-THC

The starting point in the development of CB<sub>1</sub> receptor PET radioligands was established when a <sup>18</sup>F-derivative of Δ<sup>8</sup>-THC (Fig. 12.3), one of the active constituents of marijuana, was synthesized in 1991 (Charalambous et al. 1991). The radioligand was studied in mice and baboons. The mouse study demonstrated some initial brain uptake of (–)-5′-[<sup>18</sup>F]-Δ<sup>8</sup>-THC (1% ID/g tissue, 5 min) with rapid clearance. The clearance of radioactivity from the baboon brain was also rapid and no specific binding was observed. The relatively low binding affinity of the radioligand ( $K_i = 57$  nM) (Compton et al. 1993), its potentially P-gp-mediated efflux (Bonhomme-Faivre et al. 2008b), and, most importantly, its high lipophilicity ( $\text{clogD}_{7.4} = 6.9$ ) are the likely factors behind the discouraging imaging results.

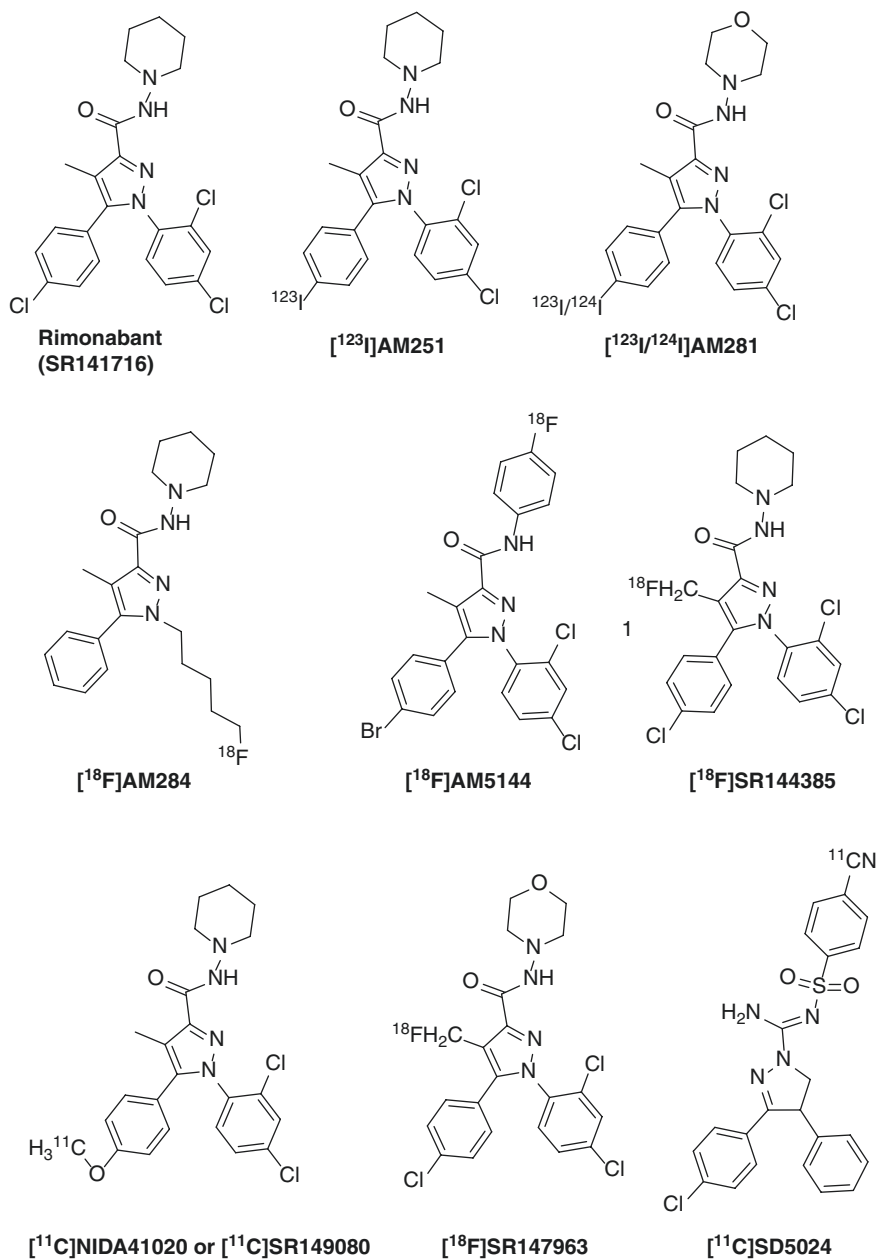
#### 12.2.1.2 Radiolabeled CB<sub>1</sub> Antagonists, Derivatives of SR141716 (Rimonabant)

In 1994 the discovery of rimonabant, a highly selective and potent CB<sub>1</sub> receptor antagonist, by Sanofi-Aventis (Rinaldi-Carmona et al. 1994) stimulated progress in the development of PET and SPECT radioligands for imaging CB<sub>1</sub> receptors. Several research groups synthesized many analogues of rimonabant and

**Fig. 12.3** Structure of (–)-5′-[<sup>18</sup>F]-Δ<sup>8</sup>-THC, the first cannabinoid PET radioligand



radiolabeled some of them for emission tomography studies (Katoch-Rouse et al. 2003; Lan et al. 1996, 1999; Gatley et al. 1996, 1998; Mathews et al. 1999, 2000, 2002; Katoch-Rouse et al. 2002; Nojiri and Ishiwata 2008) (Fig. 12.4). While more



**Fig. 12.4** Initial radiolabeled analogues of rimonabant and structurally related compounds

details on the initial stage of CB<sub>1</sub> radioligand development can be obtained from the original literature and previous reviews (Horti and Van Laere 2008; Gatley et al. 2004) a brief summary of the most representative initial radioprobes is given below.

Iodinated analogues of SR141716, AM251 and AM281, exhibited high and moderate CB<sub>1</sub> receptor-binding affinity, respectively, and good CB<sub>1</sub>/CB<sub>2</sub> selectivity (Lan et al. 1999) and were radiolabeled with <sup>123</sup>I for SPECT studies (Lan et al. 1996; Gatley et al. 1998) (Fig. 12.4). The molecular weights of AM251 and AM281 are rather high (>500) for passive BBB transport (Table 12.2). In mouse experiments, [<sup>123</sup>I]AM251 and [<sup>123</sup>I]AM281 demonstrated reasonable specific brain uptake (Table 12.2) and both radioligands were found to be suitable for further baboon SPECT studies. In baboon experiments, [<sup>123</sup>I]AM251 exhibited little brain uptake, whereas [<sup>123</sup>I]AM281 showed good permeation of the blood-brain barrier (Gatley et al. 1998). A likely explanation for these results is the high lipophilicity of [<sup>123</sup>I]AM251 and lower lipophilicity of [<sup>123</sup>I]AM281 (Table 12.2) that is within the conventional range for *in vivo* radioligands. Notably, both radioligands manifest PSAs that are within the range for passive blood-brain transport (Table 12.2). The highest regional accumulation of [<sup>123</sup>I]AM281 in the baboon brain was in the cerebellum, cortex, and striatum, and lower in the other regions. A blocking study with rimonabant demonstrated that [<sup>123</sup>I]AM281 specifically bound to CB<sub>1</sub> receptors in baboon cerebellum but the value of the signal-to-noise ratio was moderate (Table 12.2). Substantial nonspecific binding was observed in other brain regions. In spite of the modest imaging results in animals [<sup>123</sup>I]AM281 remained the best available CB<sub>1</sub> receptor radioligand for almost a decade and the only hope for *in vivo* CB<sub>1</sub> receptor imaging in humans.

The first CB<sub>1</sub> receptor SPECT imaging study in humans was performed with [<sup>123</sup>I]AM281 in six patients suffering from Tourette's syndrome before and after Δ<sup>9</sup>-THC treatment (Berding et al. 2004). The radioligand displayed some brain uptake that did not change significantly after Δ<sup>9</sup>-THC treatment. However, V<sub>3</sub>' declined in the only patient that showed a treatment response suggesting that specific binding of [<sup>123</sup>I]AM281 in human subjects is mediated by CB<sub>1</sub> receptors. Because of the low specific signal and high nonspecific binding in human subjects, it was concluded that [<sup>124</sup>I]AM281/PET could give better results than [<sup>123</sup>I]AM281/SPECT because PET provides better image statistics than SPECT.

Human PET imaging with [<sup>124</sup>I]AM281 in a single patient with schizophrenia was performed by a research group from MHH (Berding et al. 2006). The radioligand manifested limited accumulation in regions enriched with CB<sub>1</sub> receptors: the striatum and pallidum (DVR - 1 = 0.3–0.4), and cerebral cortex and cerebellum (DVR - 1 = 0.2). It was also stated that the high radiation burden of <sup>124</sup>I, a radionuclide with 4-day half-life, represents a radiation safety issue in human subjects. The low specific signal, high nonspecific binding, and safety concerns were the main reasons why [<sup>124</sup>I]AM281-PET was never used again in clinical experiments.

In summary, initial studies with radiolabeled analogues of rimonabant (Table 12.2) demonstrated that CB<sub>1</sub> receptor imaging is a tough nut to crack. Structure-imaging properties of these radioligands suggest that most of them exhibit lipophilicity that is too high (logD<sub>7.4</sub> > 4) and/or binding affinity that is too low

**Table 12.2** Physical-chemical and *in vivo* imaging characteristics of the first generation of radioligands for PET/SPECT imaging of CB<sub>1</sub>

Code name	M. W. <sup>a</sup> (Dalton)	K <sub>i</sub> <sup>CB1</sup> (nM)	K <sub>i</sub> <sup>CB2</sup> (nM)	clogD <sub>7.4</sub> <sup>b</sup>	PSA <sup>b</sup>	Target-to-nontarget region ratio – 1			References
						Mouse	Monkey	Human	
<i>Analogues of rimonabant</i>									
[ <sup>123</sup> I]AM251	555	0.6–1	2290	5.3	50	0.7 <sup>d</sup>	Low uptake	–	Lan et al. (1999), Gatley et al. (1996, 1998)
[ <sup>123</sup> I]AM281 or [ <sup>124</sup> I]AM281	557	4.5	4200	3.9	59	~0.9 <sup>d</sup>	~0.4 <sup>e</sup>	0.21 <sup>i</sup> 0.37 <sup>i</sup>	Gatley et al. (1998), Berding et al. (2004, 2006); Gifford et al. (1997)
[ <sup>18</sup> F]AM284	372	30	–	5.0	50	0.4 <sup>f</sup>	–	–	Gatley et al. (2004), Li et al. (2005)
[ <sup>18</sup> F]AM5144	519	31	–	5.7	47	–	0.5 <sup>g</sup>	–	Li et al. (2005)
[ <sup>18</sup> F]SR144385	482	14 <sup>c</sup>	1000 <sup>e</sup>	4.8	50	1.5 <sup>h</sup>	–	–	Mathews et al. (1999, 2000)
[ <sup>18</sup> F]SR147963	484	120 <sup>c</sup>	–	3.4	59	0.7 <sup>h</sup>	–	–	Mathews et al. (2000)
[ <sup>11</sup> C]NIDA41020 or [ <sup>11</sup> C]SR149080	459	4.1–16	800	4.0	59	1.2 <sup>g</sup>	0.5 <sup>g</sup>	–	Katoch-Rouse et al. (2002, 2003); Mathews et al. (2002)
[ <sup>11</sup> C]SD5024	464	0.5	>5000	2.8	120	–	0.8 <sup>i</sup>	–	Donohue et al. (2008a)
<i>Indole derivatives</i>									
[ <sup>18</sup> F]NIDA54	400	0.7	–	3.8 <sup>j</sup>	25	–	0.7	–	Willis et al. (2005)
[ <sup>18</sup> F]pJSB	492	1.5 <sup>k</sup>	–	5.0	80	–	0.9	–	Finnema et al. (2009), Donohue et al. (2008b)

<sup>a</sup>Molecular weight<sup>b</sup>Lipophilicity (clogD<sub>7.4</sub>) and polar surface area (PSA) are calculated with ACD/Structure Designer software<sup>c</sup>[C<sub>50</sub> (Mathews et al. 2000)<sup>d</sup>Cerebellum/brain stem ratio – 1<sup>e</sup>Cerebellum baseline/cerebellum block with rimonabant ratio – 1<sup>f</sup>Whole-brain baseline/whole brain blocked with rimonabant ratio – 1<sup>g</sup>Cerebellum/thalamus ratio – 1<sup>h</sup>Hippocampus/thalamus ratio – 1<sup>i</sup>Binding potential<sup>j</sup>Experimental data<sup>k</sup>In vitro functional binding activity (K<sub>b</sub>)<sup>l</sup>Striatum/pons – 1 (time-uptake curves)

(Table 12.2). As a result, animal studies with these radioligands showed either little brain uptake or some brain uptake with high nonspecific binding and relatively low target-to-nontarget ratio – 1 value (<1). It is noteworthy that within the initial series of radioligands the most reasonable imaging properties were exhibited by the least lipophilic radioligand [ $^{123/124}$ I]AM281, which also manifested a moderate CB<sub>1</sub> receptor-binding affinity (Table 12.2). [ $^{123}$ I]AM251, an analogue of AM281 with excellent binding affinity but high lipophilicity, showed little brain uptake in monkey and was clearly not suitable for PET studies.

### 12.2.1.3 Radiolabeled Aminoalkylindole Derivatives and Structurally Related Compounds

Aminoalkylindoles (AAI) that are exemplified by WIN-55,212-2 (Fig. 12.5) represent a large group of compounds that exhibit typical cannabinoid pharmacology. Because the structure-activity relationship for AAIs is well developed (Huffman et al. 2005) and many AAI display excellent CB<sub>1</sub> affinity, it is not a surprise that several compounds of this class have been radiolabeled for PET or SPECT application (Willis et al. 2005; Finnema et al. 2009; Donohue et al. 2008b; Deng et al. 2005; Makriyannis and Deng 2001; Willis et al. 2003; Tamagnan et al. 1999).

The most notable PET radioligands among AAI are [ $^{18}$ F]NIDA54 (Willis et al. 2005) and [ $^{18}$ F]PipISB (Finnema et al. 2009; Donohue et al. 2008b). [ $^{18}$ F]NIDA54 manifests greater binding affinity and more appropriate molecular weight, PSA, and lipophilicity for a CNS radioligand, but both radioligands exhibit comparable specific binding in monkey brain. This finding can be explained by the functional properties of the inverse agonist [ $^{18}$ F]PipISB that should bind with a greater population of CB<sub>1</sub> receptors than agonist [ $^{18}$ F]NIDA54. Neither [ $^{18}$ F]NIDA54 nor [ $^{18}$ F]PipISB were sufficiently good CB<sub>1</sub> radiotracers in animal experiments to initiate further clinical investigation.

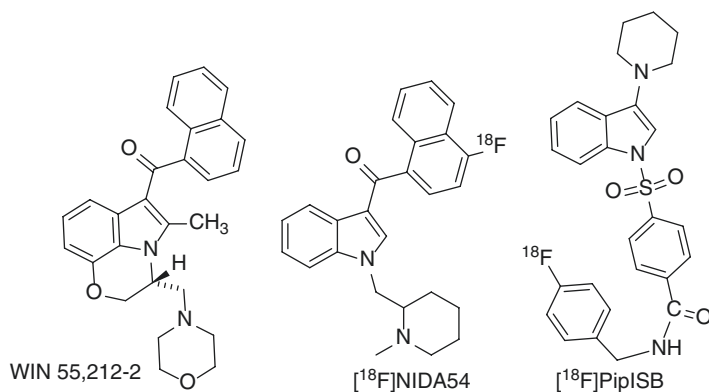


Fig. 12.5 Aminoalkylindole derivatives

## 12.2.2 Current CB<sub>1</sub> Receptor Radioligands for Human PET Imaging

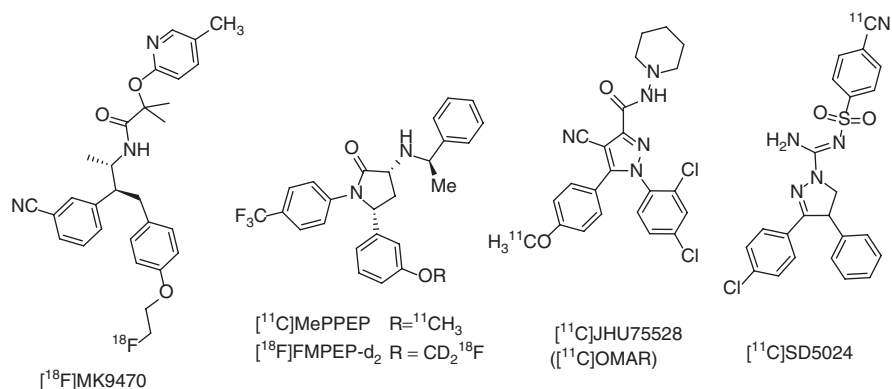
### 12.2.2.1 [<sup>11</sup>C]OMAR ([<sup>11</sup>C]JHU75528)

[<sup>11</sup>C]OMAR ([<sup>11</sup>C]JHU75528) was the first CB<sub>1</sub> selective radioligand with adequate properties for PET imaging (Horti et al. 2006; Fan et al. 2006). Structurally [<sup>11</sup>C]OMAR (Fig. 12.6) is an analog of rimonabant (Fig. 12.4) with comparable CB<sub>1</sub> receptor-binding affinity and good CB<sub>1</sub>/CB<sub>2</sub> selectivity. The lipophilicity of OMAR is considerably lower because the hydrophobic methyl and chlorine substituents of the rimonabant molecule are replaced with cyano- and methoxy-groups that exhibit negative constants of lipophilicity ( $\pi = 0.56, 0.71, -0.57, \text{ and } -0.02$ , correspondingly (Kubinyi 1995)), but the PSA value of OMAR is relatively high (Table 12.3). An *in vitro* functional assay demonstrated that OMAR is a cannabinoid antagonist (Fan et al. 2009).

Radiosynthesis of [<sup>11</sup>C]OMAR is straightforward (Fan et al. 2006) and is achieved by <sup>11</sup>C-methylation of the corresponding phenol precursor that is currently commercially available, or can be prepared as described (Gao et al. 2012).

In mice and baboon studies, [<sup>11</sup>C]OMAR manifested promising imaging results (Table 12.3). In comparison with the previously reported CB<sub>1</sub> receptor radiotracers of the first generation (Table 12.2), [<sup>11</sup>C]OMAR showed substantially greater signal-to-noise ratios. Pharmacological evaluation of [<sup>11</sup>C]OMAR in mice demonstrated that the radioligand labels CB<sub>1</sub> receptors specifically and selectively and does not bind to various other cerebral receptors including D<sub>1</sub>, D<sub>2</sub>, D<sub>5</sub>, 5-HT<sub>2A</sub>, 5-HT<sub>1C/2C</sub>, opioid, and  $\alpha$ 4 $\beta$ 2-nAChR. The specific *in vivo* binding of [<sup>11</sup>C]OMAR in CD1 mice was blocked by selective CB<sub>1</sub> antagonists in a dose-dependent manner (Horti et al. 2006). MicroPET studies with [<sup>11</sup>C]OMAR showed a 50% higher brain uptake in wild-type mice versus CB<sub>1</sub> knockout animals (Herance et al. 2011).

Baboon PET studies demonstrated that [<sup>11</sup>C]OMAR exhibits good values of BP in CB<sub>1</sub> receptor-rich brain regions (Table 12.3) and reasonable total brain uptake



**Fig. 12.6** Currently available CB<sub>1</sub> receptor PET radioligands for human studies

**Table 12.3** Physical-chemical and *in vivo* imaging characteristics of the clinical radioligands for PET imaging of CB<sub>1</sub>

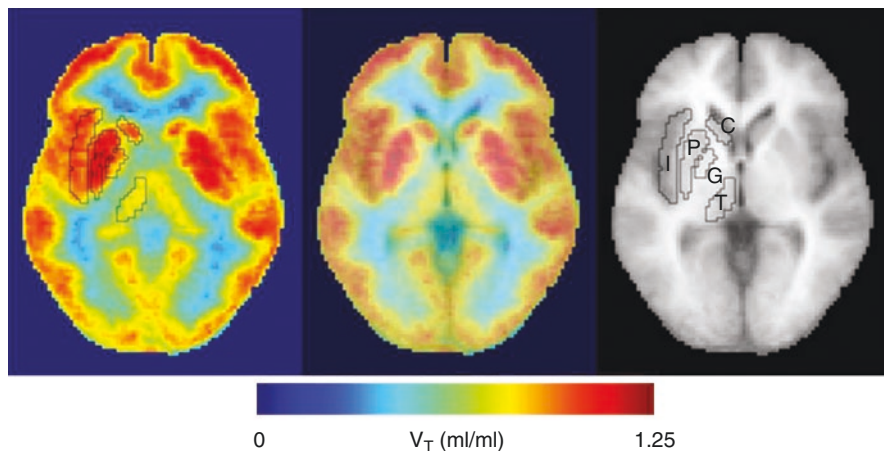
Code name	M.W. <sup>a</sup> (Dalton)	clogD <sub>7.4</sub> <sup>b</sup>	PSA	K <sub>i</sub> <sup>CB1</sup> (nM)	K <sub>i</sub> <sup>CB2</sup> (nM)	Target-to-nontarget region ratio – 1			References
						Mouse	Nonhuman primate	Human	
[ <sup>11</sup> C]OMAR	470	3.3–3.6 <sup>c</sup>	83	11; 2.05 <sup>f</sup>	2700 5250	2.4	1.4	1.0	Horti et al. (2006), Fan et al. (2006), Wong et al. (2010a), Tsujikawa et al. (2014)
[ <sup>18</sup> F] MK-9470	489	4.9	84	0.7; 0.10	44	–	~3–4	3	Burns et al. (2007), Van Laere et al. (2008a), Tsujikawa et al. (2014)
[ <sup>11</sup> C] MePEP	454	4.77 ± 0.27 <sup>c</sup>	42	0.5 <sup>d</sup> 0.11 <sup>f</sup>	363	2.3	~2	1.4	Terry et al. (2008), Yasuno et al. (2008), Tsujikawa et al. (2014)
[ <sup>18</sup> F] FMPEP-d <sub>2</sub>	474	4.24 ± 0.08 <sup>c</sup>	42	0.2 <sup>d</sup> 0.11 <sup>f</sup>	669	–	~2.2	1.8 <sup>e</sup>	Donohue et al. (2008c), Terry et al. (2009), Tsujikawa et al. (2014)
[ <sup>11</sup> C] SD5024	464	3.79 ± 0.09 <sup>c</sup>	120	0.47 <sup>f</sup>	n/a	–	~2.2	~1.1	Tsujikawa et al. (2014)

<sup>a</sup>Molecular weight<sup>b</sup>Lipophilicity (clogD<sub>7.4</sub>) and PSA are calculated with ACD/Structure Designer software<sup>c</sup>Experimental data<sup>d</sup>In vitro functional binding activity (K<sub>b</sub>)<sup>e</sup>2-h scanning<sup>f</sup>Measured in human parietal cortex

(%SUV = 150%). A blockade PET study with rimonabant showed that *in vivo* binding of [ $^{11}\text{C}$ ]OMAR in baboon brain is mediated by CB $_1$  receptors (Horti et al. 2006). Kinetic analysis suggests that [ $^{11}\text{C}$ ]OMAR rapidly reaches the steady state in the baboon brain before the end of the 90-min PET scanning session, which is a logistical advantage of [ $^{11}\text{C}$ ]OMAR versus other available PET radioligands (see below).

[ $^{11}\text{C}$ ]OMAR was used by the Johns Hopkins group under an Investigational New Drug Application, approved by the FDA, for quantification of cerebral CB $_1$  receptors in healthy subjects (Fig. 12.7) and schizophrenia patients (see Sect. 3.3) (Wong et al. 2010a). Within ten control male subjects, [ $^{11}\text{C}$ ]OMAR peaked at approximately 20 min postinjection (% SUV = 136–207% in the putamen, Table 12.4) and decreased gradually thereafter to reach a SUV between 80 and 117% at 90 min. The globus pallidus showed the highest  $V_T$  ( $1.47 \pm 0.25$ ), with a range of 0.95–1.82 in individual subjects. The cingulate cortex (mean  $1.23 \pm 0.16$ ) and the putamen (mean  $1.32 \pm 0.20$ ) also showed high binding. White matter, pons, and thalamus were the regions with the lowest accumulation of [ $^{11}\text{C}$ ]OMAR radioactivity. [ $^{11}\text{C}$ ]OMAR binding in healthy controls appeared to decline with increasing age in the globus pallidus and putamen, the regions with the highest values of  $V_T$ .

Given the relatively fast kinetics of [ $^{11}\text{C}$ ]OMAR, measurements of brain uptake can be quantified as  $V_T$  using two-tissue compartment model (2-TCM) with vascular component and multilinear analysis (MA1) modeling, which are the preferred analytical methods. Interestingly, human  $V_T$  was about 20% of that in rhesus monkey, which may have been due to low estimations of brain entry (human  $K_1$  parameter was approximately 25% of that in monkey) (Normandin et al. 2010a), and as other studies have shown human CB $_1$  receptor density to be a similar fraction compared



**Fig. 12.7** Trans-axial images of distribution volume ( $V_T$ ) of [ $^{11}\text{C}$ ]OMAR, mean of ten healthy subjects in a standard space (left panel), a standard MRI (right), and merged image (middle). Volumes of interest of selected structures in the standard space are shown on the  $V_T$  image and MRI. Regions are insula (I), putamen (P), globus pallidus (G), thalamus (T), and caudate nucleus (C) (reprinted from Wong et al. (2010a, b) with permission of Copyright Clearance Center)



to nonhuman primate. An initial modeling study in baboons tested if the simpler reference tissue model could be used for [ $^{11}\text{C}$ ]OMAR data analysis (Horti et al. 2006). When pons was used as a reference tissue the resulting values of BP fluctuated by 10–20%, further suggesting that it leads to low precision and is an unstable method to use for quantifying  $\text{CB}_1$  receptors. An alternative candidate for use as a reference tissue, white matter, did not appear to reach equilibrium in the course of the scan, and indeed had a small increase of radioactivity over the course of 90 min, eliminating it as a choice for analytic method.

Test-retest analysis of  $V_T$  in healthy human subjects demonstrated that [ $^{11}\text{C}$ ]OMAR has reasonably good precision with ~3% within-subject variability, ~9% intersubject variability, and ICC ~0.7 (Table 12.4). (Normandin et al. 2015) Analysis of potential modeling errors suggested that measurement error in the input function is the likely source for variability in  $V_T$ . Both peak brain uptake and fraction of parent tracer in plasma were found to be higher in men than women.  $V_T$  was significantly higher in women than men, presumably due to the faster plasma clearance in women than in men, and could be measured using 90 min of data (and as little as 60 min for some regions using the MA1 model) (Normandin et al. 2010b; Neumeister et al. 2013). Since brain uptake is higher and  $V_T$  is lower in men compared to women, it is recommended that  $V_T$  derived via metabolite-corrected arterial input function be used in patient population assessments with sex-matched controls, as brain uptake alone and sex differences could result in misleading or confounding results. The free fraction of [ $^{11}\text{C}$ ]OMAR is ~0.1–0.2% (Normandin et al. 2015; Ranganathan et al. 2016; D'Souza et al. 2016a).

[ $^{11}\text{C}$ ]OMAR demonstrated an age-related decline trending in nearly all regions and attaining statistical significance in the region with the highest  $V_T$  (globus pallidus) (Wong et al. 2010a; Neumeister et al. 2013).

**Table 12.4** *In vivo* properties of  $\text{CB}_1$  receptor radioligands in humans

Radioligand	[ $^{11}\text{C}$ ]OMAR	[ $^{18}\text{F}$ ]MK-9470	[ $^{18}\text{F}$ ]FMPEP- $d_2$	[ $^{11}\text{C}$ ]SD5024
Peak brain uptake (SUV)	~2	1.2	~4	~3
Retest variability	–	~7%	16%	–
Intersubject variability	–	16%	14%	–
$V_T$ (mL/cm $^3$ )	~1.5	~22	~23	~2.5–3.0
Retest variability	~3%	10–30%	14%	–
Intersubject variability	~9–15%	~55–66% (35% with FUR)	26%	22%
Effective dose ( $\mu\text{Sv}/\text{MBq}$ )	–	22.8	19.7	–
References	Wong et al. (2010a), Normandin et al. (2015, 2010b)	Burns et al. (2007), Van Laere et al. (2008a, b), Addy et al. (2008)	Terry et al. (2010a, b)	Tsujikawa et al. (2014)

Human PET studies with the CB<sub>1</sub> antagonist blocker AVE1625 demonstrated the utility of [<sup>11</sup>C]OMAR for successful occupancy evaluation of cannabinoid receptors by drugs in human brain (Wong et al. 2010b). Because of the short half-life of <sup>11</sup>C ( $t_{1/2} = \sim 20$  min), the baseline and drug challenge scans with [<sup>11</sup>C]OMAR can be done in the same subject in a single day, an advantage of [<sup>11</sup>C]OMAR over <sup>18</sup>F-labeled CB<sub>1</sub> receptor radioligands for drug development research.

The biodistribution and radiation dosimetry estimates for [<sup>11</sup>C]OMAR have not yet been reported.

### 12.2.2.2 [<sup>18</sup>F]MK-9470

Discovery of the selective CB<sub>1</sub> inverse agonist rimonabant by Sanofi-Aventis (Rinaldi-Carmona et al. 1994), available in the UK and the EU as anti-obesity drug in 2006–2009, stimulated the development of cannabinoid drugs by many pharmaceutical companies. As a result, a large number of rimonabant analogs and structurally different CB<sub>1</sub> inverse agonists have been developed. Merck disclosed a series of non-rimonabant-based CB<sub>1</sub> receptor inverse agonists (Lin et al. 2006). The best compound of the Merck series, taranabant, demonstrated a substantially higher affinity (0.4 nM) than that of rimonabant (Lin et al. 2008). Using taranabant as a lead compound Merck has developed MK-9470, a CB<sub>1</sub> receptor-selective fluoroalkyl analogue of taranabant with high CB<sub>1</sub> receptor-binding affinity, but also relatively high values of  $\log D_{7.4}$  and PSA (Burns et al. 2007) (Fig. 12.6, Table 12.3). Radiolabeling of [<sup>18</sup>F]MK-9470 was performed in two steps by preparation of [<sup>18</sup>F]fluoroethylbromide and [<sup>18</sup>F]fluoroethylation of the corresponding phenol with moderate radiochemical yield. [<sup>18</sup>F]MK-9470 can also be produced in a simplified, one-step synthesis using a GE TRACERlab FX<sub>FN</sub> with the tosylate precursor and [<sup>18</sup>F]fluoride in high yield for routine GMP production (Thomae et al. 2014).

*In vitro* autoradiography of [<sup>18</sup>F]MK-9470 in rhesus monkey brain slices (Burns et al. 2007) produced images with a high signal in cerebral cortex, cerebellum, caudate/putamen, globus pallidus, substantia nigra, and hippocampus, which is consistent with the reported localization of CB<sub>1</sub> receptors in the brain of various mammalian species. Additional *in vitro* studies showed that [<sup>18</sup>F]MK-9470 is not a P-gp substrate.

Preclinical studies using [<sup>18</sup>F]MK-9470 in rodents have provided additional information on the radioligand's characterization. [<sup>18</sup>F]MK-9470 is about 80% specific in wild-type compared to CB<sub>1</sub> receptor knockout mice. In rats, [<sup>18</sup>F]MK-9470 shows a distribution consistent with known CB<sub>1</sub> receptor distribution, a test-retest variability of <5%, and a 58% decrease of radioactivity following blockade by rimonabant (Miederer et al. 2013). Serial arterial blood sampling can permit kinetic modeling of both [<sup>18</sup>F]MK-9470 and radioactive metabolites in brain, which accounts for approximately 10–30% of radioactivity, and demonstrates that radiometabolites cause an overestimation of  $V_T$ . Measurements of  $V_T$  were better approximated after 10 h of scanning, which was independent of radiometabolites. Although it was concluded that compartmental modeling and measurements of  $V_T$  are superior to those of brain uptake alone, simplified measurements using SUV or mSUV correlated with  $V_T$  and may be preferred over the onerous arterial sampling and

extensive scan time required to measure  $V_T$  (Casteels et al. 2012). In a separate analysis, kinetic modeling with arterial blood sampling demonstrated a better fit with 2-TCM compared to 1-tissue compartment model (1-TCM), but due to micro-parameter poor identification, a coupled fitting approach improved parameter stability. However, radioactivity in brain from [ $^{18}\text{F}$ ]MK-9470 was observed to plateau from ~60 to 90 min and therefore an irreversible model was assessed and found to fit the data well (Miederer et al. 2018). [ $^{18}\text{F}$ ]MK-9470 uptake was found to increase with age when comparing adolescent and adult male rats, although this study was limited by its lack of kinetic modeling. Nevertheless, the main effect was replicated with [ $^3\text{H}$ ]CP-55,940 (a  $\text{CB}_1$ ,  $\text{CB}_2$  high-affinity agonist) *in vitro* binding, though with one-quarter the magnitude (Verdurand et al. 2011).

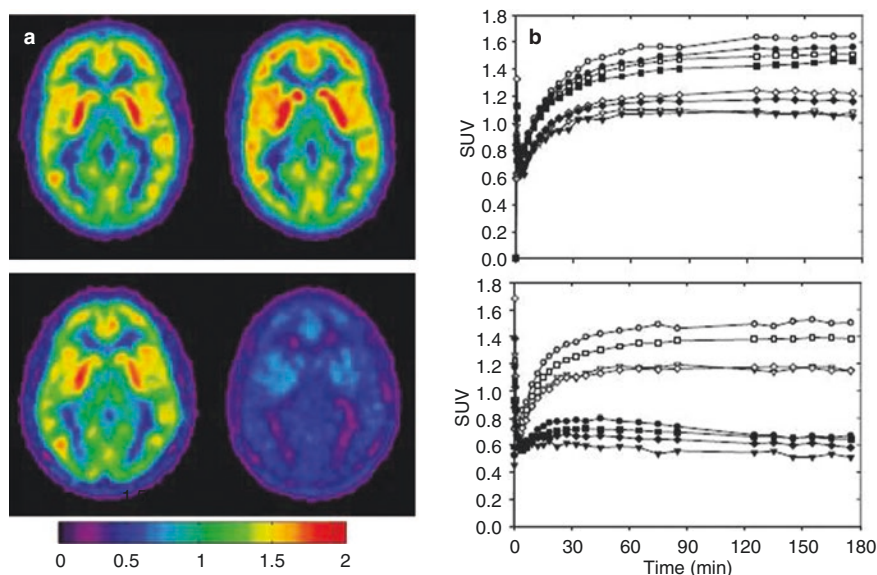
Additional rodent studies demonstrated that anesthetics affected brain uptake of [ $^{18}\text{F}$ ]MK-9470, and suggest an indirect action of  $\text{GABA}_A$  receptors on  $\text{CB}_1$  receptor binding (Casteels et al. 2010a). Following 2 weeks of treatment with anti-epileptic agents, valproic acid produced a significant increase in [ $^{18}\text{F}$ ]MK-9470 uptake while levetiracetam did not (Goffin et al. 2008). These results from rodents were not corrected for plasma concentrations of parent radioligand across conditions (*i.e.*, drug treatments) which might be expected to change with altered hepatic metabolism, and potentially confound independent measurements of brain uptake. In rats, *ex vivo* biodistribution studies were in good agreement with *in vivo* PET imaging data, with which organs known to have  $\text{CB}_1$  receptor expression and involvement in metabolism were grossly well delineated (Buchholz et al. 2017).

Baseline PET scans in the rhesus monkey showed gradual brain penetration and accumulation of [ $^{18}\text{F}$ ]MK-9470 in gray matter regions of the brain as expected for binding to  $\text{CB}_1$  receptors and consistent with autoradiography studies. The specificity of the *in vivo* binding was confirmed in a blockade study with MK-0364, a  $\text{CB}_1$ -selective inverse agonist.

After bolus injection of [ $^{18}\text{F}$ ]MK-9470 in human controls the radioactivity reached a plateau at ~120 min (Burns et al. 2007) (Fig. 12.8). The distribution of radioactivity corresponded to the known  $\text{CB}_1$  receptor distribution and regional concentrations remained nearly unchanged for 2–6 h postinjection. Despite the seemingly irreversible brain kinetics the excellent signal-to-noise characteristics of [ $^{18}\text{F}$ ]MK-9470 (Table 12.3) made it suitable for several human PET imaging studies (Burns et al. 2007; Gerard et al. 2010, 2011; Sanabria-Bohorquez et al. 2010; Van Laere et al. 2008a, b, c, 2009) that were performed by the imaging group from K.U. Leuven in collaboration with Merck.

A methoxy analogue of MK-9470, CB-119, was designed as a carbon-11 alternative for PET imaging (Hamill et al. 2009). While it showed nearly identical binding characteristics to [ $^{18}\text{F}$ ]MK-9470, it was not able to achieve equilibrium within the scanning time. [ $^{11}\text{C}$ ]CB-119 was therefore abandoned for further clinical development in lieu of [ $^{18}\text{F}$ ]MK-9470.

[ $^{18}\text{F}$ ]MK-9470 demonstrates many favorable properties for characterizing  $\text{CB}_1$  receptors with *in vivo* imaging including good brain uptake and low nonspecific uptake (Tables 12.3 and 12.4). However, its slow pharmacokinetics has proven to be a challenge for modeling accepted outcome measures such as  $V_T$ . In both monkeys



**Fig. 12.8**  $^{18}\text{F}$ -MK-9470 PET images of  $\text{CB}_1$  receptors were acquired in normal healthy male subjects before and after treatment with MK-3640 or placebo. (a) Representative transverse parametric SUV images at the level of the basal ganglia for two human subjects, acquired before (*left*) and 24 h after (*right*) treatment with either placebo (*upper*) or a 7.5 mg dose of MK-0364 (*lower*). (b) Graphs show the corresponding TACs. The open symbols denote baseline values, and the closed symbols denote the scan after treatment (reprinted from Burns et al. (2007) with permission of Copyright National Academy of Sciences, U.S.A., PNAS)

and humans, radioactivity increased in brain up to about 120 min after bolus injection with  $^{18}\text{F}$  MK-9470, and concentrations of radioactivity did not appear to decline, even though concentrations of the radioligand in arterial plasma decreased throughout the length of the scan (Burns et al. 2007). For these reasons, brain studies using  $^{18}\text{F}$  MK-9470 were analyzed using a pharmacokinetic method designed for irreversible PET radioligands (Koeppel et al. 1999) which simply uses the area under the curve (AUC) of radioactivity in brain after a period when measurements have stabilized (e.g., 120–180 min after injection), termed fractional uptake ratio (FUR). However, when challenged with another  $\text{CB}_1$  inverse agonist the radioactivity quickly washed out of brain, indicating that  $^{18}\text{F}$  MK-9470 undergoes reversible binding at the  $\text{CB}_1$  receptor, which runs counter to the rationale of using an irreversible model.

Indeed, when scanning is continued for 12 h a slow washout of the radioligand can be detected, and sufficient data is then available for analysis by 2-TCM (Sanabria-Bohorquez et al. 2010). However, even with this additional data only macroparameters from modeling could be distinguished, and the individual parameter that is proportional to receptor density ( $k_3$ ) could not be identified in any individual brain region. Using 2-TCM the test-retest variability of  $V_T$  was 10–30%, indicating that it had poor precision. After making the assumption that dissociation from the receptor

( $k_4$ ) is constant throughout the brain *in vivo*, multiple serial correlations were made linking  $V_T$  to irreversible measurements and to FUR. It was further assumed that in cases when plasma measurements are not statistically different, brain uptake measured as SUV or modified SUV (mSUV) is a sufficient outcome measurement, as they would correlate with FUR and be interpreted as CB<sub>1</sub> receptor availability. These models do not fully account for the full clearance of radioligand (*i.e.*, AUC extrapolated to infinity), and therefore do not correct the amount of radioligand in brain to that in plasma, and dissociation from receptors is not always constant *in vivo*. Thus, while the precision of these model outcomes can be assessed by test-retest studies, their accuracy remains uncertain, particularly in conditions where metabolic processes or changes in receptor density may affect radioligand concentration in plasma or local dissociation rate. Nevertheless, using this analytical method, [<sup>18</sup>F]MK-9470 has shown good precision (retest variability <7% within 24 h) and intersubject variability (16–35%) (Burns et al. 2007; Addy et al. 2008).

Gender-dependent increases of [<sup>18</sup>F]MK-9470 uptake with age were measured in women but not in men, particularly in the basal ganglia and hippocampus (Van Laere et al. 2008a). These results were obtained with and without partial volume correction of the brain images with presumably no loss of effect; partial volume effects have been minimal and likely insignificant for other studies using [<sup>18</sup>F]MK-9470. Metabolism of the radioligand was not dependent on age, although there were measured differences in metabolism between sexes. Similarly, sex-dependent differences were measured for the slow parameter of arterial input function (*i.e.*, elimination phase of clearance). However, when applied to pharmacokinetic modeling no significant difference was measured in arterial plasma input function using a three-parameter bi-exponential model. Larger intersubject variability (COV 15–20%) and region-dependent differences (up to 230%) were found than in the smaller sample size of the initial human study (Burns et al. 2007).

In part to assess this large variation within groups, additional information was obtained from subjects to find correlations between receptor availability and temperament traits, as have been shown for other receptor systems (Farde et al. 1997). Novelty-seeking traits were found to be inversely correlated to [<sup>18</sup>F]MK-9470 uptake throughout the brain, with the strongest correlation in the left amygdala (Van Laere et al. 2009). The trait subtypes of extravagance and disorderliness maintained significance even after strict limitations, while trait subtypes of exploratory excitability and impulsiveness, and trait types harm avoidance, reward dependence, and persistence, failed to attain significant correlations with CB<sub>1</sub> receptor availability. As novelty-seeking traits are associated with thrill seeking, addiction behavior (including overeating, and substance dependence), and impulsive aggression (Hiroi and Agatsuma 2005), their association with CB<sub>1</sub> receptor imaging, and particular location in the amygdala, is intriguing.

In humans, [<sup>18</sup>F]MK-9470 has shown to be useful in estimating receptor occupancy of other inverse agonists (Burns et al. 2007; Addy et al. 2008). Clinical trials of taranabant for weight loss showed that 10–40% receptor occupancy was sufficient to produce significant amounts of weight loss in a dose-dependent fashion, which is in agreement with estimates from preclinical rodent studies.

Whole-body imaging studies using [ $^{18}\text{F}$ ]MK-9470 demonstrated the radiation burden to be acceptable for completion of multiple brain studies per year ( $22.8 \pm 4.3 \mu\text{Sv}/\text{MBq}$ ), and similar to that of other commonly used PET tracers (Van Laere et al. 2008b). Brain uptake of the radioligand was about 3.2–4.9% of the total dose, and receptor occupancy was estimated to be well below 1%. [ $^{18}\text{F}$ ]MK-9470 undergoes predominantly hepatobiliary excretion, with significant radiation burden to the gallbladder which was the radiation dose-limiting organ. Although measured radioactivity in the periphery likely originates from a mixture of parent compound and its metabolites, an appreciable amount of radioactivity was measured in liver, small intestine, and red bone marrow, sites known to have  $\text{CB}_1$  receptors.

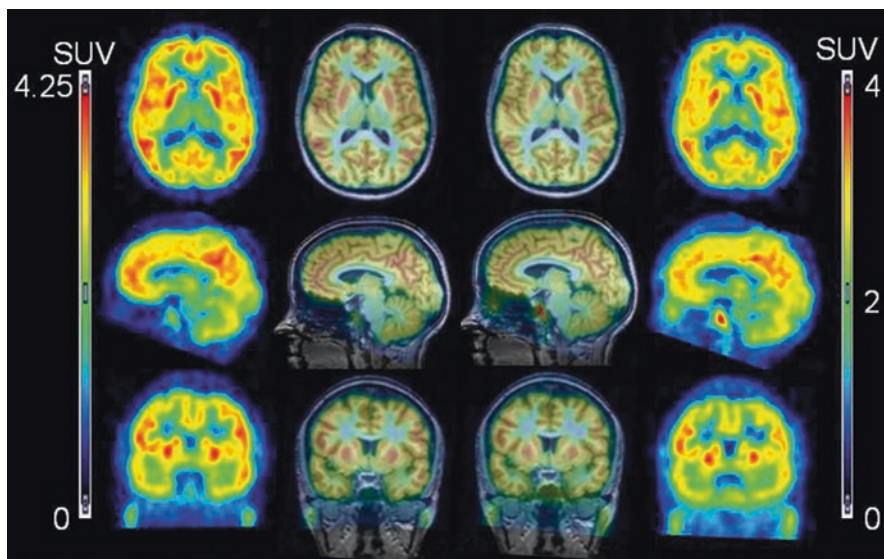
### 12.2.2.3 [ $^{11}\text{C}$ ]MePPEP and [ $^{18}\text{F}$ ]FMPEP- $d_2$

A collaborative group from Eli Lilly, NIH, and Karolinska University developed two structurally similar  $\text{CB}_1$  radioligands, [ $^{11}\text{C}$ ]MePPEP and [ $^{18}\text{F}$ ]FMPEP- $d_2$  (Fig. 12.6), for human PET studies (Terry et al. 2008; Yasuno et al. 2008; Donohue et al. 2008c; Suter et al. 2010). Both compounds are  $\text{CB}_1$  receptor inverse agonists with high  $\text{CB}_1$  receptor-binding affinity and selectivity (Table 12.3). MePPEP has about 10 times higher binding affinity compared to rimonabant, and has >700 times greater selectivity for  $\text{CB}_1$  than  $\text{CB}_2$ . The lipophilicity values of MePPEP and FMPEP- $d_2$  are high, but the PSAs are in the acceptable range for passive BBB transport and lower than those of OMAR and MK-9470 (Table 12.3).

Radiosynthesis of [ $^{11}\text{C}$ ]MePPEP is simple and can be performed by reaction of the corresponding desmethyl precursor with [ $^{11}\text{C}$ ]methyl iodide (Terry et al. 2009). In the monkey brain, [ $^{11}\text{C}$ ]MePPEP showed very high uptake with a good specific signal, reaching >89% in high-binding regions (prefrontal cortex) and ~75% in low-binding regions (pons, thalamus) (Table 12.3) (Yasuno et al. 2008). Preclinical studies in rodents further defined the *in vivo* characteristics of [ $^{11}\text{C}$ ]MePPEP (Terry et al. 2008). According to data from both genetic and pharmacologic rodent models, [ $^{11}\text{C}$ ]MePPEP is not a substrate for P-gp and has high specificity for the  $\text{CB}_1$  receptor. Approximately two-thirds of radioactivity after injection of [ $^{11}\text{C}$ ]MePPEP is specific for  $\text{CB}_1$  receptors in mouse and rat brain, and *ex vivo* mass spectrometry demonstrated >90% specific binding. Serial arterial blood sampling permitted kinetic modeling of [ $^{11}\text{C}$ ]MePPEP and measurements of  $\text{CB}_1$  receptors as  $V_T$  in rats within 70 min of scan time. A constant percentage (13%) of the radioactivity present in rat brain was in the form of metabolites, which may cause a stable, overestimated value of  $V_T$ . High doses of  $\text{CB}_1$  receptor agonists, including the endocannabinoid anandamide and the high-affinity synthetic cannabinoid CP 55,940, were unable to cause any displacement or blockade of [ $^{11}\text{C}$ ]MePPEP binding in brain; these findings were independently confirmed by *ex vivo* mass spectrometry. Despite the much higher affinity of MePPEP over lower affinity agonists (*e.g.*, endocannabinoids) for  $\text{CB}_1$  receptors, the severalfold higher concentration of agonists would be expected to compete and potentially displace or block the tracer dose of radioligand. The inability of [ $^{11}\text{C}$ ]MePPEP to be displaced by agonist *in vivo* suggests either non-overlapping binding sites of MePPEP and agonists or a large receptor reserve; that is, a small percentage of  $\text{CB}_1$  receptors are in the high-affinity, agonist-preferring,

state *in vivo*, and displacement of that small proportion might not be detected by PET. It was previously shown that *in vitro* agonism was achieved by CB<sub>1</sub> receptor occupancy of 7.5%, suggesting a >92% receptor reserve (Gifford et al. 1999).

An IND approval for the radioligand was obtained and [<sup>11</sup>C]MePPEP was initially studied in 17 human control subjects (Terry et al. 2009). The radioligand exhibited high human brain uptake in a regional distribution consistent with that of CB<sub>1</sub> receptors (Fig. 12.9), and slow washout from brain. [<sup>11</sup>C]MePPEP provided acceptable quantification of receptor density using  $V_T$  2-TCM as the outcome measure. However, the precision and accuracy of quantification of CB<sub>1</sub> receptor with [<sup>11</sup>C]MePPEP in human brain may have been diminished by slow brain kinetics and a very low fraction of free radioligand in plasma ( $f_p$ ) (Terry et al. 2009) that can be explained by the high lipophilicity of this compound (Table 12.3). Further studies determined that quantification was not limited by measurements from brain, but rather by measurements of radioligand in plasma (see Sect. 12.2.3). Despite the relatively low test-retest variability (8%) and low intersubject variability (16%) of brain uptake, when combined with concentrations of parent radioligand in arterial plasma to calculate  $V_T$ , the intersubject variability was much higher (>50%). Assuming that the slow washout of [<sup>11</sup>C]MePPEP from brain limited a sufficient amount of pharmacokinetic data from being collected, an increased scan time of 210 min demonstrated that measurements of  $V_T$  were affected not by the washout



**Fig. 12.9** [<sup>11</sup>C]MePPEP and [<sup>18</sup>F]FMPEP-*d*<sub>2</sub> in human brain. PET images from 40 to 80 min after injection of [<sup>11</sup>C]MePPEP were averaged (left column) and co-registered to the subject's MR images (middle left column). In the same individual, PET images from 30 to 60 min after injection of [<sup>18</sup>F]FMPEP-*d*<sub>2</sub> were averaged (right column) and co-registered to the subject's MR images (middle right column). Color bars on left and right correspond to SUV for [<sup>11</sup>C]MePPEP and [<sup>18</sup>F]FMPEP-*d*<sub>2</sub>, respectively

rate of [ $^{11}\text{C}$ ]MePPEP, but rather by the low concentration of [ $^{11}\text{C}$ ]MePPEP in plasma and the radioactive half-life of  $^{11}\text{C}$ . Thus, the relatively poor precision of the plasma measurements contributed to the higher within- and between-subject variabilities of  $V_T$  compared to those of brain uptake. Nevertheless, measurements of  $V_T$  within subjects were highly stable over time, and consistent values were observed 60 min after injection. The test-retest variability was acceptable ( $\sim 15\%$ ) and the intraclass correlation coefficient (ICC) of  $V_T$  (*i.e.*, the sensitivity of distinguishing variability between subjects from within subjects) was considered good to excellent (0.87), and better than that of brain uptake measured alone (0.77). Taken together, the stable measures of  $V_T$ , the high ICC of  $V_T$ , and the theoretically superior outcome measure of  $V_T$  compared to brain uptake suggest that  $V_T$  remains the more accurate index of  $\text{CB}_1$  receptor density, in spite of the poor precision of plasma measurements. In computer simulations in which  $\text{CB}_1$  receptor density was varied,  $V_T$  was much more sensitive than brain uptake alone in measuring increases, but not sensitive for decreases, of receptor density, and therefore might be expected to require fewer sample points (*i.e.*, study subjects) to detect significant differences between groups. Therefore, it was recommended using  $V_T$  as the primary outcome measure for [ $^{11}\text{C}$ ]MePPEP; however brain uptake could be used for within-subject studies (*e.g.*, receptor occupancy).

A replication test-retest study of [ $^{11}\text{C}$ ]MePPEP in 15 subjects examined multiple kinetic modeling methods (Riano Barros et al. 2014). The 1-TCM demonstrated comparable test-retest reliability ( $19.7 \pm 8.7\%$ ) and ICC ( $0.82 \pm 0.09$ ) with high pallidum-to-pons ratio (suggesting low bias), but the 2-TCM did not. This may be due in part to the difficulty with measuring radioactivity in later blood samples, with an inability to measure the parent fraction at 90 min, and therefore affecting the modeling of the input function. This in turn is likely due to injection of about  $\frac{1}{2}$  the amount of radioactivity compared to the prior studies ( $\sim 360$  MBq vs.  $\sim 650$  MBq in Terry 2009). All three spectral analyses (a “model free” approaching using an input function) demonstrated good test-retest reliability (13.5–15.5%) and ICC (0.76–0.83), though the ROI-based and voxel-based models demonstrated bias to overestimating  $V_T$  in high-binding regions, while the rank-shaping model restricted the range of  $V_T$  and underestimated those regions. While the input function free modified SUV (mSUV) method demonstrated satisfactory test-retest variability and ICC for most regions, it performed poorly for the low-binding region pons, and yielded the most regional heterogeneity of models tested. The simplified reference tissue model (with pons as reference) produced highly variable results across regions, with very poor test-retest and ICC outcomes, likely due to the presence of specific binding in pons.

Several modeling methods have been assessed in an effort to improve the parent fraction arterial plasma input function of [ $^{11}\text{C}$ ]MePPEP. Comparison of standard and convoluted (by incorporating injection time duration as a variable) exponential, Hill, and power models demonstrated improved modeling of the input function data using the convoluted Hill model compared to the standard exponential model, but this had negligible effect on  $V_T$  and its related micro- and macroparameters (Tonietto



et al. 2015). A follow-up analysis implemented an automated, user-independent pipeline on a combination of continuous (first 15 min) and discrete (up to 90 min) arterial plasma samples in comparison to the standard exponential approach. A small (3%), but statistically significant, reduction of weighted residual sum of squares for  $V_T$  suggested improvement of the input function model, though the effect on  $V_T$  itself was small ( $2 \pm 1\%$ ) (Tonietto et al. 2019). In an effort to more accurately quantify the later time points of parent compound in arterial plasma, LC-MS/MS was tested as a way to measure the nonradioactive MePPEP from the carrier of [ $^{11}\text{C}$ ]MePPEP in a single test-retest study. While the method was highly correlated with the standard radio-HPLC method and could calculate  $V_T$  with equal test-retest variability, LC-MS/MS determinations were consistently lower by 25–30%, demonstrating a promising option for input function measures when radioactivity might be difficult or impossible to detect (Victor et al. 2010).

In an effort to develop a radioligand with improved measurements of radioactivity for an extended time, several fluorinated analogues of [ $^{11}\text{C}$ ]MePPEP were developed and tested in monkeys, yielding [ $^{18}\text{F}$ ]FMPEP- $d_2$  as suitable for human studies (Terry et al. 2010a). Other fluorinated radioligands tested had either less specific binding, or showed higher amounts of radioactivity bound to bone, presumably due to free [ $^{18}\text{F}$ ]fluoride which was formed as a metabolite from the parent radioligand. Indeed, deuterium substitution on the fluoromethoxy radiolabel to inhibit defluorination resulted in about 1/3 less radioactivity uptake in bone than was observed after injection of the native hydrogen version.

The same precursor was used for the preparation of [ $^{18}\text{F}$ ]FMPEP- $d_2$  and [ $^{11}\text{C}$ ]MePPEP. The radiosynthesis of [ $^{18}\text{F}$ ]FMPEP- $d_2$  involves [ $^{18}\text{F}$ ]fluorination of dibromomethane- $d_2$ , purification of [ $^{18}\text{F}$ ]fluorobromomethane- $d_2$ , and [ $^{18}\text{F}$ ]fluoromethylation of the phenol precursor (Donohue et al. 2008c). In a rhesus monkey study, [ $^{18}\text{F}$ ]FMPEP- $d_2$  peaked in the brain at 20 min post-bolus injection (SUV 4.5–6.5) with gradual washout thereafter. The blocking experiment revealed that specific binding in the monkey brain was 80–90% of total tissue uptake. Radioactivity concentration in the mandible reached an SUV of approximately 2.0 by the end of the 3-h scan (Terry et al. 2009). Following injection with [ $^{18}\text{F}$ ]FMPEP- $d_2$  in mice, radioactivity is comprised of about 56% parent compound in plasma at 15 min, decreasing to about 19% by 240 min, and radioactivity is about 86% parent compound in cortex, with the remainder accounted by a single metabolite, and remains essentially unchanged from 60 to 240 min after injection. Similar to [ $^{11}\text{C}$ ]MePPEP, about 2/3 radioactivity was displaced by rimonabant following [ $^{18}\text{F}$ ]FMPEP- $d_2$  injection in mice (Takkinen et al. 2018).

After injection of [ $^{18}\text{F}$ ]FMPEP- $d_2$  in human, radioactivity peaked in putamen (SUV = 5) by 30 min post-bolus administration followed by slow washout (Terry et al. 2010a). The regional distribution of radioactivity matched the density of  $\text{CB}_1$  receptors in human brain (Fig. 12.9). The skull had substantial uptake of radioactivity that reached a peak SUV value of 4.5 by the end of 5 h of scanning. Pharmacokinetic modeling demonstrated that 60–120 min of human imaging gives the most accurate measurements of  $V_T$ , whereas a longer acquisition is likely vulnerable to

contamination of the brain with radiometabolites and/or spillover of radioactivity from the skull, which presumably outweighs any added accuracy from a more extended scan time.

In test-retest studies similar to those done for [ $^{11}\text{C}$ ]MePPEP, [ $^{18}\text{F}$ ]FMPEP- $d_2$  yielded similar between-subject variability for brain uptake; however [ $^{18}\text{F}$ ]FMPEP- $d_2$  demonstrated a much better test-retest variability of plasma measurements (16% vs. 58% for [ $^{11}\text{C}$ ]MePPEP), and acceptable test-retest variability of  $V_T$  (14%) (Table 12.4). This increased precision of plasma measurements led to a much reduced intersubject variability of  $V_T$  (26%), and, in conjunction with an extremely good ICC (0.89), suggests an improved accuracy of  $V_T$ . When changes of receptor density were simulated,  $V_T$  was shown to be more sensitive for detecting both increases and decreases of receptor density. The test-retest studies of both [ $^{11}\text{C}$ ]MePPEP and [ $^{18}\text{F}$ ]FMPEP- $d_2$  were conducted with at least 2-week intervals, strengthening the robustness of radioligands' precision. In conclusion, using [ $^{18}\text{F}$ ]FMPEP- $d_2$  to measure  $\text{CB}_1$  receptor density as  $V_T$  was determined to be more precise, accurate, and sensitive than [ $^{11}\text{C}$ ]MePPEP or using brain uptake alone as outcome measurement.

Two significant limitations of [ $^{18}\text{F}$ ]FMPEP- $d_2$  were identified in early human studies. First, values of  $V_T$  increased after 120 min. While the exact cause was not identified, such a trend is consistent with an accumulation of radioactive metabolites in brain. Simulations demonstrated that radioactivity originating from neighboring bone would account for <2% of the total amount. Fortunately,  $V_T$  was measured with consistent and stable values from 60 to 120 min after injection, and therefore limiting scan duration to 120 min should avoid confounding changes to the accuracy of  $V_T$ . Second, the plasma free fraction of both [ $^{11}\text{C}$ ]MePPEP and [ $^{18}\text{F}$ ]FMPEP- $d_2$  in humans is very low (approximately 0.05% and 0.63%, respectively). The test-retest variability of plasma free fraction is excellent; however, the intersubject variability was quite large. This is likely due in part to the nature of such low values, since small measured differences would be disproportionately large in percentage. Since correction of  $V_T$  with free fraction (which would be a more correct quantitation of radioligand binding in brain) added too much noise to the final outcome measurements, it was excluded from the calculation of  $V_T$ . Thus, it is recommended that potential changes of free fraction be accounted for in future studies, particularly those with pharmacological challenges, as a proportionately large amount of radioligand could be displaced from plasma proteins and made available to enter brain.

In an effort to reduce the burden related to arterial sampling, a population-based input function (PBIF) was compared to full arterial sampled input function (Zanotti-Fregonara et al. 2013). Using the Logan plot to determine  $V_T$  (which was equivalent to that determined by the 2-TCM with similar ICC of 0.88, but slightly higher test-retest variability of 16%), a PBIF function derived from 42 healthy individuals yielded nearly identical estimates of  $V_T$  compared to the individually measured input function (13.2 mL/cm $^3$  vs. 13.3 mL/cm $^3$ ;  $R^2 = 0.8765$ ,  $p < 0.001$ ). This method was able to replicate the main finding in a study of individuals with alcohol-use disorder, but underestimated global  $V_T$  from those with cannabis-use disorder at

baseline (see Sect. 12.3.1.1), suggesting that the method may be used within healthy populations, but should be avoided in the analysis of clinical groups.

The PBIF with Logan plot was utilized to explore sex differences of CB<sub>1</sub> receptor expression (Laurikainen et al. 2019).  $V_T$  of [<sup>18</sup>F]FMPEP-*d*<sub>2</sub> was consistently high in all regions (grouped bilaterally) in males compared to females, with occipital, parietal, and posterior cingulate cortex demonstrating the largest effect. The intersubject variability was similar within males and females; however females taking oral contraceptives were noted to have lower  $V_T$  than those not taking contraceptives or recent menopause. Neuropsychological testing revealed a negative correlation between  $V_T$  and visual-spatial working memory, with sex as a covariate, though there were no differences in test performance.

Whole-body imaging studies using [<sup>11</sup>C]MePPEP and [<sup>18</sup>F]FMPEP-*d*<sub>2</sub> demonstrated the radiation burden to be acceptable for both radioligands ( $4.6 \pm 0.3 \mu\text{Sv}/\text{MBq}$  and  $19.7 \pm 2.1 \mu\text{Sv}/\text{MBq}$ , respectively) for completion of multiple brain studies per year (Terry et al. 2010b). Brain uptake of both radioligands was about 7–8% of the total dose, and receptor occupancy is estimated to be well below 1% (0.3% for [<sup>11</sup>C]MePPEP, and 0.06% for [<sup>18</sup>F]FMPEP-*d*<sub>2</sub>). Both radioligands undergo hepatobiliary excretion, [<sup>11</sup>C]MePPEP exclusively, while [<sup>18</sup>F]FMPEP-*d*<sub>2</sub> undergoes also some urinary excretion. Similar to [<sup>18</sup>F]MK-9470, and as first reported for [<sup>11</sup>C]OMAR, an appreciable amount of radioactivity was measured in red bone marrow. To determine if radioactivity was concentrated in bone tissue rather than bone marrow, additional analysis showed that the pattern of accumulation in bone was consistent with areas rich in bone marrow, such as vertebrae, pelvis, and ribs. A comparison of bones rich in marrow to those devoid of marrow demonstrated that radioactivity was not largely due to accumulation in bone. In addition, studies using monkeys demonstrated that about 20% of the radioactivity in bone marrow was due to specific binding to CB<sub>1</sub> receptors.

#### 12.2.2.4 [<sup>11</sup>C]SD5024

A collaborative group from NIH and Karolinska University developed CB<sub>1</sub> radiotracer [<sup>11</sup>C]SD5024 (Fig. 12.6) (Donohue et al. 2008a; Tsujikawa et al. 2014). The tracer was evaluated *in vitro* and *in vivo* and compared with other CB<sub>1</sub> receptor ligands previously used in humans, *i.e.*, [<sup>11</sup>C]MePPEP, [<sup>11</sup>C]OMAR, [<sup>18</sup>F]MK-9470, and [<sup>18</sup>F]FMPEP-*d*<sub>2</sub>. The negative enantiomer of SD5024 exhibited intermediate binding affinity ( $K_i = 0.47 \text{ nM}$ ) between MePPEP, MK-9470, FMPEP-*d*<sub>2</sub> (0.1 nM), and OMAR (2 nM) (Table 12.3). The lipophilicity of SD5024 is lower than MePPEP and FMPEP-*d*<sub>2</sub> (Tsujikawa et al. 2014), but slightly greater than that of OMAR.

The radiosynthesis of SD5024 was performed via a palladium catalyst-assisted reaction of [<sup>11</sup>C]cyanide and corresponding Br precursor with radiochemical yield of 36% ( $n = 2$ ) (Donohue et al. 2008a), high radiochemical purity (>98%), and moderate specific activity of  $41 \pm 31 \text{ GBq}/\mu\text{mol}$  ( $n = 4$ ) (Tsujikawa et al. 2014). [<sup>11</sup>C]SD5024 demonstrates high uptake in nonhuman primate brain in a distribution consistent with CB<sub>1</sub> receptor density, peaking at ~30–60 min and decreasing thereafter, and can be displaced and blocked with the antagonists PipISB and rimonabant. The lower affinity positive enantiomer ( $K_i = 16.9 \text{ nM}$ ) demonstrated an immediate

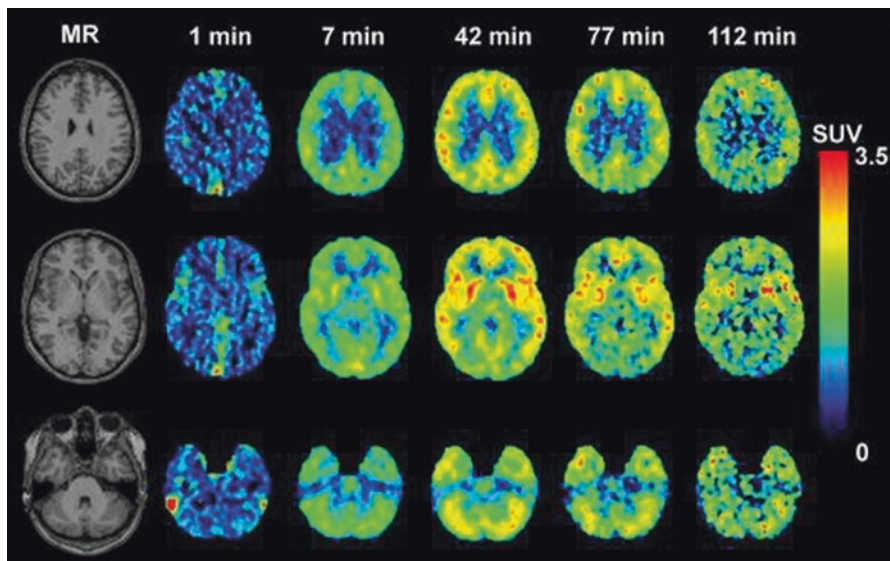
peak at 1.5 min after injection followed by rapid decline in all regions, consistent with a high proportion of nonspecific binding. Pharmacokinetic modeling with an arterial input function demonstrated best fit with the 1-TCM, and was well identified with standard error of 3.8% across brain regions, suggesting that the majority of binding is specific. Indeed, specific binding is ~80% in high-binding regions, and ~66% in low-binding regions, as determined by rimonabant preblockade.

[<sup>11</sup>C]SD5024 was applied for the comparison of rimonabant, taranabant, and three novel rimonabant-related CB<sub>1</sub> receptor antagonists, developed by AstraZeneca, in nonhuman primate (Hjorth et al. 2016). Using pons as a pseudoreference region (arterial sampling for kinetic analysis was not possible as those samples were needed for measuring test drug plasma concentration), [<sup>11</sup>C]SD5024 demonstrated a test-retest variability of 12%, and escalating displacement after 4–6 escalating doses of test drugs. Taranabant demonstrated greater potency, achieving similar receptor occupancy at lower doses and plasma concentrations than the other drugs tested. When these nonhuman receptor occupancy doses of taranabant and rimonabant were compared to clinically effective doses for weight loss, it was estimated that only 20–30% of occupancy might be needed to result in weight loss (slightly less or similar to that estimated using [<sup>18</sup>F]MK-9470, see Sect. 12.2.2.2).

In humans, [<sup>11</sup>C]SD5024 peaked at 1.5–3.0 SUV at about 40 min and decreased by 20% at the end of 120 min (Fig. 12.10). Notably, the fraction of parent [<sup>11</sup>C]SD5024 in plasma decreased slowly during the study, reaching ~50% by 75 min. Brain uptake was best fit by the 1-TCM with well-identified  $V_T$ , similar to nonhuman primate, and could be reliably measured with as little as 80 min of data. Subtracting activity from the vascular compartment in the modeling improved fitting of the data. The intersubject variability ( $n = 7$ ) was substantially lower than [<sup>11</sup>C]MePPEP at 22% (Table 12.4), likely due to its moderate binding affinity and slower metabolism, suggesting that [<sup>11</sup>C]SD5024 provides less noise (*i.e.*, greater precision) in its measurements. While a head-to-head comparison with test-retest and receptor blockade studies would better identify the preferred radioligand for human studies, the current data suggests that [<sup>11</sup>C]SD5024 might be one of the best available <sup>11</sup>C-labeled radioligand for CB<sub>1</sub> receptor neuroimaging, being superior to [<sup>11</sup>C]MePPEP (due to better plasma measures and intersubject variability) and equivalent or slightly better than [<sup>11</sup>C]OMAR (due to greater brain uptake and  $V_T$ ) (Tsujikawa et al. 2014).

### 12.2.3 Considerations in Imaging CB<sub>1</sub> Receptors and Its Interpretation

An advantage of PET over other imaging modalities is its quantitative ability, and quantitative analysis is achieved in large part through pharmacokinetic modeling. Pharmacokinetic modeling of CB<sub>1</sub> receptor PET radioligands in brain poses some particular challenges compared to more established compounds. These challenges include the high concentration of receptors throughout the brain, lack of a



**Fig. 12.10** [ $^{11}\text{C}$ ]SD5024 in human brain. Magnetic resonance anatomic images at the levels of centrum semiovale (top row), nucleus basalis (middle row), and pedunculus cerebellaris medius (bottom row) correspond to dynamic PET images obtained at 1, 7, 42, 77, and 112 min after [ $^{11}\text{C}$ ]SD5024 injection (from right to left). Each PET image represents standard uptake value (SUV) and is indicated in the SUV color scale on the right. Reprinted from Tsujikawa et al. (2014), Copyright 2014, with permission from Elsevier

receptor-devoid region, slow pharmacokinetics of the radioligands, low plasma concentrations of parent radioligand, and very high plasma protein binding. Careful consideration should be made when interpreting data from clinical PET studies, as radioligand properties and modeling methods may potentially confound results or their interpretation. To better appreciate such potential confounding variables, a brief review of modeling considerations is discussed here.

The noninvasive nature of *in vivo* PET imaging limits the ability to establish a “gold standard” method to validate endpoint measurements such as receptor density, as absolute measurements might require surgical excision followed by *in vitro* quantitation of receptors. Using a 2-TCM the radioligand can be conceptualized as being in a “non-displaceable compartment,” which represents radioligand non-specifically bound to proteins or lipids or is free in tissue, and a “specifically bound” compartment, which represents radioligand bound to a receptor and is potentially displaceable. Both compartments are detected as radioactivity by the PET camera, and so pharmacokinetic analysis with mathematical modeling serves to separate out these two compartments as discernible and quantitative measures. In doing so, estimates of receptor density and/or receptor occupancy can be calculated. Estimates of receptor density are approximated by  $V_T$  (Innis et al. 2007), which in turn can be

conceptualized as the ratio of areas under the curve of radioligand in tissue (*e.g.*, brain) divided by radioligand in arterial plasma. Over the course of the entire study the brain was exposed to the amount of radioligand available in plasma (*i.e.*, the input function), and the brain responded by taking up the radioligand. Unfortunately, we cannot always measure the complete area under the curve because of the short radioactive half-life or slow pharmacokinetics of the radioligand. Using mathematical modeling the curves can be extrapolated to infinity (or, in practice, described using rate constants) with only part of the entire curve of brain or plasma data; however enough of the curve shape must be observed for it to be successful, in particular the tissue uptake, peak, and initial decline.

The additional steps required to perform compartment modeling and calculate  $V_T$  yield theoretically more accurate outcome measures than simple methods, as it avoids potentially confounding variables. At equilibrium radioligand in brain changes proportionally by the amount of radioligand in arterial plasma, and is therefore susceptible to pharmacokinetic variables from the periphery that may affect the delivery of the radioligand to brain (*e.g.*, metabolism, distribution, excretion which could change radioligand concentration) (Liow et al. 2007). Additionally, compartment models account for radioligand in brain that is both receptor bound and unbound or free. Even for radioligands that have nearly 90% receptor specificity, not accounting for unbound radioligand may reduce the significance or obscure results.

The high density and widespread location of  $CB_1$  receptors are likely the cause for several hurdles when performing pharmacokinetic modeling of its radioligands (Hirvonen et al. 2010). First, extremely slow washout of  $CB_1$  inverse agonists may be due to a rapid process of reversible binding and rebinding to receptors, a phenomenon known as “synaptic barrier” (Frost and Wagner Jr. 1984). Similar kinetics suggestive of synaptic barrier has been seen with other radioligands targeting receptors found in high density (*e.g.*, iomazenil and benzodiazepine receptors) (Laruelle et al. 1994). Second, the widespread location of  $CB_1$  receptors in brain eliminates the use of reference tissue modeling methods, an alternative to compartment modeling using arterial plasma samples. Even areas of relatively low receptor density show up to 90% specific binding by some radioligands, and small changes in receptor binding in these regions would lead to potentially large systemic errors in outcome measurements. A potential advantage of high receptor density is that radioligands with lower specific activity could be used while maintaining the principle of tracer dosing. Tracer doses are needed to avoid inducing pharmacological effects, such as receptor internalization, and are generally accepted at 5–10% receptor occupancy or less;  $CB_1$  receptor imaging studies are well below this threshold, even in rodents, and typically maintain <1% receptor occupancy in humans.

The  $CB_1$  receptor radioligands discussed here demonstrate slow pharmacokinetics. Slow pharmacokinetics implies that ligands take a longer time to reach steady state when equilibrium has been achieved, and for reversible radioligands, after the identifiable peak in brain concentration has been achieved. This is a particular problem in PET imaging, as data acquisition time is typically limited by the radioactive half-life of the radioligand, as measured in brain and/or plasma. Thus, there is a

chance that insufficient data may be collected within the time allowed, and consequently an increased likelihood that measurement errors can be introduced. Test-retest studies are particularly useful to identify potential sources of error in kinetic modeling in measurements (brain uptake, plasma) and outcome measures (brain uptake,  $V_T$ ).

Additional potential causes of error are likely due to the high lipophilicity of  $CB_1$  receptor radioligands. First, most of the ligands have a relatively rapid distribution phase, followed by a very slow excretion phase (taken together as clearance). As a result, the excretion phase of plasma clearance typically has very low concentrations of parent radioligand, which are difficult to measure accurately. Second,  $CB_1$  receptor ligands tend to exhibit very high plasma protein binding, sometimes >99%, meaning that free ligand available to cross the BBB may be very small (*e.g.*, <1%). Thus, when comparing between groups or conditions, brain uptake or  $V_T$  should be corrected for differences in free fraction (not the fraction bound to proteins), as low absolute changes in free fraction can produce large proportional changes. For example, a change of bound fraction from 99.9% to 99.8%, which could be caused by competitive binding by a drug, would correspond to a change of free fraction from 0.1% to 0.2%, or a 100% increase; such a phenomenon may have been the cause of about 17% increased brain uptake of one radioligand when a co-administered drug caused about 10% increased free fraction (Yasuno et al. 2008). It is generally assumed that metabolites more polar than the parent tracer are unable to cross the BBB. However, ligands for cannabinoid receptors have a higher lipophilicity than more typical PET tracers, and their metabolites, though relatively more polar, may retain a lipophilicity sufficient to cross the BBB. Thus, additional consideration should be made to the radiometabolite profile of cannabinoid PET tracers to avoid confounding results from pharmacokinetic modeling. Finally, the high lipophilicity of  $CB_1$  receptor radioligands may permit passage not only through the BBB, but also through cell membranes and into intracellularly localized receptors where a substantial population may be present (*e.g.*, 85% in hippocampus) (Coutts et al. 2001). While it is assumed that the only receptors binding radioligand are membrane surface bound, the degree that differences in lipophilicity may confer for intracellular passage of these radioligands remains unknown, and could theoretically account for differences in brain uptake, washout, and estimation of receptor density of these PET tracers.

What does the data from  $CB_1$  receptor radioligand studies imply physiologically? It is generally assumed that differences or changes to receptor binding are due to changes of receptor density; however outcome measurements using PET alone cannot distinguish receptor density from receptor-binding affinity; indeed, until recently few examples of decreased binding affinity had been reported (Itoh et al. 2009). Changes observed in studies using these radioligands are likely to demonstrate changes in receptor density, whether due to direct downregulation or by indirect downregulation due to increased endocannabinoid tone (Romero et al. 1995; Sim-Selley and Martin 2002). As GPCRs, cannabinoid receptors are found in both a protein-coupled or high-affinity and agonist-preferring state and an uncoupled or low-affinity inverse agonist-preferring state. As demonstrated by *in vitro*

studies (Rinaldi-Carmona et al. 1996), high-affinity inverse agonists bind to all available receptors, and thus it is assumed that at tracer doses inverse agonist radioligand binding represents the entire population of available receptors. Regional changes relative to global  $V_T$  or brain uptake likely reflect local variations in  $CB_1$  receptor availability or density, as tracer uptake is unlikely affected by regional flow effects. However, when relative values of brain uptake are taken alone it could mask the absolute effect of receptor availability: For example, does a local increase of tracer uptake in brain with globally decreased uptake represent a true, and potentially abnormal, increase in receptor availability, or does it represent a local sparing or preservation of receptor expression with global reductions? Such relative changes may be open to wider interpretation when presented without more absolute measurements, as can be obtained with  $V_T$ . Nevertheless, relative changes can be sensitive measures, and may provide additional insight into regional alterations of  $CB_1$  receptor availability or expression (Van Laere et al. 2002).

Finally, as noted above, variabilities in CNS outcome measures of these radioligands have been described across populations of healthy individuals. In particular, sex differences (with males having lower outcomes with [ $^{11}C$ ]OMAR and greater outcomes with [ $^{18}F$ ]FMPEP- $d_2$  and [ $^{18}F$ ]MK-9470), BMI correlation (see Sect. 3.2), and  $CB_1$  receptor SNPs (see Sect. 3.1.1) suggest that care should be made when selecting comparison groups, with appropriate controls and adjustments in study design and analysis. The extent that medications may confound CNS imaging results remains incompletely known, with the exception of  $CB_1$  receptor antagonists/inverse agonists which can displace all radioligands described, and therefore may also require control in sample selection, or be appropriately adjusted for via pharmacokinetic modeling.

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## 12.3 $CB_1$ Receptor Imaging in Neuropsychiatric Disorders

### 12.3.1 Substance-Use Disorders

Multiple preclinical and clinical studies have suggested that  $CB_1$  antagonists could be used as treatments for multiple types of drug dependence (Le Foll and Goldberg 2005). Cannabis itself has the highest level of substance abuse, after alcohol and nicotine (Compton et al. 2005). Postmortem studies have shown that the density of  $CB_1$  mRNA-positive neurons is significantly lower in long-term cannabis users in the striatum, nucleus accumbens, and hippocampal regions (Villares 2007). Rimonabant blocks effects induced by  $\Delta^9$ -THC in humans and has been found to increase smoking cessation (Soyka et al. 2008; Cahill and Ussher 2007; Huestis et al. 2001).

While the synergism between alcohol and cannabinoid agonists (in various forms) creates undesirable side effects when the two are consumed together, the administration of  $CB_1$  antagonists leads to a significant reduction in alcohol consumption and cannabinoid antagonists also mitigate alcohol-withdrawal symptoms (Kleczkowska et al. 2016). Several preclinical studies have provided evidence that



CB<sub>1</sub> receptors significantly contribute to the behavioral properties of alcohol and that the chronic consumption alters CB<sub>1</sub> receptor expression in the brain nuclei associated with addiction pathways. Additionally, recent studies have further established the role of the ECS in the development of fetal alcohol spectrum disorders (FASD) (Basavarajappa et al. 2019), yet few trials have investigated cannabinergic medications for alcohol-use disorder. Although animal studies suggested that rimonabant would be a useful agent for treating alcohol dependence, the results in clinical trials were negative (Soyka et al. 2008; George et al. 2010). In animal models using both pharmacologic and genetic knockout methods, blocking or eliminating CB<sub>1</sub> receptor activity decreases the rewarding effects of opiates and alcohol, and reduces cocaine-seeking behavior (Le Foll and Goldberg 2005).

The ECS is central in the brain's reward circuitry, and interacts with the opioid and dopamine system circuitry and thus plays a role in other addictions (Nowak et al. 2006; Hungund and Basavarajappa 2004; Moranta et al. 2006; Economidou et al. 2006; Gonzalez et al. 2004; Caille et al. 2007). In both *CNR1* and *FAAH*, human genetic polymorphisms have been associated with increased risk of alcohol or drug use (Zhang et al. 2004; Hutchison et al. 2008; Sipe et al. 2002). Addiction remains a major public health concern and current evidence suggests that pharmacotherapies containing Δ<sup>9</sup>-THC, such as dronabinol and nabiximols, could still be effective treatments for a variety of disorders. Development of novel medications such as fatty acid amide hydrolase (FAAH) inhibitors and neutral CB<sub>1</sub> receptor antagonists promises to extend the range of available interventions (Sloan et al. 2017).

### 12.3.1.1 Cannabis-Use Disorder

In cannabis users, [<sup>18</sup>F]MK-9470 binding is ~12% lower in cortical, but not most subcortical, brain regions compared to non-cannabis user controls, about 5 days after last use of cannabis reaching significance in temporal lobe, anterior and posterior cortex, and nucleus accumbens (Ceccarini et al. 2015). Binding was not correlated with any clinical characteristics including amount of cannabis consumption, cannabinoid metabolite concentration, duration or initial age of cannabis use, or status of tobacco use.

[<sup>18</sup>F]FMPEP-*d*<sub>2</sub> also demonstrated a regionally selective downregulation of CB<sub>1</sub> receptors in chronic cannabis smokers, and also showed that it was reversible (Hirvonen et al. 2012a). Compared to healthy controls,  $V_T$  of [<sup>18</sup>F]FMPEP-*d*<sub>2</sub> in chronic daily cannabis smokers was ~20% lower in all regions of neocortex, but not subcortex, including basal ganglia a day after last use of cannabis (Fig. 12.11). Years of smoking correlated negatively with  $V_T$  (*i.e.*, more years of smoking correlated with smaller  $V_T$ ), as  $V_T$  of [<sup>18</sup>F]FMPEP-*d*<sub>2</sub> was independent of age in healthy males. As there was no significant difference between groups for  $V_T$  in cerebellum, this tissue was used as a reference tissue for additional analysis; it should be noted that this was performed only after the lack of difference for  $V_T$  was established between groups using the standard method correcting for radioligand in plasma, and may not be directly applicable to other between-group analyses. Following approximately 1 month of cannabis abstinence  $V_T$  increased in all affected regions except

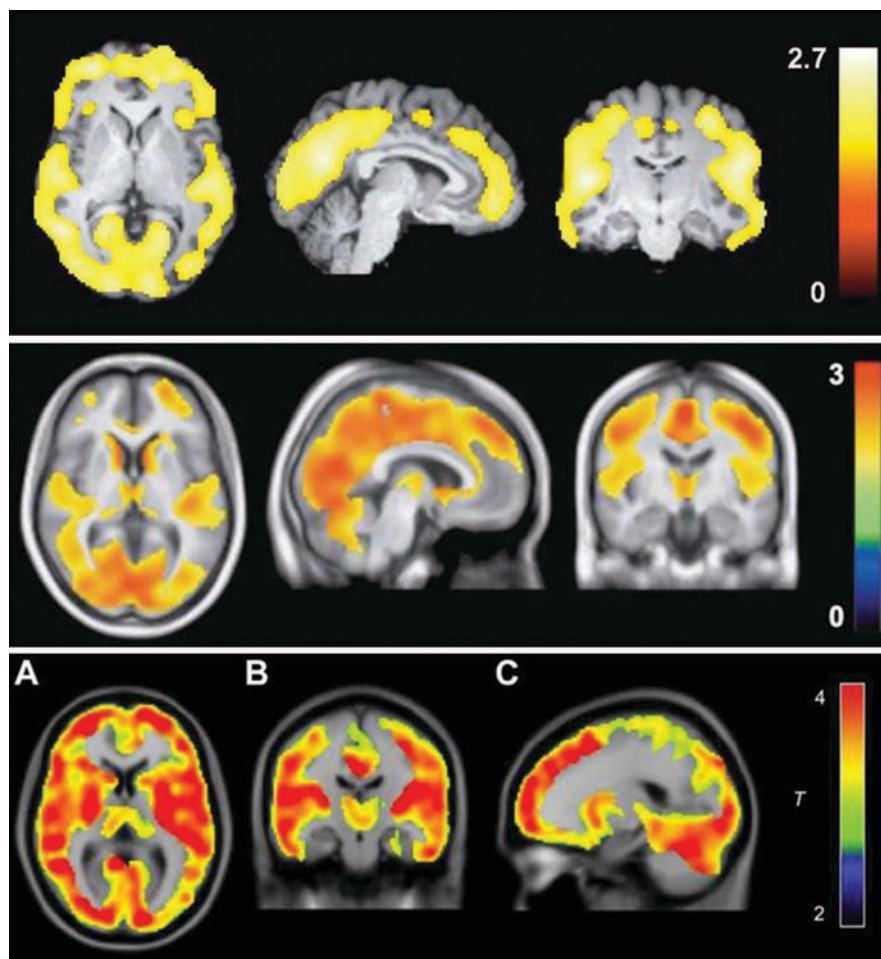
hippocampus, demonstrating that the downregulation of CB<sub>1</sub> receptors was reversible. In addition,  $V_T$  correlated negatively with BMI; therefore BMI was used as an adjustment variable in all findings. BMI or cannabis use did not correlate with free fraction of the radioligand, a potential confounding variable of the study. Neither the initial  $V_T$  nor the amount of recovery of  $V_T$  after 4-week abstinence correlated with the age of first use, amount of current daily use, or any clinical rating scales measuring withdrawal or craving. Tobacco smoking did not affect the results when included as a confounding variable in cannabis abusers (see below Sect. 12.3.1.3 for additional review). Interestingly, three control subjects of Indian descent had significantly lower  $V_T$ , but there was no difference for subjects of African or European descent, compared to the rest of the control group.

[<sup>11</sup>C]OMAR was used to examine cannabis users in a similar study; however in this instance participants used less cannabis daily and were scanned three times: at baseline while not intoxicated or in withdrawal from cannabis, after 2 days, and after 28 days of monitored abstinence (D'Souza et al. 2016a). This paradigm permitted detection of the rapid time course by which CB<sub>1</sub> receptors reverse, as the 15% lower  $V_T$  in cannabis users compared to controls at baseline were essentially reversed within 2 days and remained nonsignificantly through 28 days. In contrast to that seen using [<sup>18</sup>F]FMPEP-*d*<sub>2</sub>, [<sup>11</sup>C]OMAR binding was globally lower in cannabis users compared to controls, with the exception of thalamus and cerebellum, and there were no correlations with CB<sub>1</sub> receptor availability and cannabis use patterns. [<sup>11</sup>C]OMAR binding was negatively correlated to cannabis withdrawal symptoms on day 2 (e.g., more severe withdrawal symptoms with lower [<sup>11</sup>C]OMAR  $V_T$ ).

### 12.3.1.2 Alcohol-Use Disorder

In alcohol-dependent patients, [<sup>18</sup>F]FMPEP-*d*<sub>2</sub> demonstrated reduced CB<sub>1</sub> receptor binding that did not recover after 2–4 weeks of abstinence (Fig. 12.11) (Hirvonen et al. 2013). Compared to healthy controls ( $n = 19$ ), alcohol-dependent patients ( $n = 18$ ) had approximately 20–30% lower  $V_T$  adjusted for BMI across all brain regions when scanned during the first week of alcohol cessation. Years of alcohol use correlated negatively with  $V_T$ , which was not confounded by age. Differences between controls and alcohol-dependent patients could not be accounted for by clinical rating scales for alcohol use or withdrawal, or by tobacco use, and remained significant after partial volume correction. Repeat scans 2–4 weeks after abstinence did not result in any significant change in  $V_T$ . In both groups, those individuals carrying the *rs2023239 C* allele of *CNRI* had higher  $V_T$  (31% higher in patients, 19% in controls), although no statistical differences were detected between carriers and noncarriers based on demographic or clinical variables.

In contrast,  $V_T$  of [<sup>11</sup>C]OMAR was approximately 20% higher in alcohol-dependent patients ( $n = 8$ ) compared to healthy controls ( $n = 8$ ) when measured 4 weeks after abstinence. Anterior and posterior cingulate and orbitofrontal cortices, hippocampus, insula, and putamen were chosen *a priori* for their involvement in alcohol dependence, and analysis of other brain regions did not demonstrate any significant between-group differences. These findings were neither affected by age,



**Fig. 12.11** Statistical parametric mapping (SPM) of [ $^{18}\text{F}$ ]FMPEP- $d_2$  in three different substance-use disorders. Top: Chronic daily cannabis smokers ( $n = 30$ ) have lower  $V_T$  than control subjects ( $n = 28$ ) at baseline as a large single cluster that includes only cortical regions, and cluster-level corrected  $p$ -value of 0.043 (reprinted from Hirvonen et al. (2012b) with permission of Nature Publishing Group). Middle: Patients with alcohol dependence ( $n = 18$ ) have lower  $V_T$  than control subjects ( $n = 19$ ) at baseline as a large single cluster, with cluster-level family-wise error-corrected  $P$ -value of 0.038 (reprinted from Hirvonen et al. (2013) with permission of Nature Publishing Group). Bottom: Male tobacco smokers ( $n = 18$ ) have lower  $V_T$  than nonsmokers ( $n = 28$ ) as a large single cluster, with cluster-level family-wise error-corrected  $p$ -value of 0.001 (reprinted from Biological Psychiatry, Hirvonen et al. (2018) Copyright (2018) with permission from Elsevier). For each, corresponding color bar represents  $t$  value in each voxel within the significant cluster

body mass index (BMI), or use of tobacco, nor did they correlate with any alcohol-related or other clinical measures (Neumeister et al. 2012).

The effects of both acute and chronic alcohol exposure on  $\text{CB}_1$  receptor availability were examined with [ $^{18}\text{F}$ ]MK-9470 in rats. Acute alcohol exposure resulted

in ~7% increase of CB<sub>1</sub> receptor binding in nucleus accumbens and it highly correlated with an approximate doubling of anandamide concentration (measured by microdialysis), while chronic alcohol exposure resulted in ~5% decrease of CB<sub>1</sub> receptor binding in right hippocampus and striatum, and insular and primary sensory cortices, which correlated with the amount of ethanol consumed, and it was reversible in hippocampus and striatum after 7–14-day abstinence. Anandamide was no difference in nucleus accumbens after chronic alcohol exposure or abstinence (Ceccarini et al. 2013a). [<sup>18</sup>F]MK-9470 was used to study both acute and chronic effects of alcohol on CB<sub>1</sub> receptor availability in humans, yielding bidirectional results. In social drinkers ( $n = 20$ ), intravenous ethanol caused ~16% increase of [<sup>18</sup>F]MK-9470 binding, while alcohol-use disorder patients had 16% lower ( $n = 26$ ) [<sup>18</sup>F]MK-9470 binding immediately after chronic heavy drinking compared to nondrinking healthy controls ( $n = 17$ ), which remained lower (17%,  $n = 19$ ) after 1-month abstinence from alcohol (Ceccarini et al. 2014). In social drinkers, the change in [<sup>18</sup>F]MK-9470 binding was less in those with more previous drinking, and correlated with some subjective effects of alcohol intoxication. There were no correlations with [<sup>18</sup>F]MK-9470 binding to clinical characteristics or patterns of alcohol, tobacco, or caffeine use.

### 12.3.1.3 Tobacco Use and Nicotine Exposure

In a study examining the impact of nicotine on receptor binding, 2 weeks of nicotine exposure did not produce significant change in [<sup>18</sup>F]MK-9470 uptake in rat brain (Gerard et al. 2010).

Similar to that seen in cannabis- and alcohol-using subjects, [<sup>18</sup>F]FMPEP-*d*<sub>2</sub> demonstrated ~20% lower  $V_T$  in male tobacco smokers ( $n = 18$ ) compared to non-smoker controls ( $n = 28$ ) (Fig. 12.11) (Hirvonen et al. 2018). Decreased CB<sub>1</sub> receptor availability was observed throughout the entire brain, persisted before and after accounting for corrections of BMI and *rs2023239* genotype, and could not be accounted for by change in plasma free fraction of the radioligand. Characteristics of smoking (cigarettes per day, age of onset, years smoked, Fagerström Test for Nicotine Dependence [FTND]) were not correlated with whole-brain  $V_T$ . In an exploratory re-examination of studies using [<sup>18</sup>F]FMPEP-*d*<sub>2</sub> in cannabis- and alcohol-using subjects (see above), the prior results were not affected by tobacco use; that is, CB<sub>1</sub> receptor downregulation resulting from cannabis or alcohol use was not furthered by tobacco use.

In a study using [<sup>11</sup>C]OMAR examining patients with schizophrenia (see below, Sect. 12.3.3), tobacco smokers with schizophrenia had greater  $V_T$  than nonsmoking patients (D'Souza and Ranganathan 2015). While both groups were lower than non-smoking controls (no tobacco smoking controls were included in the study, and therefore conclusions about tobacco smoking alone cannot be drawn with regard to [<sup>11</sup>C]OMAR binding), smokers with schizophrenia were not significantly different from controls. CB<sub>1</sub> receptor availability did not correlate with FTND or other smoking-related characteristics; however more smokers were on antipsychotic medications, which increased [<sup>11</sup>C]OMAR binding in that group, and may have confounded this result.

### 12.3.2 Eating and Metabolic Disorders

There is overwhelming evidence that the ECS plays a central role in the control of food intake and in eating disorders (Gerard et al. 2011; Di Marzo et al. 2001; Cooper 2004; Maresz et al. 2007; Morton et al. 2006). The administration of cannabinoid CB<sub>1</sub> receptor agonists can promote body weight gain (Cota et al. 2003; Vickers and Kennett 2005), anandamide causes overeating in rats because of its ability to activate cannabinoid receptors (Gaetani et al. 2008), and CB<sub>1</sub> receptor agonists have been used as appetite stimulants in AIDS and cancer patients (Croxford 2003). Conversely, antagonists or inverse agonists are associated with long-term weight loss (Vickers and Kennett 2005; Smith and Fathi 2005; Hsiao et al. 2015). Mechanism-of-action studies have shown that the effect of inverse agonists is likely caused by a combination of central effects (e.g., reduction in food intake) (Nogueiras et al. 2008) and peripheral effects (increased resting energy expenditure and fat oxidation) (Addy et al. 2008; Jbilo et al. 2005). In addition, different alleles of the gene encoding the CB<sub>1</sub> receptor (*CNRI*) and FAAH (*FAAH*) have been associated with anorexia nervosa (Siegfried et al. 2004; Monteleone et al. 2009) and obesity (Benzinou et al. 2008; Sipe et al. 2005). An analysis pooling data from two studies found a negative association between [<sup>18</sup>F]MK-9470 CB<sub>1</sub> binding in regions associated with homeostasis (e.g., hypothalamus, pons, medulla) and reward (e.g., nucleus accumbens, amygdala, insula, orbitofrontal cortex, striatum) and body mass index (BMI) (Ceccarini et al. 2016), replicating an association observed in some but not all CB<sub>1</sub> PET studies (Sloan et al. 2019). This association raises the possibility that the fluctuations in BMI associated with disordered eating may explain the observed alterations in radiotracer binding, suggesting that altered CB<sub>1</sub> availability may be a result of disordered eating rather than a causal factor.

The highly selective CB<sub>1</sub> inverse agonist SR141716 (rimonabant) was developed by Sanofi-Aventis in the European Union to treat obesity (Rinaldi-Carmona et al. 1994), and has been found to improve lipid and glucose metabolism in obese humans. However, shortly after its introduction to the market, the safety of rimonabant was called into question due to the side effects that included severe depression and suicidal thoughts (Kelly et al. 2011; Lee et al. 2009; Le Foll et al. 2009). After the collection of additional clinical data, and preclinical results with taranabant showing similar side effects, rimonabant was withdrawn from the market and further development of centrally acting CB<sub>1</sub> inverse agonists was largely abandoned (Le Foll et al. 2009). Due to the structural dissimilarity of taranabant from rimonabant, and preclinical data from animal studies, the adverse effects are thought to be mechanism-dependent, and not merely compound-specific, effects. As previously discussed, the brain receptor occupancy necessary for clinical weight loss with rimonabant and taranabant is estimated at 20–30% (see Sect. 12.2.2.4).

Following the failures of rimonabant and taranabant due to anxiety, depression, and suicidal thoughts, a second generation of CB<sub>1</sub> receptor antagonists that are peripherally restricted have been developed that might confer benefit on metabolic profile but without CNS penetration and psychiatric side effects. One such compound, TM38837, was assessed for CNS penetration in nonhuman primates using

[<sup>11</sup>C]MePPEP with brain uptake as the within-subject outcome measure, as previously recommended, for calculating receptor occupancy by the Lassen plot (Takano et al. 2014). Compared to rimonabant, TM38837 required about 10 times higher dose and 100 times higher plasma concentration to achieve a similar CNS CB<sub>1</sub> receptor occupancy, and it was concluded that the clinically relevant doses of TM38837 would have low CB<sub>1</sub> receptor occupancy. This study demonstrated the potential of using CNS-penetrant CB<sub>1</sub> receptor radioligands in the assessment of peripherally restricted drugs, by screening for a lack of meaningful receptor occupancy.

There is evidence that people with anorexia nervosa do not exhibit normal endocannabinoid responses to food consumption (Monteleone et al. 2015). In a rat model of anorexia, [<sup>18</sup>F]MK-9470 cortical and subcortical binding was substantially higher in male (67%) and female (>51%) rats following weight loss by food restriction and exercise, which normalized after weight gain (Casteels et al. 2014). In an assessment of anorexia nervosa patients, [<sup>18</sup>F]MK-9470 demonstrated ~25% globally increased uptake in brain compared to healthy controls (Gerard et al. 2011). The amount of uptake was also significantly more than bulimia nervosa patients, though bulimia nervosa patients were not statistically different from controls. Uptake was relatively increased in insula for anorexia (right side) and bulimia (left side) patients, and in areas of frontotemporal cortex in anorexia patients. In addition, scores rating drive for thinness in anorexia patients correlated with [<sup>18</sup>F]MK-9470 uptake in the superior temporal region. Increased uptake was interpreted as increased CB<sub>1</sub> receptor availability, and therefore hypoactive endocannabinoid transmission. Similarly, [<sup>18</sup>F]MK-9470 was about 24% greater in multiple regions throughout brain in those with functional dyspepsia compared to BMI-matched controls (Ly et al. 2015). While no correlations could be associated with symptom severity or clinical characteristics, the regions identified are implicated in visceral nociception, homeostasis, and reward, and remained stable over the course of about 3 years.

CB<sub>1</sub> receptor expression has been identified in metabolically active brown adipose tissue (BAT), and both [<sup>11</sup>C]MePPEP/[<sup>18</sup>F]FMPEP-*d*<sub>2</sub> and [<sup>18</sup>F]MK-9470 binding can be observed in regions of human BAT distribution (Van Laere et al. 2008b; Terry et al. 2010b). In rats, CB<sub>1</sub> receptors were found to colocalize with the BAT marker uncoupling protein-1; were selectively expressed in BAT but not white adipose tissue; could be imaged with [<sup>18</sup>F]FMPEP-*d*<sub>2</sub> in BAT, with kinetic analysis conducted using an image-derived input function; and increased following β<sub>3</sub> agonist treatment to stimulate BAT activity (Eriksson et al. 2015; Lahesmaa et al. 2018). In follow-up, cold exposure increased [<sup>18</sup>F]FMPEP-*d*<sub>2</sub> binding in human BAT, and overweight subjects were found to have lower CB<sub>1</sub> receptor availability in BAT compared to lean subjects both by imaging and biopsy histochemical analysis (Lahesmaa et al. 2018). In brain, overweight subjects had about 23% lower CB<sub>1</sub> receptor availability compared to their lean counterparts, with a correlation during cold condition between midbrain and BAT, suggesting sympathetic nervous and endocannabinoid system-mediated connection during thermoregulation.

Furthering the investigation of peripheral CB<sub>1</sub> receptor-rich targets in obesity, [<sup>11</sup>C]OMAR was assessed in heart muscle in obese mice and humans (Valenta et al.

2018). Uptake of [ $^{11}\text{C}$ ]OMAR was greater in heart of genetically modified obese mice compared to wild type, which was reflected in high  $\text{CB}_1$  expression detected by PCR and RNA *in situ* hybridization. [ $^{11}\text{C}$ ]OMAR had significantly greater retention in hearts of obese vs. health weight controls.

### 12.3.3 Schizophrenia

Cannabis has long been associated with schizophrenia (Ortiz-Medina et al. 2018), and the evidence between cannabis use and development of a psychotic disorder has been mounting (D'Souza et al. 2016b). Large epidemiological studies have suggested an overall increased and potentially dose-dependent risk for developing schizophrenia after cannabis use (Fernandez-Espejo et al. 2009; Malone et al. 2010; Marconi et al. 2016), and a genetic susceptibility has been found in subjects who develop psychotic symptoms after cannabis use (Caspi et al. 2005; Ho et al. 2011). Additional genetic studies have reported an association between the  $\text{CB}_1$  receptor-encoding gene, *CNRI*, and schizophrenia, especially the disorganized subtype (Ujike et al. 2002); individuals with a 9-repeat allele of an AAT-repeat polymorphism of the gene have a 2.3-fold higher susceptibility to schizophrenia (Leroy et al. 2001). A meta-analysis has demonstrated greater CSF and blood anandamide, and blood  $\text{CB}_1$  receptor expression in those with schizophrenia compared to controls was greater during earlier stage of the illness and before antipsychotic treatment, was inversely associated with symptom severity, and normalized following successful treatment (Minichino et al. 2019).

In schizophrenia, hyperactivity of dopamine neurotransmission in the mesencephalic projections to the nucleus accumbens appears to contribute to the psychotic (positive) symptoms, and hypodopaminergic and hypoglutamatergic transmission in the prefrontal cortex to the negative symptoms (van der Stelt and Di Marzo 2003). These transmitters are under  $\text{CB}_1$  receptor signaling control, and hyperactivity of the central ECS has been shown to be involved in the pathogenesis of schizophrenia and its symptoms (Vinod and Hungund 2006a; Laviolette and Grace 2006; Ujike and Morita 2004; Zavitsanou et al. 2004; Dean et al. 2001). Since increasing evidence from the anatomical, pharmacological, and behavioral studies points to functional interactions between the cannabinoid and dopamine receptor systems (Laviolette and Grace 2006), this could be a key to the development of innovative therapy for psychotic illnesses (Ujike and Morita 2004).

Interestingly, there is growing evidence that cannabidiol acts as a negative allosteric modulator of the  $\text{CB}_1$  receptor, and reduces psychotic symptoms in patients with schizophrenia, both as a monotherapy (Leweke et al. 2012) and as an adjunctive treatment (McGuire et al. 2018). Conversely, blockade of  $\text{CB}_1$  receptors has been found to be ineffective at reducing psychotic symptoms in schizophrenia and schizoaffective disorder (Meltzer et al. 2004), suggesting the possibility that cannabidiol may exert its antipsychotic effects through other molecular targets.

Studies measuring  $\text{CB}_1$  receptor density in postmortem tissue have been inconsistent. Significantly greater  $\text{CB}_1$  receptor binding in the frontal and cingulate cortex

was found in subjects with schizophrenia (Dean et al. 2001), and has also been reported to be increased by up to 25% in the posterior cingulum (Newell et al. 2006). In contrast, other studies have shown a decrease in CB<sub>1</sub> receptor mRNA and protein by nearly 15% and 12%, respectively (Eggen et al. 2008). While discrepancies could be attributed to methodology (e.g., autoradiography generally reflecting higher CB<sub>1</sub> receptor expression in schizophrenia vs. vice versa for immunodetection), receptor trafficking and localization, and/or allosteric modulation changing affinity of ligands at the allosteric site (Price et al. 2005), confounding factors such as lifetime antipsychotic exposure which has been shown to affect CB<sub>1</sub> receptor expression (Secher et al. 2010) might also contribute. Therefore, *in vivo* imaging studies would be of great assistance in distinguishing the changes in receptor expression due to disease, and control for effects of medication.

Prior to imaging, [<sup>3</sup>H]MePPEP was assessed in postmortem human brain tissue to estimate the difference in CB<sub>1</sub> receptor binding in those with schizophrenia ( $n = 47$ ) versus controls ( $n = 43$ ), and the number of participants to enroll in subsequent PET studies (Jenko et al. 2012). Specific binding was about 20% higher in dorsolateral prefrontal cortex from those with schizophrenia compared to controls; this result was significant ( $p = 0.050$ ) when controlling numerous categorical and continuous variables (e.g., gender, age of onset, duration of illness, age of death, antipsychotic exposure, alcohol and substance abuse), and more so when the variables, all of which were not statistically significant, were omitted ( $p = 0.021$ ). Binding affinity of MePPEP was the same between groups, and the *rs2023239C* allele was not found to affect results between groups; allele carriers had higher binding than noncarriers within groups, though not reaching statistical significance. A power analysis using this *in vitro* data suggested that 53 subjects per group would be needed for an *in vivo* PET study using [<sup>18</sup>F]FMPEP-*d*<sub>2</sub> with 80% power and  $p < 0.05$ .

In a subsequent study, [<sup>11</sup>C]MePPEP and [<sup>18</sup>F]FMPEP-*d*<sub>2</sub> were used to image CB<sub>1</sub> receptors in separate cohorts of males with first-episode psychosis (i.e., diagnosis or high risk for development of schizophrenia) (Borgan et al. 2019). [<sup>11</sup>C]MePPEP was imaged in patients and controls ( $n = 20$  each) for 90 min with arterial sampling, and demonstrated lower  $V_T$  in anterior cingulate cortex, hippocampus, thalamus, and striatum in those with psychosis compared to controls; results were significant by group, but not group  $\times$  region. Significant associations were found between lower  $V_T$  in hippocampus with greater symptom severity (as measured by the Positive and Negative Syndrome Scale [PANSS]), and lower  $V_T$  in striatum and anterior cingulate cortex with worse cognitive performance. [<sup>18</sup>F]FMPEP-*d*<sub>2</sub> was imaged in patients ( $n = 7$ ) and controls ( $n = 11$ ) from 0 to 60 and 90 to 120 min with arterial sampling, and demonstrated significantly lower  $V_T$  in anterior cingulate cortex, hippocampus, thalamus, and striatum of those with psychosis compared to controls. As predicted by the previous *ex vivo* study, antipsychotic treatment did not affect the outcome in either cohort, as treatment status did not affect the results.

Autoradiography with [<sup>3</sup>H]OMAR was performed in the same schizophrenia and matched control subjects previously analyzed by *in situ* hybridization (for mRNA) and radioimmunohistochemistry (for protein) (Eggen et al. 2008; Volk et al. 2014).



[<sup>3</sup>H]OMAR was grossly 8% higher in all layers of prefrontal cortex from schizophrenia subjects compared to matched controls, and did not differ by multiple clinical characteristics (*e.g.*, psychiatric medications, cannabis use, history of suicide). [<sup>3</sup>H]OMAR correlated negatively with CB<sub>1</sub> receptor mRNA, which may reflect altered trafficking of CB<sub>1</sub> receptors with a nonlinear 1:1 relationship between transcription and translation, higher CB<sub>1</sub> receptor affinity leading to reduced expression, or reduced expression (*e.g.*, from loss of receptor reserve pools) leading to higher affinity of the remaining receptors. [<sup>3</sup>H]OMAR positively correlated with ABHD6 (which metabolizes 2-AG) mRNA expression, suggesting that lower 2-AG levels could be associated with CB<sub>1</sub> receptor upregulation.

In an initial study, patients with schizophrenia ( $n = 10$ ) trended towards higher measurements of  $V_T$  with [<sup>11</sup>C]OMAR compared to healthy controls ( $n = 10$ ). There was no significant difference in  $V_T$  between healthy controls and schizophrenia patients, with the exception of that in pons (Wong et al. 2010a); however this finding did not survive correction for multiple comparisons. Patients in the study were on either olanzapine or risperidone monotherapy, and antipsychotics have been shown to have a mixed effect on CB<sub>1</sub> receptor binding or availability; in subgroup analysis, no difference was found between patients on the two medications. Within the schizophrenia patient group, there was no association of  $V_T$  with Brief Psychotic Rating Score (BPRS); however when scores were taken as a ratio of positive over negative symptoms, there was a positive correlation with  $V_T$  in several areas of neocortex (Wong et al. 2008).

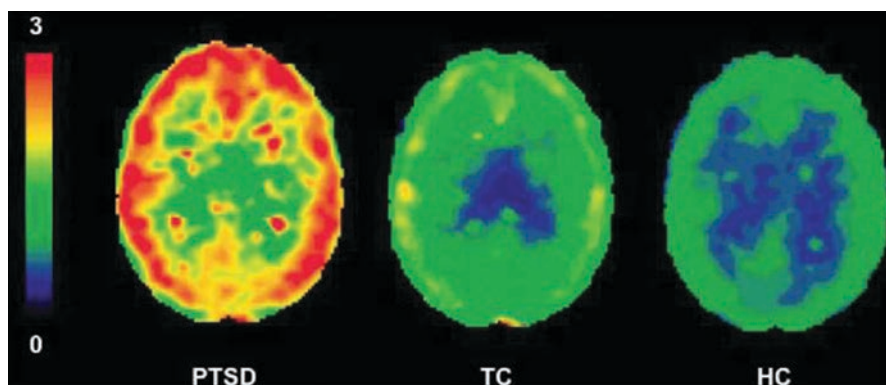
In a follow-up study, patients with schizophrenia ( $n = 25$ ) demonstrated ~12% significantly lower global  $V_T$  with [<sup>11</sup>C]OMAR compared to healthy controls ( $n = 18$ ) (Ranganathan et al. 2016). Patients with schizophrenia treated with antipsychotic medications ( $n = 18$ ) had higher  $V_T$  with [<sup>11</sup>C]OMAR compared to untreated patients ( $n = 7$ ), and while the two patient groups had no significant differences in PANSS, CB<sub>1</sub> receptor availability and PANSS were positively correlated in hippocampus, hypothalamus, and thalamus. As mentioned above, schizophrenia patients who used tobacco had CB<sub>1</sub> receptor availability that was greater than that of non-smoking patients, and it was nonsignificantly lower than that of nonsmoking controls.

In contrast to results obtained using [<sup>11</sup>C]MePPEP, [<sup>18</sup>F]FMPEP-*d*<sub>2</sub>, and [<sup>11</sup>C]OMAR, [<sup>18</sup>F]MK-9470 demonstrated 5–10% higher binding in brains of individuals with schizophrenia ( $n = 51$  with antipsychotic treatment,  $n = 6$  after antipsychotic washout,  $n = 10$  first-episode psychosis and antipsychotic naive) compared to healthy controls ( $n = 12$ ) (Ceccarini et al. 2013b). Specifically, binding was higher in regions of the mesocortical limbic circuit (nucleus accumbens, insula, cingulate cortex, inferior frontal cortex, and areas of the parietal and mediotemporal lobes), and was negatively correlated with PANSS negative symptom and depression scores in the antipsychotic free subgroup. In a rodent model of schizophrenia mimicking the *in utero* exposure to infectious agents hypothesized to contribute to risk of disease in humans, [<sup>18</sup>F]MK-9470 reflected reduced CB<sub>1</sub> receptor availability in globus pallidus of adolescent rats, and increased receptor availability in sensory cortex and hypothalamus in adult rats after prenatal exposure to a viral mimic agent, compared to controls (Verdurand et al. 2014).

### 12.3.4 Mood and Anxiety Disorders

Cannabinoid agonists demonstrate mood-altering effects (Degenhardt et al. 2003). The location of CB<sub>1</sub> receptors in limbic and cerebral cortex suggests that they are involved in mood dysfunction, and endocannabinoid stimulation does indeed have anxiolytic and antidepressant effects on animals (Gobbi et al. 2005). Inhibitors of FAAH are a hopeful and novel target for anxiety and depression, suggesting that enhancement of local endocannabinoid tone is preferred to exogenous cannabinoid agonists, which are potentially anxiogenic (more commonly at high doses) as well as anxiolytic (Moreira and Wotjak 2010). While the effect of cannabinoid agonists on depression is unclear, they generally have a negative effect on symptoms when used chronically (Lowe et al. 2019), and inverse agonists have been shown to increase depression and suicidal thoughts (as previously discussed). Several genetic polymorphisms have been identified in association with mood and anxiety disorders, and likely cause alterations to endocannabinoid signaling (Hillard et al. 2011). Genetic and postmortem studies have implicated the ECS in depression and suicide, and CB<sub>1</sub> receptors have been found to be upregulated in suicide victims (Vinod and Hungund 2006b). A recent review concluded that cannabinoid receptors, particularly CB<sub>1</sub> receptors, may become promising targets for the development of novel therapeutic tools for the treatment of suicidal behavior (Colino et al. 2018). Several studies have found reductions in circulating endocannabinoid levels in individuals with PTSD (Neumeister et al. 2013; Hill et al. 2013). Since endocannabinoids appear to influence stress, anxiety, and reversal learning related to aversive memories, cannabinoid receptor agonists have been proposed as a potential treatment for PTSD. One small randomized placebo-controlled crossover trial indicated that nabilone, a synthetic analogue of  $\Delta^9$ -THC, reduced PTSD symptoms in male military personnel compared to placebo (Jetly et al. 2015). While a follow-up study also demonstrated reduced PTSD-related nightmares (Jetly et al. 2015), several others have found that cannabis can worsen PTSD symptoms and increase the risk of violence and suicidal behaviors (Wilkinson et al. 2015; Barrett et al. 2011; Kimbrel et al. 2018; Borges et al. 2016; Moller et al. 2013). A recent review suggested that cannabidiol (CBD), while representing a less specific pharmacological approach, may be another way to treat trauma-related and anxiety disorders (Patel et al. 2017).

[<sup>11</sup>C]OMAR demonstrated ~20% higher  $V_T$  in the brain of individuals with PTSD ( $n = 25$ ) compared to trauma-exposed controls ( $n = 12$ ) and healthy controls without trauma history ( $n = 23$ ) (Fig. 12.12) (Neumeister et al. 2013). While cortisol was lower in the trauma-control and PTSD groups, peripheral anandamide, but not other endocannabinoids, was >50% lower in the PTSD group compared to either control group. As well, [<sup>11</sup>C]OMAR  $V_T$  was negatively correlated to peripheral anandamide in the PTSD group, and other demographic and clinical comorbidities and variables were not correlated with [<sup>11</sup>C]OMAR  $V_T$ . Combination of [<sup>11</sup>C]OMAR  $V_T$ , anandamide, and cortisol levels could identify ~85% PTSD individuals. The low anandamide concentration suggests compensatory CB<sub>1</sub> receptor upregulation, and together could be reflective of an ineffective modulation of stress response which results in PTSD symptoms, compared to those exposed to trauma without developing symptoms.



**Fig. 12.12** [ $^{11}\text{C}$ ]OMAR volume of distribution ( $V_T$ ) values in post-traumatic stress disorder (PTSD,  $n = 25$ ) and trauma-exposed control (TC,  $n = 12$ ) groups relative to healthy control (HC,  $n = 23$ ) group (reprinted from Neumeister et al. (2013) with permission of Nature Publishing Group)

While the role of the amygdala on the anxiogenic effects of cannabis had been examined in preclinical studies, use of PET provided an opportunity to assess this association in humans *in vivo*. Using [ $^{11}\text{C}$ ]OMAR,  $\text{CB}_1$  receptor availability in amygdala was found to be positively correlated to attentional bias to threat and severity of threat symptomatology (using the dot-probe task measuring reaction and bias to threatening versus neutral words) across a group of individuals with PTSD ( $n = 12$ ), trauma exposed without PTSD ( $n = 4$ ), and nontrauma-exposed controls ( $n = 4$ ) (Pietrzak et al. 2014). Moreover, a mediation analysis demonstrated that the endophenotype of attentional bias to threat better explains and connects the association between higher amygdala  $\text{CB}_1$  receptor availability and trauma-related psychopathology than a direct relationship alone. Similar to that seen in the PTSD study, peripheral anandamide concentrations were negatively correlated with  $\text{CB}_1$  receptor availability, as well as attentional bias to threat. In a separate study, male participants ( $n = 14$ ) with experiential use of cannabis were given THC 10 mg orally and assessed for anxiety (state-trait anxiety inventory), psychosis (PANSS), and response to fearful vs. neutral faces during fMRI imaging, and were found to have significantly greater anxiety and psychosis ratings during peak levels of THC, and activation of right amygdala and adjacent limbic and associated areas (Bhattacharyya et al. 2017).  $\text{CB}_1$  receptor  $V_T$  was measured with [ $^{11}\text{C}$ ]MePPEP (by spectral analysis), and right amygdala  $V_T$  was positively correlated with the anxiogenic (but not psychotogenic) effect of THC, and positively correlated with the activation seen during fMRI fear processing task.

### 12.3.5 Neurodegenerative Diseases

Cannabinoids exert a neuroprotective influence on some neurological diseases, including Alzheimer's disease (AD), Parkinson's disease (PD), Huntington's disease (HD), multiple sclerosis, and epilepsy. Synthetic cannabinoid receptor

agonists/antagonists or compounds can provide symptom relief or control the progression of neurological diseases. However, the molecular mechanism and the effectiveness of these agents in controlling the progression of most of these diseases remain unclear (Agar 2015).

### 12.3.5.1 Alzheimer's Disease

Alzheimer's disease (AD) is one of many neurodegenerative disorders that are associated with excitotoxicity, oxidative stress, and neuroinflammation, and certain cannabinoids have been demonstrated to inhibit these events to halt the progression of neurodegeneration. CB<sub>1</sub> and CB<sub>2</sub> receptors and FAAH have found to be co-localized with amyloid plaques in Alzheimer's disease (AD). CB<sub>2</sub>-selective agonists were shown to block amyloid-induced activation of cultured microglial cells (Ramirez et al. 2005), and would presumably decrease neuroinflammation in AD. In postmortem studies, [<sup>125</sup>I]SD7015, a CB<sub>1</sub> selective ligand, has demonstrated an inverse correlation of CB<sub>1</sub> receptor density and Braak tau pathology staging (Farkas et al. 2012), suggesting a novel *in vivo* method of staging AD progression. Independent of cannabinoid receptor activation, Δ<sup>9</sup>-THC competitively inhibits acetylcholinesterase (AChE) and prevents AChE-induced amyloid beta-peptide aggregation (Eubanks et al. 2006). Study of CB<sub>1</sub> and CB<sub>2</sub> receptors in correlation with amyloid load could lead to new treatment options for AD.

Activating CB<sub>1</sub> receptors has been demonstrated to produce certain therapeutic effects on animal models of AD (Aso et al. 2018). [<sup>18</sup>F]FMPEP-*d*<sub>2</sub> was longitudinally assessed in a mouse model of AD (Takkinen et al. 2018). Compared to wild-type littermates, APP/PS1–21 transgenic mice, which produce amyloid β<sub>42</sub> around 6 weeks and peaking at 9 months of age, were found to have lower [<sup>18</sup>F]FMPEP-*d*<sub>2</sub> uptake in parietotemporal cortex (at 9, 12, and 15 months), hippocampus (at 9 and 12 months), and cerebellum (at 9 months) relative to thalamus (used as a pseudoreference region and with no significant difference between groups). *Ex vivo* autoradiography also demonstrated reduced [<sup>18</sup>F]FMPEP-*d*<sub>2</sub> binding in amyloid-overproducing mice, but with difference by sex and time (males with lower binding in striatum at 15 months, females with lower binding in parietotemporal cortex, striatum, and posterior hippocampus). Immunohistochemical staining of CB<sub>1</sub> receptors demonstrated visible loss in amyloid-overproducing mice compared to wild type at 15 months. However, measured by Western blot, CB<sub>1</sub> receptor expression was not different by genotype at 9 months, though males were significantly lower than females.

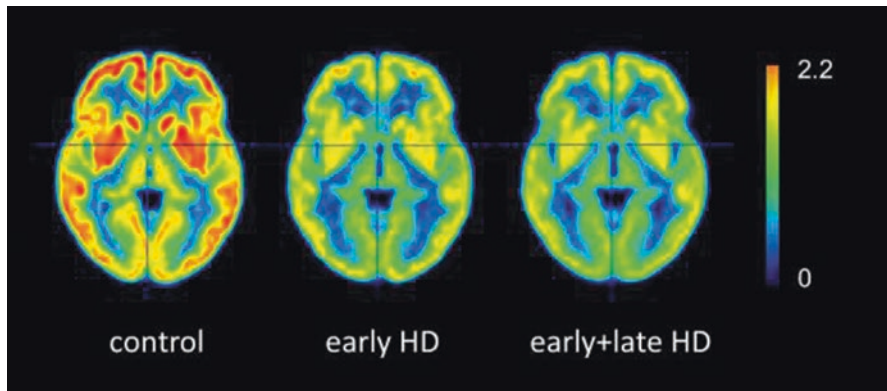
The activity of CB<sub>1</sub> receptors was studied in postmortem brain samples of AD patients during clinical deterioration. CB<sub>1</sub> receptor activity was higher at earlier AD stages in hippocampal areas and internal layers of frontal cortex, but a decrease was observed at the advanced stages. The pattern of modification appears to indicate initial hyperactivity of the endocannabinoid system in brain areas that lack classical histopathological markers at earlier stages of AD, indicating an attempt to compensate for the initial synaptic impairment, which is then surpassed by disease progression. These results suggest that initial CB<sub>1</sub> receptor stimulation might have therapeutic relevance (Manuel et al. 2014). However, another study found no

evidence for a difference in CB<sub>1</sub> receptor availability in AD compared to age-matched controls (Ahmad et al. 2014) using combined [<sup>18</sup>F]MK-9470 and [<sup>11</sup>C]PIB PET scans. When compared with reported CB<sub>1</sub> receptor changes in Parkinson's and Huntington's disease (see below), this suggests that CB<sub>1</sub> receptors may be differentially involved in neurodegenerative disorders.

### 12.3.5.2 Huntington's Disease

Huntington's disease (HD) is caused by a trinucleotide repeat expansion on the *HTT* gene, leading to a polyglutamine tail of the huntingtin protein. Although the normal physiologic function of the protein remains unclear, this pathological variant has been shown to cause disruptions in nuclear transcription factors, presumably leading to a decrease in normal expression of certain proteins. The pathological characteristic is progressive neuronal loss, preferentially affecting the caudate nucleus, resulting in motor (choreiform movements), cognitive (early-onset dementia), and psychiatric symptoms (behavioral changes, e.g., disinhibition, aggression). One of the earliest neurochemical alterations in HD is the loss of CB<sub>1</sub> receptors in the basal ganglia (Glass et al. 2000), which occurs before the D<sub>1</sub>/D<sub>2</sub> receptor dopaminergic dysfunction and characteristic choreic motor dysfunction and is strongly correlated with chorea and cognitive deficit (Laprairie et al. 2016). In animal models, the ECS is hypofunctional (Lastres-Becker et al. 2003). A huntingtin/CB<sub>1</sub> double-knockout mouse model demonstrated that endocannabinoid signaling was essential to the neuropathologic symptoms, and that CB<sub>1</sub> receptor loss may sensitize cells to additional excitotoxic damage (Blazquez et al. 2011). Oom et al. found early decreased CB<sub>1</sub> receptor binding in the bilateral caudate, putamen, globus pallidus, and thalamic nucleus of presymptomatic HD (R6/2) mice, with further progressive decline using [<sup>18</sup>F]MK-9470, suggesting that PET quantification of *in vivo* CB<sub>1</sub> binding may be a useful early biomarker for HD (Ooms et al. 2014).

In HD, there is also selective loss of spinal neurons and the appearance of activated microglia in striatum, cortex, and white matter. The number of activated microglia in the striatum and cortex has been shown to correlate with the extent and progression of neuronal loss (Sapp et al. 2001), and CB<sub>2</sub> agonists show neuroprotective effects in an HD animal model (Sagredo et al. 2009; Paldy et al. 2008). In this way, CB<sub>2</sub> receptor imaging may be useful in monitoring the progression of neuronal loss in HD. In addition, early regional dysfunctions in CB<sub>1</sub> receptor signaling, involving the lateral globus pallidus and caudate-putamen, have been found, suggesting another useful early biomarker for HD. Decreased [<sup>18</sup>F]MK-9470 brain uptake was observed in a rat genetic model of Huntington's disease when compared to wild-type littermates (Casteels et al. 2011); however, this was observed in a regionally selective fashion limited to the basal ganglia, rather than the global reduction described below in humans. A rat model of HD using unilateral quinolinic acid lesion demonstrated a disproportionately modest decrease of [<sup>18</sup>F]MK-9470 uptake compared to the severity of the lesion, and an increase in contralateral basal ganglia and cerebellum, the latter of which corresponded to improved functional outcome (Casteels et al. 2010c).



**Fig. 12.13** Cross-sectional, partial-volume-corrected, modified standardized uptake value (mSUV) parametric images of [ $^{18}\text{F}$ ]MK-9470, averaged for all controls ( $n = 14$ ), early-HD patients ( $n = 7$ ), and total group of HD patients ( $n = 19$ ) (courtesy of Dr. Koen Van Laere, KU Leuven)

Given the high concentration of  $\text{CB}_1$  receptors in the basal ganglia, neuropsychiatric and movement disorders known to affect those regions present themselves as potentially sensitive and particularly interesting targets for  $\text{CB}_1$  receptor PET imaging. In humans with HD, [ $^{18}\text{F}$ ]MK-9470 uptake was decreased in all areas of brain except white matter compared to healthy controls (Fig. 12.13) (Van Laere et al. 2010). Intriguingly, this was also observed in early symptomatic patients, even in areas with minimal volume loss. Decreased uptake was presumably due to reduced  $\text{CB}_1$  receptor availability, as huntingtin is known to interfere with protein and receptor expression. Therefore,  $\text{CB}_1$  PET imaging may be a useful tool to assess early symptomatic, and potentially presymptomatic, Huntington's disease onset. As well, decreased [ $^{18}\text{F}$ ]MK-9470 uptake was found in pre-HD subjects in the prefrontal cortex and cingulate cortex, with depression, apathy, and irritability inversely correlating with the binding (Ceccarini et al. 2019a). This suggests that a potential molecular basis of neuropsychiatric symptoms in pre-HD could lead to new therapeutic avenues. It has previously been suggested that  $\text{CB}_1$  agonists may delay the development of this disease (Lastres-Becker et al. 2003; Glass et al. 2004), and at least one clinical study has shown that they improve neuropsychiatric symptoms (Curtis et al. 2009). However, in a double-blind, randomized, placebo-controlled, crossover pilot clinical trial, no significant differences on motor, cognitive, behavioral, and functional scores were detected among HD patients treated with Sativex (a botanical extract with an equimolecular combination of  $\Delta^9$ -THC and cannabidiol) and placebo (Lopez-Sendon Moreno et al. 2016).

### 12.3.5.3 Parkinson's Disease

Several studies have reported neuroprotective effects of cannabinoids in glutamatergic excitotoxicity, oxidative damage, traumatic injury, ischemia, and neurodegeneration models (Paradisi et al. 2006; Sarne and Mechoulam 2005).  $\text{CB}_1$  receptors are densely located in the basal ganglia, and the ECS has a modulatory effect on

excitation and inhibition, as evident in movement disorders. Parkinson's disease (PD) leads to a decrease in voluntary movements, secondary to insufficient striatal dopamine, causing a subsequent increase in GABAergic activity (Pacher et al. 2006). Early on CB<sub>1</sub> receptor density is reduced, but later the density is increased to compensate for the subsequent overexcitation of GABA (Orgado et al. 2009). mRNA expression of CB<sub>1</sub> is changed and an increased binding capacity has been observed in postmortem tissue of subjects with PD (Lastres-Becker et al. 2001). It is hypothesized that alterations of endocannabinoid synthesis stimulate increased dopamine release (Giuffrida et al. 1999), and cannabinoids inhibit the excitotoxic glutamate release and counteract oxidative stress to dopaminergic neurons which is responsible for nigrostriatal damage (Levenes et al. 1998). Deficiency in endocannabinoid transmission is thought to contribute to late-stage dyskinesias. Thus, modulation of CB<sub>1</sub> receptor-mediated basal ganglia transmission can be used to suppress or prevent the development of these complications (Segovia et al. 2003; Sevcik and Masek 2000; Di Marzo et al. 2000; Sieradzan et al. 2001; Fox et al. 2002); however, the results to date have been mixed (Carroll et al. 2004).

One recent study designed to detect potential alterations in the CB<sub>1</sub> and CB<sub>2</sub> (A isoform, CB<sub>2</sub>Ar) receptors and monoacylglycerol lipase (MAGL) gene expression in the substantia nigra and putamen found them to be closely related to the neuropathological processes of PD. Therefore, the pharmacological modulation of these targets could represent a new potential therapeutic tool for the management of PD (Navarrete et al. 2018). Most interestingly, a recent study (Ceccarini et al. 2019b) found decreased CB<sub>1</sub> receptor availability in the prefrontal and midcingulate cortex in PD patients using [<sup>18</sup>F]MK-9470 PET imaging which strongly correlated with the level of disturbed executive functioning, episodic memory, and visuospatial functioning. Thus, further investigation of regional CB<sub>1</sub> receptor expression in varying levels of PD-related cognitive impairment may be useful. In recent study, upregulation of CB<sub>2</sub> receptors was shown in a PD rat model (Concannon et al. 2016). Although more studies are necessary, these findings suggest that CB<sub>2</sub> receptor-selective agonists may provide neuroprotection against the neurodegenerative processes of PD.

Rodent models of PD using 6-hydroxydopamine showed decreased [<sup>18</sup>F]MK-9470 uptake in bilateral striatum and contralateral cerebellum relative to the lesioned side. Additional correlations were observed with an inverse relationship of tyrosine hydroxylase-containing neurons (*i.e.*, dopaminergic neurons) to [<sup>18</sup>F]MK-9470 uptake in ipsilateral cerebellum, and a positive relationship of [<sup>18</sup>F]MK-9470 uptake and contralateral limb impairment (*i.e.*, disease severity) (Casteels et al. 2010b).

[<sup>18</sup>F]MK-9470 uptake was altered in a region-specific pattern in individuals with PD (Van Laere et al. 2012). In ventral midbrain (site of the substantia nigra) tracer uptake was decreased for early and late stages of PD patients compared to healthy controls, likely representing reduced CB<sub>1</sub> receptor expression in the pars reticulata, irrespective of disease progression, rather than simply a marker of neuronal loss. However in putamen, prefrontal cortex, midcingulate, anterior insula, and hippocampus (regions correlating to dopaminergic projections) tracer uptake was increased relative to total brain uptake in comparison to healthy controls. Regionally

increased CB<sub>1</sub> receptor availability could be explained by compensatory processes of reducing glutamatergic drive in the absence of normally inhibitory dopamine (in striatum) in glutamatergic neurons, and disinhibition of dopaminergic neurons in the ventral tegmental area via a feedback mechanism from mesocortical and mesolimbic regions. Comparison of PD patients with or without levodopa-induced dyskinesias did not reveal any differences in [<sup>18</sup>F]MK-9470 uptake, as was also shown in healthy rats receiving L-DOPA or bromocriptine (Casteels et al. 2010d).

#### 12.3.5.4 Multiple Sclerosis

Meta-analyses have supported the beneficial effect of cannabinoid agonists as treatment for neuropathic pain (Iskedjian et al. 2007; Rog et al. 2005) and muscle spasticity (Pryce and Baker 2007; Mori et al. 2014) in multiple sclerosis (MS). Selective CB<sub>1</sub> receptor overexpression on various glial cells and CB<sub>2</sub> receptor-positive microglia is located within active plaques and in the periphery of chronic active plaques, supporting a role for the ECS as a biomarker or therapeutic target for MS (Bifulco et al. 2007; Benito et al. 2007). Treatment of MS with cannabinoids appears to be based on two mechanisms: (i) reducing excitotoxicity by CB<sub>1</sub> receptors on glia and neurons and (ii) reducing inflammation by CB<sub>2</sub> receptors on microglia (Rossi et al. 2010; Pryce and Baker 2015). Nabiximols is a sublingual spray developed by GW Pharmaceuticals for the treatment of MS using a proprietary blend of a Δ<sup>9</sup>-THC and cannabidiol, which show differing affinities for CB<sub>1</sub> and CB<sub>2</sub> receptors, as well as other targets. Although MS has been considered mainly an inflammatory disorder, recent evidence, has revealed the importance of neurodegenerative events, opening the possibility that cannabinoid agonists, given their cytoprotective properties, may also serve to reduce oligodendrocyte death and axonal damage in MS. Thus, the treatment with WIN55,512-2, a potent CB<sub>1</sub> and CB<sub>2</sub> agonist, was reported to be effective to ameliorate tremor and spasticity in mice with chronic relapsing experimental autoimmune encephalomyelitis (EAE), a commonly used murine model of MS (de Lago et al. 2012). Growing evidence suggests that targeting the hydrolysis of 2-AG may offer a more favorable benefit-to-risk balance in MS than existing cannabinoid medicines (Manterola et al. 2018). While CB<sub>1</sub> receptor imaging might potentially be useful for probing neuroadaptations related to glutamatergic excess or other interactions between the ECS and MS pathology, no studies have yet been reported.

CB<sub>2</sub> receptor imaging may be useful to identify MS plaques of the active type. Ten days after EAE induction in mice, CB<sub>2</sub> receptor gene (*CNR2*) mRNA was ten-fold higher in activated microglia compared to microglia in a resting state. In total brain, *CNR2* mRNA was upregulated 100-fold (Maresz et al. 2005). In humans, CB<sub>2</sub>-expressing microglial cell staining was observed in spinal cord and brain sections of patients with MS and amyotrophic lateral sclerosis (Yiangou et al. 2006; Palazuelos et al. 2008). CB<sub>2</sub> receptor-positive microglial cells were homogeneously distributed within active MS plaques, but were found only in the periphery of chronic active plaques (Benito et al. 2007). However, to date, no CB<sub>2</sub> receptor imaging studies of MS patients have yet been reported.



### 12.3.5.5 Amyotrophic Lateral Sclerosis

Amyotrophic lateral sclerosis (ALS) is a rapidly progressing neurodegenerative disease accelerated by neuroinflammation. CB<sub>2</sub> receptors are dramatically upregulated in the spinal cords of mice with ALS, and CB<sub>2</sub> agonists have been shown to slow motor neuron degeneration (Shoemaker et al. 2007). Animal models have demonstrated that increased endocannabinoid tone delays disease progression (Kim et al. 2006), though the limited clinical data available to date has not shown cannabinoid agonists to be helpful in ongoing disease. Nevertheless, the potential for ECS to affect neuroinflammatory and glutamate-mediated processes in ALS, along with the sparse number of therapeutics available, should prompt additional investigation.

### 12.3.6 Pain and Migraine

Cannabinoids are antinociceptive in several types of animal models, acute tissue injury and nerve injury-induced nociception, and play a role in transmission, processing, perception, and modulation of pain (Hohmann and Suplita 2nd. 2006; Anand et al. 2009). Cannabinoid agonists and endocannabinoids act on CB<sub>1</sub> receptors in the brain, dorsal root ganglia, and peripheral nerves (Pacher et al. 2006). Activation of CB<sub>2</sub> receptors has been shown to reduce inflammatory pain (see above, Multiple Sclerosis). In addition, anti-inflammatory and analgesic medications affect proteins involved in lowering endocannabinoid levels (*e.g.*, COX-2, endocannabinoid transporter), and act synergistically with cannabinoid agonists in animal studies (Guindon et al. 2006). Interestingly, cannabidiol, which behaves as a cannabinoid receptor-negative allosteric modulator and TRPV1 agonist among other roles, has shown anti-inflammatory properties as well (Booz 2011). In humans, the mechanism of cannabinoid receptor-mediated analgesia is less clear. The exact mechanism of pain suppression is unknown, but synergistic actions and interactions with the endogenous opioid system have been reported (Corchero et al. 2004), and the ECS modulates the actions of cytokines and glucocorticoids (Hill and McEwen 2009), which play important roles in inflammation. In female migraine patients, platelets have been observed to have an increased rate of endocannabinoid metabolism, suggesting a decreased endocannabinoid tone in patients suffering from migraine headaches (Cupini et al. 2006). In addition, TRPV1 receptors are directly linked to pain transmission, and are affected by the ECS. Thus, the modulation of pain by the endocannabinoid system is likely a mixed interplay between central and peripheral mechanisms, CB<sub>1</sub>, CB<sub>2</sub>, TRPV1, and 5-HT<sub>1a</sub> receptors, and signal propagation to other neurotransmitter systems to affect the sensation and affective (and anxiety-related component) of pain. Cannabinoids are effective in suppressing pain symptoms of chemotherapy-induced and other peripheral neuropathies, but their widespread use is limited by CNS-mediated and anticholinergic side effects (Mulpuri et al. 2018). However, cannabinoid-based drugs appear to heterogeneously influence the perception of pain, causing a reduction in the affective, but not the sensory, perception of pain (Lotsch et al. 2018; Lee et al. 2013; De Vita et al. 2018).

Several synthetic cannabinomimetic drugs are being developed to treat pain and depression. However, the precise mode of action of endocannabinoids on different targets in the body and whether their effects on pain and depression follow the same or different pathways remain to be determined (Huang et al. 2016).

In women with episodic migraine, [ $^{18}\text{F}$ ]MK-9470 uptake was globally increased by about 16%, with focal increases found in anterior cingulate, mesial temporal, prefrontal, and superior frontal cortices (Van der Schueren et al. 2011). No significant clinical correlations could be made between CB<sub>1</sub> receptor availability and migraine characteristics (*e.g.*, with or without aura, age of onset, migraine frequency).

### 12.3.7 Epilepsy

In epilepsy, an uncontrolled excitatory signal causes an epileptic seizure. Endocannabinoid-mediated self-inhibition reduces excitability of neocortical interneurons (Bacci et al. 2004). The CB<sub>1</sub> receptor acts as an important endogenous locus of epileptic seizure modulation (Wallace et al. 2002) and endocannabinoids block status epilepticus in hippocampal neurons (Deshpande et al. 2007). There is evidence of a compensatory upregulation of CB<sub>1</sub> receptors in epilepsy (Wallace et al. 2003; Chen et al. 2003). Both animal and human clinical studies have shown mixed results on the use of cannabinoid agents for seizure depending on model, pathology, and dose of the agent. This is likely due to the complex interplay of multiple actions of cannabinoid receptors, and their dual association with excitatory and inhibitory neurotransmitters.

An examination of patients with mesial temporal lobe epilepsy due to hippocampal sclerosis provided an *in vivo* perspective on the role of CB<sub>1</sub> receptors in a locally hyperexcitable and excitotoxic environment. Compared to healthy controls, [ $^{18}\text{F}$ ]MK-9470 uptake was increased in the ipsilateral temporal lobe, and decreased in ipsilateral and, to a lesser extent, contralateral insula relative to the affected lobe (Goffin et al. 2011). Insufficient data is available to distinguish whether these changes are acute or chronic; however there were significant correlations to relative uptake in the seizure focus: a positive correlation with the number of seizures in the month prior to scanning, and a negative correlation to the number of days since the last seizure before scanning.

A recent study using [ $^{18}\text{F}$ ]MK-9470 PET imaging in an amygdala kindling model of epileptogenesis in rhesus monkeys (Cleeren et al. 2018) found evidence that the CB<sub>1</sub> receptors may be dynamically and progressively involved in the early development of mesial temporal lobe epileptogenesis.

### 12.3.8 Stroke

CB<sub>2</sub> receptor-positive cells occur in the brain within minutes following stroke (Ashton et al. 2007), and CB<sub>1</sub> receptor expression is increased in the ischemic zone

(Jin et al. 2000). In a CB<sub>1</sub> receptor knockout mouse model of stroke, mortality was increased and infarct size and ischemia were more severe compared to wild-type mice (Parmentier-Batteur et al. 2002). This suggests that endogenous cannabinoid signaling pathways may protect against additional anoxic and inflammatory injury following ischemic stroke. CB<sub>2</sub> receptor activation has been proven to produce protective effects against acute poststroke inflammation (Capetini et al. 2012) and a CB<sub>1</sub> receptor antagonist has been found to be protective in an ischemic mouse model (Reichenbach et al. 2016). Studies have also indicated that CB<sub>2</sub> receptor agonists reduce neurodegeneration after stroke through anti-inflammatory activity (Yu et al. 2015). Vandeputte et al. (Vandeputte et al. 2012) found that [<sup>18</sup>F]MK-9470 PET demonstrated a strong increase in CB<sub>1</sub> receptor availability 24 and 72 h after stroke surrounding the lesion, in a photothrombotic stroke model, suggesting that time-dependent and regionally increased CB<sub>1</sub>, but not CB<sub>2</sub>, binding is an early consequence of photothrombotic stroke. This implies that pharmaceutical interventions should primarily aim at CB<sub>1</sub> receptors.

### 12.3.9 Non-CNS Applications

Cannabinoid drugs have recently also become candidates for peripheral inflammatory and degenerative disorders such as atherosclerosis (Steffens et al. 2005) and in the field of oncology (Herrera et al. 2005). For example, CB<sub>2</sub> receptors have been shown to be involved in the apoptotic pathway in certain tumor cell lines, and have correlated with certain brain tumor grades (Ellert-Miklaszewska et al. 2007). In tissues with normally low receptor expression, there may be a >1000-fold increase in receptor density in the presence of certain tumor types (Oesch et al. 2009). [<sup>18</sup>F]MK-9470 has been assessed in prostate cancer and found to bind in both benign and malignant prostate tissue, but could not reliably distinguish between the two (with notably weak CB<sub>1</sub> receptor detection by immunohistochemistry), and could visualize metastases in the appendicular skeleton but not reliably in the axial skeleton (Emonds et al. 2013). Peripheral CB<sub>1</sub> receptors have been considered a target of therapy in metabolic and cardiac diseases (see above Sect. 12.3.2 for imaging applications in brown adipose tissue and myocardium). Finally, CB<sub>1</sub> and CB<sub>2</sub> receptors are upregulated in liver fibrosis as a consequence of cirrhosis (Wasmuth and Trautwein 2007; Teixeira-Clerc et al. 2008).

### 12.3.10 Considerations and Conclusions on the Use of Current CB<sub>1</sub> Receptor PET Radioligands

The development and evaluation of CB<sub>1</sub> receptor PET radioligands have presented both insight into CB<sub>1</sub> receptor pathophysiology and challenges to PET radiochemistry and pharmacokinetic modeling. The primary breakthrough in CB<sub>1</sub> receptor imaging was the development of ligands with lower lipophilicity and higher binding affinity. However, high lipophilicity continues to present anticipated challenges, and

high binding affinity and slow clearance of the radioligands have presented new ones. The results from [ $^{11}\text{C}$ ]OMAR and [ $^{11}\text{C}$ ]MePPEP exemplify the need for accurate plasma measurements when determining  $V_T$ , and show that peak uptake alone is not an appropriate outcome measure when comparisons are made between subjects. The development of additional radioligands with improved properties would be of value. To improve the measurements of parent ligand in plasma, alternative approaches with higher sensitivity could be used. Assays or spectroscopic methods with high accuracy for detecting very low concentrations of drug in plasma should be considered (*e.g.*, LC-MS).

Insights gained regarding the ECS include an understanding that CB<sub>1</sub> receptor density decreases after direct agonist stimulation. For example, cannabis abusers had lower  $V_T$  and/or radioligand binding compared to controls in all studies to date, and in diseases regarded to have increased endocannabinoid tone (*e.g.*, Huntington's disease) leading to receptor downregulation. However, these results have assumed that decreased radioligand outcome measures are the result of receptor downregulation and not occupancy; the inability of cannabinoid agonists to displace these radioligands has only been definitively demonstrated in rodents, and has been assumed via data interpretation but not yet demonstrated and reported in humans. Conversely, at least one study has suggested that decreased endocannabinoid tone is associated with upregulation of CB<sub>1</sub> receptors (Neumeister et al. 2013). Also, changes observed in CB<sub>1</sub> receptor density or availability were often region dependent, and in one case, a region-specific lesion resulted in contralateral changes (Casteels et al. 2010b); while unilateral lesions or comparisons across hemispheres are commonly performed for other tracers, findings from these CB<sub>1</sub> receptor imaging studies suggest that this may not always be an appropriate method.

Inverse discrepancy has been reported for CB<sub>1</sub> receptor imaging across several studies. For example, in alcohol-dependent patients [ $^{11}\text{C}$ ]OMAR demonstrated a 20% increase of  $V_T$  while [ $^{18}\text{F}$ ]FMPEP- $d_2$  demonstrated a 20–30% decrease, and [ $^{18}\text{F}$ ]MK-9470 demonstrated increased binding in chronic alcohol dependence but lower binding acute alcohol exposure in social drinkers. As well, [ $^{11}\text{C}$ ]MePPEP, [ $^{18}\text{F}$ ]FMPEP- $d_2$ , and [ $^{11}\text{C}$ ]OMAR all demonstrated lower  $V_T$  in schizophrenia patients compared to controls, while [ $^{18}\text{F}$ ]MK-9470 demonstrated high binding in the schizophrenia group. Whether these differences are due to study design or methods remains unclear, and therefore additional investigations are warranted.

The radioligands presented here have their particular pros and cons (Table 12.4). [ $^{11}\text{C}$ ]OMAR and [ $^{11}\text{C}$ ]SD5024, with their relatively fast kinetics and C-11 radiolabel, have the advantage that subjects could quickly complete multiple PET studies in one day, and receptor density could be quantified by  $V_T$ . Faster kinetics are attributed in part by their more modest binding affinity, which in turn likely contributes to less brain uptake and lower measures of  $V_T$ . [ $^{11}\text{C}$ ]OMAR binding was shown to decrease with age, and thus age should be analyzed as a clinical variable. [ $^{18}\text{F}$ ]MK-9470 has been studied extensively in healthy controls and in patients with several neuropsychiatric disorders, providing a wealth of data and observations. Its simplified outcome measure does not require arterial blood sampling, greatly easing subject burden and data analysis. However, this comes at the expense of potentially

less accurate measurements, as differences in metabolism and excretion are not accounted for in the final outcome measurement. These initial studies also uncovered several clinical variables that should be accounted for when analyzing [ $^{18}\text{F}$ ]MK-9470 data, including age, sex, and weight. [ $^{18}\text{F}$ ]FMPEP- $d_2$  has high brain uptake which can be quantified by  $V_T$ , and therefore is more likely to detect small changes in  $\text{CB}_1$  receptor availability, and more accurately measure an index of receptor density. It is available in C-11 form ([ $^{11}\text{C}$ ]MePPEP) in addition to F-18 form, should same-day repeat studies be preferred, and studies with either radioligand can be completed with about 90–120 min of scanning. Longer scan times with [ $^{18}\text{F}$ ]FMPEP- $d_2$  may be vulnerable to radioactivity accumulation in the brain, possibly due to radioactive metabolites. Analysis of [ $^{18}\text{F}$ ]FMPEP- $d_2$  should account for BMI as a clinical variable. Due to their lipophilicity, the free fraction of all radioligands should be measured when comparing group differences or studying the effect of drugs. To date, partial volume correction has not affected data for any of the radioligands.

Several uncertainties remain when interpreting data from  $\text{CB}_1$  receptor imaging studies. It has been assumed that changes in [ $^{18}\text{F}$ ]MK-9470 uptake or [ $^{11}\text{C}$ ]OMAR and [ $^{18}\text{F}$ ]FMPEP- $d_2 V_T$  likely represent changes in receptor availability. However, it is unknown whether changes in uptake or  $V_T$  are due to altered receptor density, internalization and/or changes in receptor trafficking (Coutts et al. 2001), or changes in receptor-binding affinity (including that occurring from allosteric modulation of other drugs), as these cannot be distinguished with PET (Martin et al. 2004). Additionally, the degree to which genetic polymorphisms of *CNRI* affect receptor density, binding affinity of the radioligands, or both has been inconsistent and remains to be more fully assessed (Zhang et al. 2004; Hutchison et al. 2008; Hirvonen et al. 2012b). Finally, most of the results presented here were all obtained either in a single study or within a relatively short time span. Therefore, little information is available regarding longitudinal changes of  $\text{CB}_1$  receptor expression over extended periods of time, and the results may suggest either a causal relationship or merely significant correlations.

Development of a  $\text{CB}_1$  receptor agonist radioligand remains an unmet need. As demonstrated by [ $^{11}\text{C}$ ]MePPEP, this class of radioligand (inverse agonist) is not displaced *in vivo* by endogenous or synthetic agonists. Therefore, these PET tracers can only be used for measuring changes or differences in receptor density, or for measuring receptor occupancy by inverse agonists. They cannot be used for measuring receptor occupancy by agonists, and estimates of endocannabinoid tone are only assessed indirectly. Therefore, an agonist radioligand that demonstrates preference for the high-affinity state and is therefore more susceptible to competition by endogenous agonists would be of great value. For example, a radioligand for high-affinity dopamine  $\text{D}_2$  receptors, [ $^{11}\text{C}$ ]MNPA, is displaced nearly twice as much by dopamine than an antagonist, such as [ $^{11}\text{C}$ ]raclopride (Seneca et al. 2006). Given that only a small proportion of receptors may be in the high-affinity state, the distribution and normal physiology of  $\text{CB}_1$  receptors may appear quite different when an agonist radioligand is used for *in vivo* PET. Therefore, development of an agonist

radiotracer may provide tremendous insight into the regulation of the endocannabinoid system.

If an agonist radiotracer for the CB<sub>1</sub> receptor were developed, several experiments probing the endocannabinoid system could be conceived. First, increasing endocannabinoid concentrations by inhibiting their degradation or reuptake (via FAAH, MAG lipase, or other proteins) would provide better understanding of this dynamic and possibly regionally selective process. Second, studies involving stimulation or inhibition of the release of primary neurotransmitters could be performed to assess the relative response of endocannabinoids to neuromodulation of GABA, glutamate, and dopamine. Third, a radioligand sensitive to dynamic changes in endocannabinoid concentration would allow for *in vivo* studies with behavioral components, such as acute stress response (Gorzalka et al. 2008).

Development of an agonist radiotracer is likely to be difficult, primarily due to their high lipophilicity. Very few CB<sub>1</sub> agonists have the moderate lipophilicity required for a PET tracer; a notable exception is the aminoalkylindole class of agonists, typified by WIN-55,212-2. In addition, most existing agonists do not have high selectivity for the CB<sub>1</sub> receptor. Indeed, one of the first attempts to develop a radioligand to image CB<sub>1</sub> receptors was [<sup>18</sup>F]Δ<sup>8</sup>-THC, which exhibited a prohibitively high lipophilicity and has nearly equal affinity for CB<sub>1</sub> and CB<sub>2</sub> receptors, as well as likely having P-gp substrate affinity. Given the relatively low expression of CB<sub>2</sub> compared to CB<sub>1</sub> receptors and the availability of selective antagonists, it is unknown if high selectivity would be an absolute requirement for an agonist radioligand. Finally, the development of an agonist radioligand may face the complexities posed by partial agonism and functional selectivity, which may present the CB<sub>1</sub> receptor with altered binding conformations and possibly differing binding affinities for agonists depending on the intracellular coupling of the receptor (Georgieva et al. 2008).

Lastly, there are many potential non-CNS applications for CB<sub>1</sub> receptor imaging, including liver fibrosis, metabolic diseases such as diabetes, and cancer. At present, insufficient data is available to determine if the existing CB<sub>1</sub> receptor radioligands would be optimal for use in the imaging of diseases outside the CNS, and, if so, what the optimal method of analysis would be (radioactivity concentration vs.  $V_T$ ). More research concerning the appropriate use of these tracers for imaging CB<sub>1</sub> receptors outside the brain needs to be performed.

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## 12.4 Imaging of CB<sub>2</sub>

The CB<sub>2</sub> receptor is expressed primarily by the immune system (Munro et al. 1993; Schatz et al. 1997; Galiegue et al. 1995), but has been found in intact CNS tissue in tiny concentrations (Van Sickle et al. 2005; Gong et al. 2006; Ashton et al. 2006). Importantly, CB<sub>2</sub> receptors are highly expressed in disorders that are associated with inflammation including cancer, pain, osteoporosis, and liver diseases (see for review (Pertwee 2009; Pisanti and Bifulco 2009; Ashton 2007)). The entire functional role of CB<sub>2</sub> receptor expression in the inflammatory processes has yet to be clarified,

though its upregulation makes CB<sub>2</sub> a unique biomarker of microglial/macrophage cell-specific activity in these neuroinflammatory processes.

Recent studies have demonstrated high expression of CB<sub>2</sub> receptors in CNS microglia in patients with various neuroinflammatory disorders (Benito et al. 2008). Postmortem tissue studies from patients with AD demonstrate dramatically increased expression of CB<sub>2</sub> receptors in areas of beta-amyloid (A $\beta$ ) plaque deposition (Benito et al. 2003). CB<sub>2</sub> receptor expression increases both in striatal microglia of transgenic mouse models of HD and in patients with HD (Palazuelos et al. 2009). Better understanding of the immune-modifying role and principal mechanisms of CB<sub>2</sub> receptor agonists in the human CNS could facilitate the development of disease-modifying treatments and preventive agents, as proposed for AD (Tolon et al. 2009). Induced expression of CB<sub>2</sub> receptors in neuroinflammatory disease is also observed in tissue samples from human and animal models of HIV-induced encephalitis and multiple sclerosis (Benito et al. 2005) (see for review (Tolon et al. 2009)). Traumatic brain injury (TBI) leads to the release of pro-inflammatory cytokines in the brain, which then cause neuronal cell death and secondary damage. Simultaneously, there is “on-demand” synthesis of cannabinoid receptors including CB<sub>2</sub>, which in turn inhibit inflammatory responses (Stella 2010; Shohami et al. 2011).

There are also unique therapeutic implications afforded by a better understanding of the endocannabinoid system and supported by recent studies of CB<sub>2</sub>-specific agonists (Tolon et al. 2009; Ehrhart et al. 2005). Recent studies demonstrated that brain CB<sub>2</sub> receptors modulate the rewarding and locomotor-stimulating effects of cocaine (Xi et al. 2011). CB<sub>2</sub> receptors are involved in some of the effects mediated by alcohol (Ishiguro et al. 2007) in rodent brains and expression of CB<sub>2</sub> in microglia increases in response to MDMA (“ecstasy”) treatment (Torres et al. 2010).

In summary, the CB<sub>2</sub> receptor that is nearly absent in the normal brain and highly expressed in activated microglia in neuroinflammation is an attractive target for PET imaging of a large variety of disorders that are associated with neuroinflammation. Although substantial progress has been made in the development of radioligands to image CB<sub>1</sub> with PET, the absence of suitable PET radioligands has hampered the noninvasive imaging of CB<sub>2</sub>. Several CB<sub>2</sub> radioligands have recently been synthesized and evaluated in animals as potential PET radioligands (Evens et al. 2008, 2009a; Fujinaga et al. 2010; Gao et al. 2010).

### 12.4.1 Development of First-Generation CB<sub>2</sub> Radioligands

Pioneer work of researchers from Canada and France revealed the first selective CB<sub>2</sub> agonists (Gallant et al. 1996) and inverse agonists (Rinaldi-Carmona et al. 1998) with nanomolar affinity, and additional review of early CB<sub>2</sub> PET imaging and radioligand development has been described in an excellent review (Evens and Bormans 2010). Several CB<sub>2</sub> receptor radioligands have been developed, including the oxoquinoline derivatives [<sup>11</sup>C]KD2, [<sup>11</sup>C]KP23, [<sup>11</sup>C]RS016, [<sup>11</sup>C]RSR-056, [<sup>11</sup>C]RS-028, and [<sup>18</sup>F]RS-126 and [<sup>11</sup>C]NE40, and the thiazole derivatives [<sup>11</sup>C]A-836339, [<sup>18</sup>F]JHU94620, and [<sup>18</sup>F]2f (Ni et al. 2019). In the last few years a large

number of selective high-affinity CB<sub>2</sub> ligands have been developed and they were a basis for the recent development of CB<sub>2</sub> PET radioligands (Tables 12.5, 12.6, and 12.7; Fig. 12.14).

The group from KU Leuven reported that [<sup>11</sup>C]1 had high CB<sub>2</sub>-binding affinity (Evens et al. 2008), but it exhibited low brain uptake (Tables 12.5 and 12.7). This is in accord with the high PSA value and high molecular weight (MW = 540) of this radioligand that are above the conventional limits for passive BBB transport. However, brain uptake increased following the administration of cyclosporine A, which inhibits several BBB efflux transporters including P-gp, indicating that the radioligand is an efflux transporter substrate. No further studies on [<sup>11</sup>C]1 have been published. Researchers from the same group presented radioligands based on the compound GW405833, [<sup>11</sup>C]2 and [<sup>18</sup>F]3, which have more moderate CB<sub>2</sub>-binding affinities (Table 12.5) (Evens et al. 2011a). Both [<sup>11</sup>C]2 and [<sup>18</sup>F]3 showed good BBB permeability (Table 12.7), in agreement with their molecular properties. Unfortunately, [<sup>18</sup>F]3 had much slower washout of radioactivity from mouse brain than [<sup>11</sup>C]2, which was due to a large fraction of radiometabolites in brain (50% at 30 min after injection), limiting its further development.

The same group (Evens et al. 2009b; Vandeputte et al. 2011) took advantage of the low expression of CB<sub>2</sub> receptors in normal brain by using the CB<sub>2</sub>/[<sup>11</sup>C]2 reporter gene/reporter probe combination as a possible way to follow suitably transfected cells. [<sup>11</sup>C]2 is capable of selective CB<sub>2</sub> imaging in rats with an induced overexpression of the CB<sub>2</sub> receptor, but due to their relatively low binding affinity both [<sup>11</sup>C]2 and [<sup>18</sup>F]3 are unlikely candidates for imaging in neuroinflammatory disorders. Also, [<sup>11</sup>C]2 exhibited low washout and high nonspecific binding in healthy monkey brain (Vandeputte et al. 2011).

A research group from Indiana synthesized a CB<sub>2</sub> ligand [<sup>11</sup>C]4 (Gao et al. 2010) with moderate CB<sub>2</sub>-binding affinity and low CB<sub>2</sub>/CB<sub>1</sub> selectivity (Table 12.5). No animal studies have been published.

The Chiba University PET center presented a series of triaryl CB<sub>2</sub> radioligands (Fujinaga et al. 2010). One of these compounds [<sup>11</sup>C]5 exhibited high binding affinity (Table 12.5). In a mouse study [<sup>11</sup>C]5 manifested good BBB penetration, but its slow brain washout suggests substantial nonspecific binding, which can be explained by the high lipophilicity of this radioligand (Tables 12.5 and 12.7).

The JHU group synthesized [<sup>11</sup>C]A836339 ([<sup>11</sup>C]6) (Horti et al. 2010), a selective CB<sub>2</sub> agonist with high binding affinity, moderate lipophilicity, and an adequate PSA value for CNS activity (Table 12.5). In healthy control CD1 mice [<sup>11</sup>C]6 shows specific binding (Horti et al. 2010) (50%, Table 12.7) in the spleen, an organ with high CB<sub>2</sub> receptor density. In control mice [<sup>11</sup>C]6 exhibited good BBB permeability, and little specific *in vivo* binding in the mouse brain, which is in agreement with low expression of CB<sub>2</sub> receptors in the intact brain (Munro et al. 1993; Schatz et al. 1997). The specific cerebral binding of [<sup>11</sup>C]6 was studied in two animal models of neuroinflammation, a lipopolysaccharide (LPS)-induced mouse model (Qin et al. 2007) and a transgenic amyloid mouse model of AD (APP<sup>swe</sup>/PS1<sup>de9</sup> mice). The dissection study (baseline and blockade (Horti et al. 2010)) showed that [<sup>11</sup>C]6 exhibits high specific binding to CB<sub>2</sub> receptors (78–84%) in LPS-treated mice.



**Table 12.5** Physical-chemical properties of first-generation CB<sub>2</sub> radioligands

Compound	M.W.	clogD <sub>7.4</sub> <sup>a</sup>	PSA	Experimental CB <sub>2</sub> K <sub>i</sub> , nM (CB <sub>2</sub> /CB <sub>1</sub> selectivity)	Reference
[ <sup>11</sup> C]1	540	2.8	158	0.4 (2263)	Evens et al. (2008)
[ <sup>11</sup> C]2	447	4.5 2.5 <sup>b</sup>	44	35 (171)	Evens and Bormans (2010), Evens et al. (2007), Baekelandt et al. (2010)
[ <sup>18</sup> F]3	356	4.7 2.8 <sup>b</sup>	44	27 (370)	Evens and Bormans (2010), Evens et al. (2011a)
[ <sup>11</sup> C]4	393	3.6	60	6.9 (28)	Gao et al. (2010)
[ <sup>11</sup> C]5	395	4.6	39	0.3 (3333)	Fujinaga et al. (2010)
[ <sup>11</sup> C]6, [ <sup>11</sup> C] A836339	310	3.37 ± 0.08 <sup>b</sup>	42	0.7 (450)	Yao et al. (2009), Horti et al. (2010)
[ <sup>11</sup> C]NE40	372	3.9 ± 0.01 <sup>b</sup>	77	9.6 (>100)	Evens et al. (2009a)
[ <sup>18</sup> F]7	398	4.4 <sup>b</sup>	77	3.4	Turkman et al. (2012)

<sup>a</sup>Lipophilicity logD<sub>7.4</sub> and PSA values were calculated with ACD/Structure Designer Suite (Toronto, Canada)

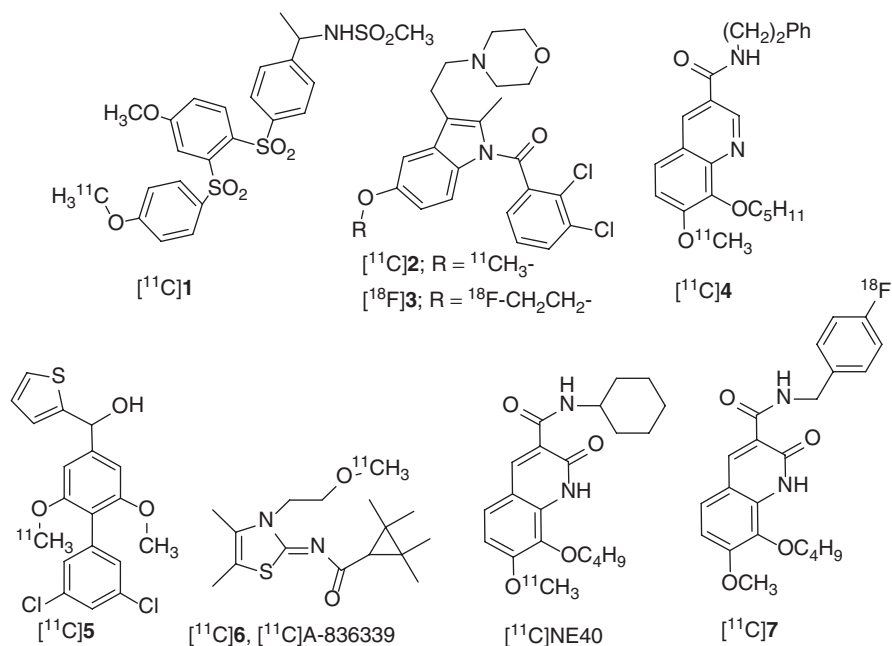
<sup>b</sup>Experimental data

**Table 12.6** Physical-chemical properties of most recent CB<sub>2</sub> radioligands

Compound	M.W.	clogD <sub>7.4</sub> <sup>a</sup>	PSA	Experimental CB <sub>2</sub> K <sub>i</sub> , nM (CB <sub>2</sub> /CB <sub>1</sub> selectivity)	Reference
[ <sup>11</sup> C]RSR-056	418	1.94	89	2.5 (>1000)	Slavik et al. (2015a)
[ <sup>11</sup> C]RS-016	424	2.8	68	0.7 (>10,000)	Slavik et al. (2015b)
[ <sup>11</sup> C]AAT-778	376	6.0	55	4.3 (256)	Haider et al. (2016)
[ <sup>11</sup> C]AAT-015	389	4.11	76	3.3 (303)	Haider et al. (2016)
[ <sup>18</sup> F]RS-126	442	2.74	68	1.2 (>10,000)	Haider et al. (2018)
[ <sup>11</sup> C]RS-028	440	0.94	88	0.8 (>10,000)	Haider et al. (2018)
[ <sup>18</sup> F] (Z)-N-(3-(4-fluorobutyl)-4,5-dimethylthiazol-2(3H)-ylidene)-2,2,3,3-tetramethyl-cyclopropanecarboxamide	325	3.22 ± 0.03	33	0.39 (>1000)	Moldovan et al. (2016)

**Table 12.7** Imaging properties of PET  $CB_2$  radioligands

Radioligand	Spleen uptake (%ID/g)	Brain uptake (%ID/g)	Brain BP (animal model)	Reference
[ $^{11}C$ ]1	–	Low	–	Evens et al. (2008)
[ $^{11}C$ ]2	1 (2 min), 1 (1 h)	1.4 (2 min), 0.1 (1 h)	>1 <sup>a</sup>	Evens and Bormans (2010)
[ $^{11}C$ ]3	0.9 (2 min), 0.7 (1 h)	1.7 (1 h)	–	Evens and Bormans (2010)
[ $^{11}C$ ]4	–	–	–	Gao et al. (2010)
[ $^{11}C$ ]5	n/a	1.9 (1 min), 0.7 (1 h)	–	Fujinaga et al. (2010)
[ $^{11}C$ ]6, [ $^{11}C$ ]A836339	1.3 (1 h)—control 0.6 (1 h)— $CB_2$ block Control/block = 2.2	0.4 (1 h)	1.4 <sup>b</sup> , 2.5 <sup>c</sup>	Horti et al. (2010)
[ $^{11}C$ ]NE40	Control/block = 1.9	0.8 (2 min) 0.1 (60 min)	~3 <sup>a</sup>	Evens et al. (2011b)
[ $^{18}F$ ]7	Insignificant	–	~1 <sup>d</sup>	Turkman et al. (2012)

<sup>a</sup>Rat model of  $CB_2$  overexpression<sup>b</sup>Mouse model of Alzheimer's disease<sup>c</sup>LPS mouse model of neuroinflammation<sup>d</sup>U87/ $CB_2$ -expressing tumor in mice**Fig. 12.14** First generation of  $CB_2$  receptor radioligands

These results were also confirmed in a small-animal PET/[<sup>11</sup>C]6 study with control- and LPS-treated mice (standardized uptake value (SUV) = 0.7 and 2.3, respectively). The cerebral specific binding of [<sup>11</sup>C]6 in the LPS-treated mice is in accordance with a previously found upregulation of CB<sub>2</sub> in this neuroinflammation model (Mukhopadhyay et al. 2006). The very high total cerebral uptake of [<sup>11</sup>C]6 in LPS-treated mice may be associated in part with dysfunction of the BBB (Sumi et al. 2010). Recent study in rat models of neuroinflammation induced by LPS or AMPA showed no specific binding of [<sup>11</sup>C]6 (Pottier et al. 2017). The inconsistency between the [<sup>11</sup>C]6 imaging results in mouse and rat models of neuroinflammation may be due to the species difference and other experimental differences. The species difference in the LPS-induced neuroinflammation models was observed previously in baboon and human PET studies. Thus, the conventional TSPO radiotracer [<sup>11</sup>C]PBR28 showed that human subjects required only a low dose of LPS (1 ng/kg, IV) for quantifiable PET imaging of neuroinflammation (Hannestad et al. 2012). Baboons, which are less sensitive to LPS than humans, required a dose of LPS that was five orders of magnitude greater (0.1 mg/kg, IV) for quantifiable PET with [<sup>11</sup>C]PBR28 (Sandiego et al. 2015).

Brain distribution of [<sup>11</sup>C]6 in the AD mouse model demonstrated that [<sup>11</sup>C]6 displays significant specific CB<sub>2</sub> binding (29–33%) in various brain regions, which is consistent with the distribution of the Aβ amyloid plaques in this mouse model of AD. Previous in vitro studies showed that CB<sub>2</sub> receptors are abundantly and selectively expressed in microglia in human brain tissue from patients with AD, whereas the expression of the CB<sub>1</sub> subtype remains unchanged (Benito et al. 2003). These results suggest the potential of the CB<sub>2</sub> radioligand [<sup>11</sup>C]6 for PET imaging of neuroinflammation in neurodegenerative disorders. The radiotracer [<sup>11</sup>C]6 is currently approved by the FDA for human PET scans under eIND, but the human PET results have not been published yet.

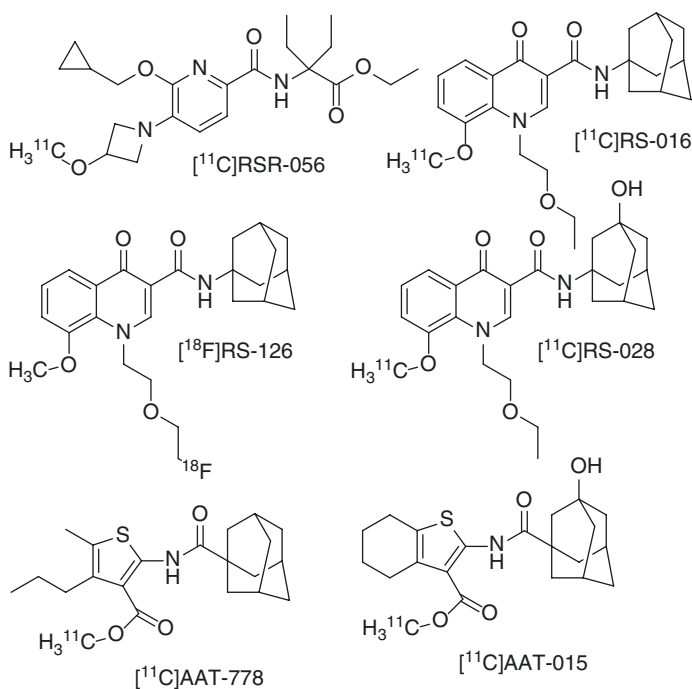
The compound of the KU Leuven group [<sup>11</sup>C]NE40 (Evens et al. 2009a) displays favorable molecular properties for BBB transport, but its binding affinity is moderate (Table 12.5). [<sup>11</sup>C]NE40 was successfully evaluated in a substantial number of preclinical safety studies (Evens et al. 2011b). [<sup>11</sup>C]NE40 showed specific CB<sub>2</sub> receptor binding in the spleen and blood of normal rats and high initial brain uptake in rhesus monkeys. In a rat model with local h-CB<sub>2</sub> receptor overexpression [<sup>11</sup>C]NE40 demonstrated specific and reversible binding to h-CB<sub>2</sub> receptors. [<sup>11</sup>C]NE40 is the first CB<sub>2</sub> receptor PET radioligand that was selected for human PET studies (see below for additional discussion).

A group from MD Anderson Cancer Center has published a series of 2-oxoquinoline derivatives (Turkman et al. 2011), and reported one as a candidate for PET studies (Turkman et al. 2012). Designed to be a more metabolically stable fluorine-18 compound, [<sup>18</sup>F]7 (Log*P* 4.4) had poor uptake in spleen and only about 50% specific binding on CB<sub>2</sub> receptor-positive tumor cells transfected into mouse. Both the preclinical applications and the clinical potential of this compound are limited by its poor solubility.

## 12.4.2 Update of CB<sub>2</sub> PET Tracer Development

The main goal of the many CB<sub>2</sub> receptor studies published after 2013 was to synthesize new PET tracers with very high CB<sub>2</sub> receptor-binding affinities and selectivity and optimal structural properties (see Fig. 12.15 and Table 12.6).

Group from Zurich synthesized [<sup>11</sup>C]RSR-056 with CB<sub>2</sub> receptor  $K_i = 2.5$  nM,  $\log D_{7,4} = 1.94$ , and SI over hCB<sub>1</sub> > 1000 (Slavik et al. 2015a). High specific binding of [<sup>11</sup>C]RSR-056 to CB<sub>2</sub>-positive tissue was demonstrated *in vitro* using rodent spleen sections. In the mouse LPS model of neuroinflammation [<sup>11</sup>C]RSR-056 demonstrated a strikingly increased brain uptake versus control animals, but the blocking with GW405833 showed only a slight reduction of the uptake. The authors concluded that to which extent binding to CB<sub>2</sub> and disruption of blood-brain barrier are involved in [<sup>11</sup>C]RSR-056 accumulation in experimental neuroinflammation remains to be elucidated. The same group also synthesized a 4-oxo-quinoline-based radiotracer [<sup>11</sup>C]RS-016 with binding affinity  $K_i$  of 0.7 nM and  $\log D_{7,4} = 2.8$  (Slavik et al. 2015b). [<sup>11</sup>C]RS-016 exhibited excellent specific binding *in vitro* in rat spleen and human ALS spinal cord. In the LPS neuroinflammation model in rodents [<sup>11</sup>C]RS-016 exhibited specific binding in the range of 10–30% suggesting that [<sup>11</sup>C]RS-016 may be suitable for PET imaging of CB<sub>2</sub> receptors in neuroinflammation. In a separate report (Haider et al. 2018) two more radiotracers of the



**Fig. 12.15** Novel radioligands for CB<sub>2</sub> receptor since 2013

4-oxo-quinoline series ( $[^{18}\text{F}]\text{RS-126}$  and  $[^{11}\text{C}]\text{RS-028}$ ) showed some specific binding by *in vitro* autoradiography in postmortem human ALS spinal cord tissue, but showed no *in vivo* specific binding in Huntington's disease model in rodents. The same group also synthesized a series of thiophene-based  $\text{CB}_2$  receptor ligands with good binding affinities ( $K_i = 3\text{--}4$  nM), but  $^{11}\text{C}$ -labeled compounds of that series failed to demonstrate *in vivo* specific binding in animals (Haider et al. 2016).

Moldovan et al. synthesized a series of fluoro analogues of A836339 with a wide range of  $\text{CB}_2$  receptor-binding affinities ( $K_i = 0.3\text{--}1100$  nM) (Moldovan et al. 2016). Compound (Z)-N-(3-(4-fluorobutyl)-4,5-dimethylthiazol-2(3H)-ylidene)-2,2,3,3-tetramethyl-cyclopropanecarboxamide was selected as the ligand with the highest  $\text{CB}_2$  receptor affinity ( $K_i = 0.39$  nM) and selectivity over those of  $\text{CB}_1$  receptor (factor of 1000). Radiolabeled  $[^{18}\text{F}](\text{Z})\text{-N-(3-(4-fluorobutyl)-4,5-dimethylthiazol-2(3H)-ylidene)-2,2,3,3-tetramethyl-cyclopropanecarboxamide}$  was tested in rodents and demonstrated high brain uptake and 20–30% specific binding in the LPS model of neuroinflammation, but rapid metabolism was a drawback of this radiotracer.

Chemists from Australia developed a series of  $^{18}\text{F}$ -labeled benzimidazole sulfones as  $\text{CB}_2$  receptor PET radioligands with binding affinities in low nanomolar range (Kallinen et al. 2019). The best compounds of the series have been radiolabeled with  $^{18}\text{F}$  and showed 32–44% specific binding in rat spleen, but neuroinflammation animal studies have not been reported.

Lately, several other research groups worked on  $\text{CB}_2$  receptor PET tracer development, but no promising *in vivo* data have been presented yet (Saccomanni et al. 2015; Yrjola et al. 2015; Heimann et al. 2018a, b).

### 12.4.3 $\text{CB}_2$ Imaging Studies

Apart from initial demonstration of *in vivo* potential and proof of target, few  $\text{CB}_2$  imaging studies have been reported. In animals,  $[^{11}\text{C}]\mathbf{6}$  has been shown to have sufficient brain uptake and receptor binding in models of substantial receptor expression; however, the preliminary findings were less favorable in rodent models of Alzheimer's related neuroinflammation (Horti et al. 2010).

In animal studies  $[^{11}\text{C}]\text{NE40}$  was shown to have rapid brain uptake and washout, and relatively rapid plasma metabolism; however, at 10 min after injection approximately 10% of radioactivity in brain was due to radiometabolites (Horti et al. 2010).  $[^{11}\text{C}]\text{NE40}$  has been used to demonstrate a PET reporter gene system in rats using an adeno-associated viral vector, a potentially useful application in gene or stem cell therapy (Vandeputte et al. 2011).

The only PET radioligand for  $\text{CB}_2$  receptors reported in human use is  $[^{11}\text{C}]\text{NE40}$ . In healthy human brain  $[^{11}\text{C}]\text{NE40}$  exhibited rapid uptake and washout (Ahmad et al. 2011). Latest human PET studies with  $[^{11}\text{C}]\text{NE40}$  demonstrated a lack of upregulation of  $\text{CB}_2$  receptors in the brain of AD patients *in vivo* compared with age-matched controls without correlation with regional amyloid deposition. In contrast,  $\text{CB}_2$  receptor availability was significantly lower in AD patients (Ahmad et al. 2016). The authors explained the results as an indication of a relative loss of

neuronal expression rather than absence of microglial expression, therefore making the CB<sub>2</sub> receptor target less promising as a biomarker for neuroinflammation in AD. In addition, the relatively low binding affinity of NE40 ( $K_i = 9.6$  nM, Table 12.5) may be the reason of this result.

[<sup>11</sup>C]NE40 has an effective dose of 4.4 μSv/MBq, as determined in three healthy males. In addition to intestine, liver, and lungs, radioactivity was observed in lymph nodes and spleen, suggestive of CB<sub>2</sub> receptor localization.

To date, no pharmacokinetic modeling from *in vivo* imaging of CB<sub>2</sub> receptors has been reported; however several predictions can be made based on preliminary studies and previous experience with CB<sub>1</sub> receptor imaging. Although CB<sub>2</sub> receptors are found at much lower density in brain than CB<sub>1</sub> receptors, image analysis and pharmacokinetic modeling of CB<sub>2</sub> receptor radioligands may have some potential sources of error. First, similar to CB<sub>1</sub> receptor ligands, CB<sub>2</sub> receptor radioligands are higher in lipophilicity than more commonly used PET tracers, and similar complications as described for CB<sub>1</sub> receptor radioligands should be anticipated. Second, many systemic or local inflammatory conditions can cause disruption of the BBB, causing brain uptake of the radioligand to be affected by flow in addition to specific binding. Third, given the low density of CB<sub>2</sub> receptors normally found in brain, simpler methods for quantifying receptor density might be employed (*e.g.*, reference tissue modeling with pseudoreference region). However, such methods should be validated using compartmental modeling with plasma measurements prior to routine use of alternative methods, with particular attention to states of systemic inflammation, as increased expression of CB<sub>2</sub> receptors may not be a region-specific phenomenon. Fourth, specific binding may occur to circulating leukocytes in addition to nonspecific binding to plasma proteins, potentially complicating measurements of plasma free fraction. As an additional consequence, competition studies with cannabinoid ligands (agonists or inverse agonists) might be expected to have a more significant effect on plasma free fraction due to preferential displacement of radioligand from specific binding sites. Finally, it is unknown if the expression of CB<sub>2</sub> receptors during neuroinflammation or pathological conditions is consistently correlated to any clinical measures, making efforts towards quantitation both a challenge and an exciting prospect.

#### 12.4.4 Potential Clinical Application of CB<sub>2</sub> Imaging

The paucity of reported results for CB<sub>2</sub> receptor imaging with PET is striking, as is its potential as a clinical biomarker. As noted, multiple inflammatory conditions could be probed by a reliable CB<sub>2</sub> receptor radioligand. Multiple sclerosis, ALS, Huntington's disease, and Alzheimer's disease have already been discussed as neuroinflammatory conditions associated with CB<sub>2</sub> receptor upregulation. In particular, *in vivo* and dynamic monitoring of MS in response to treatment with cannabinoid

agonists or other investigatory drugs may be of particular interest. Yet few clinical evaluations of CB<sub>2</sub> receptor PET imaging have been reported (Ahmad et al. 2016; Ahmad et al. 2013). The hurdles in CB<sub>2</sub> receptor radioligand development and translation include (1) differences between humans and rodents in their immune systems, microglia subtypes, and levels of CB<sub>2</sub> receptor expression (Smith and Dragunow 2014), as well as distinct age-related changes (Galatro et al. 2017); (2) low expression level of CB<sub>2</sub> receptors in the brain (Chen et al. 2017); (3) lack of specific CB<sub>2</sub> receptor antibody for postmortem validation of *in vivo* imaging results (England et al. 2015); (4) lack of full CB<sub>2</sub> receptor knockout mice (Constantinescu et al. 2011); and (5) developing a reference region for modeling (Li and Kim 2015). In addition, as age-related inflammation is also a confounding factor in mouse models, these models might not be ideal for evaluating CB<sub>2</sub> receptor tracers (Ni et al. 2019). A greater density of CB<sub>2</sub> receptor-positive microglia and macrophages has been observed in the postmortem spinal cord tissues from patients with ALS compared to healthy controls (Taylor et al. 2016). And similar increases were reported in other studies on ALS spinal cord tissues using [<sup>11</sup>C]RS-028 and [<sup>11</sup>C]KD2 (Mu et al. 2013). Emerging evidence indicates the involvement of CB<sub>2</sub> receptor and increased levels of complements, cytokines, and chemokines in the brain of patients with AD. Reactive astrocytes, activated microglia, and upregulated CB<sub>2</sub> receptor have been observed in the vicinity of Aβ plaques in postmortem AD mouse model brains and patients with AD (Ni et al. 2019).

Vasculitis and cerebritis are often difficult to diagnose, in part due to their frequent normal appearance on conventional imaging, and can be difficult to treat. Thus, CB<sub>2</sub> receptor imaging could potentially provide assistance in both diagnosis and therapy monitoring. A study on rats with a poststroke model demonstrated CB<sub>2</sub> receptor activation related to the neuroprotective responses that likely occur at an early stage of brain disorders (Hosoya et al. 2017). The advent of CB<sub>2</sub> receptor PET radiotracers has enabled the visualization of CB<sub>2</sub> receptor distribution *in vivo* in animal models of central nervous system inflammation; however translation to humans has been less successful. Certain types of cancer have demonstrated CB<sub>2</sub> receptor expression, and *in vivo* imaging could assist in identifying tumor type and grade prior to surgery, and identify tumors susceptible to treatment with cannabinoids to induce apoptosis. Peripherally, CB<sub>2</sub> receptor imaging may assist in deepening an understanding of inflammation in atherosclerotic (Steffens 2008) and cardiovascular diseases (Patel et al. 2010). As described elsewhere (Vandeputte et al. 2011), *in vivo* imaging of CB<sub>2</sub> receptors with a reporter gene/reporter probe system provides an excellent opportunity to monitor gene therapy with PET, and may have other potential applications. Finally, inflammatory states have been associated with mood disorders, such as seen in patients treated with PEGylated interferon, and a genetic polymorphism of the CB<sub>2</sub> receptor has been associated with bipolar disorder (Minocci et al. 2011). Whether CB<sub>2</sub> receptors are a suitable *in vivo* biomarker for mood disorders remains unknown.

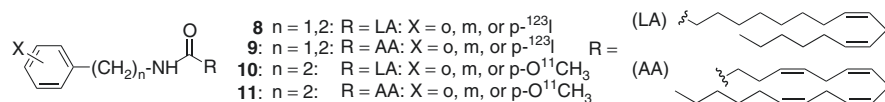
## 12.5 Imaging of FAAH

FAAH is a serine hydrolase enzyme that is responsible for the breakdown of AEA, and is found in the CNS and periphery (Egertova et al. 1998; Thomas et al. 1997). FAAH inhibitors are considered to be an important therapeutic target, allowing a means to increase endocannabinoid tone in a more regionally selective and physiologically relevant manner than administration of exogenous agonists (Otrubova et al. 2011). Clinical applications of FAAH inhibition include treatment of anxiety, PTSD, substance-use disorders (particularly cannabis-use disorder and cannabis withdrawal), depression, Alzheimer's disease, Parkinson's disease, and neuropathic pain (D'Souza et al. 2019; Pertwee 2014; Fowler 2015). As FAAH is found throughout human brain in variable concentration, it is not amenable to a reference region approach for imaging (Romero et al. 2002).

Radioligand design for enzymes is approachable from three different paradigms: reversible binding, tissue or metabolic trapping, and irreversible binding or suicide inactivation (Fowler et al. 2002; Rempel et al. 2017). Reversible binding is most similar to receptor binding and is potentially amenable to compartment kinetic modeling and could measure enzyme availability. Tissue trapping and irreversible binding via suicide inactivation could both potentially inform enzymatic activity as well as tissue distribution. In both cases, molecular design and radiolabel position should be carefully selected to ensure that the enzymatically processed trapped compound retains the radiolabel. The majority of FAAH inhibitors, including the prototypical URB597, are irreversible suicide inhibitors based on either carbamate or urea cleavage by Ser<sub>241</sub> leading to a covalent bond. It should be noted that the FAAH inhibitory potency of unlabeled compounds that was used to predict specific binding of the corresponding radiotracers does not directly correlate with specific binding of these radiotracers. The lack of binding affinity data for the most promising FAAH inhibitors is an obstacle for development of FAAH radioligands for PET. Nevertheless, the latest studies revealed radioligands for imaging FAAH with adequate properties for PET quantification (Wilson et al. 2011a; Li et al. 2011).

### 12.5.1 Tissue-Trapped FAAH Substrates

Initial attempts of a group from Belgium to develop PET- and SPECT-labeled FAAH radioligands have dealt with aryl analogues of anandamide using a tissue trapping approach (Fig. 12.16). Analogue compounds [<sup>123</sup>I]8 and [<sup>123</sup>I]9 (Fig. 12.14) exhibited poor binding to FAAH in mice, likely due to steric hinderance from the



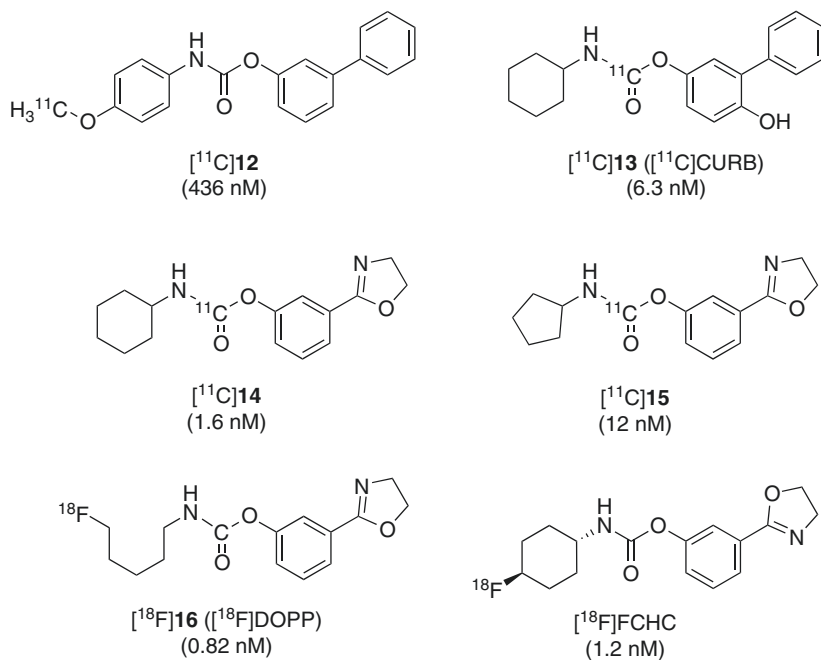
**Fig. 12.16** Radiolabeled analogues of AEA for PET and SPECT imaging of FAAH



iodine, and were abandoned (Wyffels et al. 2009). Compounds [ $^{11}\text{C}$ ]10 and [ $^{11}\text{C}$ ]11 displayed moderate inhibitory potency of AEA metabolism ( $\text{pIC}_{50}$  ranges = 4.34–5.64, 4.46–5.56, respectively) and demonstrated some brain uptake in mice (Wyffels et al. 2009), but no further *in vivo* evaluation has been presented. Another tissue-trapping radioligand reported, [ $^{18}\text{F}$ ]FHEA, was designed and intended for the lysosomal enzyme (and therefore acidic environment functioning) *N*-acylethanolamine-hydrolyzing amidase (NAAA), an enzyme functionally similar to membrane-bound FAAH (preferring basic conditions) in that it can inactivate anandamide, though it is more associated with inactivating palmitoylethanolamide (PEA). [ $^{18}\text{F}$ ]FHEA, a PEA analogue, demonstrated stable brain uptake in mice from 1 to 60 min after injection, though levels in brain were no different after administration of either an NAAA inhibitor or an FAAH inhibitor. However, [ $^{18}\text{F}$ ]FHEA metabolism was reduced following FAAH, but not NAAA, inhibition, suggesting that [ $^{18}\text{F}$ ]FHEA is at least a partial substrate for FAAH (Pandey et al. 2014).

## 12.5.2 Irreversible FAAH Inhibitors

In a follow-up attempt from the same Belgian group, [ $^{11}\text{C}$ ]-4-methoxyphenylcarbamate (12, Fig. 12.17), a radiolabeled analogue of URB597, inhibited the hydrolysis of AEA with an  $\text{IC}_{50}$  value of 436 nM (compared to 40 nM for that of URB597) and



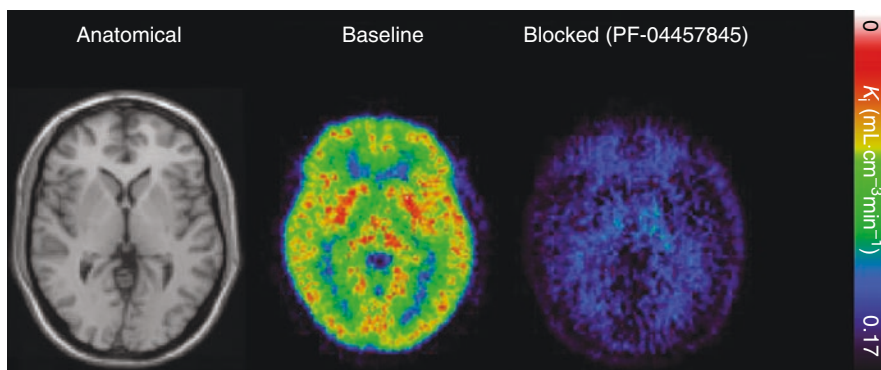
**Fig. 12.17** Radiolabeled carbamate-based FAAH irreversible inhibitors and analogues of URB597, with their  $\text{IC}_{50}$

manifested good initial brain uptake (anticipated with good lipophilicity of  $\log D = 2.27$ ), but no specific binding was observed in mouse biodistribution studies (Wyffels et al. 2010). It was concluded that this negative result is due to insufficient affinity of [ $^{11}\text{C}$ ]12.

Researchers from Toronto have reported a series of  $^{11}\text{C}$ -radiolabeled analogues of URB597 with low-nanomolar inhibitory potency values (Wilson et al. 2011a, b; Wilson et al. 2013). [ $^{11}\text{C}$ ]CURB (Fig. 12.17) (aka, URB694) showed high brain uptake in rats (SUV of 1.5–2.5 at 5 min) with no washout, which is characteristic of irreversible binding. Regional distribution of radioactivity was found to be high in the cortex, intermediate in the cerebellum, and low in the hypothalamus, which corresponds to the known distribution of FAAH in the rat brain (Egertova et al. 1998; Thomas et al. 1997). Pretreatment with unlabeled URB694 reduced binding in all brain regions by 62–86%, and URB597 reduced binding in brain by 72–88%. It was concluded that [ $^{11}\text{C}$ ]CURB represents a highly promising radiotracer for imaging FAAH using PET (Wilson et al. 2011b).

Human PET studies using [ $^{11}\text{C}$ ]CURB have demonstrated its ability to quantify FAAH activity in CNS. Following injection of [ $^{11}\text{C}$ ]CURB, radioactivity in brain peaked at  $\sim 4.3$  SUV within 2–3 min, and after a washout period of  $\sim 30$  min remained relatively stable at levels  $\sim 25$ – $33\%$  below that of peak for the remaining 90-min scan, consistent with irreversible binding. Radioactivity quickly declined in plasma, and the parent fraction quickly declined over the first several minutes and then slowly from 20 min (48%) to 90 min (37%). The free fraction of [ $^{11}\text{C}$ ]CURB is low, at  $\sim 0.9\%$ . Data were well described with good identifiability using the 2-TCM with irreversible binding (2-TCMi;  $k_4 \equiv 0$ ), with as little as 60 min of data. Though some high-uptake regions had better model fitting in a 2-TCM with reversible binding and most others in a three-tissue compartment model with irreversible binding, they were not statistically superior to the 2-TCMi and had poor identifiability in some microparameters. Binding was not limited by cerebral blood flow, and simulation studies predicted that  $\lambda k_3$  is the optimal parameter to quantify FAAH binding (with  $\lambda = K_1/k_2$ , or  $V_T$  of ligand in the free and nonspecific compartment at equilibrium, available for binding). The Patlak analysis for irreversible kinetics (*e.g.*, [ $^{18}\text{F}$ ]FDG as a prototypical application) was unsuccessful, likely due to the late time it could be implemented (40 min) with the remaining data containing too much noise (Rusjan et al. 2013). Subsequent analysis demonstrated that voxel-wise modeling was achievable for high-resolution data (HRRT camera; Fig. 12.18) using a basis function model (BAFPIC for 2-TCMi with arterial input function), though with higher bias and variability than ROI data outcomes, but  $\sim 60\%$  reduced variability compared to Patlak plot derivations (Rusjan et al. 2018). [ $^{11}\text{C}$ ]CURB has an effective dose of  $5.2 \mu\text{Sv/MBq}$  in women, and  $4.6 \mu\text{Sv/MBq}$  when estimated in both men and women, with highest radioactivity uptake in liver, gallbladder, and small intestine, within the range of other carbon-11 radioligands (Boileau et al. 2014).

[ $^{11}\text{C}$ ]CURB has been further examined in clinical studies. It has excellent reproducibility with test-retest variability of 9%, and good reliability with ICC of 0.79. The C385A single-nucleotide polymorphism (SNP) of the FAAH gene (rs324420), known to result in reduced FAAH expression (*i.e.*, loss of function) with



**Fig. 12.18** [ $^{11}\text{C}$ ]CURB in human brain. Axial slice of reference MRI (right, MNI  $z = +2$  mm), and of [ $^{11}\text{C}$ ]CURB parametric maps of  $K_1$  at baseline condition (middle, averaged of  $n = 6$ ) and 2 h after an oral dose  $\geq 1$  mg of PF-04457845 (right). Figure courtesy of Dr. Pablo Rusjan, University of Toronto

corresponding increase of anandamide, has been associated with increased risk for trait impulsiveness, cannabis sampling, and higher reward reactivity, but lower anxiety, and cannabis liking, craving, and withdrawal. [ $^{11}\text{C}$ ]CURB demonstrated lower binding in brains of A allele holders ( $\sim 20\%$  in A/C heterozygotes,  $\sim 50\%$  in the lone A/A homozygote imaged) compared to C/C homozygotes, as would be predicted with the loss of function and decreased FAAH expression (Boileau et al. 2015a). [ $^{11}\text{C}$ ]CURB was able to estimate brain FAAH occupancy of the high-potency FAAH inhibitor PF-04457845 across 1 mg, 4 mg, and 20 mg, quantified using  $\lambda k_3$  and 60 min of scan data, yielding a  $\sim 91\%$  reduction (Fig. 12.18). Following blockade, the time-activity curves had a reversible appearance, and the 1-TCM fit equally well as the 2-TCMi, confirming that the first compartment of the latter model represents non-displaceable binding. Since the non-displaceable represents  $< 10\%$  of  $\lambda k_3$ , FAAH occupancy was therefore estimated as  $> 95\%$ , confirming high potency and occupancy of FAAH by PF-04457845 at 1 mg (Boileau et al. 2015b). In cannabis users ( $n = 10$ ), compared to healthy controls ( $n = 22$ ), [ $^{11}\text{C}$ ]CURB reflected 14–20% reduced FAAH binding, which was not correlated with clinical characteristics of cannabis use (years of use, days and amount used per week, craving, withdrawal symptoms), or influenced by cigarette or alcohol use, or global cerebral blood flow. High cannabinoid and associated metabolite concentrations in blood and urine and greater impulsivity traits correlated with lower FAAH binding (Boileau et al. 2016). Among the explanations for lower FAAH binding in cannabis users 1 day after abstinence include presence of metabolites binding or modulating FAAH, related downregulation of  $\text{CB}_1$  receptors, compensation for downregulated AEA, or suppression of immune cells given identification of FAAH in microglia. However, increased translocator protein (TSPO) binding imaged in cannabis users suggests an increase of neuroinflammation, casting this last possibility in doubt (Da Silva et al. 2019).

Additional clinical studies with [ $^{11}\text{C}$ ]CURB have been briefly reported. Despite an association between C385A with obesity and metabolic syndrome, neither [ $^{11}\text{C}$ ]CURB binding nor peripheral endocannabinoid levels were different between overweight healthy subjects ( $n = 10$ ) and age-matched controls ( $n = 20$ ) (Best et al. 2017). Similarly, those with PTSD ( $n = 7$ ) had no difference in [ $^{11}\text{C}$ ]CURB binding compared to healthy controls ( $n = 30$ ); however, [ $^{11}\text{C}$ ]CURB binding in amygdala did correlate across both groups with fMRI BOLD response to fearful faces, suggesting the role of endocannabinoid tone in fear processing more generally. In contrast to the finding of reduced peripheral anandamide levels in PTSD patients (see above Sect. 12.3.4 (Neumeister et al. 2013)), there was a significant elevation in peripheral 2-AG, but no difference in anandamide levels in those with PTSD compared to controls (Isabelle Boileau et al. 2018). Across healthy subjects ( $n = 51$ ), [ $^{11}\text{C}$ ]CURB binding in prefrontal cortex was negatively correlated with impulsivity (total Barratt Impulsiveness Scale), and in whole brain, prefrontal cortex, and ventral striatum with the self-control factor (Esmaeil Mansouri et al. 2018). Young heavy drinkers (age 19–25) with ( $n = 14$ ) or without ( $n = 17$ ) a family history of alcohol-use disorder had no difference in FAAH brain levels; however across groups lower FAAH levels (and therefore increased anandamide levels) were correlated with greater Alcohol Use Disorder Identification Test (AUDIT) scores and lower subjective sedative effects of alcohol (Best et al. 2018).

Several analogues of CURB ([ $^{11}\text{C}$ ]14, [ $^{11}\text{C}$ ]15, [ $^{18}\text{F}$ ]16, or [ $^{18}\text{F}$ ]DOPP; Fig. 12.17) also demonstrated high specific FAAH uptake in the brain of rats and nonhuman primates (Wilson et al. 2011a; Wilson et al. 2013; Sadovski et al. 2013; Rotstein et al. 2014; Shoup et al. 2015). [ $^{18}\text{F}$ ]DOPP, synthesized with a multistep procedure with alkyl radiolabeling followed by coupling reaction, demonstrated good uptake in rat brain and greater than that of [ $^{11}\text{C}$ ]CURB, in a distribution consistent with FAAH, to which the  $^{18}\text{F}$ -radiolabel was bound irreversibly; notably, the compound was stable in human, but not rat, plasma, possibly due to the presence of species-specific carboxylesterases (Sadovski et al. 2013). Follow-up studies in nonhuman primate demonstrated good brain uptake (2.2–2.8 SUV), kinetic modeling best fit with 2-TCMi using  $\lambda k_3$  as an outcome measure consistent to that described with [ $^{11}\text{C}$ ]CURB, rapid metabolism, and significantly reduced binding after blocking with URB597 suggesting specificity to FAAH (Rotstein et al. 2014). No data in human subjects has yet been reported.

[ $^{18}\text{F}$ ]DOPP was utilized in a rat study to investigate the *in vivo* potency of BIA 10–2474, a putative FAAH inhibitor that caused one death and five serious neurological injuries during a Phase I dose escalation study. *Ex vivo* measures of [ $^{18}\text{F}$ ]DOPP demonstrated BIA 10–2474 to have high potency at FAAH in rats with  $\text{IC}_{50} \sim 65 \mu\text{g}/\text{kg}$ , comparable to other FAAH inhibitors (e.g., URB597 at  $0.15 \mu\text{g}/\text{kg}$ , URB694 aka CURB at  $40 \mu\text{g}/\text{kg}$ , PF-04457935 at  $30\text{--}100 \mu\text{g}/\text{kg}$ ); in contrast the pre-trial reported that  $\text{IC}_{50}$  for BIA 10–2474 was  $1.1\text{--}1.7 \text{ nM}$ , reflecting a much lower potency (Tong et al. 2017). It has since been understood that several off-target mechanisms at high doses likely led to its toxicity, possibly inhibition of other serine hydrolases or disruption of neuronal lipid metabolism (Chaikin 2017); preclinical occupancy studies might have identified FAAH saturation at lower doses and potentially prevented the tragic trial outcomes.

A second [ $^{18}\text{F}$ ]fluorine-labeled carbamate radioligand, [ $^{18}\text{F}$ ]FCHC ( $\text{IC}_{50} = 1.2 \text{ nM}$ ; Fig. 12.17), has been evaluated in preclinical studies. [ $^{18}\text{F}$ ]FCHC is radiosynthesized in a one-pot, three-step process with 10–20% radiochemical yield. Both *ex vivo* and *in vivo* studies demonstrated that [ $^{18}\text{F}$ ]FCHC uptake in rat brain was initially rapidly followed by a slow increase to  $>5 \text{ SUV}$  in high-FAAH-rich regions, which was about twice as high as low regions, and was  $>95\%$  blocked by URB597. Metabolism was initially rapid (75% parent at 2 min) with no lipophilic metabolites and 16% parent remaining by 40 min. [ $^{18}\text{F}$ ]FCHC has higher brain uptake (about two times greater than [ $^{18}\text{F}$ ]DOPP and about four times greater than [ $^{11}\text{C}$ ]CURB), greater specific binding, and faster blood clearance than other carbamate FAAH inhibitor radioligands (Shoup et al. 2015). Despite the promising characteristics of [ $^{18}\text{F}$ ]FCHC and [ $^{18}\text{F}$ ]DOPP, no fluorine-18 radioligand for FAAH has yet advanced into human studies.

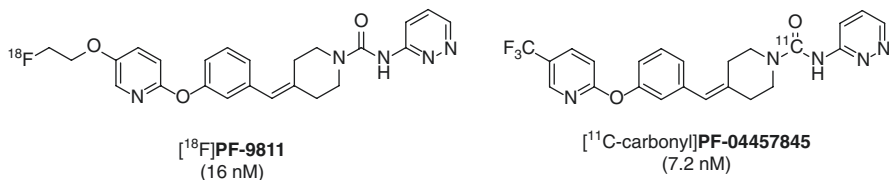
Scientists from Chiba (Japan) recently developed two high-binding-affinity FAAH PET tracers [ $^{11}\text{C}$ ]MFTC ( $\text{IC}_{50} = 0.34 \text{ nM}$ ,  $\text{clogD} = 3.6$ ) (Kumata et al. 2015) and [ $^{11}\text{C}$ ]DFMC ( $\text{IC}_{50} = 6.1 \text{ nM}$ ,  $\text{clogD} = 3.43$ ) (Shimoda et al. 2016). [ $^{11}\text{C}$ ]MFTC, a [ $^{11}\text{C}$ -carbonyl]carbamate, was synthesized via [ $^{11}\text{C}$ ]phosgene and [ $^{11}\text{C}$ ]carbonate intermediate in a two-step reaction with 20% radiochemical yield, had good and rapid uptake that plateaued in mouse and rat brain in a distribution consistent with FAAH, and was  $>80\%$  blocked by cold MFTC or URB597 in rat brain. In nonhuman primates, [ $^{11}\text{C}$ ]MFTC had rapid uptake in a FAAH regional distribution, and slow washout over 90 min that suggests nonspecific binding. However, while cold MFTC reduced radioactivity by 70–80% with homogenous regional time-activity curves, URB597 reduced radioactivity by 50–60% with heterogenous regional time-activity curves, suggesting the presence of off-target binding (Kumata et al. 2015). [ $^{11}\text{C}$ ]DFMC, a urea-based FAAH inhibitor, was synthesized via [ $^{11}\text{C}$ ]methylation with 20% radiochemical yield, and had rapid and high uptake in mouse (*ex vivo*) and rat (*in vivo* PET) brain in a FAAH distribution. However, uptake was disproportionately high in the cerebellar nucleus (2.4 SUV) compared to other FAAH-containing regions (1.6–2.0 SUV), though after blockade by URB597 60–70% of radioactivity was reduced in brain with nearly homogenous regional time-activity curves suggesting specificity for FAAH. Brain extracts largely confirmed this with  $\sim 95\%$  of radioactivity in brain at 30 min being irreversibly bound to brain tissue, which dropped to  $<2\%$  after pretreatment with URB597 (Shimoda et al. 2016).

A group from Pfizer has released a series of publications on a novel FAAH inhibitor (Ahn et al. 2011; Johnson et al. 2010) and its closely related radiotracer, [ $^{18}\text{F}$ ]PF-9811 (Fig. 12.19). Using a three-step one-pot reaction sequence [ $^{18}\text{F}$ ]PF-9811 can be attained in high yield ( $\sim 11\%$ ). Using an activity-based protein profiling probe for FAAH, [ $^{18}\text{F}$ ]PF-9811 achieved an  $\text{IC}_{50} = 16 \text{ nM}$  (compared to  $\text{IC}_{50} = 7.2 \text{ nM}$  for PF-04457845). Following injection of [ $^{18}\text{F}$ ]PF-9811, distribution of radioactivity was as previously described for FAAH and [ $^{11}\text{C}$ ]CURB in rat brain. Both *in vitro* and *in vivo* measurements of [ $^{18}\text{F}$ ]PF-9811 were reduced after pretreatment with PF-04457845 in rats, with  $\sim 40\%$  reduction of SUV as measured by PET after 90 min (Skaddan et al. 2012).

Structurally similar [ $^{11}\text{C}$ -carbonyl]PF-04457845 (Fig. 12.19) was reported by the Toronto group. Radiolabeling was accomplished by [ $^{11}\text{C}$ ]CO $_2$  fixation to the urea carbonyl position in a one-pot reaction with radiochemical yield  $\sim 4.5\%$ . It demonstrated good imaging properties with appropriate lipophilicity ( $\text{Log}P_{7.4} = 3.48$ ), good brain uptake consistent with FAAH distribution in rats and approximately double that achieved by [ $^{11}\text{C}$ ]CURB, and an estimated  $>95\%$  specific binding in brain (Hicks et al. 2013). As a therapeutic compound, PF-04457845 has demonstrated an ability to reduce cannabis withdrawal symptoms (D'Souza et al. 2019), but not pain related to osteoarthritis of the knee (Huggins et al. 2012).

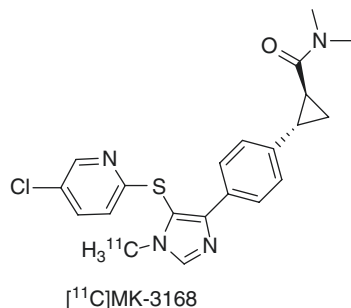
### 12.5.3 Reversible FAAH Inhibitors

The Merck PET group has presented data on [ $^{11}\text{C}$ ]MK-3168 (Fig. 12.20), a radiolabeled inhibitor of FAAH with reversible binding (Li et al. 2011; Liu et al. 2013). The compound demonstrated a favorable *in vitro* profile with  $\text{IC}_{50} = 1.0$  nM and  $K_D$  0.8 nM (both human cortex), lipophilicity by HPLC  $\log D$  of 3.3,  $>1000$ -fold selectivity over other targets, and lack of rat and human P-gp sensitivity. Autoradiographic studies in rhesus monkey and human brain slices showed highest binding in striatum, frontal cortex, and hippocampus, consistent with FAAH distribution. PET studies of [ $^{11}\text{C}$ ]MK-3168 in rhesus monkeys confirmed this uptake pattern with total to nonspecific signal  $\sim 2:1$  and pre-administration of a FAAH inhibitor provided evidence that *in vivo* binding of [ $^{11}\text{C}$ ]MK-3168 is mediated by FAAH (Liu et al. 2013). Kinetic modeling in monkeys showed 2-TCM  $V_T$  of 3–4 mL/cm $^3$ , and small



**Fig. 12.19** Radiolabeled urea-based FAAH irreversible inhibitors and analogues of PF-0445784, with their  $\text{IC}_{50}$

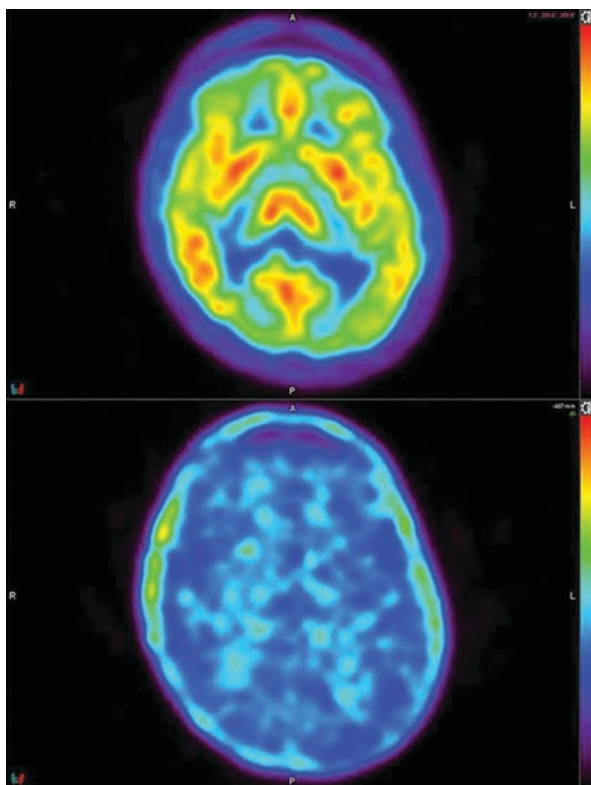
**Fig. 12.20** [ $^{11}\text{C}$ ]MK-3168, first FAAH PET tracer for imaging in human subjects



changes (<15%) in white matter between baseline and blocked scans provided  $BP_{ND}$  0.4–1 with a modest test-retest variability at <20% (Joshi et al. 2012). Human [ $^{11}C$ ]MK-3168 PET demonstrated high brain uptake (peak SUV  $\sim$ 3–3.5), rapid metabolism with parent radioligand comprising 30–40% of radioactivity at 10 min and essentially eliminated at 60 min after injection, 2-TCM  $V_T$  of 14–20 mL/cm<sup>3</sup>, good test-retest variability (<12%), and acceptable radiation burden for PET studies (effective dose of 4.65  $\mu$ Sv/MBq) (Li et al. 2011; Joshi et al. 2012; Postnov et al. 2018). [ $^{11}C$ ]MK-3168 was used to assess FAAH occupancy by another inhibitor, JNJ-42165279, first in rhesus macaque, and then in humans using a 2-TCM, demonstrating that 10 mg achieved >95% occupancy (Fig. 12.21) (Postnov et al. 2015, 2018).

A second reported reversible FAAH inhibitor, [ $^{11}C$ ]MPPO, is based on an  $\alpha$ -ketoheterocyclic scaffold that promotes binding at the serine hydrolase site without cleavage. [ $^{11}C$ ]MPPO has high binding at FAAH (*in vitro*  $IC_{50}$  = 10 nM), greater than 500-fold selectivity over triglycerol hydrolase, and acceptable lipophilicity ( $\log D_{7.4}$  = 3.43). [ $^{11}C$ ]MPPO was radiolabeled with  $^{11}C$  with 13% radiochemical yield and demonstrated moderate brain uptake (0.6–0.8 SUV) with FAAH distribution and rapid washout in *ex vivo* mouse studies and *in vivo* PET rat studies, generating time-activity curve profiles consistent with reversible binding. In rats,

**Fig. 12.21** Distribution of [ $^{11}C$ ]MK-3168 in human brain under baseline and blocked conditions. Top image: [ $^{11}C$ ]MK-3168 PET scan at baseline. Bottom image: [ $^{11}C$ ]MK-3168 PET scan in the same volunteer after receiving 10 mg JNJ-42165279. Figure reproduced from Postnov et al. (2018)



pretreatment with URB597 had little effect on [ $^{11}\text{C}$ ]MPPO brain uptake, but abolished regional uptake heterogeneity and increased washout, suggesting that uptake and/or binding is modest but specific for FAAH. The relatively low PET signal may have been due to the very rapid metabolism of the radioligand, with hydrophilic radiometabolites comprising 75% and 90% of radioactivity in brain and plasma at 15 min, respectively. [ $^{11}\text{C}$ ]MPPO was assessed for sensitivity to P-gp and Bcrp efflux transporters, and while wild-type controls had higher brain uptake compared to P-gp/Bcrp knockout mice, the amount was considered insufficient to account for the limited brain uptake (Wang et al. 2016a).

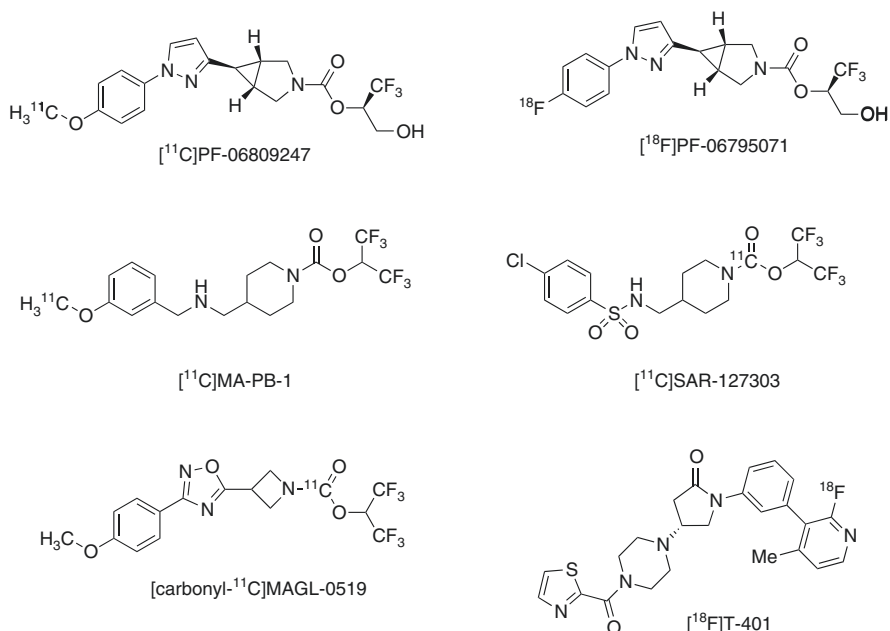
## 12.6 Imaging of MAGL

Similar to inhibition of FAAH, monoacylglycerol lipase (MAGL), an enzyme responsible for the breakdown of ~85% of 2-AG, has been the target for pharmacologic inhibitor and radioligand development. MAGL is also a serine hydrolase enzyme and is co-localized with CB<sub>1</sub> receptors in axon terminals. MAGL inhibitors may have application in anxiety disorders, traumatic brain injury, substance-use disorders, Alzheimer's disease, and Parkinson's disease (Pertwee 2014; Fowler 2015).

Several PET tracers for imaging MAGL have been published in recent years. An initial evaluation of several candidate radiotracers by the Toronto group based on known MAGL inhibitors (KML29, JJKK-048, ML30, JW642) and five novel analogues led to [ $^{11}\text{C}$ ]carbonyl radiolabeling, similar to that performed for FAAH radiotracers, of the most promising candidates based on favorable affinity ( $\text{IC}_{50}$  range 0.36–5.9 nM) and selectivity. The two hexafluoroisopropanol carbamates and three triazole urea evaluated *ex vivo* in rodents had modest nonspecific brain uptake, but only [ $^{11}\text{C}$ ]JJKK-048 had reduced uptake (by 42%) after preblockade with unlabeled compound. Several compounds also demonstrated reduced retention in blood after preblockade, possibly due to plasma and leukocyte MAGL which could confound attempts at designing brain-penetrant MAGL radiotracers (Hicks et al. 2014), and potentially affect kinetic modeling estimates that do not account for a vascular compartment.

The Leuven group modified JW642 to create [ $^{11}\text{C}$ ]MA-PB-1 ( $\text{IC}_{50} = 26$  nM,  $\text{clog}D = 3.4$ ; Fig. 12.22), which has similar MAGL affinity but lower and more acceptable lipophilicity. [ $^{11}\text{C}$ ]MA-PB-1 has high brain uptake in mice (*ex vivo* SUV 2.6, decreasing to 1.6), which was ~70% blocked by itself and MJN110, an unrelated MAGL inhibitor, with fast metabolism and no significant MAGL binding in blood. *In vivo* PET imaging in rats demonstrated similar kinetics, albeit with lower peak uptake (~1.6 SUV) and washout to a lower plateau from 30 to 90 min (~0.6), which was blocked ~60% by pre-administration of MJN110, but not by challenge administration, consistent with [ $^{11}\text{C}$ ]MA-PB-1 having irreversible binding at MAGL. The tracer was advanced to nonhuman primate imaging studies, in which it demonstrated high uptake (~4.2 SUV) with washout to plateau by 20 min (~2.5 SUV), and could be blocked by ~≥88% by MJN110 (Ahamed et al. 2017).





**Fig. 12.22** Radioligands for MAGL

A high-potency carbamate with structurally distinct sulfonamido-piperidine platform was selected as a candidate radiotracer. SAR127303 ( $IC_{50} = 36$  nM,  $\log D = 3.69$ ; Fig. 12.22) and the novel urea-based TZPU ( $IC_{50} = 39$  nM,  $\log D = 1.85$ ) were selected from a series of analogues, with SAR127303 having better selectivity for MAGL over FAAH. *Ex vivo* studies in mice demonstrated [<sup>11</sup>C]TZPU to have poor brain uptake, while [<sup>11</sup>C]SAR127303 had rapid and persisting brain uptake from 1 to 60 min after injection. *In vivo* PET imaging reflected these results, with [<sup>11</sup>C]TZPU achieving  $\sim 0.4$  SUV in rat brain which was reduced by  $\sim 48\%$  with unlabeled TZPU, and not substantially accounted for by P-gp/Bcrp. In contrast, [<sup>11</sup>C]SAR127303 achieved  $\sim 1.5$  SUV in rat brain throughout the 60-min scan, in a distribution pattern consistent with MAGL expression (cortex, striatum > cerebellum > hippocampus >> pons), and could be  $\sim 50$ – $67\%$  reduced by self-block with loss of regional heterogeneity, suggesting that  $\sim 1/3$  of radioactivity is from nonspecific binding. [<sup>11</sup>C]SAR127303 was partially reduced after pretreatment with MAGL inhibitor KML29, but not after JZL184 (consistent with prior literature) or FAAH inhibitor URB597 (as anticipated for MAGL selectivity). [<sup>11</sup>C]SAR127303 uptake was not altered by P-gp inhibitor cyclosporin A suggesting lack of P-gp substrate affinity, and could not be competed off by unlabeled SAR127303, confirming irreversible binding in rat brain (Wang et al. 2016b, c). Follow-up assessment and kinetic modeling with arterial input function in rats demonstrated that [<sup>11</sup>C]SAR127303 binding could be quantified with the 2-TCMi (similar to FAAH, described above) with good identifiability of microparameters, and provided

supporting evidence that radiometabolites do not enter brain. Results with 2-TCMi were highly correlated with those obtained by Patlak values, and  $K_i$  from both 2-TCMi and Patlak as well as  $\lambda k_3$  from 2-TCMi, but not  $k_3$ , all correlated with *in vitro* binding data; however,  $\lambda k_3$  had higher variability making  $K_i$  the preferred outcome parameter. Thus, Patlak  $K_i$  parametric images of [ $^{11}\text{C}$ ]SAR127303 in rat brain could be used to reflect MAGL availability (Yamasaki et al. 2018). Advancement of [ $^{11}\text{C}$ ]SAR127303 to nonhuman primate imaging provided encouraging results, with sufficient (peak  $\sim 1$  SUV) and persistent brain uptake from 20 to 90 min (consistent with a profile of irreversible binding) in a MAGL distribution (Wang et al. 2016c).

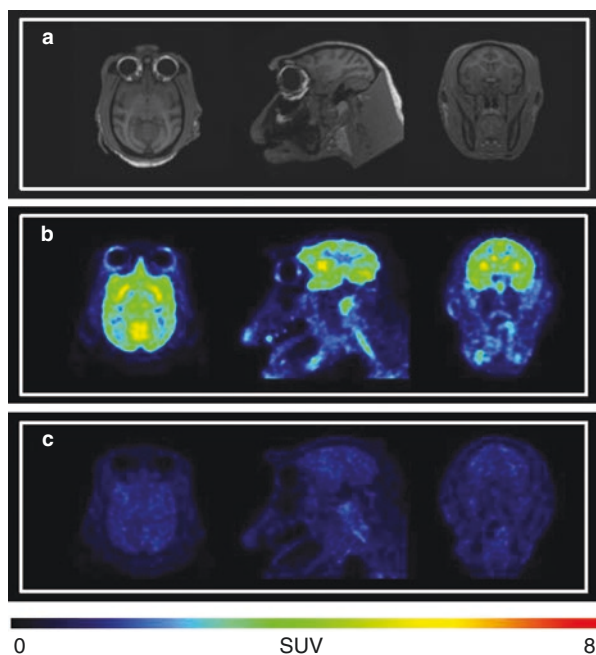
A series of azetidine scaffold compounds were developed as a novel class of MAGL radiotracers in an effort to lower lipophilicity. MAGL-0519 ( $\text{IC}_{50} = 12.7$  nM,  $\text{Log}P = 1.23$ ) was selected as the most favorable candidate, as it had the highest affinity for MAGL and selectivity over FAAH (30- to 50-fold), with no P-gp substrate affinity by *in vitro* assay. [Methyl- $^{11}\text{C}$ ] and [carbonyl- $^{11}\text{C}$ ]MAGL-0519 (Fig. 12.22) were assessed by PET in rat brain and demonstrated nearly identical kinetic profiles, with rapid heterogenous uptake (peak  $\sim 1.5$  SUV), followed by gradual washout for 90 min, and could be blocked, but not displaced, by unlabeled MAGL-0519 or KML29 ( $\sim 40$ – $50\%$  reduction), and had no significant reduction with pretreatment of FAAH inhibitor URB597. Although blocking and challenge studies were consistent with irreversible binding at MAGL, the slow washout of both [ $^{11}\text{C}$ ]methyl- and [ $^{11}\text{C}$ ]carbonyl-radiolabeled could not otherwise be explained by nonspecific binding or radiometabolites ( $<10\%$ ), and therefore likely reflected ongoing dissociation between the azetidine carbonyl and the MAGL serine residue. Initial PET studies in nonhuman primates showed [carbonyl- $^{11}\text{C}$ ]MAGL-0519 to have similar kinetics with high heterogenous uptake ( $\geq 1.5$  SUV), with gradual washout over 90 min ( $\sim 0.5$  SUV) (Cheng et al. 2018). In a separate attempt, another azetidine-based [ $^{11}\text{C}$ ]carbonyl-radiolabeled tracer (1,1,1,3,3,3-hexafluoropropan-2-yl-3-(1-benzyl-1H-pyrazol-3-yl)azetidine-1-[ $^{11}\text{C}$ ]carboxylate) was evaluated with improved affinity and reasonable lipophilicity ( $\text{IC}_{50} = 0.41$  nM,  $\text{Log}D = 3.69$ ). The radioligand was shown to be irreversibly bound in rat brain tissue, have high brain uptake in mice (*ex vivo*) and rats (*in vivo* PET) in a distribution consistent with MAGL, and be  $\sim 40\%$  reduced by pretreatment with unlabeled compound or MAGL inhibitor JW642, but not during challenge in rats. However, it also demonstrated washout from brain over time, similar to MAGL-0519 (Mori et al. 2019).

In a follow-up effort, another series of azetidines were designed for more favorable properties, including candidates as reversible binding radioligands for MAGL. An irreversible binding carbamate with high affinity (MAGL-2-11,  $\text{IC}_{50} = 0.88$  nM,  $\text{Log}D = 1.9$ ) and two reversibly binding compounds, designed by altering and stabilizing the leaving group (PAD,  $\text{IC}_{50} = 2.7$  nM,  $\text{Log}D = 3.35$ ; MAGL-4-11,  $\text{IC}_{50} = 11.7$  nM,  $\text{Log}D = 2.7$ ), were evaluated. These three new compounds are selective for MAGL, and *in silico* studies predicted MAGL-2-11 and PAD to have full inhibition of MAGL, while MAGL-4-11 had only partial docking in the binding site but would block 2-AG from entering the catalytic site. PET imaging of [carbonyl- $^{11}\text{C}$ ]PAD in rats showed high and heterogenous brain uptake (2.2

SUV) with very little washout over 90 min, and was ~50% blocked with KML29, suggesting improved irreversible kinetics compared to [ $^{11}\text{C}$ ]MAGL-0519. PET imaging of [ $^{11}\text{C}$ ]PAD and [ $^{18}\text{F}$ ]MAGL-4-11 in rats showed poor brain uptake (~0.4 and ~0.3 SUV, respectively), though substantial uptake was seen in brown adipose tissue that could be substantially blocked by KML29, suggesting peripheral specificity for MAGL. P-gp/Bcrp knockout mice had high brain uptake of both [ $^{11}\text{C}$ ]PAD and [ $^{18}\text{F}$ ]MAGL-4-11 compared to wild-type mice with plateaued kinetics, demonstrating that the inadequate blood-brain barrier permeability for CNS imaging is due to ABC efflux transporters (Chen et al. 2019a).

A Pfizer and Karolinska Institute collaborative effort identified [ $^{11}\text{C}$ ]PF-06809247 (Fig. 12.22) as an irreversible binding PET radioligand for MAGL. Initial *in silico* assessment identified PF-06809247 from reported and Pfizer library MAGL inhibitors as having optimal properties including blood-brain barrier penetration and lipophilicity ( $\text{Clog}D = 2.7$ ), and measured assays confirmed high affinity ( $\text{IC}_{50} = 13 \text{ nM}$ ), low cross-species affinity differences, >1000-fold selectivity for MAGL over FAAH, and very good selectivity overall (Zhang et al. 2019). *Ex vivo* assessments in rats demonstrated ligand uptake consistent with brain and peripheral organ MAGL distribution that was reduced by another MAGL inhibitor. PET imaging in nonhuman primates showed [ $^{11}\text{C}$ ]PF-06809247 to have high, heterogenous uptake (~3.0–3.5 SUV) peaking at 40 min and with minimal washout to 120 min, and could be ~90% blocked by MAGL inhibitor pretreatment (Fig. 12.23). [ $^{11}\text{C}$ ]PF-06809247 has entered into human testing for test-retest reliability; however results have not yet been reported (ClinicalTrials.gov Identifier: NCT03100136). Very shortly after the report of [ $^{11}\text{C}$ ]PF-06809247, another collaborative group assessed its analogue [ $^{18}\text{F}$ ]PF-06795071 (Fig. 12.22) as a [ $^{18}\text{F}$ ]fluorine-labeled PET radioligand for MAGL. Compared to its analogues also assessed, PF-06795071 possesses good affinity and lipophilicity ( $\text{IC}_{50} = 2.3 \text{ nM}$ ,  $\text{Log}D = 3.3$ ), with high selectivity, and demonstrated irreversible binding by *in vitro* assay. Autoradiography of [ $^{18}\text{F}$ ]PF-06795071 in rat brain resulted in 26–49% specific binding when blocked by itself or KML29 in a MAGL distribution. *In vivo* PET in rat brain showed good uptake (~1.75 SUV) and moderate washout over 90 min, which was 56% and 61% blocked by pretreatment with KML29 and unlabeled PF-06795071, respectively (Chen et al. 2019b).

[ $^{18}\text{F}$ ]T-401 (Fig. 12.22) is the first reversibly binding MAGL PET radioligand reported to successfully enter brain. [ $^{18}\text{F}$ ]T-401 ( $\text{IC}_{50} = 4.0 \text{ nM}$ ,  $\text{Log}D = 1.86$ ) was designed following the discovery of piperazinyl pyrrolidine-2-one scaffold of reversible MAGL compounds, and selected from a series of analogues due to its favorable binding affinity, lipophilicity, good mouse brain uptake by *ex vivo* assessment, and low P-gp affinity. [ $^{18}\text{F}$ ]T-401 immediately peaks (~0.6 SUV) followed by rapid washout with heterogenous brain uptake in mouse brain, with very little retention in MAGL knockout mice. T-401 was demonstrated to be selective and reversibly bound to MAGL by *in vitro* assays. In monkey brain, [ $^{18}\text{F}$ ]T-401 had similar kinetics as seen in mice, and could be blocked by unlabeled compound; an analogue developed alongside [ $^{18}\text{F}$ ]T-401 demonstrated poor washout and increasing brain radioactivity over time, leading to the selection of [ $^{18}\text{F}$ ]T-401 as the preferred



**Fig. 12.23** [ $^{11}\text{C}$ ]PF-06809247 in nonhuman primate brain. MRI (a) and PET images summed from 0 to 123 min under baseline (b) and blocking condition with (c) pretreatment of MAGL inhibitor. Reprinted with permission from Zhang et al. Copyright (2019) American Chemical Society (Zhang et al. 2019)

radiotracer. The estimated effective dose of [ $^{18}\text{F}$ ]T-401 as derived from monkey data is 25  $\mu\text{Sv}/\text{MBq}$ , in range for human use; [ $^{18}\text{F}$ ]T-401 has reportedly entered into human testing; however no additional details are yet available (Hattori et al. 2019).

## 12.7 Imaging of TRPV1

Although TRPV1 receptors are not currently considered part of the ECS, they function in close relation to each other as evidenced by cannabinoids and anandamide (which is sometimes even referred to as an endovanilloid) acting meaningfully as agonists. To date, four radioligands for TRPV1 have been reported. The first, [ $^{11}\text{C}$ ]SB366791, a cinnamide-derived antagonist developed at GlaxoSmithKline with good selectivity (Gunthorpe et al. 2004), had relatively poor binding affinity ( $K_D = 280 \pm 56$  nM in human TRPV1 expressing CHO), acceptable lipophilicity ( $\text{LogD} = 3.5$ ), and an increase of proportional radioactivity due to metabolites from 2 to 10 min in mouse brain (15–33% of total radioactivity), but appropriate biodistribution and good retention in target-rich trigeminal nerve (TGN) (van Veghel et al. 2013a). In follow-up, [ $^{11}\text{C}$ ]DVV24 (a trifluoromethyl- for chloro- substitution of

[<sup>11</sup>C]SB366791) had improved binding affinity ( $K_D = 163 \pm 28$  nM) and about ~2.5 times mouse TGN-to-blood binding ratio compared to blocked conditions, but rapid washout from mouse brain and a similar increase of radiometabolites from 2 to 10 min (~10–45% of total radioactivity) (van Veghel et al. 2013b). [<sup>18</sup>F]DVV54, an aminoquinazoline, demonstrated similar binding affinity ( $K_D = 171 \pm 48$  nM) and slower clearance from blood, but poorer brain and TGN uptake attributed to its high lipophilicity (cLogD 6.51) and PSA (50.70) (van Veghel et al. 2013b). Carbon-11-labeled capsaicin accumulated in rat tissues known to express TRPV1, including low, but measurable, uptake in brain, suggesting capsaicin as a potential platform for generating analogue radioligands (Goto et al. 2015). Most recently, the TRPV1 antagonist BCTC, known to have high selectivity and high CNS penetration, was labeled with carbon-11 and found to have higher ex vivo TGN-to-blood ratio and specific binding than the other radioligands (Kumata et al. 2017), though no report of brain uptake was made.

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## 12.8 Conclusion and Future Directions

Herein we have described the utility of ECS imaging with PET, its historical development, the current understanding and experience with it, and its potential applications and developments in the future. Akin to its description as a nonclassical neurotransmitter system, it has pushed the boundaries of classical PET radioligand and pharmacokinetic techniques. Given the established roles of the ECS in physiological function, it is not surprising that direct *in vivo* imaging has provided a window into basic workings of the normal brain and a wide variety of neuropsychiatric diseases. It is therefore expected to continue yielding additional new insights for those, as well as for inflammatory, immunological, and non-CNS diseases.

Progress in the medicinal chemistry of cannabinoid drugs stimulated discovery of radiotracers for imaging the ECS. Most PET imaging studies on the ECS have been done with CB<sub>1</sub> receptor radioligands. There are at least five radiolabeled CB<sub>1</sub> receptor inverse agonists ([<sup>11</sup>C]OMAR, [<sup>18</sup>F]MK-9470, [<sup>11</sup>C]MePPEP, [<sup>18</sup>F]FMPEP-*d*<sub>2</sub>, and [<sup>11</sup>C]SD5024) that are suitable for PET imaging in human subjects. All of these radioligands are not free of drawbacks and better compounds are under development. Combination of excellent specific signal, high brain uptake, appropriate brain kinetics, absence of active radiometabolites, and radiochemical availability are a must for a successful radioligand for PET imaging in humans, but this is a tough nut to crack. The high density of CB<sub>1</sub> receptors in brain, which has proved a challenge, masks the true physiology of the receptor system; a series of radioligands that can distinguish agonist-preferring coupled receptors or their associated second messenger systems would provide additional information on the pathophysiology that has already been probed. A useful radiolabeled CB<sub>1</sub> receptor agonist for PET imaging in humans remains to be discovered and is an unmet need.

Successful development and human study of CB<sub>2</sub> receptor radioligands remain a challenge. Several research groups are actively working on the development of CB<sub>2</sub>

PET radioligands as potential probes for imaging inflammation in various disorders. Very low expression of CB<sub>2</sub> in the intact CNS dictates a necessity for a CB<sub>2</sub> PET radioligand with very high binding affinity likely requiring subnanomolar to picomolar range. So far only one radioligand, [<sup>11</sup>C]NE40, has been studied and reported in humans, but its binding affinity is rather moderate considering the expected requirement of very high affinity. The major obstacle in the development of CB<sub>1</sub> and CB<sub>2</sub> receptor radioligands is a very high lipophilicity of compounds with sufficiently high binding affinities. While CB<sub>2</sub> receptor imaging remains with substantial promise for numerous CNS diseases, the lackluster and mixed results to date may call into question this target as a useful biomarker of neuroinflammation when contrasted with other well-established imaging targets such as TSPO. However, its potential as a druggable target, similar to the neuroinflammatory targets cyclooxygenases and P2X<sub>7</sub> receptor, may reveal yet unrealized dividends in developing novel treatments for debilitating neuropsychiatric disorders.

The recent report of several FAAH and MAGL radioligands advanced into human study has opened new avenues for understanding the ECS. Imaging of FAAH, a pharmacological target with exceptional promise, has the potential to broaden our understanding of the regulation of anandamide. As well, using PET to quantify the dose to percent enzyme saturation of investigational drugs demonstrates the power of this *in vivo* imaging technique, as sadly illustrated in preclinical studies following the tragedy of the BIA 10–2474 trial. Imaging MAGL, an equally promising target for drug development, may improve the understanding of 2-AG regulation, which given its ~1000-fold higher concentration than anandamide in brain (Buczynski and Parsons 2010) may provide more salient understanding to the *in vivo* regulation of ECS neurotransmission.

Non-cannabinoid targets involved in the ECS, such as TRPV1, are underexplored targets for radioligand development, and likewise represent exciting possibilities for creating tools for probing this nearly ubiquitous and far-reaching system. Radioligands for GPR18, GPR55, and the putative endocannabinoid transporter remain essentially unexplored and would add to our knowledge of endocannabinoid physiology and cannabinoid pharmacology.

The complexity of the ECS and its seemingly unending connectivity to organ and biochemical systems afford it a vast potential for future development and investigations for *in vivo* imaging. As the depths of the ECS continue to be sounded, the drive for novel radioligands remains an exciting pursuit.

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# Current Radioligands for the PET Imaging of Metabotropic Glutamate Receptors

# 13

Linjing Mu and Simon M. Ametamey

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## Abstract

Glutamate receptors are divided into two main groups: ionotropic glutamate receptors (iGluRs) and metabotropic glutamate receptors (mGluRs). Modulation of iGluRs and mGluRs has potential for the treatment of psychiatric and neurological diseases such as depression, anxiety, schizophrenia, and Parkinson's disease. Positron emission tomography (PET) might offer the possibility to visualize glutamate receptors and presents an interesting tool for studying these receptors under physiologic and pathologic conditions. MGLuR PET radioligands reported till August 2019 are summarized, and also a summary of some of the most important prerequisites for CNS PET radioligands is provided. Emphasis is given to mGluR1 and mGluR5, the two receptor subtypes for which most advances in radioligand development and human studies have been made. In addition, recent research efforts toward the development of PET radioligands for other receptor subtypes more specifically, mGluR2/mGluR3 and mGluR4 are also outlined alongside their *in vitro/in vivo* properties and clinical applications where available.

## Abbreviations

ABP688	3-(6-methyl-pyridin-2-ylethynyl)-cyclohex-2-enonemethyl-oxime
AMPA	2-amino-3-(3-hydroxy-5-methyl-isoxazol-4-yl)-propionic acid
BBB	Blood-brain barrier
$B_{\max}$	Maximal binding capacity
$BP_{\text{ND}}$	Binding potential relative to the no displaceable compartment
Bq	Becquerel
BSA	Bovine serum albumin
CAMP	Cyclic adenosine monophosphate
CNS	Central nervous system
DMF	Dimethylformamide
DMSO	Dimethyl sulfoxide
DV	Distribution volume
EOB	End of bombardment
EOS	End of synthesis
$\text{Et}_3\text{N}$	Triethylamine
GPCR	G-protein-coupled receptor
HBD	Hydrogen bond donors
HPLC	High-pressure liquid chromatography
<i>i.v</i>	Intravenous
$\text{IC}_{50}$	Inhibition constant required for displacement of 50% of radioligand binding
ID	Injected dose
IDnorm./g	Injected dose normalized to body weight per tissue weight

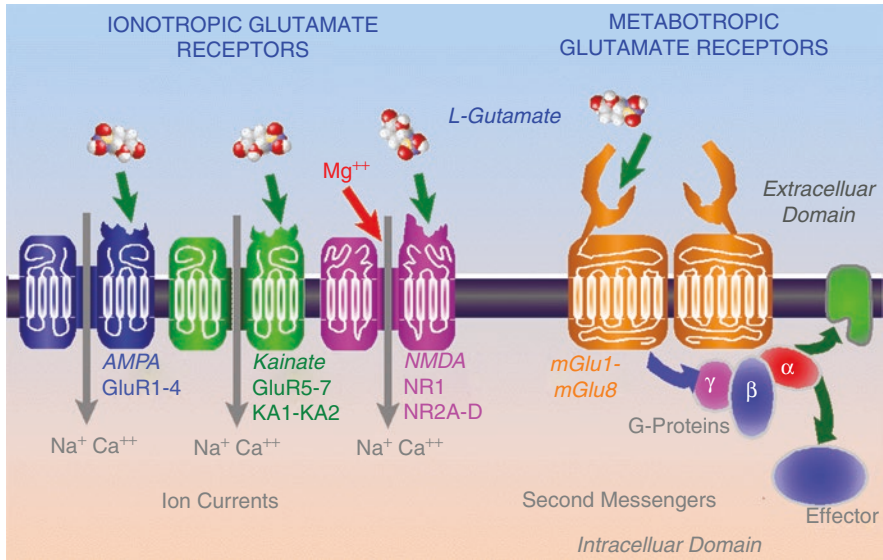
iGluR	Ionotropic glutamate receptor
$K_d$	Dissociation constant
$K_i$	Inhibition constant
ko	Knockout
LPS	Lipopolysaccharide
LTP	Long-term potentiation
MeI	Methyl iodide
M-FPEP	2-methyl-6-(3-fluoro-phenylethynyl)-pyridine
mGluR	Metabotropic glutamate receptor
mGluR5	Metabotropic glutamate receptor subtype 5
min	Minute(s)
M-MPEP	2-methyl-6-((methoxyphenyl)ethynyl)-pyridine
MPO	Multi-parameter optimization
MW	Microwave
NAC	<i>N</i> -acetylcysteine
NMDA	<i>N</i> -methyl-D-aspartate
NMDAR	<i>N</i> -methyl-D-aspartate receptor
p.i.	Postinjection
PBS	Phosphate-buffered saline
PET	Positron emission tomography
P-gp	P-glycoprotein
PI	Phosphoinositol
PLC	Phospholipase C
ROI	Region of interest
RT	Room temperature
SAR	Structure-activity relationship
SUV	Standard uptake value
TAC	Time activity curve
TEMPO	2,2,6,6-Tetramethylpiperidine-1-oxyl
THF	Tetrahydrofuran
TM	Transmembrane domain
TPSA	Topological polar surface area
wt	Wild-type

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## 13.1 Introduction

### 13.1.1 Glutamate Receptors

Glutamate (L-glutamic acid) serves as the neurotransmitter at the majority of excitatory synapses in the mammalian central nervous system (CNS). Glutamate receptors are divided into two main groups, namely, ionotropic glutamate receptors (iGluRs) and metabotropic glutamate receptors (mGluRs) (Fig. 13.1).



**Fig. 13.1** Ionotropic and metabotropic glutamate receptors (Source: <http://www.ucl.ac.uk/ioo/research/salt/glutam.htm> (Prof. Tom Salt))

The iGluRs are the principal mediators of excitatory neurotransmission. These are ligand-gated cationic-selective channels which are permeable to Na<sup>+</sup>, K<sup>+</sup>, and Ca<sup>2+</sup> ions. They are subdivided into three major classes based on structural similarities and named according to the type of synthetic agonist that activates them. These three selective agonists are NMDA (*N*-methyl-D-aspartate), AMPA ( $\alpha$ -amino-3-hydroxy-5-methyl-isoxazole-4-propionate), and kainate (Bettler and Mulle 1995; Mori and Mishina 1995). Native receptors of all of these families are likely heteromeric assemblies comprising more than one type of subunit. *N*-methyl-D-aspartate (NMDA) tetrameric receptors consist of subunits NR1 and NR2. There are four NR2 subunit types, NR2A, B, C, and D. AMPA receptors are composed of a four-subunit family (GluR1–4) with members that are products of separate genes and are believed to assemble as functional tetramers. Kainate receptors are composed of two related subunit families, GluR5–7 and KA-1 and 2.

Metabotropic glutamate receptors (mGluRs) are G-protein-coupled receptors (GPCR), which activate intracellular secondary messenger systems when bound by the physiological ligand glutamate. From a structural viewpoint, mGluRs belong to the seven-transmembrane protein family and have been classified into three subgroups. Group I mGluRs (mGluR1 and mGluR5) are coupled to phospholipase C and upregulate or downregulate neuronal excitability (Gereau and Conn 1995). Group II (mGluR2 and mGluR3) and group III (mGluR4 and mGluR6–8) inhibit adenylate cyclase and hence reduce synaptic transmission (Bradley et al. 2000; Ritzen et al. 2005). The mGluRs are expressed on neuronal and glial cells, with each receptor subtype exhibiting distinct spatial and temporal expression profiles in the

brain (Shigemoto et al. 1997; Thomas et al. 2001), with the exception of mGluR6 existing in the retina. In neurons, group I mGluRs are mainly localized in somatodendritic domains and postsynaptically regulate neuronal excitability and synaptic transmission via several intracellular second messenger systems, whereas group II and III mGluRs are predominantly localized in axonal domains and axon terminals to presynaptically regulate neurotransmitter release (Cartmell and Schoepp 2000). mGluRs function as dimers, with two glutamate molecules being required for full receptor activation (Shin et al. 2008; Tsuchiya et al. 2002).

### 13.1.2 PET

Positron emission tomography (PET) is a noninvasive imaging technology, allowing for the quantitative imaging of physiologic and pathophysiologic processes in vivo (Ametamey et al. 2008; Piel et al. 2014). High resolution of about 1–2 mm can be achieved with small animal imaging systems and 4–6 mm for clinical PET scanners. The visualization of biological targets and systems is of special interest in this research field. Specific PET tracers for labeling such targets are potential future diagnostics of various diseases and provide elegant tools for drug development (Fowler 1999), especially for the development of drugs with less side effects. Besides oncologic and cardiac applications, there is a growing demand for specific and selective radiopharmaceuticals in the field of neuroscience. PET can be used to image receptor distribution, concentration, and functions in normal and pathological states and has proven to be a powerful technique in neurological research (Passchier et al. 2002). Due to the convenient half-lives of carbon-11 and fluorine-18 and their chemical properties, most effort has been put into the development of tracers labeled with one of these isotopes. In general, carbon-11 and fluorine-18 are produced by nuclear reactions which require the bombardment of convenient stable isotopes with high energetic protons provided by a cyclotron. Carbon-11 can easily be introduced into biologically interesting compounds without changing their biochemical and pharmacological properties. The relatively short physical half-life of  $^{11}\text{C}$  (20 min) allows for multiple imaging studies in a short period of time, which makes it advantageous for sequential investigations with short time intervals in the same individual (animal or human), thereby allowing the subject to serve as its own control. Therefore  $^{11}\text{C}$ -labeled compounds can be used for instance in a receptor occupancy paradigm or in cognitive intervention studies where several PET investigations are usually needed within a few hours apart from each other. The short half-life of  $^{11}\text{C}$ , however, is disadvantageous for commercial production. Fluorine-18 has long physical half-life of 110 min allowing more complex synthesis, extended in vivo investigation, and, most importantly, satellite distribution of the radiotracer to clinical PET centers that do not have radiochemistry facilities. In addition, it shows better imaging characteristics than carbon-11 due to its lower positron energy (0.64 MeV vs. 0.96 MeV for  $^{11}\text{C}$ ). The larger the positron energy, the larger the average distance the positron travels

before annihilating and as such more loss in spatial resolution will be expected. Thus, efforts have been put into developing  $^{18}\text{F}$ -labeled radiotracers. Radiolabeling methods with either carbon-11 or fluorine-18 will not be discussed in this chapter. Several review papers on this topic can be found in the literature (Dahl et al. 2017; Deng et al. 2018; van der Born et al. 2017).

### 13.1.3 General Requirements for CNS Radiotracers

The general requirements for successful PET radiotracers for the imaging of specific brain targets such as transporters, receptors, and enzymes also hold true for glutamate receptors. Several excellent review papers have been published on this topic (Eckelman et al. 2006; Van de Bittner et al. 2014; Waterhouse 2003). A summary of some of the most major points is provided below.

#### 13.1.3.1 High Affinity and Selectivity

The affinity of the radiotracer relative to the concentration of binding site present in the target tissue is an important characteristic of site-specific radiotracers (Eckelman et al. 2006). The ability to image a receptor is determined by delivery, by retention at the specific binding site, and by washout of nonspecifically associated radiotracer. Two terms are related to the affinity of a ligand toward its target, and depending on how the value is obtained, the  $K_d$  (equilibrium dissociation constant) and the  $K_i$  (inhibition constant) express the affinity of a ligand to the target. Usually, binding affinities should be in the low nanomolar to subnanomolar range. With regard to selectivity, the radioligand in question should have the highest affinity for the actual target and only low affinity for any other binding site.

#### 13.1.3.2 Concentration of Target Sites ( $B_{\max}$ )

A high ratio of binding site concentration toward radiotracer concentration is a requirement for a valid response to changes in binding site concentration caused by diseases or drug occupancy during a PET experiment. Therefore, it is important that the  $B_{\max}$  value exceeds the affinity ( $K_d$ ) of the radioligand, and this positively affects the target-to-nontarget ratio and the imaging quality. It is considered that PET tracers with an in vitro  $B_{\max}/K_d$  ratio (also called binding potential) of at least ten have a high chance of providing a specific signal in vivo. A high level of nondisplaceable binding in vivo may require an in vitro  $B_{\max}/K_d$  much greater than ten to provide a useful specific signal in vivo (Eckelman and Mathis 2006).

#### 13.1.3.3 BBB Permeability

Lipophilicity is a measure of how much of a compound partitions into a lipid environment compared to an aqueous environment. CNS compounds have to pass the blood-brain barrier (BBB) and ideally should therefore have  $\log P$  values well above 1 to allow free diffusion across the BBB. Nonspecific binding is known to be higher with increasing lipophilicity, and high nonspecific binding in target tissues reduces the signal-to-noise ratio and, thus, the ability to detect the specific signal (Pike 2009). It has been recommended that the  $\log P$  values of potential CNS compounds

should be between 1 and 3.5 (Waterhouse 2003; Clark 2003). It should also be pointed out that lipophilicity values (normally  $\log P$  or  $\log D$ ) are highly variable, depending on the methodology used for their determination. Values obtained using the traditional “shake-flask” method tend to be lower than those obtained by HPLC, C18, and calculation methods.  $\log P$  values determined using *in silico* methodology are also highly variable, depending on the algorithm used. Therefore, it has been suggested that  $\log P$  should not be relied upon as a sole predictor of BBB penetration during brain radiotracer discovery. HPLC measurement of permeability, percentage of plasma protein binding, and membrane interactions may be potentially useful in predicting a radioligand’s *in vivo* performance (Tavares et al. 2012). Additionally, the BBB possesses efflux pumps which can prevent compounds from effectively accumulating in the brain, of which P-glycoprotein (P-gp) is a key efflux pump (Ohe et al. 2003). Therefore, a PET tracer should not be a substrate for P-gp. There are *in vitro* assays to determine whether a compound is a P-gp substrate or not (Wanger-Baumann et al. 2011).

#### 13.1.3.4 In Vivo Stability

The formation of radioactive metabolites which might enter into the brain and confound the brain signals is undesired. Unfavorable is also the metabolic elimination of the radioactive nuclide from the radiotracer. Thus, the radioligand should be resistant to unfavorable metabolism during the acquisition time, and radioactive metabolites should not be taken up in the target area.

#### 13.1.3.5 Efficient Radiosynthesis

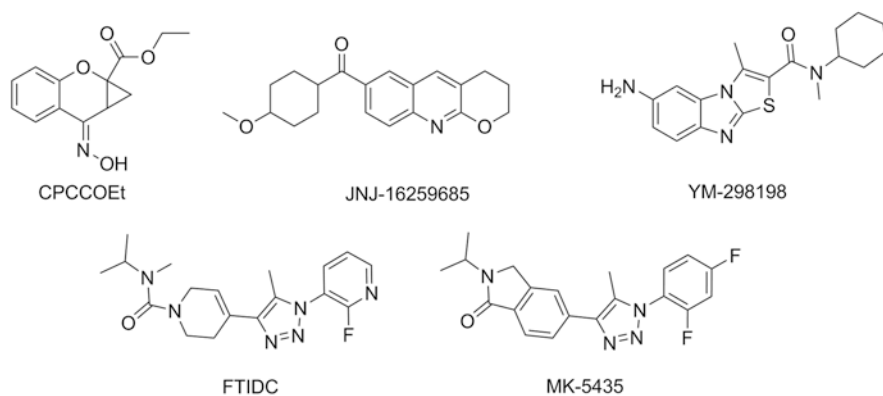
Due to the short half-lives of common PET nuclides, a rapid and efficient radiolabeling process is crucial. The best radiosynthetic scheme should incorporate the radionuclide at the last synthetic step. Typically, the reaction time for synthetic schemes with multiple steps should be accomplished within two half-lives of the radioisotope. The establishment of a one-step radiosynthesis is desired for obtaining high radiochemical yields with high molar activities. The ratio between labeled and unlabeled tracer is defined by the molar activity. Low molar activities result in low-quality PET images especially for saturable systems such as enzymes, transporter proteins, and receptors. For receptors with low binding sites (low  $B_{\max}$ ), very high molar activity is required in order to exclude substantial occupation of target sites by unlabeled ligand and the saturation of the biological system of interest.

It should be stressed that these criteria do not necessarily guarantee a successful glutamate radiotracer. They are merely guidelines to assist in the selection of potentially useful CNS imaging ligands. Recently, multiparameter optimization methods have been used to assess ligands which have central nervous drug-like properties. The parameters are (1) lipophilicity, calculated distribution coefficient ( $\text{Clog}P$ ); (2) calculated distribution coefficient at  $\text{pH} = 7.4$  ( $\text{Clog}D$ ); (3) molecular weight (MW); (4) topological polar surface area (TPSA); (5) number of hydrogen bond donors (HBD); and (6)  $\text{p}K_a$  (Wager et al. 2010a; Wager et al. 2010b; Zhang et al. 2013). This multiparameter optimization method has successfully been used to develop CNS PET tracers (Zhang et al. 2013; Chen et al. 2016; Zhang et al. 2017a).

When a potential radiotracer has been selected and a radiolabeling method has been developed, some preclinical evaluations and acute toxicity studies have to be performed prior to PET studies in humans. The characterization of a new radioligand includes autoradiography, biodistribution, and metabolite studies as well as pretreatment and displacement studies, using *in vitro*, *in vivo*, and *ex vivo* techniques. With the increasing ability to carry out PET studies in rodents, using dedicated small animal PET cameras, the evaluation of PET tracers is simplified and serves as an important bridge between laboratory and clinical science (Myers 2001; Xi et al. 2011; Xu and Li 2019).

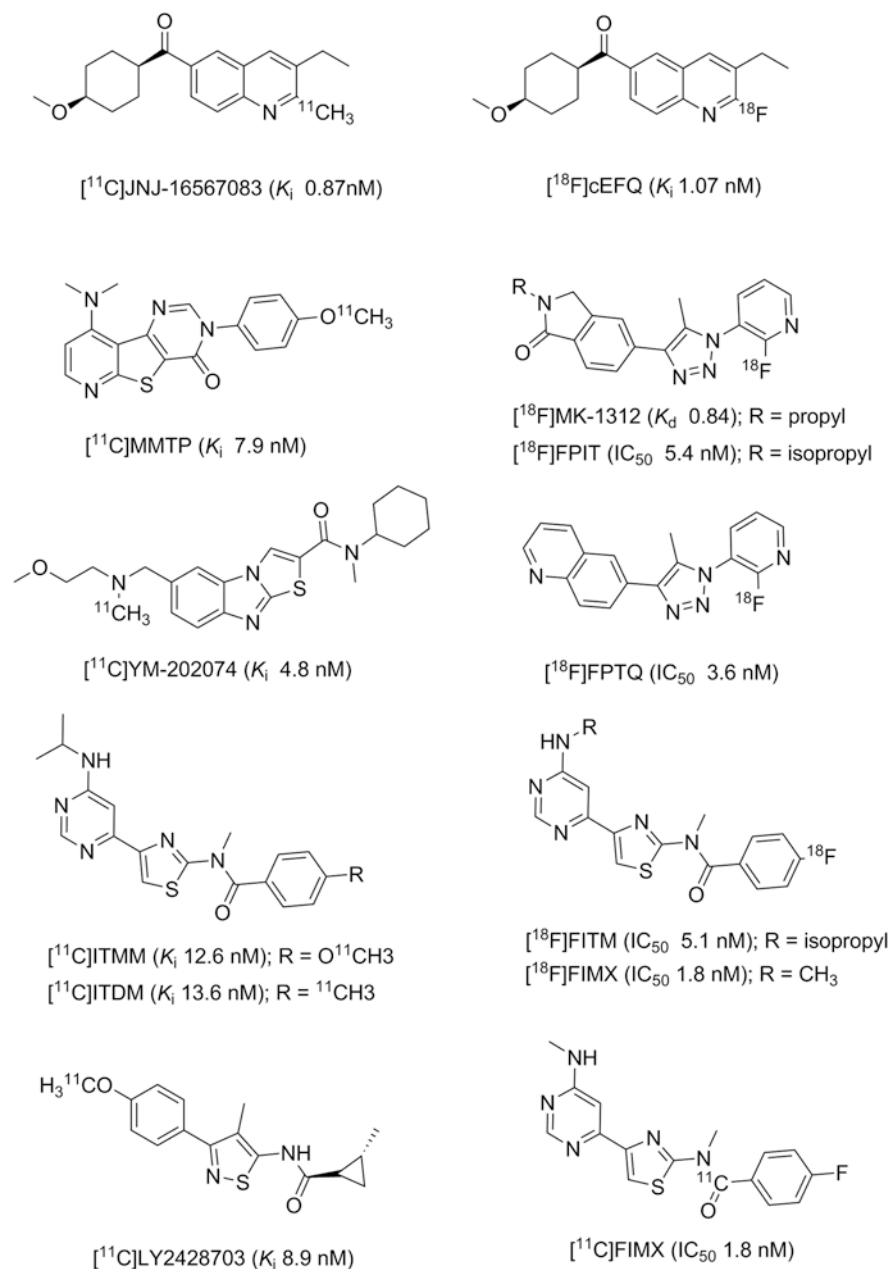
### 13.2 mGluR1 PET Tracers

mGluR1 has been implicated in neuroprotection, chronic and neuropathic pain, multiple sclerosis, motor dysfunction, epilepsy, cerebral ischemia, and cerebellar long-term depression (Ito et al. 2009; Kohara et al. 2008; Yamasaki et al. 2017). Hence, monitoring mGluR1 *in vivo* and noninvasively would lead to a better understanding of the role of this receptor in the pathophysiology of biological processes mediated through glutamatergic pathways. It has been reported that the highest expression of mGluR1 in rat brain is in the cerebellum; moderate or low level in the thalamus, striatum, hippocampus, and cerebral cortex; and very low level in the brain stem (Fotuhi et al. 1993; Lavreysen et al. 2004a). Since noncompetitive ligands do not bind to the evolutionarily conserved glutamate binding site, they offer the potential for improved selectivity for mGluR subtypes (Parmentier et al. 2000). Therefore, potent and selective allosteric mGluR1 antagonists such as CPCCOEt (Annoura et al. 1996), JNJ-16259685 (Lavreysen et al. 2004b), YM-298198 (Kohara et al. 2005), FTIDC (Suzuki et al. 2007), and MK-5435 (Ito et al. 2009) have been developed (Fig. 13.2). Subsequently, several radiolabeled



**Fig. 13.2** Structures of some mGluR1 allosteric antagonists





**Fig. 13.3** Structures of some important mGluR1 PET tracers

**Table 13.1** Summary of affinity, lipophilicity values, and in vivo specificity of mGluR1 radioligands

Radioligand	Binding affinity (nM)	Lipophilicity (LogD)	Specificity in the brain
[ <sup>11</sup> C] JNJ-16567083	K <sub>i</sub> 0.87 (4 °C); 3.49 (37 °C) <sup>a</sup> K <sub>i</sub> 0.57 (4 °C); 4.41 (37 °C) <sup>b</sup> K <sub>i</sub> 4.75 (4 °C); 13.3 (37 °C) <sup>c</sup>	3.38	No (rat, baboon)
[ <sup>11</sup> C]MMTP	K <sub>i</sub> 7.9 <sup>b</sup> ; IC <sub>50</sub> 9.5 <sup>c</sup>	3.15	Not clear
[ <sup>11</sup> C] YM-202074	IC <sub>50</sub> 4.8 <sup>c</sup>	2.7	Yes (rat), no (monkey and human)
[ <sup>11</sup> C] LY2428703	IC <sub>50</sub> 8.9 <sup>c</sup>	4.02	Yes (rat), no (monkey and human)
[ <sup>11</sup> C]ITMM	K <sub>i</sub> 12.6 <sup>b</sup>	2.57	Yes (rat and human)
[ <sup>11</sup> C]ITDM	K <sub>i</sub> 13.6		Yes (monkey)
[ <sup>18</sup> F]FPQT	IC <sub>50</sub> 3.6 <sup>c</sup>	2.53	Confound by radiometabolites (rat)
[ <sup>18</sup> F]FPIT	IC <sub>50</sub> 5.4 <sup>c</sup>	2.53	Confound by radiometabolites (rat, monkey)
[ <sup>18</sup> F]cEFQ	K <sub>i</sub> 0.87(4 °C); 3.38 (37 °C) <sup>a</sup> K <sub>i</sub> 1.07 (4 °C); 1.77 (37 °C) <sup>b</sup> K <sub>i</sub> 3.74 (4 °C); 24.7 (37 °C) <sup>c</sup>	3.54	Yes (rat), no (baboon)
[ <sup>18</sup> F]MK-1312	IC <sub>50</sub> 0.84 <sup>c</sup>	2.3	Yes (monkey)
[ <sup>18</sup> F]FITM	IC <sub>50</sub> 5.1 <sup>c</sup>	1.5	Yes (rat, monkey)
[ <sup>18</sup> F]FIMX	IC <sub>50</sub> 1.8 <sup>c</sup>	2.52	Yes (monkey, human)

<sup>a</sup>Determined in ligand competition binding assays using cloned rat mGluR1 membrane

<sup>b</sup>Using rat cerebellum homogenate

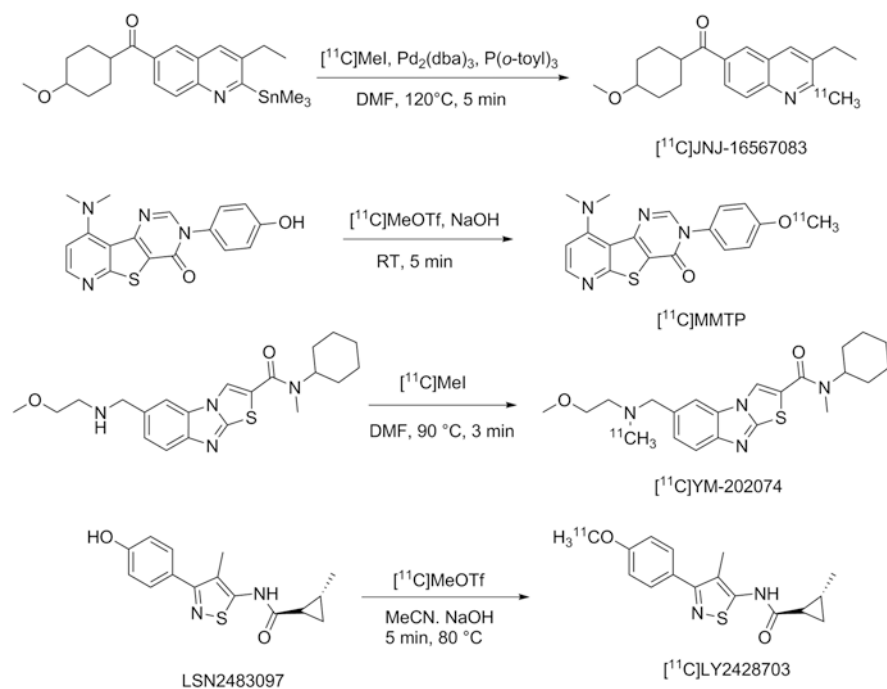
<sup>c</sup>Using cloned human mGluR1

ligands for PET studies were synthesized and evaluated for their utility as potential imaging agents for mGluR1 (Xu and Li 2019; Li and Huang 2014) (Fig. 13.3). Their binding affinities, lipophilicities, and in vivo specificity to mGluR1 are summarized in Table 13.1.

### 13.2.1 <sup>11</sup>C-Labeled mGluR1 Tracers

The most important carbon-11 radioligands for mGluR1 are described below.

(3-Ethyl-2-[<sup>11</sup>C]-methyl-6-quinolinyloxy)(cis-4-methoxycyclohexyl)-methanone, also denoted as [<sup>11</sup>C]JNJ-16567083, was the first reported allosteric antagonist PET tracer for imaging mGluR1 in the rodent brain (Huang et al. 2005). High binding affinity of JNJ-16567083 toward rat mGluR1 receptor was discovered with a K<sub>i</sub> value of 0.87 nM. [<sup>11</sup>C]JNJ-16567083 was synthesized via palladium-mediated



**Scheme 13.1** Radiosyntheses of [ $^{11}\text{C}$ ]JNJ-16567083, [ $^{11}\text{C}$ ]MMTP, [ $^{11}\text{C}$ ]YM-202074, and [ $^{11}\text{C}$ ]LY2428703

Stille coupling of the trialkyltin precursor with [ $^{11}\text{C}$ ] methyl iodide (Scheme 13.1). Biodistribution studies in rats under baseline and blockade conditions showed that [ $^{11}\text{C}$ ]JNJ-16567083 is selective and binds specifically to the mGluR1. Among the brain regions examined, the cerebellum, a region with the highest expression of the mGluR1, exhibited the highest uptake of radioactivity and highest specific binding of 81%. Further studies with [ $^{11}\text{C}$ ]JNJ-16567083 in rodents and nonhuman primates revealed species differences in the binding affinities toward mGluR1 (Table 13.1). A higher specificity of binding was obtained in rat brain, compared to baboon brain (Huang et al. 2012).

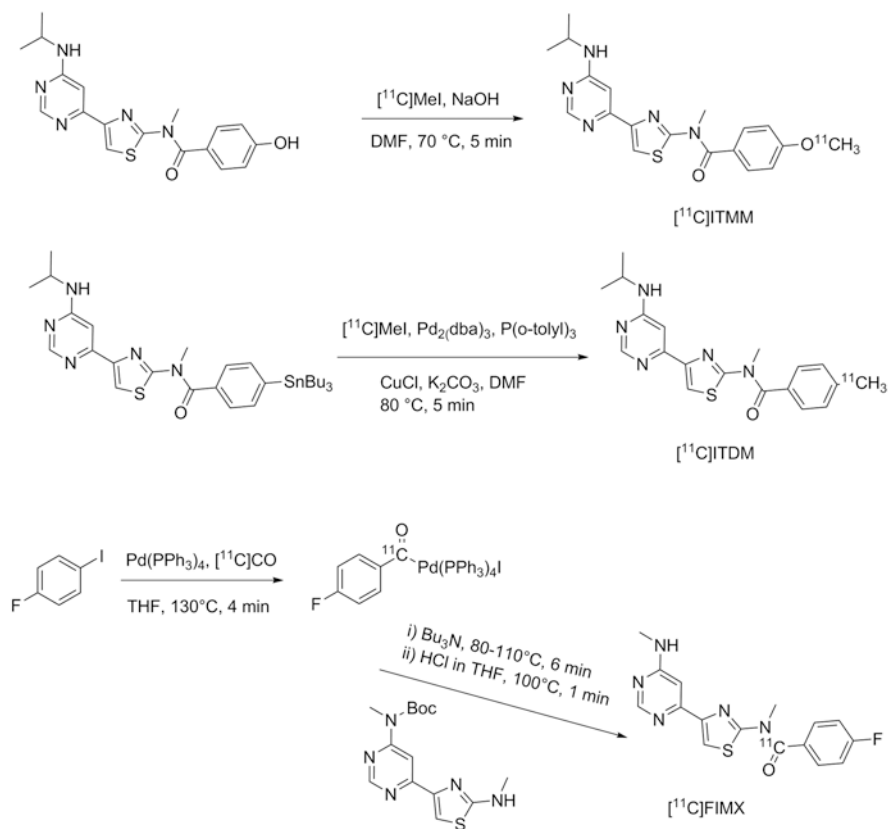
[ $^{11}\text{C}$ ]MMTP, [*O*-methyl- $^{11}\text{C}$ ]dimethylamino-3(4-methoxyphenyl)-3H-pyrido[3',2':4,5]thieno[3,2-d]pyrimidin-4-one is another promising carbon-11-labeled mGluR1 PET radioligand, which was synthesized by methylation of the corresponding desmethyl precursor using [ $^{11}\text{C}$ ]MeOTf in the presence of NaOH at room temperature (Scheme 13.1) (Prabhakaran et al. 2010). [ $^{11}\text{C}$ ]MMTP selectively labeled mGluR1 binding sites in postmortem human brain slices containing the cerebellum, hippocampus, frontal cortex, and striatum using phosphor imaging autoradiography. The densities of binding sites ( $B_{\text{max}}$ ) for the various brain regions estimated from these autoradiography studies were 140, 110, 85, and 20 fmol/cm<sup>2</sup> for cerebellum, hippocampus, frontal cortex, and striatum, respectively. The authors

performed PET studies in a baboon and showed that [ $^{11}\text{C}$ ]MMTP penetrates the BBB and accumulated in mGluR1-rich regions. The highest uptake was observed in the cerebellum, a region known to have the highest level of mGluR1 expression. However, it was not clear whether the binding in the cerebellum was specific, given that no blocking experiment was performed.

*N*-cyclohexyl-6- $\{[N-(2\text{-methoxyethyl})\text{-}N\text{-methylamino}]\text{methyl-}N\text{-methylthiazolo}[3,2\text{-}a]\text{benzimidazole-2-carboxamide}$  (YM-202074) is a derivative of YM-298198 (Fig. 13.2) and exhibits an inhibition constant ( $K_i$ ) of 4.8 nM toward mGluR1 and no inhibition up to 1  $\mu\text{M}$  for mGluR5 (Kohara et al. 2008). [ $^{11}\text{C}$ ]YM-202074 was synthesized by *N*-[ $^{11}\text{C}$ ] methylation of its desmethyl precursor with [ $^{11}\text{C}$ ] methyl iodide in DMF at 90  $^\circ\text{C}$  for 3 min (Scheme 13.1). [ $^{11}\text{C}$ ]YM-202074 exhibited optimal lipophilicity with a  $\log D$  value of 2.7. High specific binding of [ $^{11}\text{C}$ ]YM-202074 for mGluR1 in rat brain was confirmed by in vitro autoradiography under blockade conditions with nonradioactive mGluR1 antagonists YM-202074, JNJ-16259685, and CPCCOEt. However, ex vivo autoradiography and PET study showed that the in vivo radioactivity distribution of [ $^{11}\text{C}$ ]YM-202074 in rat brain region did not reflect the distribution pattern obtained from in vitro autoradiographic results. It was suggested that the discrepancy between in vitro and in vivo distributions might be caused by radiometabolite(s) of [ $^{11}\text{C}$ ]YM-202074 in the brain, and less than 10% intact compound was found in mouse plasma at 30 min p.i. Therefore, it was concluded that [ $^{11}\text{C}$ ]YM-202074 may not be a useful PET ligand for the in vivo imaging of mGluR1 (Yanamoto et al. 2010).

(1*S*,2*R*)-*N*-(3-(4-(methoxy)phenyl)-4-methylisothiazol-5-yl)-2-methylcyclopropane-1-carboxamide (LY2428703) exhibits an  $\text{IC}_{50}$  value of 8.9 nM to human mGluR1. [ $^{11}\text{C}$ ]LY2428703 was prepared by reacting the phenolic precursor, LSN2483097, with [ $^{11}\text{C}$ ]methyl triflate (Scheme 13.1) (Zanotti-Fregonara et al. 2013a). PET studies showed that [ $^{11}\text{C}$ ]LY2428703 readily penetrated the rat brain and distributed preferentially in the target-rich cerebellum. Its in vivo specificity and selectivity were confirmed by using mGluR1 and mGluR5 blocking agents, respectively. To verify species differences, a saturation binding assay was used to estimate its in vitro binding potential. The values for the binding potential expressed as  $B_{\text{max}}/K_d$  were 25.3, 18.2, and 16.9 for rat, monkey, and human, respectively (Zanotti-Fregonara et al. 2013b). PET imaging studies in rhesus monkeys under baseline and blocking conditions with and without nonradioactive LY2428703 (2.0 mg/kg) showed that although the brain uptake in the cerebellum was slightly higher than other brain regions, the uptake was not specific. In humans, a low and a homogeneous brain uptake was observed with [ $^{11}\text{C}$ ]LY2428703. The unspecific and low brain uptake was suspected to be due to high plasma protein binding given that the plasma free fraction of [ $^{11}\text{C}$ ]LY2428703 was 0.094% as measured by ultrafiltration. In summary, despite the promising in vitro and in vivo results in rodents, [ $^{11}\text{C}$ ]LY2428703 is unsuitable for imaging mGluR1 expression in monkey and human brain.

*N*-(4-(6-(isopropylamino)pyrimidin-4-yl)-1,3-(thiazol-2-yl)-4-[ $^{11}\text{C}$ ]methoxy-*N*-methylbenzamide ([ $^{11}\text{C}$ ]ITMM, Fig. 13.3) was prepared by reacting the corresponding phenolic precursor with [ $^{11}\text{C}$ ]MeI (Scheme 13.2) (Fujinaga et al. 2012c). A  $K_i$



**Scheme 13.2** Radiosyntheses of  $[^{11}\text{C}]\text{ITMM}$ ,  $[^{11}\text{C}]\text{ITDM}$  and  $[^{11}\text{C}]\text{FIMX}$

value of 12.6 nM was determined toward mGluR1 using rat brain homogenates in competition binding experiments.  $[^{11}\text{C}]\text{ITMM}$  was demonstrated to have high selectivity over mGluR5 ( $K_i$  mGluR5/mGluR1, >306) and low binding affinity to 28 major neurotransmitter receptors, uptake systems, and ion channels (Toyohara et al. 2013a). The  $\log D$  value of  $[^{11}\text{C}]\text{ITMM}$  was 2.57 using shake-flask method in octanol/phosphate buffer at pH 7.4. In vitro autoradiography on sagittal section of rat brain slices showed heterogeneous distribution of  $[^{11}\text{C}]\text{ITMM}$ , with highest uptake in the cerebellum, moderate radioactivity in the thalamus, and low in the striatum. The lowest radioactivity was detected in the brain stem. The in vivo PET results confirmed the expected distribution pattern of  $[^{11}\text{C}]\text{ITMM}$  in rat brain with the highest uptake in the cerebellum, followed by thalamus and striatum. Pretreatment with unlabeled ITMM (1 mg/kg) significantly reduced the uptake compared to the control group, and the distribution of radioactivity became fairly uniform throughout the brain. The maximum reduction of radioactivity uptake exceeded 85% in the thalamus and striatum, demonstrating that  $[^{11}\text{C}]\text{ITMM}$  binds specifically to mGluR1 in the rat brain. The specificity of binding of  $[^{11}\text{C}]\text{ITMM}$  was also

confirmed in mGluR1-knockout mouse brain. Although rapid metabolism was observed in the plasma, > 80% total radioactivity in the rat brain homogenate was intact [ $^{11}\text{C}$ ]ITMM (Fujinaga et al. 2012b). PET imaging studies were performed in eight healthy male volunteers (aged 21–27y) with [ $^{11}\text{C}$ ]ITMM (Toyohara et al. 2013b). Five of the eight subjects underwent dynamic brain PET study, and the other three subjects a whole-body distribution study. The uptake of [ $^{11}\text{C}$ ]ITMM was consistent with the known distribution of mGluR1 in the human brain. Brain uptake kinetics was slow, with a continuing increase of activity in the cerebellar cortex, and little washout in other brain regions during the 90 min dynamic PET scan. Small individual differences of [ $^{11}\text{C}$ ]ITMM distribution and brain TACs were detected in the five healthy young male subjects. A two-tissue-compartment model was used to analyze the regional TACs with the input function based on the metabolite-corrected arterial plasma concentration curve. Brain mGluR1 densities were estimated as distribution volume ( $V_T$ ), and the values for the cerebellar cortex and thalamus were 2.7 and 1.1 mL/cm<sup>3</sup>, respectively. The whole-body distribution studies revealed that the urinary bladder wall absorbed the highest dose, followed by the small intestine, liver, and heart wall. No adverse or clinically detectable pharmacologic effects were detected in any of the eight subjects. The authors concluded that [ $^{11}\text{C}$ ]ITMM is suitable for imaging cerebellar mGluR1 with acceptable dosimetry and pharmacologic safety at the dose required for adequate PET imaging (Toyohara et al. 2013b).

N-(4-(6-(isopropylamino)pyrimidin-4-yl)thiazol-2-yl)-N-methyl-4- $^{11}\text{C}$ ]methyl benzamide ([ $^{11}\text{C}$ ]ITDM), a structurally similar compound to ITMM, was developed by the same research group by replacing the O-[ $^{11}\text{C}$ ]methyl group in [ $^{11}\text{C}$ ]ITMM with C-[ $^{11}\text{C}$ ]methyl group. ITDM displayed a binding affinity ( $K_i$ ) value of 13.6 nM toward mGluR1, which is similar to ITMM ( $K_i$  value of 12.6 nM). The [ $^{11}\text{C}$ ]methyl group was introduced to the benzene ring via cross-coupling reaction of arylstannane precursor with [ $^{11}\text{C}$ ]MeI in the presence of palladium(0) complex (Scheme 13.2) (Fujinaga et al. 2012b). [ $^{11}\text{C}$ ]ITDM showed high specific binding in the cerebellum and thalamus in in vitro autoradiographic studies using rat brain sections. PET studies with [ $^{11}\text{C}$ ]ITDM in monkeys confirmed the high brain uptake and mGluR1 specificity in vivo. Metabolite analysis showed more than 95% intact [ $^{11}\text{C}$ ]ITDM at 60 min postinjection in rat brain extracts.

Yamasaki et al. used R6/2 mice, the most widely used animal model of Huntington's disease, for the measurement of nondisplaceable binding potential ( $BP_{ND}$ ) of [ $^{11}\text{C}$ ]ITMM (Yamasaki et al. 2014). It was established that the pons, a part of the brainstem with negligible mGluR1 expression, could be used as the reference tissue for the in vivo measurement of mGluR1 levels without the need for arterial blood sampling. Reduced radioactivity uptake was observed in cerebellum, thalamus, striatum, and cingulate cortex brain regions in R6/2 mice compared with control mice. The  $BP_{ND}$  measurements based on the Logan reference (Logan Ref) method indicated decreased  $BP_{ND}$  values in several R6/2 mice brain regions with highest in the striatum of roughly 35% followed by thalamus, cingulate cortex, and cerebellum at 20% to 30%. These percentage decreases in various of R6/2 mice brain regions were similar to the decreased mRNA levels reported previously in these brain regions (Cha et al. 1998). A close correlation between changes in

radioactive signal intensity and the degree of mGluR1 expression was found by comparison of the autoradiograms of R6/2 mouse brain sections with immunohistochemical images. Similar to the PET results, autoradiographic and immunohistochemical images of R6/2 mouse brain sections showed a heterogeneous decrease of signal in striatum, thalamus, and cerebellum, suggesting that the radioactive signal intensity corresponded with mGluR1 expression. The roles of mGluR1 were further investigated in chronic PD pathology by performing longitudinal [ $^{11}\text{C}$ ]ITDM PET imaging in A53T transgenic (A53T-Tg) rats expressing an abnormal human  $\alpha$ -synuclein (ASN) gene (Yamasaki et al. 2016). A53T-Tg rats showed a dramatic decline in general motor activities with age, along with abnormal ASN aggregation and striatal neuron degeneration. In longitudinal PET imaging, dynamic changes associated with PD progression were observed. Using the same reference tissue as mentioned above (Yamasaki et al. 2014), striatal  $BP_{\text{ND}}$  values of [ $^{11}\text{C}$ ]ITDM were higher in A53T-Tg rats than control rats prior to PD pathology onset but dramatically decreased afterward with age. [ $^{11}\text{C}$ ]ITDM was further assessed in pilocarpine-induced status epilepticus (PISE), a rat model of neuroinflammation (Yamasaki et al. 2017). PET with [ $^{11}\text{C}$ ]ITDM showed significant decreases of mGluR1 availability ( $BP_{\text{ND}}$ ) in the thalamus and hippocampus after PISE over the chronic period. PET with [ $^{11}\text{C}$ ]ITDM could successfully visualize hippocampal and thalamic declines of mGluR1 related to neuroinflammation. These results demonstrated that monitoring mGluR1 in vivo may provide beneficial information to further understand central nervous system disorders.

[Carbonyl- $^{11}\text{C}$ ]4-fluoro-N-methyl-N-(4-(6-(methylamino)pyrimidin-4-yl)thiazol-2-yl) benzamide ([ $^{11}\text{C}$ ]FIMX) is the C-11 version of [ $^{18}\text{F}$ ]FIMX, which has been demonstrated as an effective radioligand for imaging brain mGluR1 with PET (Hong et al. 2015; Xu et al. 2013). [ $^{11}\text{C}$ ]FIMX with a short physical half-life was developed to allow PET imaging of mGluR1 in the same animal subject on the same day, for example, in receptor occupancy studies. [ $^{11}\text{C}$ ]FIMX was prepared via a two-step synthetic approach (Scheme 13.2). The intermediate [ $^{11}\text{C}$ ]acyl-palladium complex was prepared by Pd-promoted carbonylation of 1-fluoro-4-iodobenzene with [ $^{11}\text{C}$ ]carbon monoxide, which was prepared online from cyclotron-produced [ $^{11}\text{C}$ ]carbon dioxide. THF was the preferred solvent because of its low boiling point, allowing it to be easily removed during the next step. To achieve optimal [ $^{11}\text{C}$ ]carbon monoxide insertion and avoid the formation of [ $^{11}\text{C}$ ]4-fluorobenzoic acid, the authors recommend the use of fresh and anhydrous THF. Boc-protected [ $^{11}\text{C}$ ]FIMX was formed by reacting [ $^{11}\text{C}$ ]acyl-palladium complex with the Boc-protected pyrimidine-thiazol amine; the aminolysis reaction went smoothly upon the complete evaporation of THF. Removal of the Boc-protecting group with 1.1 M HCl at 100 °C for 1 min led to the formation of the final product [ $^{11}\text{C}$ ]FIMX, which was separated from the reaction mixture by reversed-phase HPLC and formulated as a sterile solution for intravenous injection. The radiochemical yield was about 5% (d.c.) with a molar activity of about 100 GBq/ $\mu\text{mol}$ . The total synthesis time was ca. 40 min. [ $^{11}\text{C}$ ]FIMX was evaluated in anesthetized rhesus monkey. High radioactivity uptake (SUV 8.5) was observed in mGluR1-rich cerebellum after intravenous injection. The signal was abolished after pretreatment with the selective antagonist

JNJ16259685 (3 mg/kg, i.v.), confirming the specificity of uptake in rhesus monkey brain.

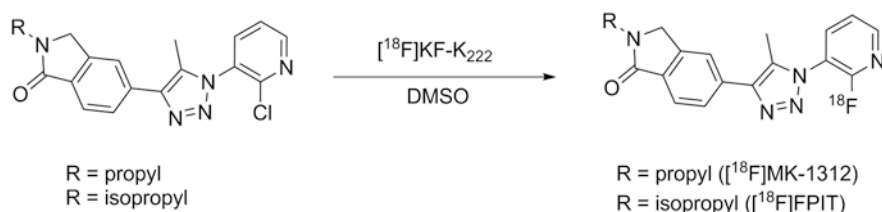
### 13.2.2 $^{18}\text{F}$ -Labeled mGluR1 Tracers

A number of fluorine-18-labeled compounds have been evaluated as potential PET ligands for imaging mGluR1. Some of the most important  $^{18}\text{F}$ -labeled mGluR1 radioligands reported so far are described below.

#### 13.2.2.1 $^{18}\text{F}$ -Labeled Triazole Analogues

Based on the structure of MK-5435, a potent and selective mGluR1 allosteric antagonist (Ito et al. 2009), several pyridyl triazole derivatives have been developed for imaging mGluR1. Depicted in Fig. 13.3 are the structures of three triazole analogues  $^{18}\text{F}$ MK-1312,  $^{18}\text{F}$ FTIDC, and  $^{18}\text{F}$ FPIT.

$^{18}\text{F}$ MK-1312 was synthesized via reaction of  $^{18}\text{F}$ KF with the corresponding chloropyridine in DMSO with microwave heating (Scheme 13.3) (Hostetler et al. 2011). Saturation binding studies in cerebellum homogenates with  $^{18}\text{F}$ MK-1312 revealed a  $K_d$  value of 0.40 and 0.84 nM for rhesus monkey and human cerebellum, respectively. Autoradiography was performed with both human and monkey brain slices. The cerebellum exhibited the highest accumulation of activity for both species. Baseline PET studies in a monkey showed that this radiotracer rapidly penetrates the BBB, which is in line with its optimal lipophilicity value of 2.3. The distribution of  $^{18}\text{F}$ MK-1312 was heterogeneous, with highest uptake observed in the cerebellum and moderate uptake in the thalamus and striatum. Displacement studies with an mGluR1 allosteric antagonist, MK-5435, indicated specific binding of  $^{18}\text{F}$ MK-1312 to mGluR1 in the cerebellum of rhesus monkey, and blockade was concentration dependent (Hostetler et al. 2011). Measurement of in vivo tracer metabolism in plasma revealed that 17% of plasma radioactivity was due to  $^{18}\text{F}$ MK-1312 at the end of a 90-min baseline PET scan.  $^{18}\text{F}$ MK-1312 shows the desired characteristics as a potential mGluR1 PET tracer with respect to its lipophilicity, affinity, and specificity toward mGluR1. However, a direct comparative study with  $^{18}\text{F}$ FITM indicated a lower brain uptake and specificity for  $^{18}\text{F}$ MK-1312 (Yamasaki et al. 2012b).



**Scheme 13.3** Radiosynthetic scheme for  $^{18}\text{F}$ MK-1312 and  $^{18}\text{F}$ FPIT

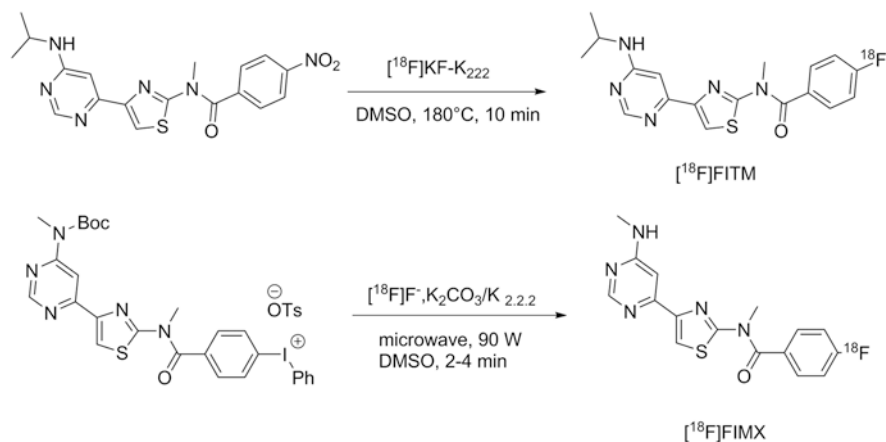


[<sup>18</sup>F]FPTQ, 6-[1-(2-[<sup>18</sup>F]fluoro-3-pyridyl)-5-methyl-1H-1,2,3-triazol-4-yl]quinolone, was synthesized by heating the bromo precursor with [<sup>18</sup>F]KF in anhydrous DMSO at 150 °C for 10 min. High binding affinities (IC<sub>50</sub> values) of FPTQ toward mouse and human mGluR1 of 1.4 and 3.6 nM, respectively, have been reported (Suzuki et al. 2009). In vitro autoradiographic results showed that the distribution pattern of [<sup>18</sup>F]FPTQ in the brain of rats was consistent with the distribution of mGluR1 reported previously (Lavreysen et al. 2004a). The specific binding of [<sup>18</sup>F]FPTQ accounted for higher than 97% of total binding, as determined by incubating with unlabeled FPTQ or mGluR1-selective JNJ-16259685 in rat brain slices. PET imaging showed that the distribution and uptake of [<sup>18</sup>F]FPTQ in the rat brain was consistent with the results obtained by dissection as well as by in vitro autoradiography. The maximum standard uptake values (SUV) were 2.3 in the cerebellum, 2.1 in the thalamus, and ranged from 1.1 to 1.5 in the hippocampus, striatum, cerebral cortex, and medulla. However, the presence of a radiolabeled metabolite in the brain may confound the specific binding of this radioligand in vivo. Thus, [<sup>18</sup>F]FPTQ may be of limited utility for the in vivo investigation of mGluR1 (Fujinaga et al. 2011).

[<sup>18</sup>F]FPIT, 1-(2-[<sup>18</sup>F]fluoro-3-pyridyl)-4-(2-isopropyl-1-oxoisindoline-5-yl)-5-methyl-1H-1,2,3-triazole, an analogue of [<sup>18</sup>F]MK-1312, was synthesized by reacting the bromo precursor with [<sup>18</sup>F]KF in anhydrous DMSO at 150 °C for 10 min (Scheme 13.3). FPIT was shown to have an IC<sub>50</sub> value of 5.4 nM for mGluR1 and 2500 nM for mGluR5 (Hostetler et al. 2011). The experimentally determined log*D* value of [<sup>18</sup>F]FPIT was 2.53, suggesting good BBB penetration. Autoradiography showed that the binding of [<sup>18</sup>F]FPIT is consistent with the reported distribution of mGluR1 with high specific binding in the cerebellum and thalamus of rat and monkey brains. [<sup>18</sup>F]FPIT displayed a similar distribution of radioactivity in both monkey and rat brains. The sequence of radioactivity level in the brain regions was cerebellum >thalamus >hippocampus >cerebral cortex >striatum >pons-medulla. Pretreatment with FPIT or JNJ-16259685 showed substantial inhibition of [<sup>18</sup>F]FPIT binding in these brain regions to a level close to or lower than that in the pons-medulla and almost diminished the difference in radioactivity among all brain regions. As indicated by the authors, a disadvantage of [<sup>18</sup>F]FPIT is its slow kinetics in the monkey brain and also the generation of a radiolabeled metabolite in the brain that could hamper accurate modeling and quantification with PET (Fujinaga et al. 2012a).

### 13.2.2.2 <sup>18</sup>F-Labeled Thiazole Analogues

[<sup>18</sup>F]FITM (4-[<sup>18</sup>F]fluoro-*N*-[4-[6-(isopropylamino)pyrimidin-4-yl]-1,3-thiazol-2-yl]-*N*-methylbenzamide) has a novel chemical structure that does not contain a triazole ring. Moreover, instead of the 2-[<sup>18</sup>F]fluoropyridine ring, [<sup>18</sup>F]FITM contains a [<sup>18</sup>F]fluorobenzene ring, which is expected to be stable against defluorination. [<sup>18</sup>F]FITM was synthesized by [<sup>18</sup>F]fluorination of its corresponding nitro precursor with [<sup>18</sup>F]KF in the presence of Kryptofix 222 (Scheme 13.4). FITM was shown to have potent antagonistic activity against human mGluR1 with an IC<sub>50</sub> value of 5.1 nM. Excellent selectivity over other subtypes has been reported whereby the IC<sub>50</sub> values were 7000 nM for human mGluR5, >10,000 nM for mGluR2, and



**Scheme 13.4** Radiosynthetic scheme for  $[^{18}\text{F}]\text{FITM}$  and  $[^{18}\text{F}]\text{FIMX}$

$>10,000$  nM for mGluR8 (Sato et al. 2009). The distribution coefficient ( $\log D$ ) of  $[^{18}\text{F}]\text{FITM}$  was 1.46 using the shake-flask method at pH 7.4. Strong signals of radioactivity were found in known mGluR1-rich regions such as the cerebellum and thalamus in *in vitro* autoradiographic studies using rat brain sections. By co-incubation with the nonradioactive reference FITM or mGluR1 antagonist JNJ-16259685, radioactivity throughout the brain section decreased significantly compared with those in the control section. Upon co-incubation with MPEP, an antagonist for mGluR5, no change in radioactivity accumulation was observed. The signals of radioactivity in *ex vivo* autoradiography in the brain sections of rat co-injected with JNJ-16259685 decreased by 86–91% in mGluR1-rich regions compared with those of the control, indicating the tracer was specifically bound to mGluR1 (Yamasaki et al. 2011). Very recently, the same research group reported on the further evaluation of  $[^{18}\text{F}]\text{FITM}$  in a monkey brain. High *in vivo* stability was also observed. The uptake of  $[^{18}\text{F}]\text{FITM}$  in the monkey brain was not only high in mGluR1-rich cerebellum but also in brain regions such as the thalamus and hippocampus which have moderate expression levels of mGluR1. Specific binding of  $[^{18}\text{F}]\text{FITM}$  could be demonstrated in all the aforementioned brain regions by pretreatment with JNJ-16259685 (Yamasaki et al. 2012a).

In order to circumvent the harsh radiolabeling conditions and also to improve the brain washout of  $[^{18}\text{F}]\text{FITM}$ , a new thiazole derivative code named  $[^{18}\text{F}]\text{FIMX}$  (4- $[^{18}\text{F}]\text{Fluoro-N-methyl-N-(4-(6-(methylamino)pyrimidin-4-yl)thiazol-2-yl)benzamide}$ ) was developed (Yamasaki et al. 2012a). Structurally, the *N*-isopropyl group in FITM was replaced by the less lipophilic *N*-methyl group in FIMX. FIMX displayed a threefold higher human mGluR1 affinity ( $\text{IC}_{50} = 1.8$  nM) than FITM and high selectivity over mGluR5 and a wide range of other receptors, transporters, and binding sites (Xu et al. 2013; Sato et al. 2009). For the F-18 radiolabeling, a diaryliodonium salt served as the precursor (Scheme 13.4). Various radiolabeling conditions were investigated including heating sources (conventional heating and

microwave), reaction time and temperature, solvent, and the influence of free-radical scavenger TEMPO. High radiolabeling yield (40%) was obtained using both conventional and microwave heating; however, microwave heating could remove the Boc-protecting group without the addition of acid which simplified the radiolabeling process. The most suitable solvent was DMF or DMSO. The addition of TEMPO improved the incorporation of [ $^{18}\text{F}$ ]fluoride ion from 26 to 42% under thermal conditions (100 °C bath temperature, 5 min) in DMF (Scheme 13.4). [ $^{18}\text{F}$ ]FIMX was produced in high radiochemical purity (96–100%) with molar activities of ca. 105 GBq/ $\mu\text{mol}$ . The decay-corrected radiochemical yield was 20%. The specificity and selectivity of [ $^{18}\text{F}$ ]FIMX were confirmed by imaging the mGluR1 expression in a rhesus monkey brain. Radioactivity signal distribution in monkey brain was consistent with the known mGluR1 expression. The highest uptake was in the receptor-rich cerebellum (5.3 SUV at 12 min). Moderate uptake values were observed in the thalamus (2.9 SUV), frontal cortex (2.7 SUV), hippocampus (2.5 SUV), putamen (2.5 SUV), and anterior cingulate (2.4 SUV). The regional time-activity curves were almost indistinguishable after i.v. injection of the mGluR1 selective antagonist, JNJ-16259685 (3 mg/kg) 30 min postinjection of the radiotracer. In sharp contrast, no changes of radioactivity distribution were observed using the mGluR5-selective ligand MTEP (5 mg/kg). Encouraged by the promising results, the same research group translated [ $^{18}\text{F}$ ]FIMX into humans. Twelve healthy volunteers were included for dynamic PET scan over 120 min, and four other subjects underwent whole-body scan to estimate radiation exposure (Zanotti-Fregonara et al. 2016). PET imaging revealed high [ $^{18}\text{F}$ ]FIMX uptake (SUV = 4–6) in mGluR1-rich cerebellum, and washout was faster than that of [ $^{11}\text{C}$ ]ITMM. The high density of mGluR1 in the cerebellum was quantified based on an unconstrained two-tissue-compartment model, and values of distribution volume ( $V_T$ ) ranged from 1.5 in the caudate to 11 mL $\cdot\text{cm}^{-3}$  in the cerebellum. Stable  $V_T$  values could be obtained in 120 min scan in all regions except the cerebellum, for which an acquisition time of at least 170 min was necessary. Regional  $V_T$  values of [ $^{18}\text{F}$ ]FIMX correlated well with mGluR1 transcript density, indicating that [ $^{18}\text{F}$ ]FIMX could be used for mapping mGluR1 in the human brain. The estimated effective dose of [ $^{18}\text{F}$ ]FIMX was 23  $\mu\text{Sv}/\text{MBq}$  and compares favorably with other  $^{18}\text{F}$ -labeled ligands (Zanotti-Fregonara et al. 2013c).

### 13.2.2.3 [ $^{18}\text{F}$ ]cEFQ

(3-Ethyl-2-[ $^{18}\text{F}$ ]fluoroquinolin-6-yl) cis-(4-methoxycyclohexyl)methanone ([ $^{18}\text{F}$ ]cEFQ), the  $^{18}\text{F}$ -labeled analogue of JNJ-16567083, was prepared by nucleophilic substitution of chloroquinoline precursor with [ $^{18}\text{F}$ ]KF in DMSO in the presence of Kryptofix 222 at 100 °C for 5 min (Huang et al. 2012; Lee et al. 2017). It was found that even when a pure cis-form precursor was used, both cis and trans isomers [ $^{18}\text{F}$ ]cEFQ and [ $^{18}\text{F}$ ]tEFQ were obtained under the abovementioned radiolabeling conditions. Separation of the isomers was possible using a reversed-phase HPLC column. Higher amount of the trans isomer, [ $^{18}\text{F}$ ]tEFQ, was obtained compared to the cis-isomer [ $^{18}\text{F}$ ]cEFQ (60% vs. 40%) due to the more stable thermodynamic nature of the trans product. It was reported that the trans isomers bearing methoxy-

methyl, and chloro-quinoline functionalities were 20–100-fold less potent toward GluR1 compared to their cis counterparts (Mabire et al. 2005). For biological evaluations, it is therefore essential to separate [ $^{18}\text{F}$ ]cEFQ from [ $^{18}\text{F}$ ]tEFQ by HPLC. In vitro stability studies indicated no cis-trans isomerization in human serum at body temperature. [ $^{18}\text{F}$ ]cEFQ was shown to be stable at 4 °C but slowly degraded into free [ $^{18}\text{F}$ ]fluoride and other radioactive metabolites over time at 37 °C. The binding affinity of [ $^{18}\text{F}$ ]cEFQ to mGluR1 was also shown to be temperature-dependent (Table 13.1). Species differences were found for [ $^{18}\text{F}$ ]cEFQ in vitro and in vivo studies. [ $^{18}\text{F}$ ]cEFQ was more potent for cloned rat mGluR1 than human mGluR1 (3.38 nM vs. 24.7 nM at 37 °C). PET images and TACs with [ $^{18}\text{F}$ ]cEFQ in rat brain indicated highest radioactivity uptake in the cerebellum. Moderate radioactivity uptake was found in the thalamus, hippocampus, and striatum. The pattern of radioactivity distribution was in line with the expected mGluR1 expression in the rat brain. The amount of the intact [ $^{18}\text{F}$ ]cEFQ in the brain was 88.9% and 59.5% at 5 and 30 min, respectively. Polar radiometabolites did not confound with the radiosignals in the brain tissue. Specificity and selectivity of binding were confirmed by blocking studies using the known mGluR1 antagonist JNJ16259685 and mGluR5 selective antagonist ABP688, respectively. [ $^{18}\text{F}$ ]cEFQ, however, did not show any detectable specific binding in the baboon brain, which impeded its further evaluation in human subjects.

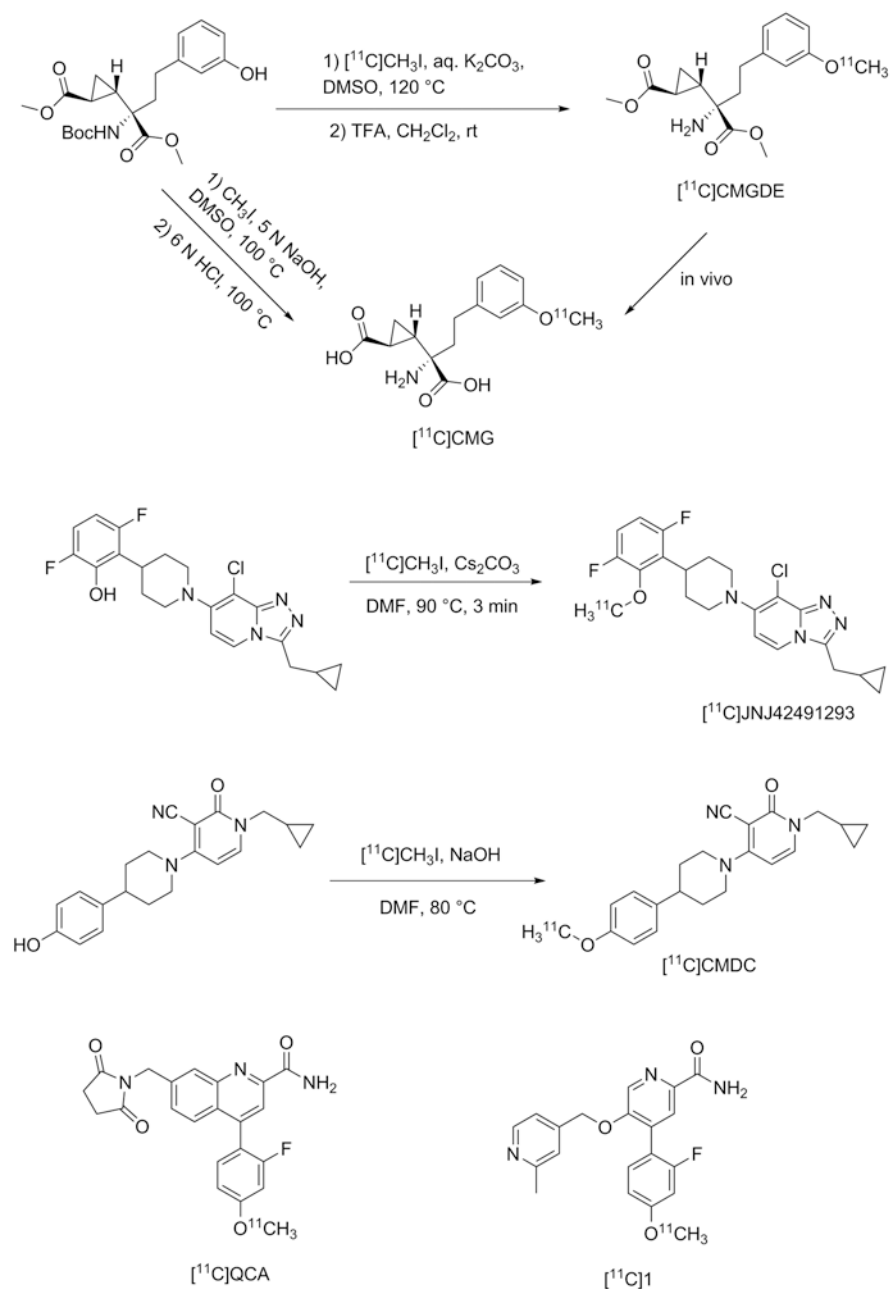
So far, three mGluR1 PET ligands, [ $^{11}\text{C}$ ]YM-202074, [ $^{11}\text{C}$ ]ITMM, and [ $^{18}\text{F}$ ]FIMX, were evaluated in human subjects. [ $^{11}\text{C}$ ]YM-202074 failed to visualize mGluR1 in humans, whereas both [ $^{11}\text{C}$ ]ITMM and [ $^{18}\text{F}$ ]FIMX demonstrated potential for mapping mGluR1 in the human brain.

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### 13.3 mGluR2/3 PET Tracers

Group II mGluRs consists of mGluR2 and mGluR3. They couple negatively to adenylyl cyclase through Gi/Go proteins. Emerging preclinical and clinical data suggested that group II mGluRs are implicated in various neurological and psychiatric disorders including anxiety disorders (Swanson et al. 2005), schizophrenia (Ghose et al. 2008; Muguruza et al. 2016; Patil et al. 2007), drug addiction (Moussawi and Kalivas 2010), Parkinson's disease (Dickerson and Conn 2012), and pain states (Chiechio 2016; Mazzitelli et al. 2018; Neugebauer et al. 2009). Specifically, high potent mGluR2 allosteric modulators have advanced to phase 2 clinical trials in patients with schizophrenia and MDD (Trabanco et al. 2019). In the brain, mGluR2 is highly expressed in the cerebral cortex, striatum, hippocampus, and amygdala (Richards et al. 2005). So far, several PET ligands targeting mGluR2 have been developed, and one of them has been evaluated in human subjects.

(S,S,S)-2-(2-carboxycyclopropyl)-2-(3-[ $^{11}\text{C}$ ]methoxyphenethyl) glycine dimethyl ester ([ $^{11}\text{C}$ ]CMGDE) was the first PET tracer developed as a prodrug for imaging group II metabotropic glutamate receptors in vivo. Its hydrolyzed product cyclopropane amino-diacid CMG was designed based on an mGluR2/3 antagonist LY341495 ( $\text{IC}_{50}$  = 21 nM for mGluR2; 14 nM for mGluR3) (Ornstein et al. 1998). CMG



**Scheme 13.5** Radiosynthetic scheme for mGluR2 radioligands and structures of  $[^{11}\text{C}]\text{QCA}$  and  $[^{11}\text{C}]\text{1}$

displayed a  $IC_{50}$  value of 94 nM toward mGluR2/3 with the mixture of its four constituent isomers (R,S,S-, S,S,S-, S,R,R- and R,R,R-isomer). As the binding affinities of the S,S,S-isomer of several CMG derivatives surpassed that of the corresponding isomer mixture by factors of 3.4–6.9 (Ornstein et al. 1998), the S,S,S-isomers of CMG and CMGDE were radiolabeled with carbon-11 as illustrated in Scheme 13.5 (Wang et al. 2012). As expected, the high hydrophilic tracer [ $^{11}C$ ]CMG showed no brain penetration after injection into male Sprague-Dawley rats. However, its pro-drug [ $^{11}C$ ]CMGDE readily accumulated into mGluR2/3-rich rat brain regions such as striatum, hippocampus, and cortical areas. The maximum accumulation (2.5–3.5% of the injected dose per  $cm^3$ ) was observed at 2 min after the tracer application. Pretreatment with LY341495, a highly potent mGluR2/3 ligand, induced 29.5% decrease of [ $^{11}C$ ]CMGDE accumulation in the whole brain, indicating relatively low *in vivo* specificity.

1-(Cyclopropylmethyl)-4-(4-[ $^{11}C$ ]methoxyphenyl)piperidin-1-yl-2-oxo-1,2-dihydro pyridine-3-carbonitrile ([ $^{11}C$ ]CMDC) was synthesized by O-[ $^{11}C$ ]methylation of its corresponding phenolic precursor with [ $^{11}C$ ]methyl iodide under basic conditions (Ma et al. 2017). CMDC showed potent functional activity ( $EC_{50}$ : 98 nM) for human mGluR2 *in vitro* (Cid et al. 2014). The logD value of [ $^{11}C$ ]CMDC as measured by the shake flask method was  $3.1 \pm 0.2$ . This value lies in the range normally considered favorable for CNS PET radiotracers. *In vitro* autoradiography indicated that the distribution pattern of radioactivity was heterogeneous with high expression in mGluR2-rich brain regions of cerebral cortex, striatum, hippocampus, and granular layer of the cerebellum. Co-incubation with unlabeled CMDC demonstrated its specificity in rat brain sections. However, limited brain penetration (peak brain uptake ca. 0.6 SUV) and low *in vivo* specific binding of [ $^{11}C$ ]CMDC in rat brain suggest that [ $^{11}C$ ]CMDC has limited potential as a PET tracer for mapping brain mGluR2 *in vivo*. Replacing the C-11 labeled methoxy group with F-18 labeled fluoroethoxyl and shifting the methoxy group from para- to meta- or ortho-position of the benzene ring resulted in similar or slightly increased binding affinities compared to CMDC (Kumata et al. 2017). *In vitro* specificities were demonstrated by autoradiographic studies using rat brain sections. No *in vivo* studies have been disclosed so far.

7-((2,5-dioxopyrrolidin-1-yl)methyl)-4-(2-fluoro-4-[ $^{11}C$ ]methoxyphenyl) quinoline-2-carboxamide ([ $^{11}C$ ]QCA) was prepared in a similar way as [ $^{11}C$ ]CMDC (Zhang et al. 2017b). The  $EC_{50}$  value reported for QCA was 8 nM. Heterogeneous brain distribution and moderate-to-high level specific binding were obtained *in vitro* autoradiographic studies using rat brain sections applying [ $^{11}C$ ]QCA. Dynamic PET imaging using [ $^{11}C$ ]QCA showed limited radioactivity uptake (ca. 0.3 SUV) in the rat brain. Pretreatment with unlabeled QCA (1 mg/kg) failed to show significant reduction of brain uptake. PET imaging studies in PgP/Bcrp knockout mice indicated that [ $^{11}C$ ]QCA had intensive interactions with brain efflux pumps at the murine blood-brain barrier. These results suggest that [ $^{11}C$ ]QCA is not a suitable tracer for imaging mGluR2 in rodent brain.

4-(2-Fluoro-4-[ $^{11}C$ ]methoxyphenyl)-5-((2-methylpyridin-4-yl)methoxy)picolinamide ([ $^{11}C$ ]1) was synthesized by O-[ $^{11}C$ ]methylation of the phenolic precursor with [ $^{11}C$ ]methyl iodide (Kumata et al. 2019). This compound displayed nanomolar

binding affinity for mGluR2 ( $IC_{50} = 26$  nM). Although this radioligand showed high in vitro specific binding to mGluR2 in autoradiographic studies, its in vivo properties (low binding and substrate of ATP binding cassette efflux transporters) were very similar to the lead compound [ $^{11}C$ ]QCA; therefore it could not be taken into consideration as a potential PET radiotracer candidate.

8-Chloro-3-(cyclopropylmethyl)-7-(4-(3,6-difluoro-2- $[^{11}C]$ (methoxyphenyl)piperidin-1-yl)-[1,2,4]triazolo[4,3-a]pyridine ( $[^{11}C]$ JNJ42491293) was selected as a potential PET tracer from a series of C-11 labeled 7-(phenyl-piperidinyl)-1,2,4-triazolo[4,3-a]pyridine derivatives (Andres et al. 2012). JNJ42491293 binds to an allosteric modulator site of the human mGluR2 with 9.6 nM binding affinity and has a calculated logP value of 4.84. PET studies in rats indicated that [ $^{11}C$ ]JNJ42491293 binds specifically and reversibly to mGluR2. The intensity of the PET signals were in the order of cerebellum > striatum > thalamus > whole brain > hippocampus > cortex. The maximum radioactivity concentration (SUV 1.7) in the brain was reached at 12 min p.i. and remained constant until 27 min followed by a slow washout. A clear displacement of radioactivity in all the brain regions was observed after injection of JNJ42153605, a high affinity, and selective ligand for mGluR2. Ex vivo radiometabolite studies revealed that the radioactivity signal in the rat brain was attributed to the intact parent compound (> 95%). Given these promising results in rat, [ $^{11}C$ ]JNJ42491293 was further studied in mGluR2 knockout rats and humans (Leurquin-Sterk et al. 2017). The brain uptake of [ $^{11}C$ ]JNJ42491293 in human subjects peaked at around 30 min after bolus injection and showed a gradual brain washout over the next hour. However, unexpected high cardiac retention was observed in a human subject, which prompted the authors to re-examine the tracer's specificity by extending preclinical research including the mGluR2 knockout rat and rhesus monkey. Similar [ $^{11}C$ ]JNJ42491293 uptake was obtained in both mGluR2 knockout and wild-type rat brain, suggesting non-specific binding. Using JNJ-46356479, another mGluR2 selective positive allosteric modulator, also failed to show specific binding of [ $^{11}C$ ]JNJ42491293 in rat and monkey brains. The unknown in vivo off-target binding in the knockout rat and the lack of in vivo specificity precluded [ $^{11}C$ ]JNJ42491293 as a useful mGluR2 PET tracer.

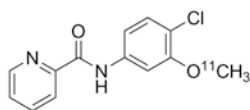
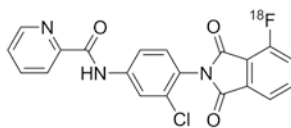
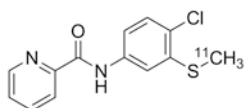
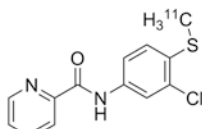
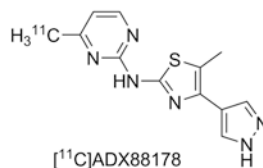
Apart from [ $^{11}C$ ]JNJ42491293, two structurally unknown PET tracers, one was labeled with F-18 ( $[^{18}F]$ ER-000604699), and the other with C-11 have been reported as mGluR2 PET tracers (Li et al. 2017; Lohith et al. 2016). However, limited information is available on these two radioligands as only conference abstracts are available.

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### 13.4 mGluR4 PET Tracers

mGluR4 belongs to the group III metabotropic glutamate receptors along with mGluR6, mGluR7, and mGluR8. Based on mRNA studies, mGluR4 has high expression in cerebellar cortex, moderate in thalamus and hippocampus (Corti et al. 2002). Compared to other groups of the mGlu receptor family, less attention has been received for group III mGluRs in the field of drug discovery. The identification

## mGluR4

[<sup>11</sup>C]ML128rEC<sub>50</sub> = 110 nM; hEC<sub>50</sub> = 240 nM[<sup>18</sup>F]KALB001hEC<sub>50</sub> = 50.4 nM; IC<sub>50</sub> = 5.1 nM[<sup>11</sup>C]PXT012253IC<sub>50</sub> = 3.4 nM[<sup>11</sup>C]iso-PXT012253IC<sub>50</sub> = 3.1 nM[<sup>11</sup>C]ADX88178EC<sub>50</sub> = 4 nM**Fig. 13.4** Structures of mGluR4 radioligands

of a partially selective mGluR4 positive allosteric modulator (PAM), N-phenyl-7-(hydroxyimino)cyclopropa[b]-chromen-1a-carboxamide (PHCCC), and its antiparkinsonian activity in animal models of PD has prompted intense interest for this target (Marino et al. 2003). Compound PXT002331 was the first mGluR4 PAM to enter into a clinical trial study (Charvin et al. 2018). Several PET tracers for imaging mGluR4 were developed since 2013 (Fig. 13.4); however, as of today none of them has advanced through to human studies.

N-(4-chloro-3-[<sup>11</sup>C]methoxyphenyl)picolinamide ([<sup>11</sup>C]ML128) was synthesized by reacting its corresponding phenolic precursor with [<sup>11</sup>C]CH<sub>3</sub>I (Kil et al. 2013). ML128, a positive allosteric modulator for mGluR4 (rEC<sub>50</sub> = 110 nM and hEC<sub>50</sub> = 240 nM), showed efficacy in a preclinical rodent model of motor impairments associated with PD (Engers et al. 2009; Lindsley et al. 2009). Fast accumulation of [<sup>11</sup>C]ML128 (peak activity between 1–3 min) was observed in several rat brain regions including striatum, thalamus, hippocampus, and cerebellum, which reflected the distribution pattern of mGluR4 in rodent brain. Partial blocking effect (22–28%) was demonstrated by using N-(3-(difluoromethoxy)phenyl)picolinamide, a potent mGluR4 modulator, suggesting low in vivo specificity. [<sup>11</sup>C]ML128 is the first reported mGluR4 PET tracer and may serve as a lead compound for future mGluR4 PET imaging agents.

N-(4-chloro-3-(<sup>11</sup>C)methylthiophenyl)picolinamide ([<sup>11</sup>C]PXT012253) was developed as an analog of [<sup>11</sup>C]ML128. (Kil et al. 2016). It emerged from a series of ML128 derivatives which were synthesized and evaluated for their binding affinities toward mGluR4. PXT012253 and its regioisomer exhibited high in vitro binding affinity for mGluR4 with IC<sub>50</sub> values of 3.4 nM and 3.1 nM, respectively. Both compounds were radiolabeled with C-11 and characterized in vivo using PET. [<sup>11</sup>C]PXT012253 outperformed its regioisomer and the lead compound



[<sup>11</sup>C]ML128 as it showed the highest uptake in rat brain and 25% increased signal-to-noise ratio compared to [<sup>11</sup>C]ML128. [<sup>11</sup>C]PXT012253 was therefore selected for further evaluation in cynomolgus monkeys (Takano et al. 2019). The uptake of [<sup>11</sup>C]PXT012253 peaked at 4 min and decreased to half in around 40 min on average. The mean %ID (injected dose) of the whole brain was 3.6% with the highest binding in the thalamus and putamen, and the lowest in the cortexes, which was similar to that previously reported in the rat brain with the same PET ligand (Kil et al. 2013). A two-tissue compartment model (2TC) fitted well to the brain regional TACs of [<sup>11</sup>C]PXT012253. Competition with PXT002331 showed that the SUV values at the peak under the blocking conditions were higher than those at baseline conditions. The authors suggested that blocking compounds other than PXT002331 should be evaluated.

N-(3-chloro-4-(4-[<sup>18</sup>F]fluoro-1,3-dioxoisindolin-2-yl)phenyl)picolinamide ([<sup>18</sup>F]KALB001) was prepared by nucleophilic substitution of the nitro group with [<sup>18</sup>F]fluoride in the presence of Kryptofix 222 and potassium carbonate at 120 °C for 5 min (Kil et al. 2014). KALB001 was developed from picolinamide ML128 by introducing the phthalimide group at the 4-position on the phenyl ring. It exhibited enhanced activity (hEC<sub>50</sub> = 50.4 nM) compared to ML128 (hEC<sub>50</sub> = 240 nM) and excellent binding affinity (IC<sub>50</sub> = 5.1 nM) to mGluR4. The characterization of [<sup>18</sup>F]KALB001 was conducted in both rats and monkeys. It was found that [<sup>18</sup>F]KALB001 occupied brain areas known to express mGluR4 and showed better permeability in the monkey brain than in the rat brain. Blocking studies revealed 10–20% reduced uptake in the rat brain pretreated with mGluR4-specific blocking agent N-(3-(difluoromethoxy)phenyl)-2-picolinamide (10 mg/kg). Negligible blocking effect was observed when mGluR5 antagonist MTEP was used, indicating good selectivity to mGluR4. [<sup>18</sup>F]KALB001 is the first <sup>18</sup>F-labeled mGluR4 radioligand, and further structural modification to improve its specificity may lead to the discovery of a suitable radioligand for imaging mGluR4 in human subjects.

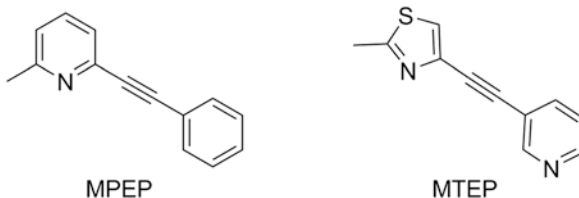
5-Methyl-N-(4-[<sup>11</sup>C]methylpyrimidin-2-yl)-4-(1H-pyrazol-4-yl)thiazol-2-amine ([<sup>11</sup>C]ADX88178) was reported in 2016 (Fujinaga et al. 2016). ADX88178 potentiated glutamate-mediated activation of human mGluR4 with an EC<sub>50</sub> value of 4 nM without significant effects on other mGluRs (EC<sub>50</sub> > 30 μM). The [<sup>11</sup>C]methyl group at position 4 in pyrimidine was introduced by Stille coupling. Protecting the pyrazole ring with p-methoxybenzyl (PMB) was required for the C-C cross-coupling reaction. The accumulation pattern of [<sup>11</sup>C]ADX88178 was consistent with the reported biological distribution of mGluR4 in the rat brain. Although low in vitro and in vivo specificity to mGluR4 was observed, compound ADX88178 may serve as a new chemical scaffold for structural optimization.

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### 13.5 mGluR5 PET Tracers

mGluR5 is an important target for PET imaging with potential for application in drug development, diagnosis, and therapy monitoring. Metabotropic glutamate subtype 5 receptor (mGluR5) was cloned in 1992 and exists as a covalently bound

**Fig. 13.5** Chemical structures of MPEP and MTEP

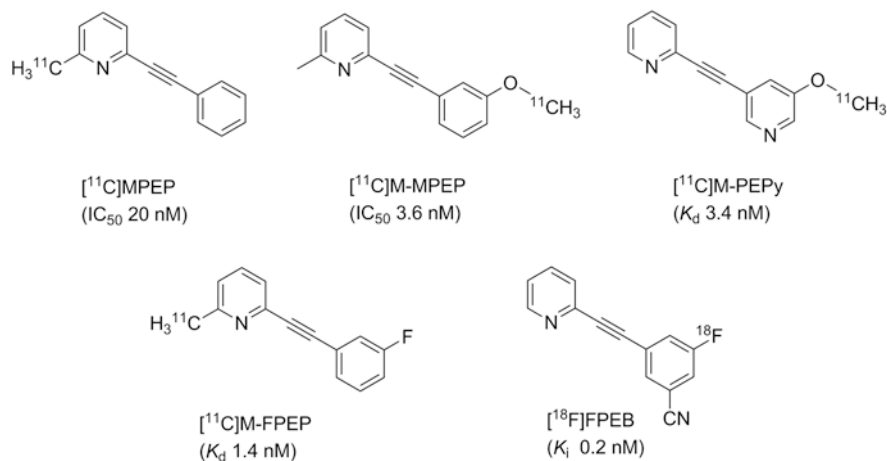


homodimer in heterologous expression system as well as in native tissue (Abe et al. 1992; Romano et al. 1996). It is expressed in brain regions such as the hippocampus, basal ganglia (striatum and nucleus accumbens), amygdala, and cortex (Romano et al. 1996; Gupta et al. 2005). mGluR5 has been implicated in a number of central nervous system disorders including depression, anxiety, addiction, schizophrenia, Parkinson's disease, and fragile X syndrome (Bear 2005; Carroll 2008; Ohnuma et al. 1998; Palucha et al. 2005; Pietraszek et al. 2007; Rouse et al. 2000; Spooren et al. 2003; Spooren and Gasparini 2004; Spooren et al. 2000; Tatarczynska et al. 2001; Tsai et al. 2005). As such, this receptor provides a novel target for the development of therapeutic drugs for the aforementioned neurological disorders. A number of noncompetitive mGluR5 ligands have been developed during the past years, and their potential as therapeutic drugs for some of the abovementioned CNS disorders has been investigated (Augelli-Szafran and Schwarz 2003; Chen and Conn 2008; Chua et al. 2005; Emmitte 2011; Gasparini et al. 2008; Jaeschke et al. 2008; Kulkarni et al. 2009; Lindemann et al. 2011; Lindsley and Emmitte 2009; Porter et al. 2005; Ritzen et al. 2009; Varney et al. 2002). In addition to CNS disorders, mGluR5 was found to be upregulated in CNS tumors and some non-CNS tumors (Brocke et al. 2010; Choi et al. 2011), participating in tumor growth and aggression.

The first potent and selective noncompetitive mGluR5 antagonists reported were the diaryl alkynes, 2-methyl-6-(phenylethynyl)pyridine (MPEP) (Gasparini et al. 1999; Lea and Faden 2006), and 3-[(2-methyl-1,3-thiazol-4-yl)ethynyl]pyridine (MTEP) (Fig. 13.5) (Cosford et al. 2003a; Cosford et al. 2002; Cosford et al. 2003b). Since their publication, MPEP and MTEP have been used as templates for the design of noncompetitive mGluR5 antagonists. Also a series of PET radioligands based on their core structures have been labeled with either carbon-11 or fluorine-18 and evaluated in in vitro and in vivo studies. Three reviews providing an overview on available mGluR5 PET ligands have been published (Mu et al. 2010; Pillai and Tipre 2016; Yu 2007).

### 13.5.1 MPEP-Derived mGluR5 Tracers

MPEP and its methyl analogue M-MPEP have been identified as potent, highly selective, noncompetitive antagonists for mGluR5. Derivatives of MPEP that have been labeled with either carbon-11 or fluorine-18 are depicted in Fig. 13.6.

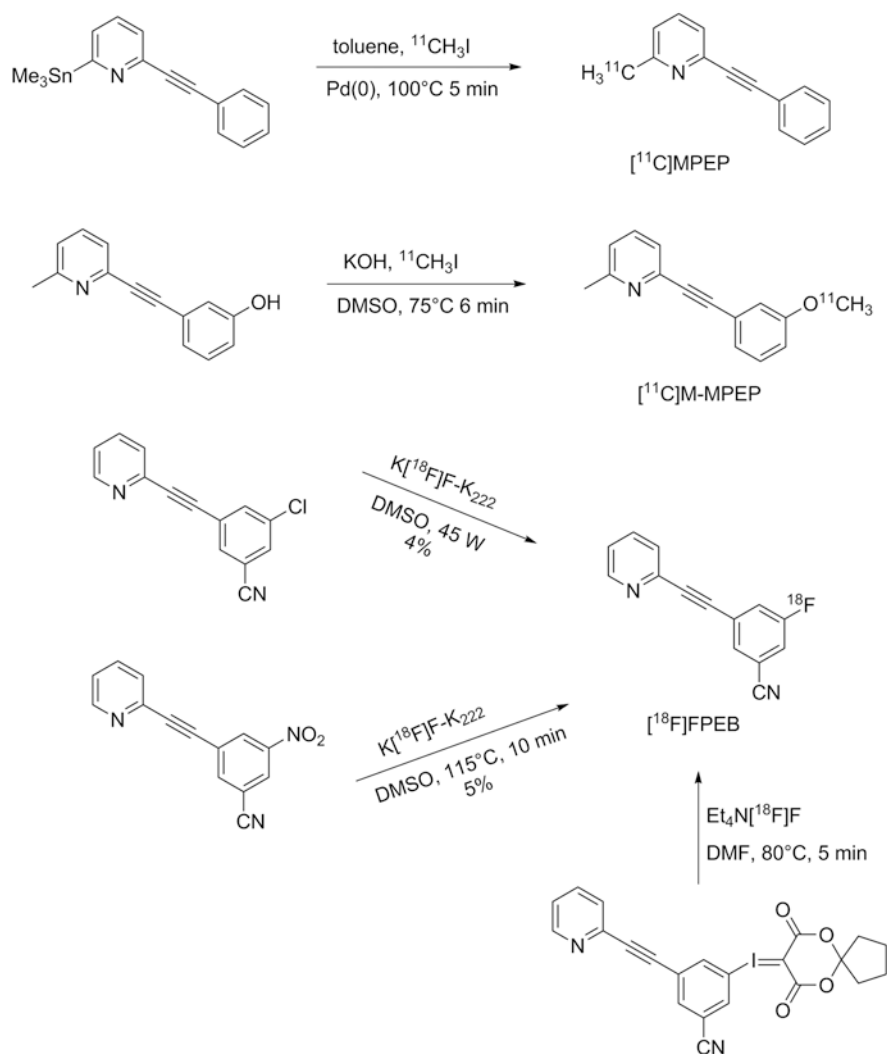


**Fig. 13.6** Chemical structures of <sup>11</sup>C- and <sup>18</sup>F-labeled MPEP derivatives

The high binding affinity of MPEP ( $IC_{50}$  20 nM) (Gasparini et al. 1999; Gasparini et al. 2002) to the heptahelical domain mGluR5 initiated the first radiolabeling of this compound with tritium. The identification of high-affinity analogues of M-MPEP (2-(2-(3-methoxyphenyl)ethynyl)pyridine,  $IC_{50}$  3.6 nM) (Gasparini et al. 2002), and M-PEPy (3-methoxy-5-pyridin-2ylethynyl)pyridine,  $K_d$  3.4 nM) (Cosford et al. 2003a; Anderson et al. 2003; Patel et al. 2003) prompted the radiolabeling of these derivatives with carbon-11 for evaluation as potential candidates for imaging mGluR5.

[<sup>11</sup>C]MPEP was prepared via Stille coupling reaction of 2-(phenylethynyl)-6-(trimethylstannyl)pyridine with [<sup>11</sup>C]MeI (Scheme 13.6) in high radiochemical purity and good molar activity (33–44 GBq/ $\mu$ mol) (Yu et al. 2005). In vivo PET imaging studies of [<sup>11</sup>C]MPEP in Sprague-Dawley rats showed high accumulation of [<sup>11</sup>C]MPEP in brain areas such as the olfactory bulb, striatum, hippocampus, frontal cortex, and cerebellum. Pre-administration of unlabeled MPEP (10 mg/kg) 5 min prior to administration of [<sup>11</sup>C]MPEP led to a decreased [<sup>11</sup>C]MPEP binding in the olfactory bulb. In contrast, an increased uptake was noted for the hippocampus and striatum, suggesting that [<sup>11</sup>C]MPEP binds nonspecifically to these brain regions. No reasons have been provided by the authors to explain this phenomenon. For the olfactory bulb, the extent of specific binding was 45 and 61% at 5 and 40 min postinjection, respectively.

The radiosynthesis of [<sup>11</sup>C]M-MPEP was accomplished by reacting [<sup>11</sup>C]MeI with phenolic precursor in the presence of solid KOH powder in 85% radiochemical yield and 98% radiochemical purity (Scheme 13.6) (Yu et al. 2005). Moderate uptake was observed after tail vein injection of [<sup>11</sup>C]M-MPEP into rats. Pre-administration of MPEP (10 mg/kg) induced a 60% decrease in [<sup>11</sup>C]M-MPEP binding in the olfactory bulb 5 min postinjection. In other brain regions, the mGluR5 antagonist, MPEP, decreased [<sup>11</sup>C]M-MPEP binding only during the early time



**Scheme 13.6** Radiosyntheses of [ $^{11}\text{C}$ ]MPEP, [ $^{11}\text{C}$ ]M-MPEP, and [ $^{18}\text{F}$ ]FPEB

points: 45% in the cortex, 14% in the hippocampus, and 29% in the striatum at 5 min postinjection. At 40 min postinjection, a binding ratio of olfactory bulb to cerebellum was 4.2. Kocic et al. also reported on the radiosynthesis and the *in vivo* evaluation of [ $^{11}\text{C}$ ]M-MPEP albeit in an abstract (Kocic et al. 2002). In their study, Kocic and colleagues found an increased uptake of [ $^{11}\text{C}$ ]M-MPEP radioactivity 30 min postinjection in the rat brain after co-injection with unlabeled M-MPEP (1 mg/kg). However, no information was provided on radioactivity uptake in the olfactory bulb.

[<sup>11</sup>C]M-PEPy was obtained by reacting the desmethyl precursor with either [<sup>11</sup>C]MeI (Yu et al. 2005) or [<sup>11</sup>C]CH<sub>3</sub>OTf (Severance et al. 2006) in the presence of KOH or NaOH in moderate radiochemical yield. In vitro phosphor imaging with [<sup>11</sup>C]M-PEPy using human and rat brain tissues demonstrated high specific binding in the hippocampus, striatum, and cortex with minimal specific binding in the cerebellum. The hippocampus-to-cerebellum ratio found in this study was 37 in rats and 8 in humans. However, further in vivo micro-PET studies in rats using urethane anesthesia, PET studies in baboons using isoflurane anesthesia, and ex vivo studies in rats showed little specific binding in the brain and very fast tracer washout (Severance et al. 2006). One possible explanation is that [<sup>11</sup>C]M-PEPy is actively extruded from the brain by P-gp or some other multidrug-resistant like transport protein in the blood-brain barrier. Despite the promising in vitro results, the low signal-to-noise ratio found in vivo did not justify the use of [<sup>11</sup>C]M-PEPy as a PET radiotracer in humans (Severance et al. 2006).

Kessler labeled 2-methyl-6-(3-fluoro-phenylethynyl)-pyridine (M-FPEP), another analogue of MPEP, with <sup>11</sup>C (Kessler 2014). [<sup>11</sup>C]M-FPEP was prepared in good radiochemical yields and high molar activity starting from its bromo-pyridine precursor. Saturation assays of [<sup>11</sup>C]M-FPEP binding resulted in a single high-affinity binding site with a  $K_d$  of  $1.4 \pm 0.1$  nM and a  $B_{max}$  value of  $563 \pm 190$  fmol/mg protein. Dynamic PET studies in a rat indicated rapid uptake of [<sup>11</sup>C]M-FPEP in rat brain, followed by a fast clearance. Classical biodistribution studies in rats showed a homogeneous distribution of [<sup>11</sup>C]M-FPEP in all brain regions examined, suggesting nonspecific uptake in the brain, which was also confirmed in blocking studies.

Based on the data summarized in Table 13.2, although moderate to high binding affinities toward mGluR5 for all the <sup>11</sup>C-labeled MPEP derivatives have been observed, their utility as mGluR5 imaging agents has been limited due to their unfavorable in vivo properties. The radiosynthesis of [<sup>18</sup>F]FPEP ([<sup>18</sup>F]3-fluoro-5-[(pyridin-3-yl)ethynyl] benzonitrile), a structural analogue of M-FPEP, was accomplished via a nucleophilic aromatic substitution reaction using either nitro or Cl<sup>-</sup> as a leaving group under conventional or microwave heating conditions (Belanger et al. 2008; Hamill et al. 2005; Patel et al. 2007; Wang et al. 2007). Low radiochemical yields were obtained under both conditions (Scheme 13.6). The low radiochemical yields may be due to the position of the leaving group which is not activated enough for aromatic nucleophilic substitution. Increased radiochemical yield ( $20\% \pm 5\%$ ,  $n = 3$ ) was achieved when using a hypervalent spirocyclic iodonium(III) ylide precursor (Stephenson et al. 2015). High binding affinities of

**Table 13.2** Lipophilicity, in vitro binding affinity, and specificity of <sup>11</sup>C-labeled MPEP derivatives

Compd	CLogP <sup>a</sup>	IC <sub>50</sub>	Specificity (brain)
[ <sup>11</sup> C]MPEP	3.6	20 nM	No
[ <sup>11</sup> C]M-MPEP	3.7	3.6 nM	Low
[ <sup>11</sup> C]M-PEPy	1.5	3.4 nM ( $K_d$ ) <sup>b</sup>	Middle
[ <sup>11</sup> C]M-FPEP	2.5	1.4 nM ( $K_d$ ) <sup>b</sup>	No

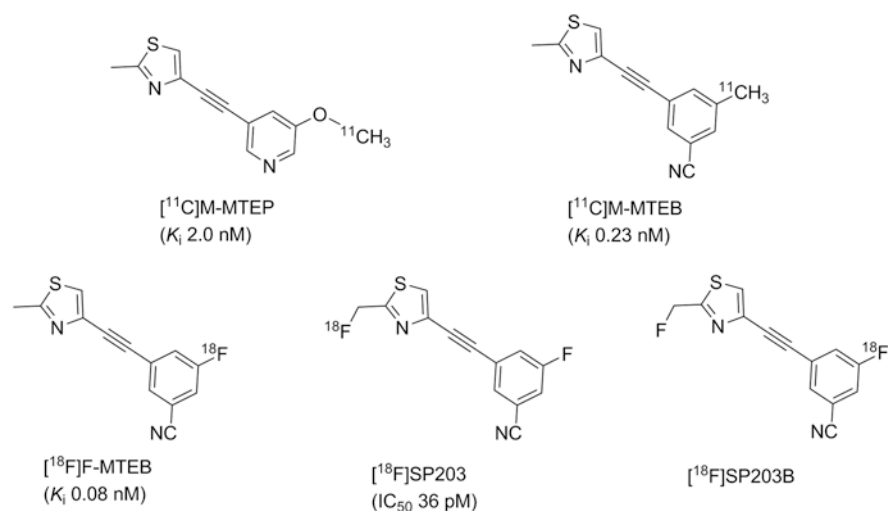
<sup>a</sup>CLogP values were calculated based on ChemBioDraw software

<sup>b</sup>Obtained from saturation assays using [<sup>3</sup>H]M-FPEP or [<sup>11</sup>C]M-FPEP

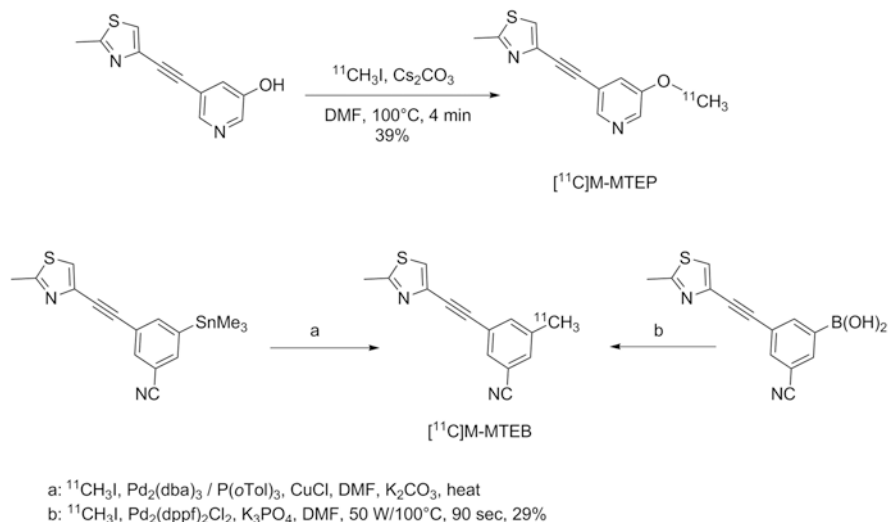
[ $^{18}\text{F}$ ]FPEB toward mGluR5 in rat, rhesus, and human brain sections have been reported ( $K_d = 0.1\text{--}0.15\text{ nM}$ ,  $n \geq 3$ ) (Patil et al. 2007). [ $^{18}\text{F}$ ]FPEB showed an approximately 70% specific signal in the cortex and caudate-putamen of rat, rhesus, and human brain (Patil et al. 2007). Small animal PET imaging studies using Sprague-Dawley rats demonstrated that [ $^{18}\text{F}$ ]FPEB accumulated in mGluR5-rich regions of the brain such as striatum and hippocampus with 77% specific binding after 20 min postinjection (Patel et al. 2005). In blocking studies of [ $^{18}\text{F}$ ]FPEB, pretreatment with 3 mg/kg, MTEP did not result in identical residual uptake in all regions. When [ $^{18}\text{F}$ ]FPEB was blocked with 10 mg/kg MTEP, nearly homogeneous uptake of tracer in all regions was observed, indicating full blockade had been achieved. It was also found that blocking mGluR5 is more efficient with MTEP than MPEP under the same conditions (Wang et al. 2007). PET imaging studies in a monkey also showed that [ $^{18}\text{F}$ ]FPEB readily enters the brain and provides mGluR5-specific signal in all gray matter regions, including the cerebellum (Hamill et al. 2005). Initial human studies have been reported by Tamagnan (Tamagnan et al. 2009). Dynamic PET images of six human subjects examined so far revealed excellent penetration of [ $^{18}\text{F}$ ]FPEB into human brain with initial visualization of anterior cingulate, thalamus, caudate, and midbrain, while cerebellum showed low uptake.

### 13.5.2 MTEP-Derived mGluR5 Tracers

Structural modification of MTEP (Fig. 13.5) led to M-MTEP (4-(5-methoxypyridin-3-yl)ethynyl-2-methyl-4,5-dihydrothiazole,  $K_i$  2.0 nM) (Cosford et al. 2003a; Patel et al. 2005) and M-MTEB (3-methyl-5-[(2-methyl-1,3-thiazol-4-yl)



**Fig. 13.7** Chemical structures of  $^{11}\text{C}$ - and  $^{18}\text{F}$ -labeled MTEP derivatives



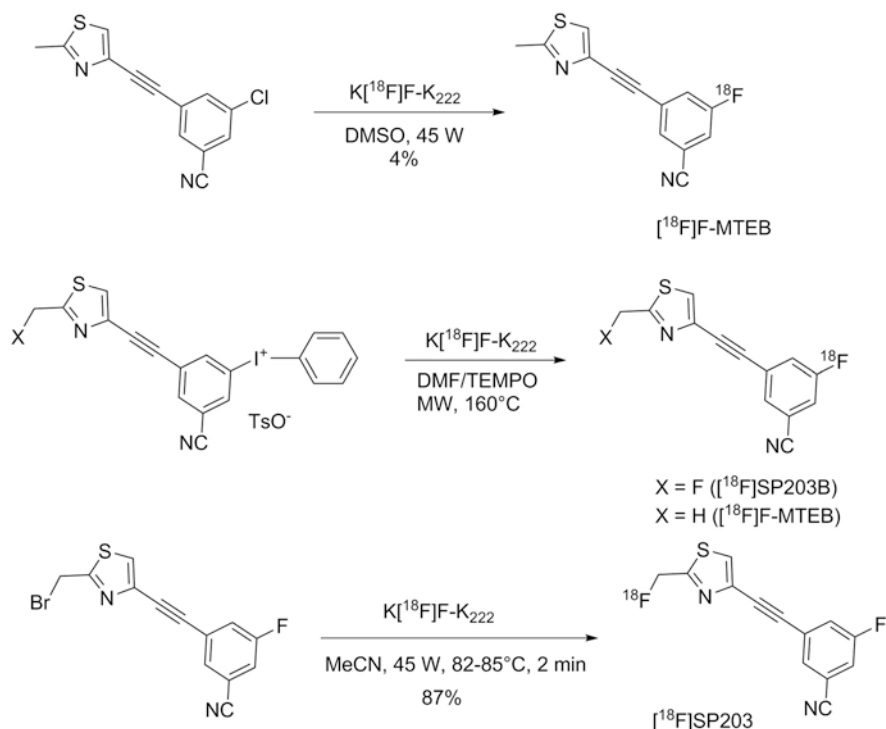
**Scheme 13.7** Radiosynthetic routes to  $[^{11}\text{C}]\text{M-MTEP}$  and  $[^{11}\text{C}]\text{M-MTEB}$

ethynyl]benzotrile,  $K_i$  0.23 nM) with improved binding affinity to mGluR5. The corresponding carbon-11- and fluorine-18-labeled PET radioligands derived from MTEP are shown in Fig. 13.7.

$[^{11}\text{C}]\text{M-MTEP}$  was synthesized by reacting the cesium salt of 5-((2-methyl-4,5-dihydrothiazol-4-yl)ethynyl)pyridin-3-ol with  $[^{11}\text{C}]\text{MeI}$  in 39% radiochemical yield (Patel et al. 2005). The  $[^{11}\text{C}]\text{methyl}$  group in  $[^{11}\text{C}]\text{M-MTEB}$  can be introduced by using either Stille (Suzuki et al. 1997) or Suzuki (Hostetler and Burns 2003) coupling reaction with  $[^{11}\text{C}]\text{MeI}$  as shown in Scheme 13.7. However, due to the ease of precursor synthesis and radiolabeling, the Suzuki route (Scheme 13.7, pathway b) with aryl boronic acid as the precursor was chosen (Hamill et al. 2005). A radiochemical yield of 29% was obtained for  $[^{11}\text{C}]\text{M-MTEB}$  using DMF as solvent. The total synthesis time was ~20–23 min, and the molar activity achieved was 70 GBq/ $\mu\text{mol}$ .

PET imaging studies with  $[^{11}\text{C}]\text{M-MTEP}$  in a rat and a rhesus monkey showed a uniform distribution across most cerebral regions, and its clearance from the brain was also rapid, suggesting little retention by mGluR5 (Hamill et al. 2003; Krause et al. 2003). More promising results were obtained with  $[^{11}\text{C}]\text{M-MTEB}$ .  $[^{11}\text{C}]\text{M-MTEB}$  showed rapid uptake into the brain of rhesus monkey and provided large, long-lasting specific signals in mGluR5-rich regions such as the striatum, lower uptake in the frontal cortex, but surprisingly also specific uptake in the cerebellum, a region known to contain low densities of mGluR5. Species differences in the distribution of mGluR5 may account for this uptake in the cerebellum (Hamill et al. 2005).

$[^{18}\text{F}]\text{F-MTEB}$  ( $[^{18}\text{F}]\text{3-Fluoro-5-}[(2\text{-methyl-1,3-thiazol-4-yl)ethynyl]benzotrile$ ) was synthesized (Scheme 13.8) by the same synthetic route as shown for  $[^{18}\text{F}]\text{FPFB}$  (Scheme 13.6) in low radiochemical yields (Hamill et al. 2005; Patel et al. 2005). The binding affinity ( $K_i$ ) of F-MTEB toward mGluR5 was in the



**Scheme 13.8** Radiosynthetic routes to  $[\text{}^{18}\text{F}]\text{F-MTEB}$ ,  $[\text{}^{18}\text{F}]\text{SP203}$ , and  $[\text{}^{18}\text{F}]\text{SP203B}$

subnanomolar range (80 pM). High-quality mGluR5-specific images in rat and rhesus monkey PET studies were obtained with  $[\text{}^{18}\text{F}]\text{F-MTEB}$ . The low radiochemical yield of 4% would limit its application in human subjects; however, the recently reported high yielding method that uses iodonium salt as a precursor in the presence of TEMPO could increase the RCY of  $[\text{}^{18}\text{F}]\text{F-MTEB}$  threefold to 20% (Telu et al. 2011). This is a step forward which would make possible the application of  $[\text{}^{18}\text{F}]\text{F-MTEB}$  in humans.

3-Fluoro-5-(2-(2-(fluoromethyl)thiazol-4-yl)ethynyl) benzonitrile (designated SP203) an analogue of F-MTEB was found to have unusually high affinity ( $\text{IC}_{50} = 36$  pM) and potency in a phosphoinositol hydrolysis assay ( $\text{IC}_{50} = 0.71$  pM) for mGluR5 (Simeon et al. 2007). SP203 was labeled with fluorine-18 by treatment of its bromomethyl analogue with  $[\text{}^{18}\text{F}]\text{-fluoride}$  in high radiochemical yield (87%, Scheme 13.8).

The PET evaluation of  $[\text{}^{18}\text{F}]\text{SP203}$  in a monkey demonstrated that a high proportion of radioactivity in brain was bound to mGluR5. However, radioactivity also accumulated in the bone, suggesting *in vivo* defluorination (Simeon et al. 2007). *Ex vivo* analysis of metabolites in rat brain extracts indicated that defluorination of  $[\text{}^{18}\text{F}]\text{SP203}$  occurs by glutathionylation at the 2-fluoromethyl group in  $[\text{}^{18}\text{F}]\text{SP203}$  (Shetty et al. 2008).  $[\text{}^{18}\text{F}]\text{SP203}$  was further evaluated in human subjects by the same group (Brown et al. 2008; Kimura et al. 2010). The results showed that

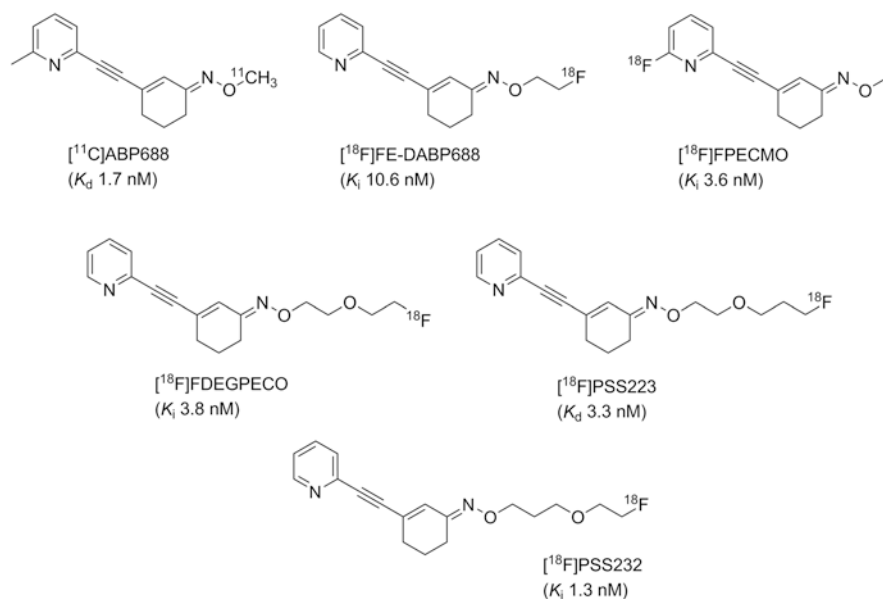


defluorination of [ $^{18}\text{F}$ ]SP203 was relatively small and that brain uptake could be robustly quantified. A recent study in nine healthy subjects using a bolus plus constant infusion protocol (equilibrium method) showed that this method is a viable alternative method and is slightly more precise than the standard bolus injection and kinetic modeling method (Kimura et al. 2012). To circumvent the problem of in vivo radiodefлуorination, the aryl fluorine part of SP203 was recently labeled with  $^{18}\text{F}$  at the meta-position on the aromatic ring (Telu et al. 2011). Because this position is not activated enough toward aromatic nucleophilic substitution, iodonium salt was used as a precursor. A radiochemical yield of 33% was obtained (Scheme 13.8). Compared to [ $^{18}\text{F}$ ]SP203, it is expected that [ $^{18}\text{F}$ ]SP203B will show a higher in vivo stability with respect to in vivo  $^{18}\text{F}$ -defluorination.

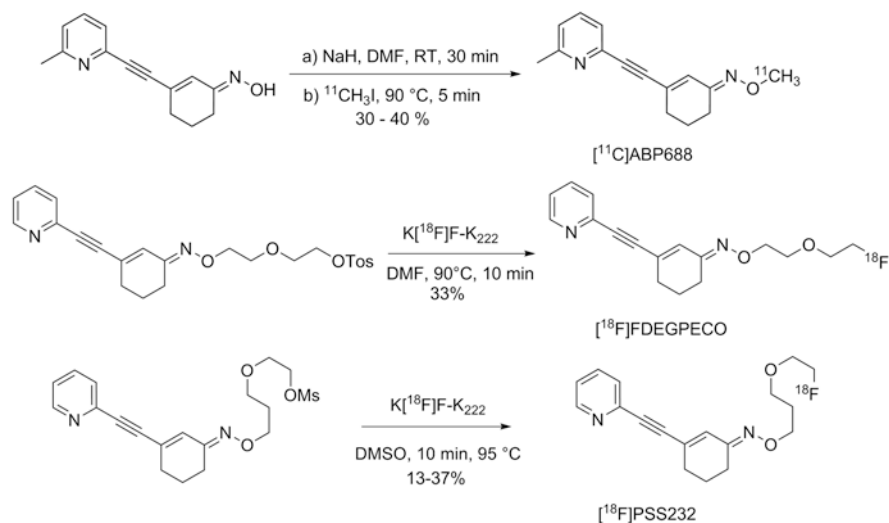
### 13.5.3 ABP688-Related mGluR5 Tracers

Chemical modification of the original MPEP series allowed the identification of ABP688, a derivative in which the aromatic ring of the MPEP series is replaced by a functionalized cyclohexanone moiety (Fig. 13.8).

[ $^{11}\text{C}$ ]ABP688(3-(6-methyl-pyridin-2-ylethynyl)-cyclohex-2-enone-*O*- $^{11}\text{C}$ -methyl-oxime) was described in 2006 by Ametamey et al. (Ametamey et al. 2006). The radiosynthesis of ABP688 was accomplished by reacting desmethyl-ABP688 in anhydrous DMF with [ $^{11}\text{C}$ ]MeI at 90 °C for 5 min using sodium hydride as a base (Scheme 13.9). [ $^{11}\text{C}$ ]ABP688 showed an optimal log*D* value of 2.4 and high binding



**Fig. 13.8** Chemical structures of  $^{11}\text{C}$ - and  $^{18}\text{F}$ -labeled ABP derivatives

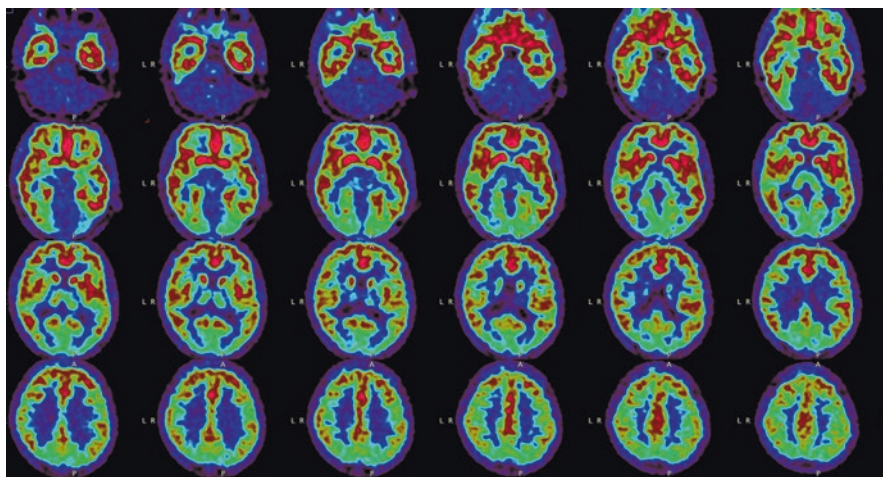


**Scheme 13.9** Radiosyntheses of  $[^{11}\text{C}]\text{ABP688}$ ,  $[^{18}\text{F}]\text{FDEGPECO}$ , and  $[^{18}\text{F}]\text{PSS232}$

affinity to mGluR5 ( $K_d$  1.7 nM in rat) with a  $B_{\text{max}}$  value of  $231 \pm 18$  pmol/mg protein. Ex vivo autoradiography in wild-type mice and rats showed that the brain uptake of  $[^{11}\text{C}]\text{ABP688}$  was highly selective and specific, with high heterogeneous accumulation in known mGluR5-rich regions such as the hippocampus, caudate putamen, and cortex. The specificity of  $[^{11}\text{C}]\text{ABP688}$  binding was confirmed by blockade studies with M-MPEP (1 mg/kg), an antagonist for mGluR5. Up to 80% reduction in radioactivity uptake in mGluR5-rich regions (hippocampus, striatum) was observed; however, no blocking effect was observed in the cerebellum. The observed heterogeneity of tracer uptake is consistent with the known distribution of mGluR5 in the brain. The specificity of binding of  $[^{11}\text{C}]\text{ABP688}$  was also confirmed in mGluR5-knockout mouse brain. Using the cerebellum as a reference region, radioactivity uptake ratios of  $6.6 \pm 0.1$ ,  $5.4 \pm 0.1$ , and  $4.6 \pm 0.1$  were obtained for the striatum, hippocampus, and cortex, respectively.

PET studies in rats and mice using a small animal PET scanner also demonstrated receptor-specific uptake in the brain regions known to be rich in mGluR5.

The first human PET studies on six healthy male volunteers using  $[^{11}\text{C}]\text{ABP688}$  as the PET tracer were reported by the same group (Ametamey et al. 2007; Treyer et al. 2008). Relatively high radioactivity concentrations were observed in mGluR5-rich brain regions such as the anterior cingulate, medial temporal lobe, amygdala, caudate, and putamen, whereas radioactivity uptake in the cerebellum and white matter, regions known to contain low densities of mGluR5, was low (Fig. 13.9). Specific distribution volume as an outcome measure of mGluR5 density in the various brain regions ranged from  $5.45 \pm 1.47$  (anterior cingulate) to  $1.91 \pm 0.32$  (cerebellum), and the rank order of the corresponding specific distribution volumes of  $[^{11}\text{C}]\text{ABP688}$  in cortical regions was temporal >frontal >occipital >parietal. The rank order of specific distribution volume obtained was in



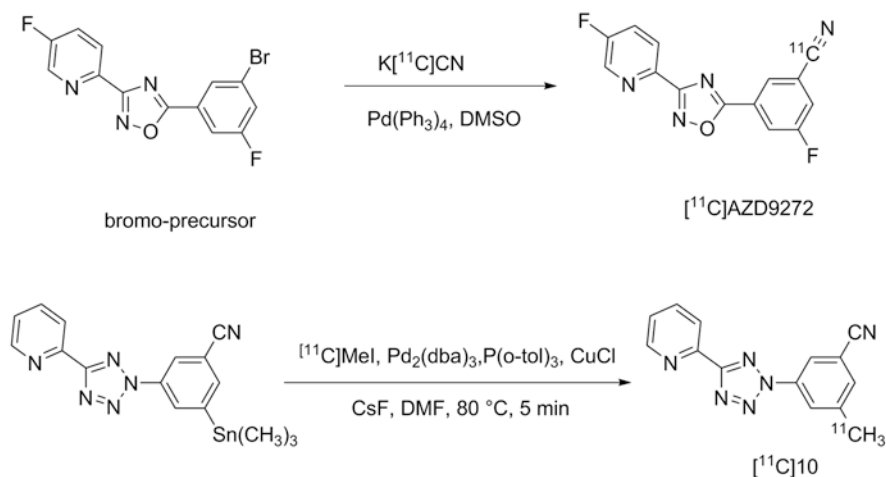
**Fig. 13.9** Uptake of [ $^{11}\text{C}$ ]ABP688 in a healthy volunteer

accordance with the reported distribution pattern of mGluR5 in rodents and humans. These results indicated that [ $^{11}\text{C}$ ]ABP688 has suitable characteristics in humans and is a promising PET ligand for imaging mGluR5 distribution in humans.

Encouraged by the animal and human PET imaging results of [ $^{11}\text{C}$ ]ABP688, several novel  $^{18}\text{F}$ -labeled ABP688 derivatives have been evaluated as potential mGluR5 imaging agents by the group that first reported on [ $^{11}\text{C}$ ]ABP688 (Wanger-Baumann et al. 2011; Baumann et al. 2010; Honer et al. 2007; Lucatelli et al. 2009; Sephton et al. 2012; Sephton et al. 2015).

[ $^{18}\text{F}$ ]FE-DABP688 was produced by reacting 2-[ $^{18}\text{F}$ ]-fluoroethyl tosylate with the sodium salt of 3-(pyridin-2-ylethynyl)-cyclohex-2-enone-oxime in dry DMF. The isolated radiochemical yield was  $25 \pm 8\%$  (Honer et al. 2007). Saturation assays of [ $^{18}\text{F}$ ]FE-DABP688 revealed a single high-affinity binding site with a  $K_d$  of 1.6 nM and a  $B_{\text{max}}$  value of  $119 \pm 24$  fmol/mg protein. Biodistribution and PET studies of [ $^{18}\text{F}$ ]FE-DABP688 in rats indicated radioactivity uptake in mGluR5-rich regions such as the hippocampus, striatum, and cortex, while radioactivity accumulation in the cerebellum was significantly low. Although the obtained PET image quality of [ $^{18}\text{F}$ ]FE-DABP688 was comparable to that of [ $^{11}\text{C}$ ]ABP688, [ $^{18}\text{F}$ ]FE-DABP688 displayed unfavorable pharmacokinetics in the anesthetized rat with a fast washout from the forebrain resulting in a relatively short-lived signal. Consequently, [ $^{18}\text{F}$ ]FE-DABP688 was abandoned and was not further exploited in primates.

[ $^{18}\text{F}$ ]-(*E*)-3-((6-Fluoropyridin-2-yl)ethynyl)cyclohex-2-enone *O*-methyl oxime ([ $^{18}\text{F}$ ]FPECMO) is another derivative of [ $^{11}\text{C}$ ]ABP688 (Lucatelli et al. 2009). It was synthesized in a one-step reaction sequence by reacting  $\text{K}[^{18}\text{F}]\text{F-K}_{222}$  complex with (*E*)-3-((6-bromopyridin-2-yl)ethynyl)cyclohex-2-enone *O*-methyl oxime in dry DMSO. The radiochemical yield after semi-preparative HPLC was 35%, and molar



**Scheme 13.10** Radiosynthesis of  $[^{11}\text{C}]\text{AZD9272}$  and  $[^{11}\text{C}]\text{10}$

activity was greater than  $240 \text{ GBq}/\mu\text{mol}$ .  $[^{18}\text{F}]\text{FPECMO}$  exhibited optimal lipophilicity ( $\log D = 2.1$ ) and high metabolic stability in vitro. Displacement studies revealed a  $K_i$  value of  $3.6 \text{ nM}$  for FPECMO. Biodistribution studies and ex vivo autoradiographies confirmed this specificity revealing highest radioactivity accumulation in mGluR5-rich brain regions. Despite high metabolic stability in various in vitro test systems, a rapid defluorination of the radiolabeled substance was observed in rats in vivo.

$[^{18}\text{F}]\text{FDEGPECO}$  (*E*)-3-(pyridin-2-ylethynyl)cyclohex-2-enone *O*-2-(2-fluoroethoxy)ethyl oxime is one of the most recently developed ABP688 derivatives (Wanger-Baumann et al. 2011; Baumann et al. 2010). A  $K_i$  value of  $3.8 \pm 0.4$  ( $n = 3$ ) nM was obtained for FDEGPECO, and the radioligand was synthesized in one-step reaction sequence by reacting  $\text{K}[^{18}\text{F}]\text{F-K}_{222}$  complex with the corresponding tosyl precursor in anhydrous DMF (Scheme 13.10) in good radiochemical yield and high molar activity. Scatchard plot analysis indicated two affinity binding sites with dissociation constants of  $0.6 \pm 0.2 \text{ nM}$  and  $13.7 \pm 4.7 \text{ nM}$  ( $n = 3$ ) and  $B_{\text{max}1}$  and  $B_{\text{max}2}$  values ranging from 470 to 1870 fmol/mg and from 4.0 to 15.6 pmol/mg protein, respectively. The in vitro autoradiography on rat brain slices showed a heterogeneous uptake consistent with the distribution of mGluR5 in the rat brain. PET imaging on rats revealed specific uptake of radioactivity in the mGluR5-rich brain regions. Postmortem biodistribution studies in rats that received intravenous injections of 8–16 MBq  $[^{18}\text{F}]\text{FDEGPECO}$  at 15 min p.i. confirmed the distribution pattern observed in PET. The highest radioactivity concentrations were observed in mGluR5-rich brain regions. The hippocampus/cerebellum and striatum/cerebellum ratios were 2.8 and 3.3, respectively.  $[^{18}\text{F}]\text{FDEGPECO}$  is one of the most promising fluorine-18-labeled ABP688 analogues so far although its background activity is relatively high and the radioactivity ratios when compared to  $[^{11}\text{C}]\text{ABP688}$  (Ametamey et al. 2006) are lower.

[<sup>18</sup>F]PSS223 ((*E*)-3-(pyridin-2-ylethynyl)cyclohex-2-enone *O*-(2-(3-<sup>18</sup>F-fluoropropoxy)ethyl) oxime), an [<sup>18</sup>F]FDEGPECO analogue, was designed with the aim to increase the lipophilicity of the ligand, in which the side chain was extended by one methylene group (Sephton et al. 2012). [<sup>18</sup>F]PSS223 was obtained in 20% decay-corrected radiochemical yield. Experimentally determined log*D* value of [<sup>18</sup>F]PSS223 was 1.89, which is 0.2 log units higher than that of [<sup>18</sup>F]FDEGPECO (Baumann et al. 2010) as expected. The in vitro binding affinity of [<sup>18</sup>F]PSS223 was measured directly in a Scatchard assay to give a *K<sub>d</sub>* value of  $3.34 \pm 2.05$  nM. In vitro autoradiography on horizontal rat brain slices showed heterogeneous distribution of [<sup>18</sup>F]PSS223 with the highest accumulation in brain regions where mGluR5 is highly expressed (hippocampus, striatum, and cortex). Autoradiography in vitro under blockade conditions with ABP688 also confirmed the high specificity of [<sup>18</sup>F]PSS223 for mGluR5. Under the same blocking conditions but using the mGluR1 antagonist, JNJ-16259685, no blockade was observed demonstrating the selectivity of [<sup>18</sup>F]PSS223 for mGluR5 over mGluR1. [<sup>18</sup>F]PSS223 was stable in PBS and rat plasma but was significantly metabolized by rat liver microsomal enzymes and to a lesser extent by human liver microsomes. Within 60 min, 90 and 20% of [<sup>18</sup>F]PSS223 were metabolized by rat and human microsome enzymes, respectively. PET studies showed that [<sup>18</sup>F]PSS223 has a fast clearance and only a weak accumulation in mGluR5-rich brain regions. A significant accumulation of radioactivity in the skull and jaws was also observed suggesting in vivo defluorination of [<sup>18</sup>F]PSS223. Consequently a clear-cut visualization of mGluR5-rich brain regions in vivo in rats was not possible.

The difference in the in vivo behavior of [<sup>18</sup>F]PSS223 and [<sup>18</sup>F]FDEGPECO could be attributed to the β-heteroatom effect (French et al. 1993), by which primary aliphatic <sup>18</sup>F-atoms in a β-position to heteroatom (e.g., [<sup>18</sup>F]FCH<sub>2</sub>CH<sub>2</sub>OR) are found to be metabolized at a slower rate. This is in line with the absence of in vivo radio-defluorination of [<sup>18</sup>F]FDEGPECO and also (*E*)-3-(pyridin-2-ylethynyl) cyclohex-2-enone *O*-(3-(2-[<sup>18</sup>F]-fluoroethoxy)propyl) oxime ([<sup>18</sup>F]PSS232). [<sup>18</sup>F]PSS232 was designed by our group to improve the metabolic stability of [<sup>18</sup>F]PSS223 by switching the oxygen atom located between the ethylene and the fluoropropyl moieties by one CH<sub>2</sub> unit to the right in the side chain (Fig. 13.8). From the two geometrical *E* and *Z* isomers of PSS232, the *E*-geometrical isomer was shown to exhibit a higher binding affinity when compared to the *Z*-isomer (1 nM for *E*-isomer vs. 15 nM for *Z*-isomer). The *E*-isomer was radiolabeled in one step via nucleophilic substitution of a mesylate leaving group and evaluated in vitro and in vivo (Sephton et al. 2015; Sephton et al. 2013). As expected, increased in vitro stabilities were observed with rat and human microsomal enzymes in 60 min (20% hydrophilic radiometabolites for rat, 4% for human, respectively). No hydrophilic decomposition products or radiometabolites were found in PBS or plasma. The specificity of [<sup>18</sup>F]PSS232 was confirmed in in vitro autoradiography on rat brain slices and in vivo PET images. Radioactivity distribution pattern was consistent with the known distribution of mGluR5 in rat brain with high binding to hippocampal and cortical regions and negligible radioactivity in the cerebellum. Under displacement conditions with MMPEP, reduced [<sup>18</sup>F]PSS232 binding was found in all brain regions except the

cerebellum. Test-retest studies were reproducible with a variability ranging from 6.8% to 8.2%. An extended single-dose toxicity study in Wistar rats showed no compound-related adverse effects (Sephton et al. 2015). Encouraged by the promising preclinical data obtained with [<sup>18</sup>F]PSS232 in rodents, its potential for imaging mGluR5 density in humans was investigated by the same research group (Warnock et al. 2018). [<sup>18</sup>F]PSS232 was evaluated in ten healthy male volunteers aged 20–40 years old, and seven of the volunteers were additionally scanned with [<sup>15</sup>O]water for cerebral blood flow determination. Arterial blood activity was measured using an online blood counter. Brain uptake of [<sup>18</sup>F]PSS232 was very similar to its analog [<sup>11</sup>C]ABP688 and matched the distribution pattern of mGluR5. The uptake kinetics of [<sup>18</sup>F]PSS232 followed a two-tissue compartment model (Warnock et al. 2018). [<sup>18</sup>F]PSS232 is thus the only fluorinated derivative of [<sup>11</sup>C]ABP688 which has so far shown utility for imaging mGluR5 in humans, and it is hoped that this new fluorinated version of ABP688 will find a wider application in clinical studies in the near future. Recently, [<sup>18</sup>F]PSS232 was evaluated in our group for its usefulness to measure mGluR5 expression levels in LPS-induced murine neuroinflammation model and postmortem tissue samples from AD and amyotrophic lateral sclerosis (ALS) patients (Muller Herde et al. 2019). The results indicated that LPS-induced neuroinflammation lead to increased mGluR5 expression in mouse brain. Levels of mGluR5 in human AD and ALS postmortem brain tissues were also found to be several-fold increased compared to control brains. These results suggest that mGluR5 is a useful target to consider in neuroinflammation diseases and that diagnosis and the therapy monitoring of AD and ALS patients could be accomplished via mGluR5 imaging.

### 13.5.4 Other Type of mGluR5 Radioligands Without Alkyne Moiety in MPEP

[<sup>11</sup>C]AZD9272 (3-fluoro-5-(3-(5-fluoropyridin-2-yl)-1,2,4-oxadiazol-5-yl)benzo-[<sup>11</sup>C]nitrile) was labeled using palladium mediated <sup>11</sup>C-cyanation (Andersson et al. 2013). Reacting the bromo-precursor 5-(3-bromo-5-fluorophenyl)-3-(5-fluoropyridin-2-yl)-1,2,4-oxadiazol with K[<sup>11</sup>C]CN in the presence of Pd(PPh<sub>3</sub>) (5.5 mg, 4.8 μmol) afforded [<sup>11</sup>C]AZD9272 over 50% incorporation yield from hydrogen [<sup>11</sup>C]cyanide in a total synthesis time of 45–50 min (Scheme 13.10). AZD9272 showed high affinity ( $K_D = 3.8 \pm 0.7$  nM) toward human mGluR5 and shared the same allosteric binding site as MPEP (Raboisson et al. 2012). In PET studies using *Cynomolgus* monkeys, high and heterogeneous brain uptake were demonstrated. The highest radioactivity was seen in the caudate, cingulate gyrus, and thalamus, whereas uptake was moderate in temporal cortex and lower in the cerebellum. Co-injection of unlabeled AZD9272 led to significantly decreased radioactivity in many brain regions (Andersson et al. 2013). Further studies revealed unique binding sites of [<sup>11</sup>C]AZD9272 in primate brain (Varnas et al. 2018). The binding of [<sup>11</sup>C]AZD9272 in NHP was almost completely inhibited by the structurally unique mGluR5 compound fenobam (2.0 mg/kg; 98% occupancy). In contrast, unlabeled

MTEP and ABP688 (2.0 mg/kg) inhibited [ $^{11}\text{C}$ ]AZD9272 binding only partially by 20% and 46%, respectively. The authors concluded that additional PET studies employing multiple doses of fenobam should be done to further assess the binding of fenobam at AZD9272 binding sites (Varnas et al. 2018).

3-(–[ $^{11}\text{C}$ ]methyl)-5-(5-(pyridin-2-yl)-2H-tetrazol-2-yl)benzotrile designated as [ $^{11}\text{C}$ ]10 in the publication of Shimodu and co-workers (Shimoda et al. 2016) has similar structural attributes as AZD9272, i.e., a tetrazole ring instead of the alkyne group (in MPEP) between two the aromatic rings. A binding affinity ( $K_i$ ) of 9.4 nM and moderate lipophilicity (cLogD, 2.4) have been reported for this compound. The radiolabeling was performed by reacting the arylstannyl precursor with [ $^{11}\text{C}$ ]methyl iodide (Scheme 13.10). PET imaging with [ $^{11}\text{C}$ ]10 in rats visualized the regional distribution of mGluR5 and demonstrated high reliability for quantification of mGluR5 density in test-retest PET studies. This radioligand might serve as a useful lead compound for development new PET ligands with higher affinity and more specific binding to mGluR5 (Shimoda et al. 2016).

Among all the abovementioned mGluR5 PET ligands, [ $^{11}\text{C}$ ]ABP688 and [ $^{18}\text{F}$ ]FPEB are the most successful and widely used PET tracers for imaging mGluR5 in humans. The first-in-man PET studies with [ $^{18}\text{F}$ ]PSS232 confirmed its utility for mapping mGluR5 in humans. The longer physical half-life of F-18 (110 min) and its easy/high yield radiolabeling process will make [ $^{18}\text{F}$ ]PSS232 a very attractive mGluR5 tracer for PET centers without cyclotron facilities.

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### 13.6 Application of mGluR Tracers

With the availability of suitable mGluR5 radioligands, the utility of mGluR5 PET radioligands for drug development and the study of psychiatric and neurological diseases have been demonstrated by various studies in humans and rodents as illustrated in Table 13.3. As mentioned earlier, [ $^{11}\text{C}$ ]ABP688 and [ $^{18}\text{F}$ ]FPEB are the most widely used tracers for imaging mGluR5 densities in the brain. Compared to healthy individuals, altered mGluR5 availability was observed in the brain of stressed subjects, epilepsy, Parkinson's disease, and adults with autism and behavioral variant frontotemporal dementia patients (DuBois et al. 2016a; Fatemi et al. 2018; Holmes et al. 2017; Kang et al. 2019; Lam et al. 2019; Leuzy et al. 2016). Downregulation of mGluR5 was demonstrated in smokers and patients with depression, while sleep-deprived subjects and alcohol use disorders showed an increased mGluR5 availability compared to the healthy controls (Akkus et al. 2013; Akkus et al. 2018; Hefti et al. 2013; Holst et al. 2017). Species difference was realized in the studies of drug induced changes in extracellular glutamate levels on mGluR5 tracer binding in the brains of humans and rodents. In humans, lower [ $^{11}\text{C}$ ]ABP688 binding was observed after administration of ketamine, known to increase glutamate levels (DeLorenzo et al. 2015; Esterlis et al. 2018a), while contradictory results were obtained in rat studies. Increased mGluR5 availability was visualized in rats after administration of ceftriaxone, which decreases glutamate (Zimmer et al. 2015); however, no change in mGluR5 availability was indicated when [ $^{11}\text{C}$ ]ABP688 or [ $^{18}\text{F}$ ]PSS232 was used

**Table 13.3** Application of mGluR5 PET tracers and main findings

PET tracer	Application	Subject	Main findings	Reference
[ <sup>11</sup> C] ABP688	Major depressive disorder	Human	Severity of depression correlated negatively with mGluR5 binding in the hippocampus	Deschwanden et al. (2011)
[ <sup>11</sup> C] ABP688	Nicotine addiction	Human	Marked global reductions in mGluR5 binding in smokers and ex-smokers	Akkus et al. (2013)
[ <sup>11</sup> C] ABP688	Pilocarpine-induced chronic epilepsy	Rat	mGluR5 availability changes during epileptogenesis in the rat models	Choi et al. (2014)
[ <sup>11</sup> C] ABP688	Ketamine administration	Human	Decreased mGluR5 binding after ketamine administration	DeLorenzo et al. (2015)
[ <sup>18</sup> F] FPEB	Receptor occupancy	Rat	Useful as a biomarker for drug development, but in vitro/in vivo difference exists	Rook et al. (2015)
[ <sup>18</sup> F] FPEB	ALS model expressing SOD1- G93A gene	Mouse	Enhanced expression of mGluR5	Brownell et al. (2015)
[ <sup>11</sup> C] ABP688	Epilepsy (ocal cortical dysplasia, FCD)	Human	Reduced mGluR5 availability in FCD	DuBois et al. (2016a)
[ <sup>11</sup> C] ABP688	Age/sex difference	Human	No sex/age effect in healthy subjects	DuBois et al. (2016b)
[ <sup>11</sup> C] ABP688	Behavioral variant frontotemporal dementia (bvFTD)	Human	Decreased availability of mGluR5 in bvFTD	Leuzy et al. (2016)
[ <sup>18</sup> F] FPEB	Novelty-seeking temperament	Human	High novelty-seeking temperament was robustly associated with increased mGluR5 availability	Leurquin-Sterk et al. (2016)
[ <sup>18</sup> F] FPEB	Posttraumatic stress disorder (PTSD)	Human	Higher mGluR5 availability in PTSD	Holmes et al. (2017)
[ <sup>18</sup> F] FPEB	Major depressive disorder	Human	No significant difference of mGluR5 availability	Abdallah et al. (2017)
[ <sup>11</sup> C] ABP688	Schizophrenia	Human	mGluR5 binding has no difference in schizophrenia from healthy control	Akkus et al. (2017)
[ <sup>11</sup> C] ABP688	Huntington disease (Q175 mouse model)	Mouse	Reduced mGluR5 availability in Q175 mouse	Bertoglio et al. (2018)
[ <sup>18</sup> F] FPEB	Parkinson's disease (6-OHDA); levodopa-induced dyskinesia (LID)	Rat	Altered mGluR5 availability in PD and LID rats	Crabbe et al. (2018)
[ <sup>18</sup> F] PSS232	Ketamine or ceftriaxone	Rat	No change of mGluR5 availability	Muller Herde et al. (2018)



**Table 13.3** (continued)

PET tracer	Application	Subject	Main findings	Reference
[ <sup>18</sup> F] FPEB	Adults with autism	Human	mGluR5 binding is altered in critical brain areas of subjects with autism	Fatemi et al. (2018)
[ <sup>18</sup> F] FPEB	Cocaine self-administration	Rat	Reduced mGluR5 availability in rats	de Laat et al. (2018)
[ <sup>11</sup> C] ABP688	Ketamine infusion	Rat	No change of mGluR5 availability	Kosten et al. (2018)
[ <sup>11</sup> C] ABP688	Alcohol use disorder	Human	Altered mGluR5 signaling in the amygdala in alcohol use disorder	Akkus et al. (2018)
[ <sup>11</sup> C] ABP688	Major depressive disorder	Human	Ketamine-induced reduction in mGluR5 availability is associated with an antidepressant response	Esterlis et al. (2018b)
[ <sup>18</sup> F] PSS232	LPS-induced Neuroinflammation	Mouse	Increased mGluR5 levels in LPS-treated mouse brain	Muller Herde et al. (2019)
[ <sup>18</sup> F] FPEB	Alcohol exposure	Rat	Decreased mGluR5 availability in the hippocampus	de Laat et al. (2019)
[ <sup>18</sup> F] FPEB	Parkinson's disease (PD)	Human	Slightly increased mGluR5 availability	Kang et al. (2019)
[ <sup>11</sup> C] ABP688	Sex differences	Human	PET [ <sup>11</sup> C]ABP688 is higher in healthy men than women	Smart et al. (2019b)
[ <sup>11</sup> C] ABP688	Mesial temporal lobe epilepsy (MTLE)	Human	Reduced mGluR5 availability	Lam et al. (2019)

to assess endogenous glutamate fluctuations induced by either ketamine or ceftriaxone (Kosten et al. 2018; Muller Herde et al. 2018). [<sup>11</sup>C]ABP688 binding to brain tissue was reported to be higher in healthy men than in women (Akkus et al. 2013; Smart et al. 2019b); however, some other studies found no effect of gender on radioligand binding (Abdallah et al. 2017; DuBois et al. 2016b). Given that the (*Z*)-isomer of [<sup>11</sup>C]ABP688 has a low affinity and affects binding potential values (Smart et al. 2019a), efforts should be undertaken to minimize the amount of the (*Z*)-isomer in the formulated solution of [<sup>11</sup>C]ABP688. [<sup>11</sup>C]ABP688 has shown good reproducibility of mGluR5 quantification in rodents and non-human primates (DeLorenzo et al. 2011b; Elmenhorst et al. 2012). In humans, high within-subject variability of  $V_T$  and  $BP_{ND}$  in the same-day scans was observed in test-retest studies with [<sup>11</sup>C]ABP688 (DeLorenzo et al. 2017; DeLorenzo et al. 2011a); however, variability was low when the time interval between the scans was greater than 7 days (Burger et al. 2010). Test-retest variability for [<sup>18</sup>F]FPEB was also high when scans were performed on the same day. Because similar high same-day within-subject variability was observed for both [<sup>11</sup>C]ABP688 and [<sup>18</sup>F]FPEB, it is suggested that this phenomenon may be related to alterations in glutamatergic activity which cause

up- and downregulation of mGluR5 (DeLorenzo et al. 2017; DeLorenzo et al. 2011a). It is further suggested that studies using these radiotracers should be carefully controlled and that further work should be done to determine sources and mechanisms of mGluR5 PET tracer variability in humans (Smart et al. 2018).

The mGluR1 PET tracer [ $^{11}\text{C}$ ]ITMM has been used for several clinical studies since its first-in-man study in 2013 (Toyohara et al. 2013b). Decreased mGluR1 availability was observed in a patient with spinocerebellar ataxia type 6 compared to age-matched healthy controls (Ishibashi et al. 2015). In comparison with the most often used [ $^{18}\text{F}$ ]FDG, [ $^{11}\text{C}$ ]ITMM demonstrated higher specificity for evaluating cerebellar ataxia in human subjects (Ishibashi et al. 2017). Recently, [ $^{11}\text{C}$ ]ITMM was also used in a PET study to measure mGluR1 in relatively early stage of Alzheimer's disease patients; however, no significant changes in mGluR1 availability were found in these patients (Ishibashi et al. 2019). Future studies in patients with the fluorinated derivative, [ $^{18}\text{F}$ ]FIMX, for imaging mGluR1 are highly expected given its demonstrated utility in first-in-man studies (Zanotti-Fregonara et al. 2016).

In summary, a number of group I mGluRs ligands have been successfully used to label mGluR1 and mGluR5 *in vitro*. The development of these ligands into useful PET tracers *in vivo*, however, in most cases failed due to high nonspecific binding, unfavorable brain uptake kinetics, and/or limited metabolic stability. So far, only three mGluR5 and two mGluR1 PET radioligands have demonstrated their usefulness in human studies. Several chemical scaffolds as lead structures for targeting mGluR2 and mGluR4 subtypes are currently under evaluation as potential PET radioligands. We are optimistic that the translation of the pre-clinical results to the clinic will be feasible in the near future for radioligands targeting mGluR2 and mGluR4. There are currently no PET radioligands for mGluR subtypes 6, 7, and 8. Different mGluR subtype PET radioligands would have high utility in drug development programs by supporting the selection of appropriate doses of clinically relevant drugs and the determination of receptor occupancy of drugs, guiding proof of concept studies, and characterizing various disease states related to the type of receptor involved. Furthermore, they will have great implications for the identification of potential molecular therapeutic targets, in the development of new treatment strategies, and in helping to understand psychiatric and neurological diseases.

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# PET and SPECT Imaging of Steroid Hormone Receptors in the Brain

# 14

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## Abstract

Steroid hormones, like sex hormones and corticosteroids, are involved in normal brain function. They can influence behavior and higher functions via binding to the corresponding hormone receptors. Steroid hormones also play a crucial role in psychiatric disorders by interacting with different neurotransmitter systems in the brain. Most of the studies in this field have been performed on postmortem

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tissue of animals and humans. However, noninvasive techniques could provide new insight into the role that steroid hormones play in normal and pathological brain function. PET and SPECT are techniques that allow noninvasive quantitative analysis of the expression of steroid hormone receptors. This chapter will briefly explain the role of steroid hormones in physiological and pathological conditions and provide an overview of the available PET and SPECT imaging methods that could be used to study the role steroid hormones and their receptors in the brain.

## Abbreviations

(Z)-[ <sup>123</sup> I]MIVE	(20Z)-11beta-methoxy-17alpha-[ <sup>123</sup> I]iodovinylestradiol
[ <sup>123</sup> I]IES	16alpha-[ <sup>123</sup> I]iodo-17beta-estradiol
[ <sup>123</sup> I]IVMMNT	17alpha-[ <sup>123</sup> I]iodovinyl-18-methyl-11-methylene-19-nortestosterone
[ <sup>123</sup> I]TAM	[ <sup>123</sup> I]iodotamoxifen
[ <sup>125</sup> I]IVNT	(20Z)-17alpha-[ <sup>125</sup> I]iodovinyl-19-nortestosterone
[ <sup>125</sup> I]MIE	16alpha-[ <sup>125</sup> I]iodo-11beta-methoxy-17beta-estradiol
[ <sup>18</sup> F]betaFMOX	17alpha-ethynyl-16beta-[ <sup>18</sup> F]fluoro-11beta-methoxy-estradiol
[ <sup>18</sup> F]FDHT	16beta-[ <sup>18</sup> F]fluorodihydrotestosterone
[ <sup>18</sup> F]FENP	21-[ <sup>18</sup> F]fluoro-16alpha-ethyl-19-norprogesterone
[ <sup>18</sup> F]FES	16alpha-[ <sup>18</sup> F]fluoro-17beta-estradiol
[ <sup>18</sup> F]FFNP	21-[ <sup>18</sup> F]fluoro-16alpha,17alpha-(R)-(1'alpha-furylmethylidene)dioxy]-19-norpregn-4-ene-3,20-dione
[ <sup>18</sup> F]-FHNP	2-[ <sup>18</sup> F]fluoro-6-[6-hydroxynaphthalen-2-yl]pyridin-3-ol
[ <sup>18</sup> F]FMDHT	7alpha-[ <sup>18</sup> F]fluoro-17-methyl-5-dihydrotestosterone
[ <sup>18</sup> F]FMNP	21-[ <sup>18</sup> F]fluoro-16alpha-methyl-19-norprogesterone
[ <sup>18</sup> F]FPTP	4-[ <sup>18</sup> F]fluoropropyl-tanaproget
[ <sup>18</sup> F]FTX	[ <sup>18</sup> F]Fluoromethyl- <i>N,N</i> -dimethyltamoxifen
16alpha-[ <sup>18</sup> F]FMNT	16alpha-[ <sup>18</sup> F]fluoro-7alpha-methyl-19-nortestosterone
16beta-[ <sup>18</sup> F]FMNT	16beta-[ <sup>18</sup> F]fluoro-7alpha-methyl-19-nortestosterone
16beta-[ <sup>18</sup> F]FT	16beta-[ <sup>18</sup> F]fluorotestosterone
20-[ <sup>18</sup> F]Fmib	20-[ <sup>18</sup> F]fluoromibolone
20-[ <sup>18</sup> F]R1881	20-[ <sup>18</sup> F]fluorometribolone
4F-M[ <sup>18</sup> F]FES	11beta-methoxy-4,16alpha-[ <sup>18</sup> F]difluoroestradiol
7alpha-[ <sup>125</sup> I]IDHT	7alpha-[ <sup>125</sup> I]iodo-5alpha-dihydrotestosterone
8beta-[ <sup>18</sup> F]FEE	8beta-(2-[ <sup>18</sup> F]fluoroethyl)estradiol
AR	Androgen receptors
CNS	Central nervous system
DES	Diethylstilbestrol
ER	Estrogen receptors
GR	Glucocorticoid receptors
HPA axis	Hypothalamic-pituitary-adrenal axis

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HRE	Hormone response elements
MR	Mineralocorticoid receptors
PR	Progesterone receptors
SHBG	Sex hormone-binding globulin
SHR	Steroid hormone receptors

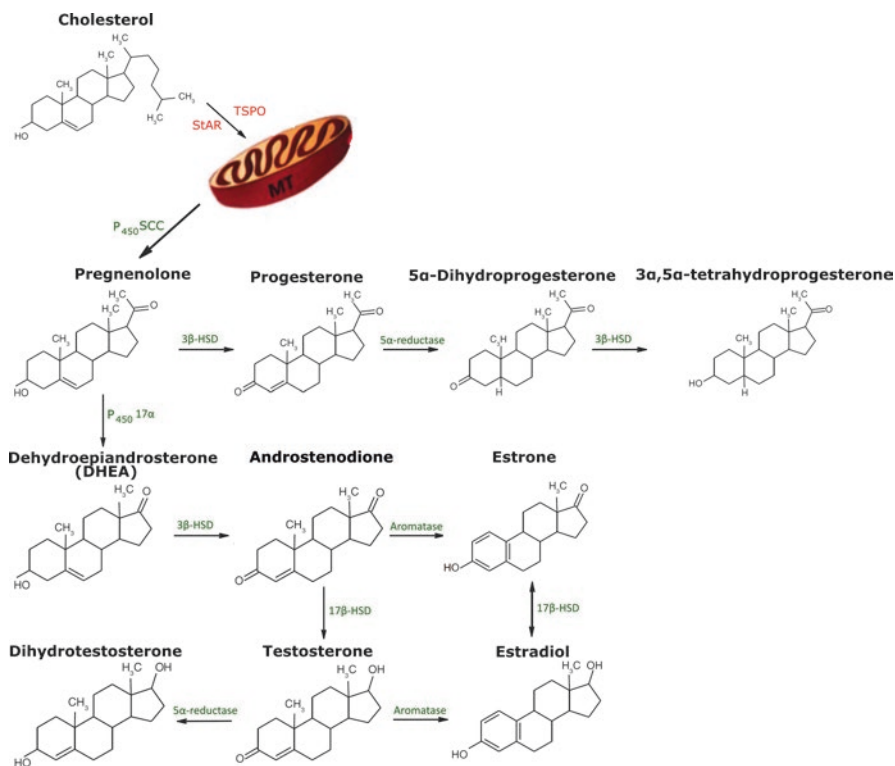
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## 14.1 Introduction

Steroid hormones are messenger molecules that are produced and secreted by endocrine glands, such as ovaries, testes, and adrenal cortex. Structurally all steroid hormones consist of a steroidal scaffold, which is composed of three six-membered rings and one five-membered ring; nonsteroidal hormones lack this steroid scaffold. Secreted steroid hormones can travel through the blood vessels toward their receptors and can easily cross cell membranes, including the blood-brain barrier, because of their high lipophilicity. Steroidal hormones can be classified into two main families: sex hormones and corticosteroids. In general, they are responsible for biological responses in a wide range of endocrine processes, including sexual differentiation, reproductive physiology, glucose metabolism, and maintenance of cell homeostasis. The sex hormones can be further classified into three subfamilies: progestins, estrogens, and androgens.

The brain is an important target organ for circulating steroid hormones. It is known that these molecules can affect several brain functions (McEwen and Milner 2017; Zalachoras et al. 2013). Steroid hormones can either bind directly to their respective membrane receptors or cross the cell membrane and bind to intracellular receptors (Gray et al. 2017; Kawata et al. 2008; Mani et al. 2012). Although circulating steroid hormones can diffuse across the blood-brain barrier, the brain itself is also capable of synthesizing steroid hormones *de novo* from cholesterol (Fig. 14.1), as was first discovered in studies, in which peripheral organs that produce steroid hormones were surgically removed (Porcu et al. 2016). Neuroactive steroids (neurosteroids) produce their effects by autocrine or endocrine pathways (Baulieu 1998) and act mostly as allosteric modulators of different neurotransmitter receptors (for a review on the functions of neurosteroids, see Rupprecht (2003)).

Neuroactive steroids can act as allosteric modulators of ligand-gated ion channels (GABA<sub>A</sub>), NMDA, and sigma receptors. Because of these interactions, neuroactive steroids have sleep-inducing, anticonvulsant, anesthetic, nootropic, and antipsychotic properties, among others (Rupprecht 2003). They also play a crucial role in neuronal development and plasticity, including processes like learning, memory, emotion, behavior, synaptic transmission, and neuroprotection (Giatti et al. 2012). Because of these neuroprotective properties, neurosteroids have been proposed as adjuvant treatment for neurodegenerative diseases and neuropsychiatric disorders like Alzheimer's disease, multiple sclerosis, Parkinson's disease, depression and anxiety-related disorders, among others (Borowicz et al. 2011; Eser et al. 2006).



**Fig. 14.1** Scheme of neurosteroidogenesis. Steroidogenesis follows a sequential, highly compartmentalized procedure, starting with the translocation of cholesterol from cytoplasm to mitochondria in the cells of the CNS, which is mediated by steroidogenic acute regulatory protein (StAR) and an 18 kDa translocator protein (TSPO). In the mitochondria, P450 side-chain cleavage (P450SCC) cleaves the side chain from cholesterol, resulting in the formation of pregnenolone. Pregnenolone is subsequently converted to progesterone and dehydroepiandrosterone in endoplasmic reticulum. The lipophilic nature of these compounds allows them to diffuse from one cell to another. Progesterone and dehydroepiandrosterone are further metabolized to form other neuroactive metabolites like testosterone, estrone, and estradiol

In this chapter, we will provide an overview of the available literature about PET and SPECT imaging techniques that can be applied to study the role of steroid hormones and their receptors in the brain. We will start discussing the role of steroidal hormones in the brain and subsequently provide an overview of the available PET and SPECT tracers for imaging of steroid hormone receptors and their applicability in brain research.

## 14.2 Steroid Hormones and Their Receptors in Brain Disorders

Steroid hormones exert their biological effects through specific steroid hormone receptors (SHR) that are expressed by the target cells. To date, two estrogen receptor (ER) subtypes (ER $\alpha$  and ER $\beta$ ) with several isoforms (Ogino et al. 2018; Tonn

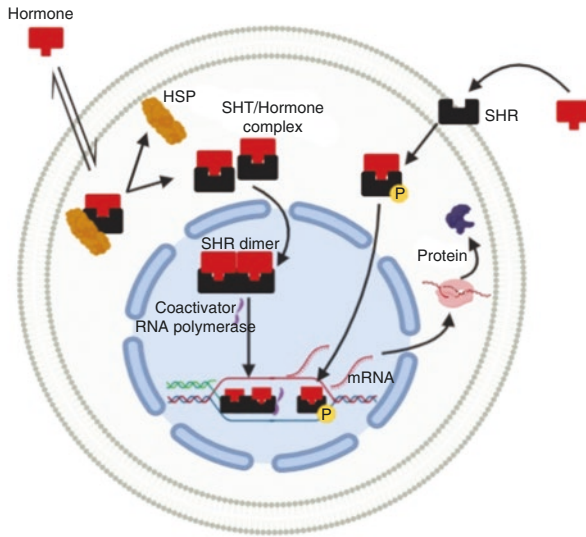


**Fig. 14.2** Functional domains of steroid hormone receptors (SHR). All the SHRs share common structural domains; the N-terminal transcription activation-1 domain (A/B), which regulates the ligand-independent transcription; a DNA-binding domain (C) which recognizes and binds with the steroid hormone-responsive elements on DNA; a flexible hinge domain (D) which carries information for the posttranscriptional modification; and a multifunctional ligand-binding domain (E) which recognizes and binds the ligand and acts as a ligand-dependent transcription activation function. ER $\alpha$  contains an extra region (F). Numbers represent the number of amino acids in each receptor. ER, estrogen receptor; PR, progesterone receptor; AR, androgen receptor; MR, mineralocorticoid receptor; GR, glucocorticoid receptor; AF, activation function; DBD, DNA-binding domain; LBD, ligand-binding domain. Figure modified from (Griekspoor et al. 2007)

Eisinger et al. 2018), two progesterone receptors (PR) subtypes (PR $\alpha$  and PR $\beta$ ) with several splice variants (Brinton et al. 2008; Knutson and Lange 2014), and two types of androgen receptor (AR) subtypes have been identified (AR $\alpha$  and AR $\beta$ ) (Ogino et al. 2018; Pelletier et al. 2000). All SHR share similar functional domains but differ in the length of the amino acid chain (Fig. 14.2).

In steroid-responsive cells, SHR are mainly present in the cytoplasm and the nucleus, although SHR are also found on the cell membrane (Schwartz et al. 2016). When SHR are activated by the corresponding steroid hormone, they dimerize and move into the nucleus of the cell, where they bind to the hormone-responsive element in the promoter region of specific target genes. SHR may act as transcriptional activators or transcriptional suppressors, resulting in the induction or suppression of the expression of hormone-responsive genes. These responsive genes may evoke a wide variety of physiological responses. Some rapid actions of estrogens and progestins are directly mediated by cell surface receptors in a non-genomic manner (Fig. 14.3). Although the involvement of steroid hormones in various brain disorders has been demonstrated, the mechanisms by which they exert their effects are still largely unknown.

Animal experiments and postmortem brain analyses using molecular biology techniques, such as western blot and immunohistochemistry, have shown that sex hormone receptors are expressed in various regions of the brain. ER, PR, and AR are expressed in those brain areas that are associated with emotion, cognition, and behavior, such as the hypothalamus, amygdala, cerebral cortex, hippocampus, and brainstem (Marrocco and McEwen 2016). This expression pattern is in agreement with the association of sex hormones with psychiatric disorders. Literature suggests that the neuroprotective effects of estrogens are mainly due to ER $\alpha$ -mediated signaling (Ooishi et al. 2012), whereas the effect on mood and cognitive functions is



**Fig. 14.3** Generalized mechanisms of action of steroid hormones in the cell. All the steroid hormone receptors mediate their action by a common genomic pathway, upon entering the cell by passive diffusion. Hormones can bind to either cytoplasmic, nuclear, or membrane receptors. In the cytoplasm, they exist as complexes with heat shock proteins (HSP) in an inactive form. When a steroid hormone binds, the receptor complex releases the HSP and becomes activated. The activated receptor forms a homodimer or heterodimer, which subsequently enters into the nucleus. The dimer binds to the specific hormone-responsive element in DNA and initiates or inhibits the process of gene transcription. Hormones with receptors in the nucleus get activated by their ligands and follow the same signaling pathway as the cytoplasmic receptors. In the case of membrane receptors, binding of the target molecule will induce a signaling cascade or the translocation of the receptor to the nucleus. In the nucleus, the receptor complex will activate the transcription to mRNA of certain genes and their following translation into functional proteins

mainly mediated by ER $\beta$  (Foster 2012). A study on patients with depression or schizophrenia, who committed suicide, showed downregulation of ER $\beta$  expression in the limbic system (Östlund et al. 2003). To our knowledge, no data exist on altered expression of PR and AR in the human brain in mood and behavioral disorders. Alternatively, steroid hormones may be acting indirectly through regulation of neurotransmission of serotonin and dopamine among others (Bristol et al. 2014; Fink et al. 1999; MacKenzie and Maguire 2014).

Not only sex hormone receptor expression but also polymorphisms in their genes were found to be implicated in psychiatric diseases. Sex hormone receptor polymorphisms have been associated with an increased risk for schizophrenia, depression, anxiety traits, and cognitive impairment. The association of polymorphisms in estrogen and androgen receptors with psychiatric disorders has been reviewed recently (Zhang et al. 2017).

The corticosteroid receptors may be divided into two classes: mineralocorticoid receptors (MR) and glucocorticoid receptors (GR) (Barabás et al. 2018). MR are mainly expressed in the limbic system, hypothalamus, and circumventricular

regions and to a lesser extent in other parts of the brain. GR expression is mainly observed in subregions of the cerebral cortex, olfactory cortex, hippocampus, amygdala, dorsal thalamus, hypothalamus, cerebellar cortex, trapezoid body, locus coeruleus, and dorsal raphe nucleus in rats (Morimoto et al. 1996; Owen 2002). Both the corticosteroid receptors are co-expressed in the hippocampus, amygdala, inferior frontal gyrus, cingulate gyrus, and nucleus accumbens in the human brain, whereas predominantly MR were found in the hippocampus (Klok et al. 2011).

Corticosteroids have the ability to bind to the corresponding MR or GR, via which they can exert behavioral changes. In clinical studies, a reduction in MR and GR expression and an increase in corticosteroid levels are suggested to be associated with suicide in different brain disorders (López et al. 1998; Xing et al. 2004; Young et al. 2003). Prolonged (chronic) stress has been reported to downregulate the expression of GR in the prefrontal cortex in rats (Chiba et al. 2012) and to reduce GR mRNA in the basolateral/lateral nuclei in patients with schizophrenia or bipolar disorder (Perlman et al. 2007, 2004). Increased production of cortisol causes a downregulation of GR receptors and desensitization of the GR signaling pathway, which in turn leads to an increase in the levels of cortisol (Sapolsky 2000).

### 14.2.1 Estrogens in Brain Disorders

Estrogens are a family of steroidal hormones that regulate reproductive and nonreproductive functions in both males and females. These hormones exert their pleiotropic effects through a combination of nuclear and membrane-associated receptors, which are often associated with different latencies of action (Cornil 2018; Micevych et al. 2015). Binding of estrogens to intracellular nuclear ER (mostly ER $\alpha$  but also ER $\beta$ ) leads to a transcriptional regulation of estrogen-related genes, through the activation of the hormone response element (HRE). On the other hand, estrogens can also bind to membrane receptors to induce fast responses through non-genomic signaling, affecting various aspects of brain function and behavior. For a review on the molecular mechanisms, see (Kawata et al. 2008).

Although both male and female subjects produce estrogens, the major source of estrogens are the ovaries, which is responsible for the difference in estrogen quantities and functions between both sexes. Changes of circulating estrogens in women, e.g., due to pregnancy or menopause, have been associated with an increased prevalence of brain disorders like reduced cognitive function, depression, and anxiety-related and eating disorders (Holsen et al. 2011). This hypothesis is supported by the observation that several mental changes occur during the transition of women from pre- to postmenopause, including reduced sexual drive and premenstrual or perimenopausal dysphoria induced by oral contraceptives or hormone replacement therapy (Eriksson et al. 2002; Rubinow and Schmidt 2006; Steiner et al. 2006). Other studies showed that during menopause, when circulating levels of estrogens are strongly reduced, there is a higher prevalence of social and psychological changes in women, like anxiety, irritability, stress, memory loss, lack of concentration, and loss of libido, which eventually may culminate into depression (Bryant

et al. 2012). Moreover, prolonged deficiency of estrogens also increases the risk to develop dementia (Sherman et al. 2005), including Alzheimer's disease (Pike et al. 2009). One possible explanation for these phenomena is that changes in circulating estrogens can induce changes in neurotransmitters like acetylcholine, dopamine, noradrenaline, and serotonin (McEwen and Milner 2017).

Maintenance of estrogen levels seems essential for normal physiological and mental status. In fact, estrogen replacement therapies have shown to have neuroprotective effects. Studies in humans and animal models have shown positive effects on perimenopausal and postmenopausal depression (Grigoriadis and Kennedy 2002), a reduction of ischemia-induced brain injury (Carswell et al. 2000), improvement of cognition (McCarrey and Resnick 2015), and neuroprotection against other neurodegenerative diseases like multiple sclerosis, Parkinson's disease and Alzheimer's disease (Chakrabarti et al. 2014). This protective effect was partly ascribed to the anti-inflammatory activity of estrogens (Resnick and Henderson 2002; Vegeto et al. 2008). Neurosteroids produced by activated microglia are able to shift a pro-inflammatory immune response into an anti-inflammatory phenotype. This effect seems to be dependent on both the expression level and the extent of stimulation of ER in the CNS (Giatti et al. 2012).

### 14.2.2 Progestins in Brain Disorders

Progestins have been demonstrated to play an important role in neuroprotection in experimental models and clinical trials in patients with stroke and traumatic brain injury. Progestins are able to decrease the lesion volume progression in the brain of stroke patients. A few studies were performed to assess the role of PR in experimental animal models of stroke, traumatic brain injury, and spinal cord injury (for a review see (Gibson et al. 2008)). For example, Liu et al. found that ischemia to the brain for 6 h resulted in a rapid increase in the progesterone and 5 $\alpha$ -dihydroprogesterone levels both in wild-type and PR knockout mice, suggesting a possible role of progestins in the salvage of neurons at risk (Liu et al. 2012). Progestins are also involved in cellular processes related to maintenance of myelin integrity, regulation of spinogenesis, synaptogenesis, neuronal survival, and dendritic growth (McEwen and Woolley 1994; Mellon 2007; Schumacher et al. 2012; Zhang et al. 2009). These observations suggest that neuroprotective effects of progestins may be mainly due to cellular maintenance processes.

Changes in steroid hormone levels affected the expression of membrane PR, especially PR $\alpha$ , in rats and mice. Upon treatment with estradiol or progesterone, significant expression of PR $\alpha$  was observed in neurons of the olfactory bulb, striatum, cortex, thalamus, hypothalamus, septum, hippocampus, and cerebellum, but not on oligodendrocytes or astrocytes. Moreover, traumatic brain injury induced the expression of PR $\alpha$  not only on neurons but also on oligodendrocytes, astrocytes, and reactive microglia, suggesting a role of progestins and PR in inflammation in the injured brain (Meffre et al. 2013). The neuroprotective mechanisms and

anti-inflammatory effects of progestins have been reviewed by several authors (De Nicola et al. 2009; Giatti et al. 2012; Grandi et al. 2017; Luoma et al. 2012; Singh and Su 2013). Progestins can inhibit the activation of microglia, which prevents NO and TNF-alpha production and the release of other inflammatory cytokines, such as IL-1beta, IL-6, complement factor C3 and C5, and macrophage-inducing factor-1.

Other mechanisms of action for the neuroprotective effects of progestins have been suggested. Progestins may prevent brain damage by controlling edema formation (vasogenic or cytogenic) via modulation of the expression of the aquaporin-4 water transporter, moderating  $\text{Ca}^{2+}$  flux caused by excitotoxicity, and reconstitution of the blood-brain barrier (O'Connor et al. 2005). Additionally, it has been reported that the antioxidant properties of progestins can prevent cellular insults by oxidative stress induced by free radicals (Ishihara et al. 2015). Besides the aforementioned roles in neuroprotection, progestins also play a role in neuronal remodeling by upregulating several neurotrophic factors, such as brain-derived neurotrophic factor (BDNF), Na/K ATPase, microtubule-associated protein 2 (MAP-2), choline acetyltransferase (ChAT), and glial-derived neurotrophic factor (GDNF) (Singh and Su 2013).

To investigate progestins as neuroprotective agents in patients, a clinical trial (phase IIa) was conducted in 100 male and female patients with blunt head trauma with moderate-to-severe damage. Treatment with progesterone was associated with a significant reduction in mortality compared to the vehicle group (Wright et al. 2007). However, the follow-up phase III clinical study could not demonstrate any neuroprotective effect of progestins in moderate-to-severe traumatic brain injury (Goldstein et al. 2017; Skolnick et al. 2014). Discussions about this lack of efficacy are suggesting a dose calibration problem (Howard et al. 2017). On the other hand, administration of progestins in combination with estrogens showed efficacy as treatment for multiple sclerosis in both animal models and in patients. The beneficial effect was mediated by modulation of peripheral and brain-intrinsic immune responses and regulation of local growth factor supply, oligodendrocytes, and astrocytes (Kipp et al. 2012). The combination of progestins and estrogens was also found to be an effective treatment for postmenopausal symptoms.

### 14.2.3 Androgens in Brain Disorders

Androgens are known to affect various brain functions and behavior. The most common central effect of androgens is the induction of aggression. Several studies have shown that excessive levels of testosterone induce aggression in both males and females of various species (Carré et al. 2011). Additionally, studies have shown that androgens play a role in depression-like and anxiety-like disorders, especially after menopause in women and during hypogonadism (andropause) in men. Changes in circulating androgens have been suggested to increase the risk of mood disorders,



depression, and psychosis (McHenry et al. 2014; Talih et al. 2007). Other studies reported that changes in circulating androgen levels have effects on cognition and cell survival (Lašaitė et al. 2017; Spritzer and Galea 2007).

Testosterone is converted to estradiol in the CNS, which plays a pivotal role in the feedback regulation in the hypothalamus. The hypothalamus can induce hormone release by the endocrine system through the pituitary. When men show hypogonadism, there are very low circulating androgen concentrations, which causes impairment of the higher functions in the brain, as described above (Ciocca et al. 2016). Testosterone replacement therapy (THT) showed either positive effects or no improvements in higher brain function (Ciocca et al. 2016; Corona et al. 2017; Davis et al. 2013). For this reason, the use of THT as a therapy for brain-related dysfunctions due to androgen hypogonadism is still under debate (Corona et al. 2015; Scovell and Khera 2018).

#### 14.2.4 Corticosteroids in Brain Disorders

Corticosteroids, such as glucocorticoids and mineralocorticoids, are produced by the adrenal glands and liver and by the placenta and maternal glands during pregnancy. In stressful conditions, the glucocorticoid cortisol is rapidly synthesized and secreted in response to adrenocorticotrophic hormone, released from the pituitary, and corticotrophin-releasing hormone, secreted by the hypothalamus. Cortisol stimulates the production of energy-rich compounds such as glucose, free fatty acids, and amino acids in different tissues. On the other hand, the mineralocorticoid aldosterone is produced in response to angiotensin II and promotes sodium reabsorption and fluid retention. Both cortisol and aldosterone release are controlled by the hypothalamic-pituitary-adrenal (HPA) axis. The HPA axis is a neuroendocrine system that has complex interactions with the serotonergic, noradrenergic, and dopaminergic systems in the brain. This axis regulates our body's responses to stress, and its dysregulation may lead to brain disorders (Stratakis and Chrousos 1995).

Excessive activity of the HPA axis or enlargement of the pituitary or adrenal gland cause an increase in cortisol levels, which in turn leads to hypercortisolemia. The overactivity of the HPA axis in stressful conditions may lead to dysregulation of the serotonergic system and is one of the most important predictors of suicide attempts in depressed patients (Pompili et al. 2010). Depression, on the other hand, is one of the major causes of hypercortisolemia, which may lead to neurotoxicity and reduced neurogenesis in the hippocampus in depressed patients (Axelson et al. 1993). Besides their role in glucose and mineral metabolism, corticosteroids are also implicated in the regulation of brain functions like sleep, ingestive behavior, behavioral adaptation, learning, and memory. In addition, corticosteroids play a significant role in brain damage, aging, mood, mental performance, and the pathogenesis of neuropsychiatric disorders, such as depression and Alzheimer's disease (de Kloet et al. 1986; McEwen and Sapolsky 1995; Sapolsky 2004).

## 14.3 Radiopharmaceuticals for Imaging of Steroid Hormone Receptors

Up to now, almost all information about the role of steroid hormones and their receptors in the healthy and diseased brain is obtained from experimental animals and postmortem human studies, because brain biopsy in patients is generally not feasible or highly undesirable. In order to facilitate research on the role of steroid hormones in neurodegenerative and psychiatric diseases, it is therefore of importance to have noninvasive techniques available to measure the expression of SHR in the brain. PET and SPECT are noninvasive nuclear imaging techniques that allow measurement of receptor expression and receptor occupancy in the living brain. Until now, several PET and SPECT tracers have been developed to image the SHR. Most of these radiopharmaceuticals, however, were developed for applications in oncology, in particular for imaging of receptor expression and occupancy in steroid hormone-sensitive tumors like breast and prostate cancer (de Vries et al. 2007; Hospers et al. 2008).

So far, only a few studies on imaging of SHR in the brain have been reported. However, it is expected that most radiopharmaceuticals that have been developed for tumor imaging can also be applied in brain imaging. Although many tracers have been described for each SHR, most of these tracers did not provide satisfactory results and consequently did not enter clinical studies. In the following paragraphs, we will not provide a complete overview of all tracers for SHR that have been reported but only discuss the most promising candidate tracers for PET and SPECT imaging of SHR in the brain.

### 14.3.1 Estrogen Receptor Imaging

The ER is the most widely studied SHR. Although literature exists on western blotting and in situ hybridization studies to determine the expression of the ER in the rodent brain, hardly any data exists on ER expression in the human brain. For this purpose, imaging methods to measure ER expression could certainly be of added value. Several tracers have been developed for PET and SPECT imaging of ER, in particular for imaging breast cancer.

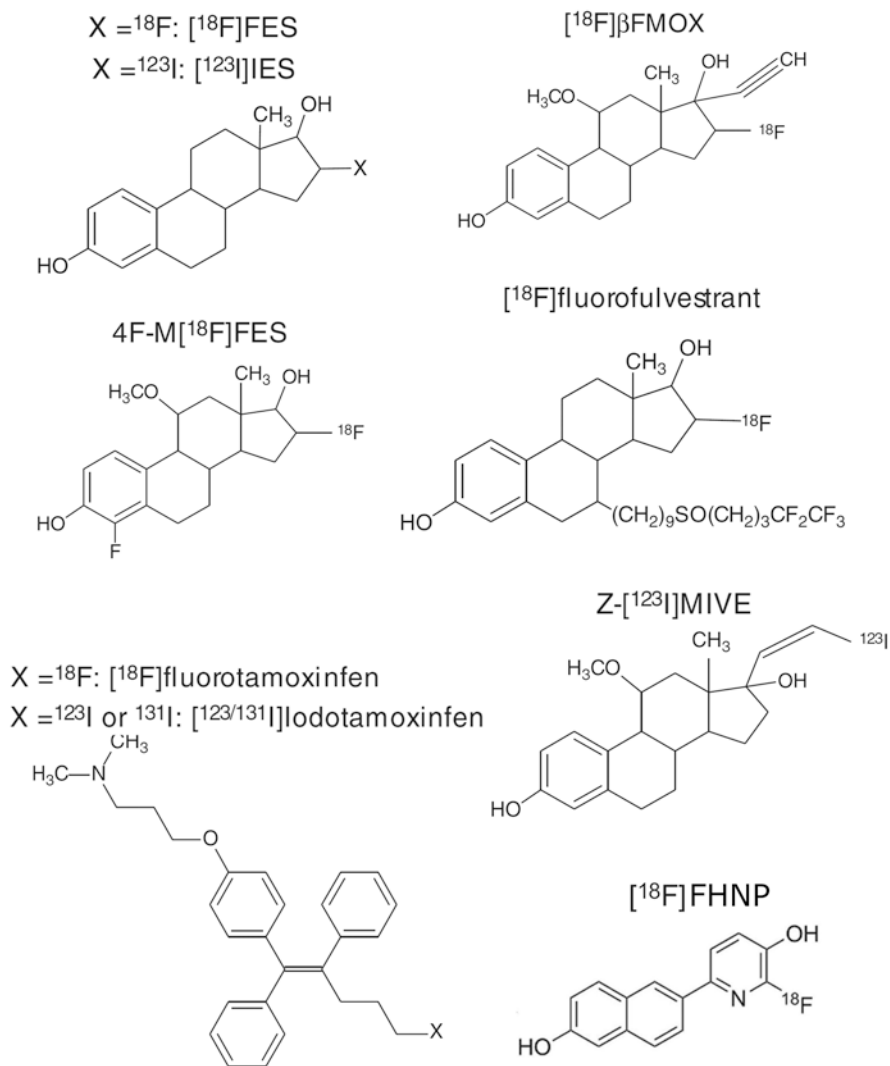
#### 14.3.1.1 PET Tracers for Estrogen Receptors

[<sup>18</sup>F]FES (16alpha-[<sup>18</sup>F]fluoro-17beta-estradiol) was one of the first PET tracers introduced for imaging of ER in the 1980s (Kiesewetter et al. 1984), and it is still used in the clinic today. In rats and mice, [<sup>18</sup>F]FES presented highest uptake in tissues with high ER expression, such as the uterus and ovaries (Aliaga et al. 2004; Kiesewetter et al. 1984; Sasaki et al. 2000; Seimbille et al. 2004; Yoo et al. 2005). [<sup>18</sup>F]FES uptake in ER-rich organs could be blocked with unlabeled estradiol in a dose-dependent manner, indicating that the tracer uptake is ER-mediated (Katzenellenbogen et al. 1993). Treatment of mice bearing ER-positive breast tumors with fulvestrant, an irreversible ER antagonist, also reduced the uptake of

[<sup>18</sup>F]FES (Fowler et al. 2012). [<sup>18</sup>F]FES binds to both ER $\alpha$  and ER $\beta$ , but the affinity of [<sup>18</sup>F]FES is 6.3-fold higher for ER $\alpha$  than for ER $\beta$  (Yoo et al. 2005). Both ER subtypes are expressed in breast tumors, but ER $\alpha$  is usually the major type expressed (Fox et al. 2008). ER $\alpha$  is associated with tumor growth and development (Dotzlaw et al. 1997; Järvinen et al. 2000), while the function of ER $\beta$  is not well understood yet but is suggested to be involved in the inhibition of ER $\alpha$  signaling (Fox et al. 2008). In breast cancer patients, [<sup>18</sup>F]FES PET has been applied successfully for the diagnosis and monitoring of anticancer treatment efficacy in ER-positive breast cancer. Previous studies showed ER expression in primary breast tumors and in metastases, in which the accumulation of [<sup>18</sup>F]FES in these tumors correlated well with ER density, as determined by immunohistochemistry (Dehdashti et al. 1995; Mintun et al. 1988). In ER-positive breast cancer patients, [<sup>18</sup>F]FES PET displays an excellent specificity (98%) and fairly good sensitivity (85%) for tumor detection (van Kruchten et al. 2013). In human plasma, [<sup>18</sup>F]FES binds to sex hormone-binding globulin (SHBG) (Tewson et al. 1999), which improves the plasma stability of the tracer, although substantial metabolism is still observed. The radiation burden associated with [<sup>18</sup>F]FES PET is within the normal range of other nuclear medicine procedures: a typical dose of 200 MBq (6 mCi) causes a radiation burden to the patient of 4.4 mSv (Mankoff et al. 2001; Mortimer et al. 1996; Peterson et al. 2008). [<sup>18</sup>F]FES PET has now been used in many clinical studies in ER-positive cancer patients and is nowadays even used in routine clinical practice in our institution.

In order to increase the binding affinity and in vivo stability of the tracer, several analogues of [<sup>18</sup>F]FES have been developed. One of these candidates, 11beta-methoxy-4,16alpha-[<sup>18</sup>F]difluoroestradiol (4F-M[<sup>18</sup>F]FES), showed high uptake and high target-to-background ratios in ER-rich organs like the uterus (VanBrocklin et al. 1994). 4F-M[<sup>18</sup>F]FES has a low affinity toward the SHBG, but studies in humans showed a significant, potentially ER-mediated uterus uptake in both pre- and postmenopausal women (Beauregard et al. 2008). A preclinical study performed by Paquette and colleagues compared both 4F-M[<sup>18</sup>F]FES and [<sup>18</sup>F]FES, showing a higher specific tumor uptake and better contrast for 4F-M[<sup>18</sup>F]FES (Paquette et al. 2013). Some years later, the same research group performed a comparative phase II clinical study in ER+ breast cancer patients and showed that 4F-M[<sup>18</sup>F]FES yielded a lower nonspecific signal and better tumor contrast than [<sup>18</sup>F]FES (Paquette et al. 2018). More studies should be performed to determine if 4F-M[<sup>18</sup>F]FES is indeed more suitable than [<sup>18</sup>F]FES for diagnosis and treatment evaluation of ER+ breast cancer patients.

Besides [<sup>18</sup>F]FES and 4F-M[<sup>18</sup>F]FES, several other radiolabeled estradiol derivatives and ER-targeting anticancer drugs have been evaluated as candidate PET tracers for imaging ER density and occupancy (Fig. 14.4). The potent estrogen 17alpha-ethynyl-16beta-[<sup>18</sup>F]fluoro-11beta-methoxy-estradiol ([<sup>18</sup>F]betaFMOX) showed promising results in rodents with a fourfold higher uptake in immature rat uterus than [<sup>18</sup>F]FES. [<sup>18</sup>F]betaFMOX was considered to be a highly sensitive tracer for imaging of the ER, as [<sup>18</sup>F]betaFMOX also exhibited specific binding in organs with low ER density, such as the kidney, muscle, and thymus (VanBrocklin et al. 1993a, b). Despite the promising results in rodents, [<sup>18</sup>F]betaFMOX PET failed to



**Fig. 14.4** Chemical structures of tracers that have been evaluated for PET or SPECT imaging of estrogen receptors

detect ER-positive lesions in breast cancer patients. This lack of sensitivity in humans may be due to the fast metabolism of the tracer, because it exhibited low binding to SHBG, which protects steroids from metabolic degradation (Jonson et al. 1999). Other ethynyl analogues of FES were tested for breast cancer imaging. These tracers also showed faster degradation in vivo, due to low SHBG binding, although they showed better target-to-background ratios (VanBrocklin et al. 1992).

In addition to estradiol derivatives, radiolabeled derivatives of anticancer drugs, such as tamoxifen and fulvestrant, were evaluated for imaging of ER.  $^{18}\text{F}$ -labeled

tamoxifen, [ $^{18}\text{F}$ ]fluoromethyl-*N,N*-dimethyltamoxifen ([ $^{18}\text{F}$ ]FTX), exhibited specific uptake in the uterus and mammary tumors, which could be blocked with an excess of estradiol (Yang et al. 1994) and appeared to correlate with tamoxifen treatment outcome (Inoue et al. 1996). So far, no additional studies have been performed to prove the clinical utility of [ $^{18}\text{F}$ ]FTX. Labeling of fulvestrant, a full antagonist of the ER, resulted in a reduced binding affinity due to the addition of  $^{18}\text{F}$ . Consequently,  $^{18}\text{F}$ -labeled fulvestrant is not suitable for PET imaging of ER (Seimille et al. 2004).

The discovery of ER $\beta$  in breast cancer initiated the search for ER $\beta$ -selective PET tracers. Lee and colleagues developed two beta-selective tracers: 8beta-(2-[ $^{18}\text{F}$ ]fluoroethyl)estradiol (8beta-[ $^{18}\text{F}$ ]FEE), a derivative of the potent ER $\beta$ -selective steroid 8beta-vinylestradiol, and [ $^{76}\text{Br}$ ]BrERB-041 derived from the nonsteroidal ER $\beta$  ligand ERB-041 (Lee et al. 2012). Although both 8beta-[ $^{18}\text{F}$ ]FEE and [ $^{76}\text{Br}$ ]BrERB-041 showed high binding affinities in a radiometric assay, ex vivo biodistribution studies in rats and mice could not demonstrate ER $\beta$ -mediated uptake (Lee et al. 2012). On the other hand, Antunes and colleagues recently synthesized a fluorinated pyridinic tracer [ $^{18}\text{F}$ ]FHNP for selective binding to ER $\beta$ . [ $^{18}\text{F}$ ]FHNP showed a higher affinity toward ER $\beta$  than ER $\alpha$ , showing selective binding of the tracer to ER $\beta$  in tumor-bearing mice, which makes this tracer a possible candidate for the measurement of ER $\beta$  in vivo (Antunes et al. 2017a, b). However, more studies are needed for the validation of this tracer.

In summary, at this moment [ $^{18}\text{F}$ ]FES remains the best validated tracer for molecular imaging of ER expression and is currently used in clinical studies. Even though its characteristics are not ideal, [ $^{18}\text{F}$ ]FES is an adequate tracer for PET imaging of ER in humans (van Kruchten et al. 2013). Efforts to develop better alternatives for [ $^{18}\text{F}$ ]FES are still made (e.g., 4F-M[ $^{18}\text{F}$ ]FES), but no phase III clinical studies have been performed yet. So far, [ $^{18}\text{F}$ ]FHNP is the only PET tracer showing ER $\beta$ -selectivity in vivo, but this ligand needs further evaluation.

#### 14.3.1.2 SPECT Tracers for Estrogen Receptors

In addition to PET tracers, some SPECT tracers have been developed for imaging of ER (Fig. 14.4) (for a review see (Liu et al. 2016)). Among the first successful SPECT tracers for ER was 16alpha-[ $^{125}\text{I}$ ]iodo-11beta-methoxy-17beta-estradiol ([ $^{125}\text{I}$ ]MIE) (Zielinski et al. 1986). In rodents, [ $^{125}\text{I}$ ]MIE was found to accumulate in ER-rich areas, such as the uterus. [ $^{125}\text{I}$ ]MIE showed higher uterus-to-blood ratios than 16alpha-[ $^{125}\text{I}$ ]iodoestradiol. 16alpha-[ $^{123}\text{I}$ ]iodo-17beta-estradiol ([ $^{123}\text{I}$ ]IES) and (20Z)-11beta-methoxy-17alpha-[ $^{123}\text{I}$ ]iodovinylestradiol (Z-[ $^{123}\text{I}$ ]MIVE) have not only been applied in animals but also in clinical studies. [ $^{123}\text{I}$ ]IES showed ER-mediated uptake in the rabbit reproductive system and in ER-positive tumors in breast cancer patients, but no specificity or sensitivity data were reported (Kenady et al. 1993; Pavlik et al. 1990; Scheidhauer et al. 1991; Schober et al. 1990). Both stereoisomers of [ $^{123}\text{I}$ ]MIVE were evaluated for ER imaging. The (Z)-isomer of [ $^{123}\text{I}$ ]MIVE showed highest ER-mediated specific uptake in rodents and in clinical studies in breast cancer patients. The tracer uptake could be blocked by saturation of the ER with tamoxifen (Bennink et al. 2004, b; Rijks et al. 1997a, b). Although both [ $^{123}\text{I}$ ]IES and (Z)-[ $^{123}\text{I}$ ]MIVE showed ER-mediated specific uptake, these

tracers also displayed fast metabolism and low SHBG binding (Nachar et al. 1999; Rijks et al. 1998a; Rijks et al. 1998b).

Attempts have been made to generate a SPECT tracer for ER by labeling derivatives of the anticancer drug tamoxifen. In rodents, the uptake of [ $^{123}\text{I}$ ]iodotamoxifen ([ $^{123}\text{I}$ ]TAM) was found to be high in ER-rich organs (Biber Muftuler et al. 2008). In clinical studies, [ $^{123}\text{I}$ ]TAM showed ER-specific uptake in breast tumors (Hanson and Seitz 1982; Van de Wiele et al. 2001). Recently tamoxifen was also labeled with  $^{131}\text{I}$  ([ $^{131}\text{I}$ ]TAM). Biodistribution studies in rodents proved that the uptake of this tracer was ER-mediated in the uterus and breast tissue (Biber Muftuler et al. 2008). No further studies on iodine-labeled tamoxifen have been published lately.

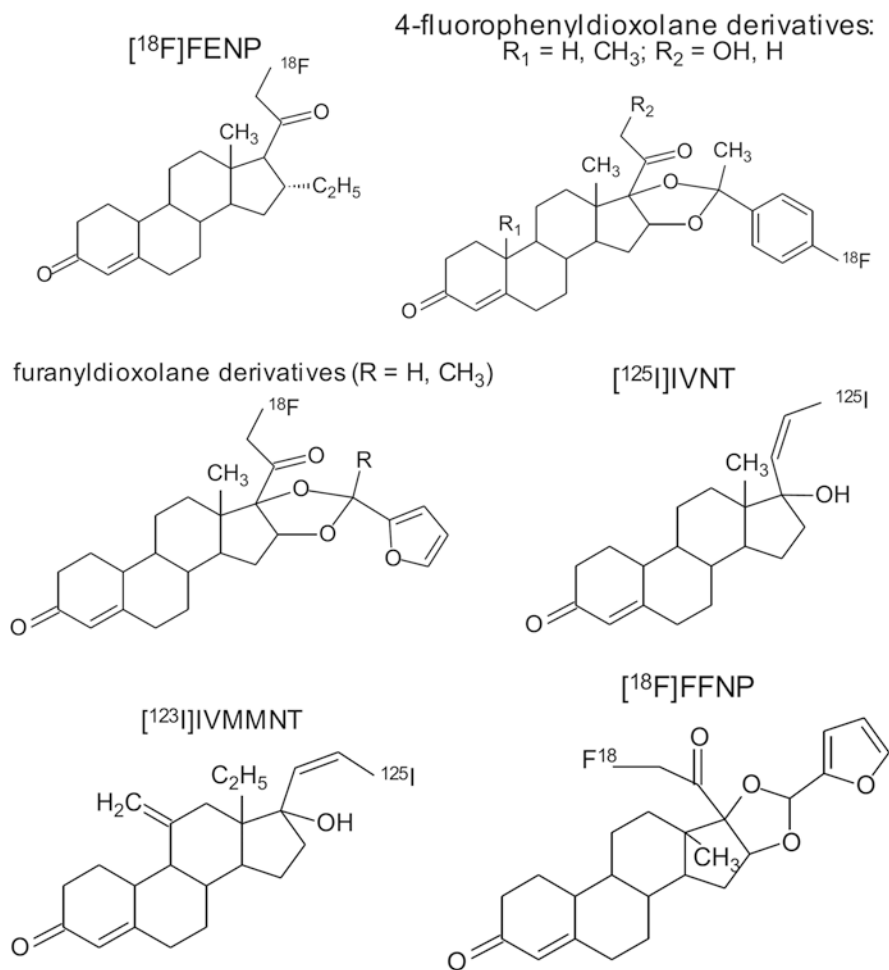
Estradiol and anticancer drug derivatives have also been labeled with  $^{99\text{m}}\text{Tc}$ . A few of them have even been successfully evaluated in rodents (Arterburn et al. 2003; Nayak et al. 2008, 2014; Ramesh et al. 2006; Takahashi et al. 2007a, b; Xia et al. 2016; Yurt et al. 2009; Zhu et al. 2010). However, the usefulness of these tracers for brain imaging is questionable, because of the high hydrophilicity of the  $^{99\text{m}}\text{Tc}$  conjugates, which likely precludes efficient brain penetration.

## 14.3.2 Progesterone Receptor Imaging

### 14.3.2.1 PET Tracers for Progesterone Receptors

A few substrates of PR have been labeled with  $^{18}\text{F}$ ,  $^{76}\text{Br}$ ,  $^{123}\text{I}$ , and  $^{125}\text{I}$  for imaging of PR with PET and SPECT (Fig. 14.5) (for a review, see Linden et al. 2018). The  $^{18}\text{F}$ -labeled candidate PET tracers for PR include 21-[ $^{18}\text{F}$ ]fluoro-16 $\alpha$ -ethyl-19-norprogesterone ([ $^{18}\text{F}$ ]FENP) (Dehdashti et al. 1991; Jonson and Welch 1998; Pomper et al. 1988; Verhagen et al. 1991a), 21-[ $^{18}\text{F}$ ]fluoro-16 $\alpha$ -methyl-19-norprogesterone ([ $^{18}\text{F}$ ]FMNP) (Verhagen et al. 1991a), and 21-[ $^{18}\text{F}$ ]fluoro-16 $\alpha$ ,7 $\alpha$ -[(R)-(1'- $\alpha$ -furylethylidene)dioxy]-19-norpregn-4-ene-3,20-dione (Buckman et al. 1995). [ $^{18}\text{F}$ ]FMNP exhibited PR-mediated uptake in the uterus and tumors in rats (Verhagen et al. 1991b), but no further studies in humans were reported. [ $^{18}\text{F}$ ]FENP showed highly selective PR-mediated uptake in the uterus of estrogen-primed rats (Pomper et al. 1988) and in PR-positive carcinoma in mice (Verhagen et al. 1991b). In eight breast cancer patients, however, [ $^{18}\text{F}$ ]FENP could only detect 50% of PR-positive lesions, and tracer uptake did not correlate with PR expression (Dehdashti et al. 1991). The major reason for failure of [ $^{18}\text{F}$ ]FENP in clinical studies was its extensive metabolism in humans (Verhagen et al. 1994).

To overcome this problem, several ketals of 16 $\alpha$ ,17 $\alpha$ -dihydroxyprogesterone were labeled with positron-emitting isotopes (Buckman et al. 1995; Vijaykumar et al. 2002; Zhou et al. 2006), with 21-[ $^{18}\text{F}$ ]fluoro-16 $\alpha$ ,17 $\alpha$ -[(R)-(1'- $\alpha$ -furylmethylidene)dioxy]-19-norpregn-4-ene-3,20-dione ([ $^{18}\text{F}$ ]FFNP) being the most studied one. In a study performed in mice with mammary tumors, treatment with estradiol, letrozole, or fulvestrant showed treatment-induced changes in the uptake of [ $^{18}\text{F}$ ]FFNP, suggesting that early evaluation of response to the anticancer treatment could be feasible (Fowler et al. 2012; Salem et al. 2018, 2019). A study in breast cancer patients showed that [ $^{18}\text{F}$ ]FFNP PET is



**Fig. 14.5** Chemical structures of tracers that have been evaluated for PET or SPECT imaging of progesterone receptors

a safe, noninvasive method to evaluate the tumor's PR status in vivo (Dehdashti et al. 2012). Brain uptake of [<sup>18</sup>F]FFNP seems to be acceptable, but further studies are needed to prove its suitability for brain imaging.

Another series of candidate PET tracers for PR imaging comprised the derivatives of the nonsteroidal PR agonist tanaproget. Several derivatives of tanaproget were evaluated (Zhou et al. 2010), and 4-[<sup>18</sup>F]fluoropropyl-tanaproget ([<sup>18</sup>F]FPTP) demonstrated highest uptake in the target tissues like uterus and ovaries. The biodistribution pattern of [<sup>18</sup>F]FPTP in rats was comparable with other PR tracers, such as FENP and FFNP (Lee et al. 2010). Furthermore, synthesis of tanaproget-based compounds labeled with <sup>11</sup>C is currently ongoing. Haywood and colleagues successfully tested the synthesis of different <sup>11</sup>C-based molecules, but further

preclinical studies are needed to characterize the pharmacokinetics of these compounds (Haywood et al. 2015).

At the moment, [ $^{18}\text{F}$ ]FFNP seems to be the most promising candidate tracer for imaging PR, but more studies are required to validate the utility of this tracer in humans.

### 14.3.2.2 SPECT Tracers for Progesterone Receptors

A few iodinated tracers for PR imaging have been developed, but none of these candidate tracers showed promising results. (20Z)-17alpha- $^{125}\text{I}$ iodovinyl-19-nortestosterone ( $^{125}\text{I}$ IVNT) had interesting binding properties, but lacked selectivity in vivo (Ali et al. 1994). Later, both isomers of 17alpha- $^{123}\text{I}$ iodovinyl-18-methyl-11-methylene-19-nortestosterone ( $^{123}\text{I}$ IVMMNT) were investigated in rats and rabbits. (Z)- $^{123}\text{I}$ IVMMNT displayed highest in vivo binding in target organs, such as in uterus and ovaries. The uptake of (Z)- $^{123}\text{I}$ IVMMNT was found to be PR-mediated (Rijks et al. 1998b). So far, no human data exist to prove the feasibility of PR imaging with these tracers.

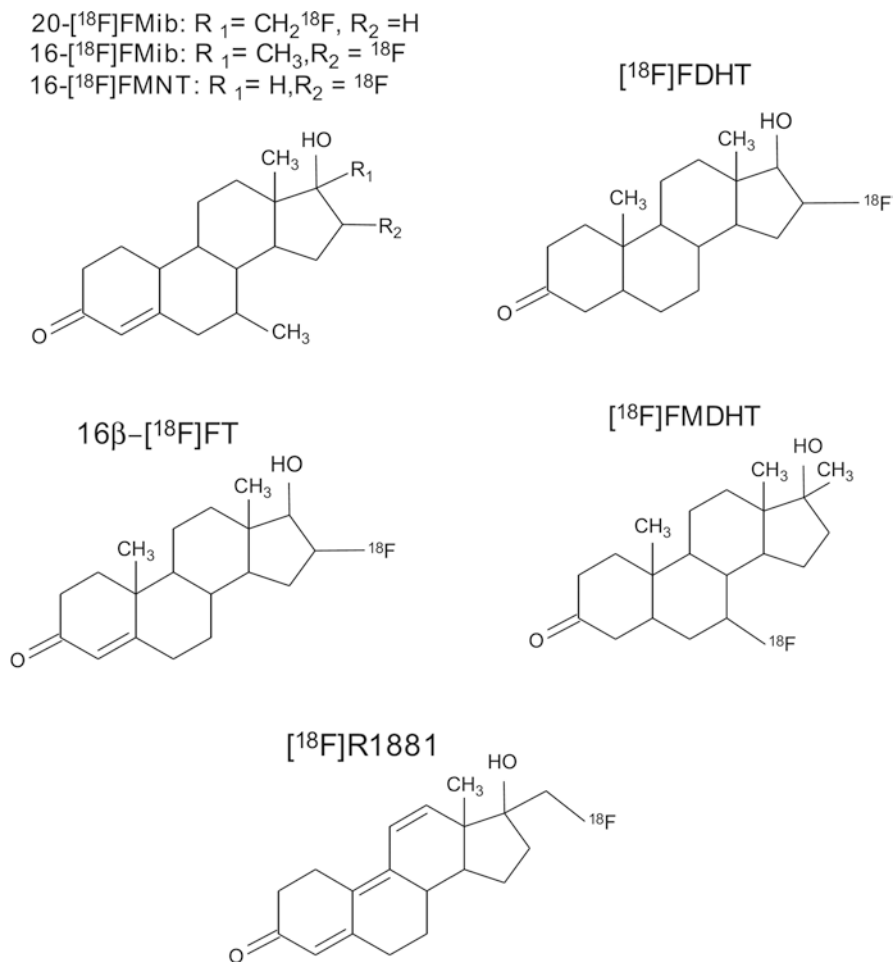
## 14.3.3 Androgen Receptor Imaging

### 14.3.3.1 PET Tracers for Androgen Receptors

Several promising PET tracers for imaging of androgen receptors have been successfully developed (Fig. 14.6). 20- $^{18}\text{F}$ fluoromibolone (20- $^{18}\text{F}$ Fmib) was the first radiolabeled PET tracer for AR that showed promising uptake in the prostate of rats and in baboons (Bonasera et al. 1996; Liu et al. 1991). Several other fluorinated compounds have been developed, and some of them showed promising results, such as 16beta- $^{18}\text{F}$ fluorodihydrotestosterone ( $^{18}\text{F}$ FDHT), 16beta- $^{18}\text{F}$ fluorotestosterone (16beta- $^{18}\text{F}$ FT), 16beta- $^{18}\text{F}$ fluoro-7alpha-methyl-19-nortestosterone (16beta- $^{18}\text{F}$ FMNT), 16alpha- $^{18}\text{F}$ fluoro-7alpha-methyl-19-nortestosterone (16alpha- $^{18}\text{F}$ FMNT), and 20- $^{18}\text{F}$ fluorometribolone (20- $^{18}\text{F}$ R1881) (Liu et al. 1992). Most of these tracers showed quick clearance from the body and rapid metabolism.

$^{18}\text{F}$ FDHT is so far the most promising tracer, showing the best results in rats and baboons. Uptake of  $^{18}\text{F}$ FDHT in the prostate was specific and AR-mediated. Metabolism of  $^{18}\text{F}$ FDHT was slower than metabolism of the other candidate tracers (Bonasera et al. 1996; Choe et al. 1995).  $^{18}\text{F}$ FDHT showed specific binding to AR and was able to detect changes in AR availability in preclinical studies (Larimer et al. 2018). Currently,  $^{18}\text{F}$ FDHT is the only PET tracer that has proceeded into the clinical evaluation phase (Beattie et al. 2010; Dehdashti et al. 2005; Kramer et al. 2019; Larson et al. 2019; Zanzonico et al. 2004). The radiation burden associated with  $^{18}\text{F}$ FDHT PET is within the normal range of other clinical nuclear medicine procedures: 0.018 mSv/MBq; for the maximum administered dose of 331 MBq, the total radiation burden is 6.0 mSv. Furthermore, recent studies aimed to position this tracer as a biomarker for metastatic castration-resistant prostate cancer patients have shown positive results (Cysouw et al. 2019; Fox et al. 2018; Jansen et al. 2019). In addition,  $^{18}\text{F}$ FDHT has also been





**Fig. 14.6** Chemical structures of tracers that have been evaluated for PET or SPECT imaging of androgen receptors

investigated as a PET tracer for detection of AR expression in breast cancer (Venema et al. 2017) and for detection of intra-abdominal testicles (Glaudemans et al. 2015).

The major drawback of  $^{18}\text{F}$ FDHT is its rapid metabolism. This led to the investigation of nonsteroidal derivatives with better in vivo stability, like  $^{11}\text{C}$ -labeled derivatives of propanamide, a selective androgen receptor modulator (Gao et al. 2011). Some nonsteroidal antagonists of AR have been radiolabeled, such as a  $^{11}\text{C}$ -labeled diethylamine flutamide derivative (Jacobson et al. 2006), a  $^{18}\text{F}$ -labeled hydroxyflutamide derivative (Jacobson et al. 2005), 3- $^{76}\text{Br}$ ]bromohydroxyflutamide (Parent et al. 2006),  $^{18}\text{F}$ ]bicalutamide, 4- $^{76}\text{Br}$ ]bromobicalutamide, and  $^{76}\text{Br}$ ]bromo-thiobicalutamide (Parent et al. 2007). However, none of these compounds showed promising results that warranted further evaluation.

Another interesting candidate PET tracer is 7 $\alpha$ -[<sup>18</sup>F]fluoro-17-methyl-5-dihydrotestosterone ([<sup>18</sup>F]FMDHT), although the first results reported for this tracer were not very promising. The uptake of [<sup>18</sup>F]FMDHT in the prostate was low; results were not reproducible and not comparable with other labeled steroids like [<sup>18</sup>F]FDHT. In these first experiments, [<sup>18</sup>F]FMDHT was investigated in rats treated with DES to suppress endogenous testosterone production (Garg et al. 2001). The same authors reevaluated the suitability of [<sup>18</sup>F]FMDHT for AR imaging in chemically castrated rats. In this animal model, uptake in the prostate was proven to be AR-mediated. Prostate uptake was comparable to other <sup>18</sup>F-labeled steroids and twofold higher than [<sup>18</sup>F]FMDHT uptake in DES-treated rats (Garg et al. 2008).

The most recent attempt for the synthesis of fluorinated androgen analogues was made by Okamoto and colleagues (2017). They could successfully synthesize two <sup>18</sup>F-labeled derivatives of 7 $\alpha$ -(3-[<sup>18</sup>F]fluoropropyl)-testosterone that showed proper affinity for AR. However, further studies are needed to determine if these tracers could be useful in determining AR changes in the brain.

So far, [<sup>18</sup>F]FDHT is the only AR tracer that has proceeded into the clinical evaluation phase. Clinical studies confirmed that this tracer is suitable for AR imaging in cancer patients. However, it still remains to be evaluated whether [<sup>18</sup>F]FDHT PET is also able to monitor AR expression in the human brain. In addition, preclinical data of [<sup>18</sup>F]FMDHT suggest that it may have favorable characteristics for AR imaging as well, but further evaluation is required to establish the merit of this tracer.

#### 14.3.3.2 SPECT Tracers for Androgen Receptors

The radioiodinated steroid 2 $\alpha$ -[<sup>125</sup>I]dihydrotestosterone was the first SPECT tracer showing high uptake in AR-rich organs like the prostate, epididymis, and testis in rats. Pretreatment with dihydrotestosterone reduced the uptake in these organs, suggesting that tracer uptake is specific and AR-mediated (Tarle et al. 1981). 7 $\alpha$ -[<sup>125</sup>I]iodo-5 $\alpha$ -dihydrotestosterone (7 $\alpha$ -[<sup>125</sup>I]IDHT) is another analogue of testosterone that was labeled with <sup>125</sup>I. 7 $\alpha$ -[<sup>125</sup>I]IDHT showed AR-mediated uptake in the prostate of rats. In vitro autoradiography of 7 $\alpha$ -[<sup>125</sup>I]IDHT produced excellent autoradiograms with low nonspecific binding in the prostate of rats (Labaree et al. 1997), but no further studies were published to demonstrate the usefulness of this tracer in vivo.

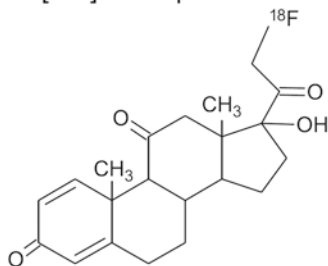
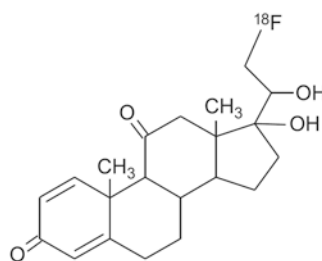
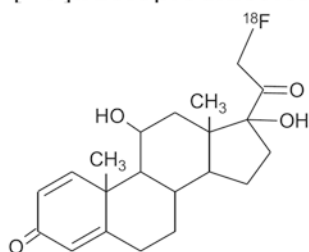
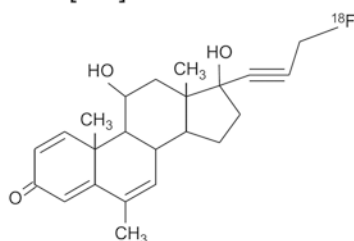
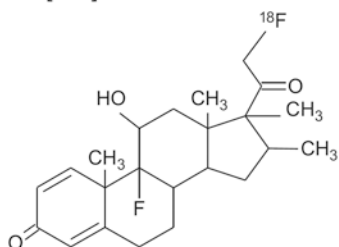
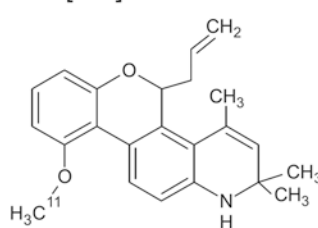
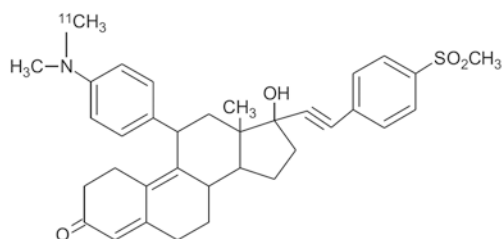
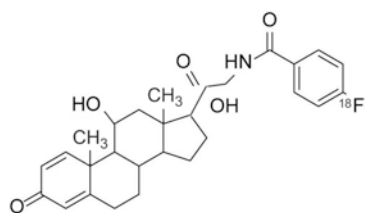
Some steroid and flutamide derivatives have been labeled with <sup>99m</sup>Tc and tested as SPECT tracers for AR. However, in vivo evaluation in rats did not show any AR-mediated specific binding for any of these compounds (Dallagi et al. 2010; Dhyani et al. 2011).

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## 14.4 Corticoid Receptor Imaging

### 14.4.1 PET Tracers for Glucocorticoid Receptors

So far, only a few radiolabeled GR ligands have been synthesized and evaluated for the potential use as PET tracer (Fig. 14.7). Most of these candidate tracers showed disappointing results in rodents and baboons. The first labeled compound that was

21-[<sup>18</sup>F]fluoroprednisone20-dihydro-21-[<sup>18</sup>F]fluoroprednisone21-[<sup>18</sup>F]fluoroprednisolone[<sup>18</sup>F]RU 52461[<sup>18</sup>F]ORG 6141[<sup>11</sup>C]AL-438[N-methyl-<sup>11</sup>C]ORG 34850[<sup>18</sup>F]-GR02

**Fig. 14.7** Chemical structures of tracers that have been evaluated for PET imaging of glucocorticoid receptors

introduced as ligand for the imaging of GR was 21- $^{18}\text{F}$ fluoroprednisone. 21- $^{18}\text{F}$ fluoroprednisone was rapidly metabolized, leading to low uptake in rat brain (Feliu and Rottenberg 1987). Additionally, an analogue of the selective GR agonist RU28362 was tested. The biodistribution of  $^{18}\text{F}$ RU52461 showed high GR-mediated uptake in the adrenals and pituitary in rats. PET imaging studies in baboons, however, showed low uptake of  $^{18}\text{F}$ RU 52461 in the brain (Dasilva et al. 1992). In another study in rats,  $^{18}\text{F}$ RU52461 uptake in the hippocampus could only be partially blocked with an excess of the unlabeled ligand, whereas complete blocking of tracer uptake in peripheral organs was observed (Pomper et al. 1992). The potent GR ligand  $^{18}\text{F}$ ORG6141 was evaluated in adrenalectomized and sham-operated rats. Ex vivo biodistribution 3 h postinjection revealed higher uptake of tracer in the hippocampus and brain stem of adrenalectomized animals as compared to sham-operated controls. However, GR-mediated specific retention of activity in these brain areas could not be demonstrated (Visser et al. 1995).

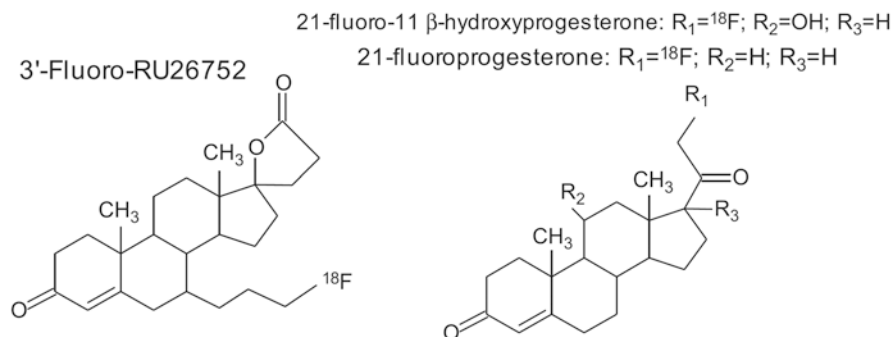
Wuest et al. synthesized a series of novel 4-fluorophenylpyrazolo steroids and tested their binding affinities to GR. Some of these compounds showed binding affinities up to 56% relative to dexamethasone (100%) (Wüst et al. 2003). One of these compounds, 2'-(4-fluorophenyl)-21- $^{18}\text{F}$ fluoro-20-oxo-11 $\beta$ ,17 $\alpha$ -dihydroxy-pregn-4-eno[3,2-*c*]pyrazole, was evaluated in rats using autoradiography and small animal PET imaging. Brain uptake of this tracer was found to be constant between 5 and 60 min after tracer injection. However, brain uptake was not specifically GR-mediated (Wüst et al. 2005) but only due to nonspecific binding. Another high-affinity GR antagonist Org34850 was labeled with  $^{11}\text{C}$ . [N-methyl- $^{11}\text{C}$ ]Org34850 was found to rapidly metabolize in rats. Ex vivo biodistribution and small animal PET studies demonstrated that [N-methyl- $^{11}\text{C}$ ]Org34850 was not able to penetrate the blood-brain barrier (Wuest et al. 2009).

Additional attempts were made with nonsteroidal molecules. The selective GR modulator AL-438 was labeled with  $^{11}\text{C}$ . The biodistribution of [ $^{11}\text{C}$ ]AL-438 showed high uptake in the pituitary and the brain, but treatment with a high dose of the GR antagonist corticosterone did not result in any blocking of tracer uptake, suggesting that the uptake was not GR-mediated (Wuest et al. 2007).

Thus, none of the aforementioned studies were successful, and consequently suitable tracers for imaging GR expression are currently not available. Nonetheless, in a recent work performed by Truillet and colleagues, a set of GR ligands was synthesized and labeled with  $^{18}\text{F}$  (Truillet et al. 2018). One of these tracers,  $^{18}\text{F}$ GR02, showed promising results. In vitro and in vivo testing in rats showed sufficient binding affinity to GR in GR-rich tissue. Additionally, they found uptake in the brain, which was increased substantially in adrenalectomized animals. Further studies are needed to validate this compound as a tracer for GR.

#### 14.4.2 PET Tracers for Mineralocorticoid Receptors

There is only one report that describes a series of MR-specific compounds that have been labeled with fluorine-18 for PET imaging. These compounds include



**Fig. 14.8** Chemical structures of tracers that have been evaluated for PET imaging of mineralocorticoid receptors

21- $[^{18}\text{F}]$ fluoroprogestosterone, 21- $[^{18}\text{F}]$ fluoro-11beta-hydroxyprogesterone, and 3'- $[^{18}\text{F}]$ fluoro-RU26752 (Fig. 14.8). Unfortunately, none of these compounds displayed the characteristics that are required for PET imaging of MR (Pomper et al. 1992).

## 14.5 Imaging of Steroid Hormone Receptors in the Brain

Although several tracers for imaging of SHR have now been applied in humans, hardly any studies on imaging of these receptors in the brain have been published. So far, mostly tracers for ER have been successfully used in rodents for imaging SHR in the brain.

### 14.5.1 Imaging of Estrogen Receptors in the Brain

So far, there is one study testing the usefulness of 16alpha- $[^{125}\text{I}]$ iodoestradiol for the detection of ER in the brain. Stumpf and colleagues (1987) used ex vivo autoradiography of the brain of prepubertal Sprague Dawley rats injected with 16alpha- $[^{125}\text{I}]$ iodoestradiol and showed high concentrations of radioactivity in the marginal zone of the mid-region anterior lobe of the pituitary and in preoptic neurons 1 h after tracer injection. The intermediate lobe and most cells in the anterior lobe, together with the dorsocaudal paraventricular nucleus and the medial amygdaloid region showed only a weak signal. Long-term exposure of the photographic plate allowed topographical recognition of tracer binding to cells in the medial amygdala, medio-dorsal and cortical nuclei, and piriform cortex. However, 16alpha- $[^{125}\text{I}]$ iodoestradiol uptake in the brain could not be significantly blocked with an excess of unlabeled estradiol, which was ascribed to deiodination of the tracer, leading to nonspecific binding of  $^{125}\text{I}$  within the myelin sheaths.

In addition to 16alpha- $[^{125}\text{I}]$ iodoestradiol, the SPECT tracer 11beta-methoxy-16alpha- $[^{125}\text{I}]$ iodoestradiol ( $[^{125}\text{I}]\text{MIE}_2$ ) has also been evaluated as tracer for ER in binding and displacement studies in rat, rabbit, and human brain slices. A

dose-dependent uptake of [ $^{125}\text{I}$ ]MIE<sub>2</sub> in pituitary and brain cell nuclei was observed in 25-day-old female rats (Brown et al. 1989), suggesting specific binding to the ER with no cross-reactivity with the other SHR. Additional experiments by Toran-Allerand and colleagues (1992) combined autoradiography with in situ hybridization and revealed a correlation between SPECT tracer uptake and binding of the mRNA probe in the cerebral cortex of female rats. In another study, ER expression during brain development was studied with [ $^{125}\text{I}$ ]MIE<sub>2</sub> in mice of both sexes. This study demonstrated the presence of ER-positive cells in the early postnatal cortex and showed a profound change in the topography and number of ER-positive cells during development (Shughrue et al. 1990). The most recent study used ex vivo autoradiography with [ $^{125}\text{I}$ ]MIE<sub>2</sub> to evaluate the effect of long-term treatment (21–60 days) with estradiol and tamoxifen on ER occupancy in the brain. The distribution of [ $^{125}\text{I}$ ]MIE<sub>2</sub> uptake was in agreement with the expected ER distribution in ovariectomized rats: highest uptake in the preoptic and hypothalamic nuclei, followed by the amygdala, hippocampus, midbrain, and frontal cortex. In the tamoxifen-treated rats, a reduction in [ $^{125}\text{I}$ ]MIE<sub>2</sub> binding was mainly seen in ER $\alpha$ -dominated regions like the cortical amygdala, central gray, and anterior paraventricular thalamus. A moderate inhibition of tracer uptake was found in the regions where both ER subtypes are expressed, like the ventrolateral-ventromedial hypothalamus, medial preoptic area, medial amygdala, and arcuate nucleus. After estradiol treatment, [ $^{125}\text{I}$ ]MIE<sub>2</sub> showed a low and homogenous uptake in the brain, with an overall reduction in ER binding of 87% as compared to untreated controls (Pareto et al. 2004).

Fluoroestradiol ([ $^{18}\text{F}$ ]FES) is the only PET tracer that has been used to assess ER expression in rat brain. In a very detailed study, the suitability of [ $^{18}\text{F}$ ]FES for in vivo quantification of ER was tested by both equilibrium and dynamic kinetic analysis (Moresco et al. 1995). Uptake of [ $^{18}\text{F}$ ]FES was mainly seen in ER-rich regions in the brain, such as in the pituitary and hypothalamus, with a maximum uptake observed between 25 and 30 min postinjection, reaching a pseudoequilibrium at 30 min. Blocking with increasing doses of unlabeled estradiol reduced the accumulation in the pituitary and hypothalamus in a dose-dependent manner, whereas no effect was observed in the hippocampus, cortex, and striatum. The specific binding of [ $^{18}\text{F}$ ]FES was calculated by equilibrium analysis considering the striatum as a reference tissue where the expression of the ER was expected to be very low. The equilibrium analysis showed saturable binding in the pituitary and hypothalamus with highest binding potential ( $B_{\text{max}}/K_{\text{d}}$ ) in the pituitary (16.11) and hypothalamus (1.97). The  $B_{\text{max}}$  and the  $K_{\text{d}}$  were found to be much higher in the pituitary ( $124.01 \pm 12.36$  pmol/g and  $7.70 \pm 1.07$ ) and then in the hypothalamus ( $11.85 \pm 0.16$  pmol/g and  $6.01 \pm 0.41$ ). The in vivo quantification of ER gave similar results in the pituitary and hypothalamus when either equilibrium or graphical analysis was used (Moresco et al. 1995). In another study, the effect of long-term treatment of estradiol and tamoxifen on ER occupancy in rat brain was studied by ex vivo autoradiography with [ $^{18}\text{F}$ ]FES. Although this study showed marked reduction in the uptake of [ $^{18}\text{F}$ ]FES in ER-rich areas, such as the hypothalamus, preoptic area, amygdala, and frontal cortex in the treatment groups, no differences in cerebellar uptake was observed (Pareto et al. 2004).

All the aforementioned studies were performed by *ex vivo* measurement of tracer uptake, without the use of *in vivo* imaging. We studied the feasibility of [ $^{18}\text{F}$ ]FES to monitor ER in the rat brain by small animal PET, and the results were in agreement with the ones found by Moresco et al. (1995). The brain time-activity curves of [ $^{18}\text{F}$ ]FES showed a quick peak uptake immediately after intravenous tracer injection, which was followed by a washout leading to a state of pseudoequilibrium after 25 min. Highest uptake was seen in the pituitary gland, followed by the hypothalamus. The rest of the brain areas showed low uptake. Blocking of the receptor with an excess of estradiol resulted in a statistically significant reduction in tracer uptake only in the pituitary gland and hypothalamus. A statistically significant higher volume of distribution and binding potential was found in the pituitary, when compared to other brain regions. [ $^{18}\text{F}$ ]FES uptake was not significantly affected by differences in plasma estrogen levels (Khayum et al. 2014).

So far, [ $^{18}\text{F}$ ]FES is the only tracer that was tested for *in vivo* visualization of ER in the brain of rats. [ $^{18}\text{F}$ ]FES showed promising results, as it displayed highest uptake in the regions of the brain with high expression of ER. It has also been shown that using this tracer, ER-positive tumors can be identified in the brain (Moresco et al. 1997). Nonetheless, further research is necessary to validate this tracer for ER imaging in the human brain. We recently used [ $^{18}\text{F}$ ]FES PET to determine the receptor occupancy of a novel ER-binding drug in postmenopausal women (Conlan et al. 2020). Highest [ $^{18}\text{F}$ ]FES uptake was observed in pituitary, with low tracer uptake in all other brain regions. After treatment with the ER-binding drug, [ $^{18}\text{F}$ ]FES uptake was significantly reduced in pituitary only, suggesting that [ $^{18}\text{F}$ ]FES PET can only show ER-specific binding in this brain region.

Moreover, the availability of a new ER $\beta$ -selective tracer would be highly desirable, since the different isoforms of ER have different expression patterns and different functions in neuroprotection, mood, and memory (Foster 2012; Ooishi et al. 2012). [ $^{18}\text{F}$ ]FHNP seems to be a good candidate for selective binding to ER $\beta$  in the brain (Antunes et al. 2017a; Antunes et al. 2017b). However, further studies are needed to validate this tracer.

### 14.5.2 Imaging of Other Hormone Receptors in the Brain

The only tracer that has been tested for AR imaging in the brain is [ $^{18}\text{F}$ ]FDHT. Although this tracer is already used in the clinical studies in cancer patients, it proved not suitable for the detection of AR in the brain of rats. Fast metabolism, low and nonspecific binding are the main issues that were encountered (Khayum et al. 2015). [ $^{18}\text{F}$ ]FPTP is the only tracer that could visualize PR both in humans and in experimental animals so far, but further evaluation of the tracer for brain imaging is required. Additionally, [ $^{18}\text{F}$ ]FFNP PET was found to be a safe, noninvasive method to evaluate tumor PR *in vivo* in patients with breast cancer, positioning this tracer as a candidate to be used for brain research (Fowler

et al. 2012; Salem et al. 2018, 2019). Unfortunately, no selective tracers for the different isoforms of the PR and AR are available yet. No suitable tracers for GR and MR imaging are available either. Although some studies explored the use of some PET tracers for GR and MR receptors in the brain, none of them proved to be suitable.

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## 14.6 Conclusion and Perspectives

Despite the large amount of research focusing on SHRs and their functions in the brain, many questions still remain unanswered. PET and SPECT imaging could be attractive tools for studying SHR function in the brain in living subjects. Several PET and SPECT tracers have been already evaluated for detection of SHR in oncology, and some of them have already entered into the clinical setting. Tracers that are used for diagnosis and monitoring of therapy response in hormone-sensitive cancers can easily be tested for applications in neurological and psychiatric disorders as well. So far, [ $^{18}\text{F}$ ]FES, [ $^{18}\text{F}$ ]FFNP, and [ $^{18}\text{F}$ ]FDHT have been used successfully in cancer patients for imaging of ER, PR, and AR, respectively. Only [ $^{18}\text{F}$ ]FES and [ $^{18}\text{F}$ ]FDHT have been tested to detect ER and AR in the brain of rodents. More studies using these tracers are needed to validate their use in brain research. The development of new tracers with higher affinity and specificity can help to increase the use of molecular imaging of sex hormone receptors in the brain.

Development of tracers for corticoid receptors has long been attempted, but these efforts have not yet produced any promising results. When tracers for corticoid receptors finally become available, however, they will be valuable assets for research in stress, mood, and neuropsychiatric disorders that are known to be associated with dysregulation of the HPA axis.

Aging and lifestyle-associated stress are known to change SHR expression. Because SHR receptors are associated with many brain functions, imaging of SHR in the brain could be a valuable tool to study disease mechanisms associated with, e.g., changes in expressional levels of SHR, which can ultimately culminate into neurological and psychiatric disorders. Understanding disease mechanisms could provide a rationale for treatment regimens based on pharmacological compounds associated with SHR. PET and SPECT imaging of SHR could also be helpful in drug discovery for individual SHR-associated disorders. PET and SPECT may provide information about the amount of a test drug that binds to the target and about the residence time of the drug in the brain. Such information about a new drug is highly valuable for decision-making in drug development and optimizing the dose and route of administration.

Thus, PET and SPECT imaging of SHR may not only be helpful in the diagnosis of neurological and psychiatric disorders associated with altered SHR expression but could also be instrumental in providing a better understanding of disease mechanisms and the development of new intervention strategies.



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## Abstract

The monoamine oxidase B (MAO-B) enzyme has a prominent role in brain function, being involved in the breakdown of monoaminergic neurotransmitters. Abnormally high MAO-B levels in the brain, found across different neurological and neurodegenerative disorders including Alzheimer's disease (AD) and Parkinson's disease (PD), are associated with deleterious effects involving reduc-

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tions in dopamine levels and increased oxidative stress, astrocytosis, and neuroinflammation. Recent advances in positron emission tomography (PET) using radiotracers specific to MAO-B have allowed the *in vitro* and *in vivo* investigation of the brain regional and temporal patterns of MAO-B in normal aging and brain disorders including depression, AD, PD, and amyotrophic lateral sclerosis (ALS). Multiple MAO-B tracers, characterized by either irreversible or reversible binding kinetics, have been under development and validation over the past ~30 years. Recent multimodal MRI-PET studies have allowed investigating MAO-B in relation to other pathophysiological changes including amyloid- $\beta$  and tau deposition, glucose metabolism, and brain structural changes. PET imaging of MAO-B is also useful for the monitoring of clinical trials of MAO-B inhibitors. This chapter presents a review of *in vivo* PET imaging studies of MAO-B from preclinical to clinical applications as well as *in vitro* autoradiography investigations, highlighting recent progress in the field and future prospects.

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## Abbreviations

AD	Alzheimer's disease
ALS	Amyotrophic lateral sclerosis
CJD	Creutzfeldt-Jakob disease
DWI	Diffusion-weighted imaging
FDG	Fluorodeoxyglucose
GABA	$\gamma$ -aminobutyric acid
GFAP	Glial fibrillary acidic protein
MAO-A	Monoamine oxidase A
MAO-B	Monoamine oxidase B
MRI	Magnetic resonance imaging
PD	Parkinson's disease
PET	Positron emission tomography
PiB	Pittsburgh compound B
SUVR	Standardized uptake value ratio
TSPO	Translocator protein

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## 15.1 Neurobiology and Clinical Relevance of MAO-B

Monoamine oxidases (MAOs) are flavin-containing enzymes that regulate the levels of monoaminergic neurotransmitters, with critical implications for the maintenance of healthy brain function. From a neurochemical perspective, the MAO enzymes catalyze the oxidative deamination of endogenous monoamine neurotransmitters as well as of amines of exogenous origin. As a by-product of this oxidation reaction, reactive oxygen species such as hydrogen peroxide are released, and their excess

has been linked to oxidative stress and neuroinflammation (Ramsay 2016; Youdim et al. 2006). There are two types of MAO, MAO-A and MAO-B, both located in the mitochondrial outer membrane, but with distinct substrate specificities and regional distributions in the brain and peripheral organs (Saura et al. 1996).

### 15.1.1 Neurobiology of MAO-B

MAO-B has high clinical relevance, being the major enzyme responsible for metabolizing dopamine in the human brain, a neurotransmitter that is essential to multiple brain functions including movement, mood, motivation, and cognition. The accumulation of data from *in vitro* and *in vivo* studies, as reviewed in Sections 15.4 and 15.5, has shown that brain MAO-B levels are elevated in a range of neurodegenerative and other neurological and neuropsychiatric disorders, including depression, Parkinson's disease (PD) and Parkinsonian disorders, amyotrophic lateral sclerosis (ALS), and Alzheimer's disease (AD) (Fowler et al. 2015; Ramsay 2016). In these disorders, the abnormal upregulation of MAO-B leads to declining levels of dopamine and other neurotransmitters, together with excess neurotoxic byproducts, oxidative stress, and neuroinflammation, all of which promote neuronal dysfunction and neurodegeneration (Arranz and De Strooper 2019; De Strooper and Karran 2016). MAO-B is abundantly located in astrocytes, and overexpressed under certain pathological conditions associated to astrocytosis (De Strooper and Karran 2016). Preclinical studies in a transgenic mouse model of AD reported MAO-B upregulation in astrocytes, resulting in excess  $\gamma$ -aminobutyric acid (GABA) production and cognitive deficits (Jo et al. 2014). Suppressing GABA synthesis by inhibiting MAO-B improved synaptic plasticity and reduced astrocytosis in the brains of transgenic AD mice (Park et al. 2019). Therefore, there is a growing interest in the investigation of reactive astrocytes as a promising therapeutic target (Arranz and De Strooper 2019; Boche et al. 2019), and PET imaging of MAO-B could play a major role toward the quantification of astrocytosis in the context of the pathological cascade in neurodegenerative and other brain disorders. Further studies have shown that MAO-B overexpression is associated to increased  $\gamma$ -secretase-mediated amyloid- $\beta$  production in the hypothalamus and frontal cortex in AD, being found both in astrocytes and in neurons (Schedin-Weiss et al. 2017). Building on all the available preclinical evidence, there has been a long-standing effort to unravel the biological mechanisms underlying MAO-B overexpression and to explore its regulation as a potential therapeutic target in a range of brain disorders.

### 15.1.2 Molecular Neuroanatomy of MAO-B

Knowledge about the neuroanatomical distribution and cellular localization of MAO-B has been gained through *in vitro* studies in human and animal models using biochemical techniques such as immunoblotting, immunohistochemistry, and immunocytochemistry. While animal models have provided a wealth of valuable



data, interspecies differences are a limitation of translational research. For example, the human brain has substantially higher tissue concentrations of MAO-B than are found in rodents (Saura et al. 1996). This means that while animal models are helpful for initial testing of radiotracers, *in vitro* and *in vivo* studies in human brain are needed before tracers can be validated for use in clinical studies.

Early studies determined that MAO-B is located both in neurons and astrocytes, but with different predominance across brain regions. Studies in normal aging and across neurodegenerative diseases are converging to the finding that MAO-B is abundantly expressed in glial cells, particularly in the outer wall of mitochondria in astrocytes (Saura et al. 1996). The astrocytic localization of MAO-B has been revealed experimentally using immunohistochemical staining for glial fibrillary acidic protein (GFAP), an intermediate filament expressed by reactive astrocytes, as well as using markers for other astrocytic filaments such as vimentin and nestin. Early immunocytochemistry studies in the rat brain (Levitt et al. 1982) using a double immunofluorescence technique by means of antisera to MAO-B and either GFAP as a marker of astrocytes or serotonin as a marker of serotonergic neurons, determined that MAO-B was overexpressed in astrocytes. In certain areas of the rat brain, MAO-B was also expressed in serotonin-containing neurons, in particular in the brainstem raphe nuclei and the hypothalamus, *i.e.*, regions rich in serotonergic cell groups. In postmortem spinal cord tissue from ALS patients, double-staining studies using MAO-B and GFAP determined that MAO-B was co-located with astrocytes as measured by GFAP; high MAO-B was also observed in serotonergic and histaminergic neurons (Ekblom et al. 1993, 1994). The astrocytes displayed a high level of phenotypic heterogeneity, whereby not all types of astrocytes overexpressed MAO-B, so that the correlation between MAO-B and GFAP was only moderate.

More recently, quantitative immunoblotting analyses of MAO-B antibodies in postmortem human control brain tissue with an average age of 48 years have shown that the MAO-B distribution in the brain is heterogeneous. Highest levels of MAO-B were observed in the hypothalamus, nucleus basalis, and hippocampal uncus, followed by rather high levels in the striatum and globus pallidus, and somewhat lower levels in other cortical regions including frontal and cingulate cortices; lowest levels were measured in the cerebellum (Tong et al. 2013). When focusing on the frontal cortices, MAO-B was positively associated with age showing an ~18% increase in MAO-B per decade from 18 to 99 years (Tong et al. 2013). Besides age, a number of other factors influence MAO-B expression in the human brain, including smoking, depressive symptoms, neuroinflammation, and medications such as the use of MAO-B inhibitors (Fowler et al. 2015).

Compared to control subjects, MAO-B levels were high in PD and Parkinsonian disorders including multiple system atrophy and progressive supranuclear palsy (Tong et al. 2017). Also, MAO-B levels were positively correlated with standard astrocyte markers including GFAP, vimentin, and nestin across PD and Parkinsonian disorders. These findings provide additional evidence that levels of MAO-B protein may serve as a biochemical marker of astrocyte activation, with the limitation that

MAO-B may not be entirely specific for astrocytosis (Tong et al. 2017). Overall, the elevation in MAO-B levels was less prominent than that measured with standard astrocytic markers (Tong et al. 2017), which highlights that MAO-B and GFAP, while moderately correlated, are not measuring exactly the same processes. More detailed findings from *in vitro* autoradiography studies are reviewed in Sect. 15.4.

### 15.1.3 MAO-B as Therapeutic Target

From a therapeutic point of view, there are a number of treatment strategies and clinical trials aiming at inhibiting MAO-B levels in the brain. Based on accumulated neurobiological knowledge about the role of MAO-B in brain function, the inhibition of this enzyme would dampen the metabolism of dopamine, leading to increased levels of endogenous dopamine and reduced neuroinflammation. To date, many clinical trials of MAO-B inhibitors have been conducted or are underway toward the treatment of depression, PD, and more recently AD, partly due to the growing interest in neuroinflammation as a therapeutic target (Ramsay 2016; Schapira et al. 2006; Youdim et al. 2006). The FDA (US Food and Drug Administration) approved the irreversible MAO-B inhibitors selegiline and rasagiline for the symptomatic treatment of motor dysfunction in PD. More recently, reversible MAO-B inhibitors are undergoing clinical trials, but their efficacy is still uncertain.

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## 15.2 Requirements to Develop PET Tracers for Imaging MAO-B

Positron emission tomography (PET) is a molecular imaging technique that allows measuring molecular changes in the brain with very high sensitivity. With a growing recognition of the role of neuroinflammation across multiple brain disorders, PET imaging of MAO-B is a promising tool to quantify astrocytosis and as an outcome measure in clinical trials for evaluating the efficacy and optimize the dosage of MAO-B inhibitors (Fowler et al. 2015, 2005; Narayanaswami et al. 2019). The main challenges in the development of radiotracers for MAO-B have been to produce chemical compounds with high specificity and sensitivity for MAO-B and with appropriate pharmacokinetics so that the quantitative PET binding measures are reliable estimates of the actual concentrations of MAO-B enzyme in the brain. The radiotracers should be sufficiently lipophilic to cross the blood-brain barrier, but their lipophilicity should be limited to minimize their non-specific binding to white matter. It is also important to minimize the amounts of radioactive metabolites entering the brain, which may confound the MAO-B PET signal.

Despite active research into MAO-B tracers for the past ~30 years, to date only a limited number of radiotracers have met the required qualities for potential translation to human. A summary of the radiotracers that will be reviewed in this chapter is presented in Table 15.1; the corresponding chemical structures of these tracers are

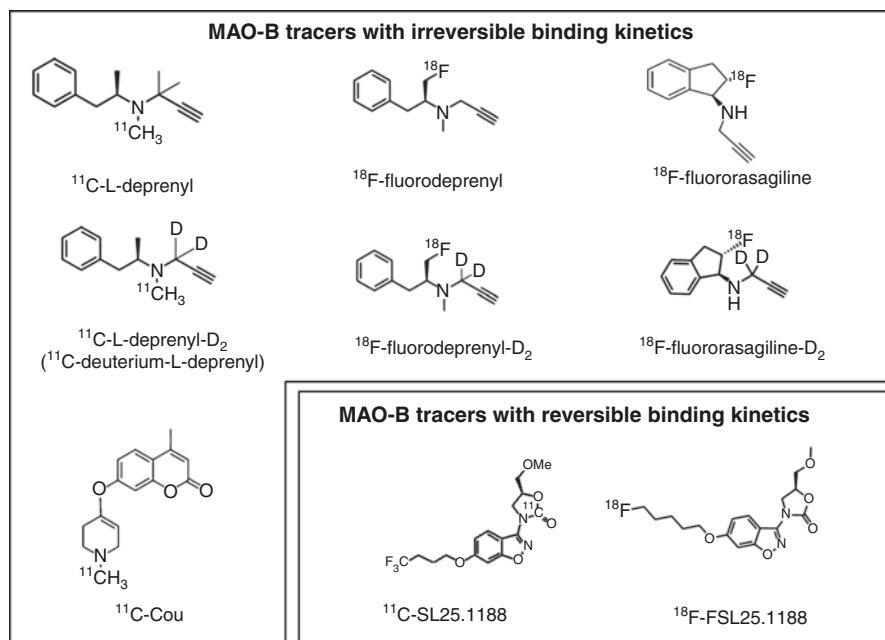
**Table 15.1** Schematic summary of MAO-B PET tracers applied in in vivo imaging studies

PET tracers	Pharmacokinetic modelling studies	In vivo PET imaging studies in animal models	In vivo PET imaging studies in human
<i>Irreversible binding tracers</i>			
<sup>11</sup> C-L-deprenyl	<i>2-tissue-compartment model with plasma input:</i> Lammertsma et al. (1991) (human) Fowler et al. (1995) (human) <i>Graphical Patlak analysis with plasma input:</i> Fowler et al. (1987) (human) Lammertsma et al. (1991) (human)	<i>Mouse:</i> MacGregor et al. (1985) <i>Baboon:</i> Fowler et al. (1988)	<i>Test-retest studies:</i> Logan et al. (2000) <i>Clinical trials of MAO-B inhibitors:</i> Bench et al. (1991) Fowler et al. (1994) Freedman et al. (2005)
<sup>11</sup> C-L-deprenyl-D <sub>2</sub> (also known as <sup>11</sup> C-deuterium-L-deprenyl)	<i>2-tissue-compartment model with plasma input:</i> Fowler et al. (1988) (baboon) Fowler et al. (1995) (human) <i>Graphical Patlak analysis with plasma input:</i> Fowler et al. (1995) (human) <i>Modified reference Patlak analysis:</i> Bergström et al. (1998) (human) Johansson et al. (2007) (human)	<i>Baboon:</i> Fowler et al. (1988) <i>Transgenic mouse models of Alzheimer's disease:</i> Rodriguez-Vieitez et al. (2015) Olsen et al. (2018)	<i>Test-retest studies:</i> Logan et al. (2000) Arakawa et al. (2017) <i>Aging:</i> Fowler et al. (1997) <i>Effects of smoking:</i> Fowler et al. (1996) <i>Alzheimer's disease:</i> Carter et al. (2012), Carter et al. (2019a, 2019b) Rodriguez-Vieitez et al. (2016a, 2016b) Vilaplana et al. (2020) <i>Amyotrophic lateral sclerosis:</i> Johansson et al. (2007) <i>Creutzfeldt-Jakob disease:</i> Engler et al. (2003, 2012) <i>Epilepsy:</i> Bergström et al. (1998) <i>Fibromyalgia:</i> Albrecht et al. (2019) <i>Clinical trials of MAO-B inhibitors:</i> Hirvonen et al. (2009) Sturm et al. (2017)
<sup>18</sup> F-fluorodeprenyl	Not available	<i>Cynomolgus monkey:</i> Nag et al. (2012b)	Not available

**Table 15.1** (continued)

PET tracers	Pharmacokinetic modelling studies	In vivo PET imaging studies in animal models	In vivo PET imaging studies in human
<sup>18</sup> F-fluorodeprenyl-D <sub>2</sub>	<i>2-tissue-compartment model with plasma input:</i> Nag et al. (2016) (cynomolgus monkey)	<i>Cynomolgus monkey:</i> Nag et al. (2016)	Not available
<sup>18</sup> F-fluororasagiline	Not available	<i>Cynomolgus monkey:</i> Nag et al. (2012a)	Not available
<sup>18</sup> F-fluororasagiline-D <sub>2</sub>	Not available	<i>Cynomolgus monkey:</i> Nag et al. (2013)	Not available
<sup>11</sup> C-Cou	<i>2-tissue-compartment model with plasma input:</i> Drake et al. (2018) (rhesus monkey)	<i>Rhesus monkey:</i> Drake et al. (2018)	Not available
<i>Reversible binding tracers</i>			
<sup>11</sup> C-SL25.1188	<i>1-tissue-compartment model with plasma input:</i> Saba et al. (2010) (baboon) Rusjan et al. (2014) (human) <i>2-tissue-compartment model with plasma input:</i> Saba et al. (2010) (baboon) Rusjan et al. (2014) (human) <i>Graphical Logan analysis with plasma input:</i> Saba et al. (2010) (baboon) Rusjan et al. (2014) (human)	<i>Baboon:</i> Saba et al. (2010)	<i>Healthy human brain:</i> Rusjan et al. (2014) <i>Depression:</i> Moriguchi et al. (2019)
<sup>18</sup> F-FSL25.1188	<i>Graphical Logan analysis with plasma input:</i> Dahl et al. (2019) (cynomolgus monkey)	<i>Cynomolgus monkey:</i> Dahl et al. (2019)	Not available

(continued)



**Fig. 15.1** Chemical structures of PET tracers for imaging MAO-B

illustrated in Fig. 15.1. From a pharmacokinetic point of view, tracers are classified into those that are best described by either irreversible or reversible binding kinetics. Most MAO-B tracers that have reached application in human PET studies have been labelled with carbon-11 (half-life of 20.4 min), which is useful when repeated scans are needed in the same subjects over a short period of time, for example, in drug dosing studies. More recently, there is an intense research effort to develop fluorine-18 (half-life of 110 min)-labelled radiotracers, which have other advantages, in particular greater clinical utility for administration in facilities that do not have a cyclotron on site. Full pharmacokinetic modelling with arterial sampling is used for newly developed MAO-B tracers, followed by validation of simplified models using reference regions that obviate the need for arterial sampling, and therefore have greater potential for clinical application.

## 15.3 Development and Pharmacokinetic Characterization of MAO-B PET Tracers

### 15.3.1 Tracers with Irreversible Binding Kinetics

#### 15.3.1.1 Carbon-11-Labelled Tracers

Originally developed in the 1980s, the most widely investigated PET tracers for MAO-B have been the irreversible binding tracers  $^{11}\text{C}$ -L-deprenyl and its deuterated analog,  $^{11}\text{C}$ -L-deprenyl- $\text{D}_2$ —also referred to as  $^{11}\text{C}$ -deuterium-L-deprenyl (Fowler

et al. 2015); these tracers are stereoselective, so that only the L enantiomers (but not the D) are active toward MAO-B. In these tracers, also known as “suicide enzyme inhibitors,” the propargylamine chemical structure forms a covalent bond to the flavin cofactor of the MAO-B enzyme. This covalent bond is formed via the cleavage of the carbon-hydrogen or carbon-deuterium bonds on the methylene carbon of the propargyl group. As the tracers bind to the catalytically active MAO-B enzyme molecules, the signal intensity recorded by the PET scanner reflects the concentration of MAO-B in brain tissue. The carbon-deuterium bonds are stronger and thus require more energy to be cleaved compared to the carbon-hydrogen bonds. As a result, the deuterated tracers have lower binding rates to MAO-B compared to the respective non-deuterated forms—also known as the “deuterium effect”—and thus the deuterated tracers have enhanced sensitivity to MAO-B.

Early *in vivo* PET investigations showed that  $^{11}\text{C}$ -L-deprenyl has a high level of specificity to MAO-B in mouse brain (MacGregor et al. 1985), as well as in human brain (Fowler et al. 1987)—where it was blocked by the administration of unlabelled L-deprenyl and the non-selective MAO inhibitor phenelzine. A subsequent pharmacokinetic study of  $^{11}\text{C}$ -L-deprenyl using experimental data with arterial plasma input in normal subjects compared irreversible vs. reversible 2-tissue-compartment models (Lammertsma et al. 1991) where the brain tissue is divided into two compartments: (1) free and non-specific binding in tissue and (2) specific binding in tissue. The  $^{11}\text{C}$ -L-deprenyl binding data were well fitted by the 2-tissue-compartment model, using rate constants for the transfer from plasma to tissue ( $K_1$ ) and viceversa ( $k_2$ ), and with  $k_3$  as the rate of irreversible binding to MAO-B. The inclusion of a  $k_4$  constant to simulate reversible binding kinetics to MAO-B did not significantly improve the model fit, thus supporting the irreversible model. The data were also well fitted by a graphical Patlak model for irreversible binding (Patlak and Blasberg 1985), where the concentration of MAO-B in brain is quantified by  $K_i$ , the rate of net influx to the specifically bound compartment. Of note, the influence of cerebral blood flow on the Patlak-derived  $K_i$  net influx rate and on  $k_3$  as obtained from the 2-tissue-compartment model was examined by using data from one subject with both  $^{11}\text{C}$ -L-deprenyl and  $^{15}\text{O}$ -water PET scans available. The analyses showed that the regional Patlak-derived net influx rates ( $K_i$ ) for  $^{11}\text{C}$ -L-deprenyl were positively correlated to cerebral blood flow. Using various doses of the reversible MAO-B inhibitor Ro 19–6327, Lammertsma et al. (1991) showed that the inhibition of MAO-B cannot be reliably measured using the Patlak-derived  $K_i$  constant because of variations in cerebral blood flow. In contrast, the rate of binding to MAO-B derived from the 2-tissue-compartment model ( $k_3$ ) was not strongly influenced by cerebral blood flow, and therefore this study suggests that the full kinetic modelling ( $k_3$  constant) is a more reliable method for MAO-B quantification compared to the Patlak analysis ( $K_i$ ) for the  $^{11}\text{C}$ -L-deprenyl tracer, especially for the evaluation of drug trials.

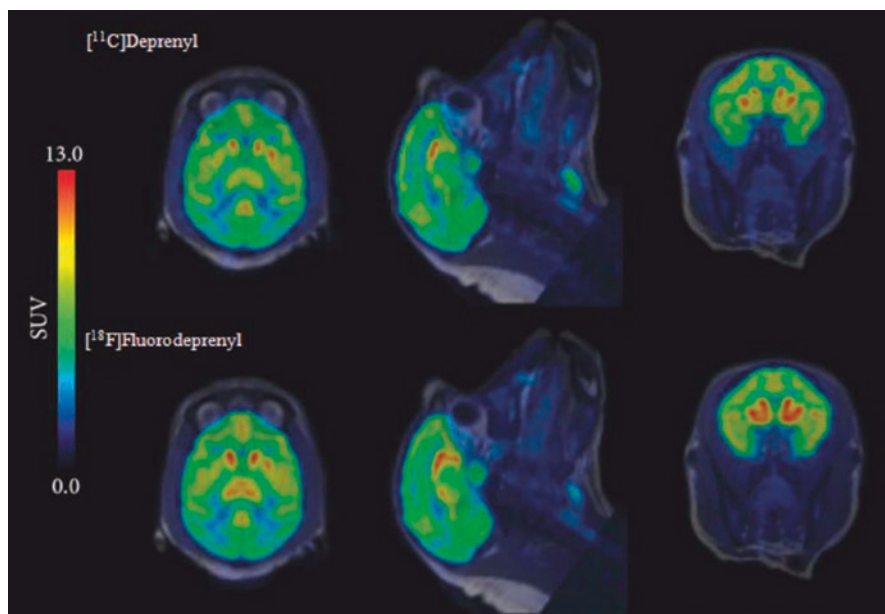
The pharmacokinetic properties of  $^{11}\text{C}$ -L-deprenyl- $\text{D}_2$  and  $^{11}\text{C}$ -L-deprenyl were compared to investigate the deuterium effect in the baboon brain (Fowler et al. 1988) and in human volunteers (Fowler et al. 1995). Both tracers were modelled by a 2-tissue-compartment model with arterial plasma input as previously described (Lammertsma et al. 1991), as well as by graphical analysis for irreversible tracers in

the study in humans (Fowler et al. 1995). Using the compartmental model, it was found that  $\lambda k_3$  (with  $\lambda = K_1/k_2$ ) is a more robust measure of radiotracer binding than  $k_3$ , being less influenced by cerebral blood flow. Importantly, the deuterated compound ( $^{11}\text{C}$ -L-deprenyl- $\text{D}_2$ ) had slower rate of trapping in human brain as measured by  $\lambda k_3$  compared to  $^{11}\text{C}$ -L-deprenyl, leading to increased sensitivity to changes in MAO-B, and lower influence of brain perfusion on tracer binding. Test-retest investigations with  $^{11}\text{C}$ -L-deprenyl- $\text{D}_2$  indicated that  $\lambda k_3$  had a low test-retest variability with an absolute value of less than  $\sim 10\%$  in healthy subjects (Logan et al. 2000; Arakawa et al. 2017); further test-retest investigations in patients would be valuable. Due to its improved pharmacokinetic properties, subsequent studies focused on the  $^{11}\text{C}$ -L-deprenyl- $\text{D}_2$  tracer only, where a modified reference Patlak model using the cerebellum gray matter as reference allowed estimating tracer binding by means of parametric Patlak slope images, a simplified approach with greater potential for clinical application as it obviates the need for arterial sampling (Bergström et al. 1998; Johansson et al. 2007; Rodriguez-Vieitez and Nordberg 2018; Patlak and Blasberg 1985).

During the past few years, a new family of metabolically trapped compounds with irreversible binding kinetics has been under development (Narayanaswami et al. 2019). A metabolically trapped tracer for MAO-B is a compound that serves as substrate for the MAO-B enzyme, and which is selectively metabolized and trapped within the brain tissue, but without forming a covalent bond (the most well-known metabolically trapped tracer is  $^{18}\text{F}$ -FDG, an analog of glucose that is trapped in highly metabolic regions in the brain). Preliminary studies on tetrahydropyridine compounds showed that they are metabolically trapped by MAO-B enzyme in brain tissue of rhesus monkey brain, but not substantially trapped in rat brain, pointing to species differences and the need to perform tests in human brain. The tracer  $^{11}\text{C}$ -Cou is a tetrahydropyridine, which is a metabolically trapped PET tracer for MAO-B, showing rapid and irreversible retention in the primate brain. Since  $^{11}\text{C}$ -Cou demonstrated quite rapid rate of trapping, a deuterated analog was developed, but it did not show suitable pharmacokinetics in rhesus monkey brain (Drake et al. 2018). MAO-B mouse models of PD (Chamoli et al. 2018; Mallajosyula et al. 2008; Siddiqui et al. 2011) are being considered for the testing of newly developed compounds prior to potential translation to human PET application.

### 15.3.1.2 Fluorine-18-Labelled Tracers

Recent advances in the chemical organic synthesis of fluorinated L-deprenyl compounds have allowed developing various tracers that incorporate fluorine-18 in different positions on the aliphatic chain rather than in the aromatic ring (Nag et al. 2011), resulting in compounds with high MAO-B affinity and selectivity as shown by autoradiography in postmortem human brain tissue. PET studies with  $^{18}\text{F}$ -fluorodeprenyl in the cynomolgus monkey brain showed the expected in vivo MAO-B distribution in the brain as illustrated in Fig. 15.2, and irreversible pharmacokinetic behavior (Nag et al. 2012b). Subsequently, the same research group developed the respective deuterated analog,  $^{18}\text{F}$ -fluorodeprenyl- $\text{D}_2$ , which showed the



**Fig. 15.2** Illustrative PET scans using MAO-B tracers  $^{11}\text{C}$ -L-deprenyl and  $^{18}\text{F}$ -fluorodeprenyl co-registered with MRI and averaged between 9 and 120 min in cynomolgus monkey brain. Reprinted from Nag et al. (2012b), with permission from John Wiley & Sons Inc. © 1999–2020. *SUV* standardized uptake value

expected *in vivo* MAO-B regional distribution in cynomolgus monkey brain, and decreased tracer uptake after administration of MAO-B inhibitor L-deprenyl (Nag et al. 2016). Kinetic modeling of the *in vivo*  $^{18}\text{F}$ -fluorodeprenyl- $\text{D}_2$  uptake in cynomolgus monkey brain using a 2-tissue-compartment model with 4 rate constants fitted the data better than the corresponding model with 3 rate constants, providing evidence for a certain degree of reversibility of this tracer. Also, a head-to-head comparison in cynomolgus monkey brain showed that  $^{18}\text{F}$ -fluorodeprenyl- $\text{D}_2$  has more reversible kinetic behavior than  $^{11}\text{C}$ -L-deprenyl- $\text{D}_2$  (Nag et al. 2016); despite these encouraging findings and that  $^{18}\text{F}$ -fluorodeprenyl- $\text{D}_2$  has shown the expected MAO-B regional distribution by *in vitro* autoradiography in human brain tissue, this tracer has not yet been translated to *in vivo* human PET studies.

Continuing on the path toward developing fluorinated MAO-B tracers for clinical use, the same research group developed and evaluated fluorine-18-labelled derivatives of rasagiline, also known as  $^{18}\text{F}$ -fluororasagiline (Nag et al. 2012a), but this tracer showed significant levels of radioactive metabolites entering the brain. With the motivation to develop a tracer with a certain degree of reversibility, the deuterated analog  $^{18}\text{F}$ -fluororasagiline- $\text{D}_2$  was subsequently synthesized (Nag et al. 2013). The  $^{18}\text{F}$ -fluororasagiline- $\text{D}_2$  tracer showed suitable kinetics in cynomolgus monkey brain; however, its translation to human *in vivo* studies is still pending.



### 15.3.2 Tracers with Reversible Binding Kinetics

While the irreversibly binding tracers reviewed above (Sect. 15.3.1) have provided good correlations with *in vitro* MAO-B protein concentrations in the brain (Tong et al. 2013), their rate of binding is partly influenced by cerebral blood flow and some radiotracers present with undesirable radioactive metabolites entering the brain. To overcome these limitations, there has been a recent effort to develop reversible tracers.

#### 15.3.2.1 Carbon-11-Labelled Tracers

A reversible carbamate inhibitor of MAO-B,  $^{11}\text{C}$ -SL25.1188, was synthesized via  $^{11}\text{C}$ -phosgene as a carbon-11 precursor. *In vivo* data on  $^{11}\text{C}$ -SL25.1188 tracer binding in baboon brain were appropriately fitted to 1-tissue-compartment or 2-tissue-compartment reversible models as well as by the graphical Logan approach (Logan 2000), and tracer binding could be blocked by pretreatment with L-deprenyl (Saba et al. 2010). Despite these promising findings, the need for  $^{11}\text{C}$ -phosgene as a precursor poses a limitation toward clinical application. To avoid the use of  $^{11}\text{C}$ -phosgene, a new technique using a direct  $^{11}\text{CO}_2$  fixation method was recently developed (Vasdev et al. 2011); the newly synthesized  $^{11}\text{C}$ -SL25.1188 tracer was subsequently tested in healthy human brain (Rusjan et al. 2014). In this paper, the pharmacokinetic evaluation of  $^{11}\text{C}$ -SL25.1188 in human brain revealed that the tracer has excellent properties including high reversibility, selectivity for MAO-B, and absence of metabolites entering the brain. The time-activity data including arterial sampling were well fitted by a 2-tissue-compartment model and by the graphical Logan approach for reversible binding. Being the first reversible tracer to be used in human for imaging MAO-B, the  $^{11}\text{C}$ -SL25.1188 tracer binding as measured by the distribution volume  $V_T$  derived from the Logan graphical approach provided a robust measure of MAO-B binding in the human brain, which was well correlated with the known regional distribution of MAO-B from postmortem studies (Rusjan et al. 2014). The  $^{11}\text{C}$ -SL25.1188 tracer was recently applied for the first time in a human brain disorder, being evaluated in a cohort of patients with major depressive disorder (Moriguchi et al. 2019), as further described in Sect. 15.5.

#### 15.3.2.2 Fluorine-18-Labelled Tracers

Current efforts are being directed at developing fluorinated reversible tracers, which would have clear advantages from a clinical perspective. A fluorinated version of SL25.1188,  $^{18}\text{F}$ -FSL25.1188, showed reversible *in vivo* binding kinetics in cynomolgus monkey brain with highest uptake in the thalamus and striatum, moderate in cortical regions, and lowest in the cerebellum; the expected MAO-B regional distribution was also observed by autoradiography in postmortem human brain sections (Dahl et al. 2019). As a potential limitation, the  $^{18}\text{F}$ -FSL25.1188 tracer showed some level of uptake in the skull at late time frames (from 50 to 90 min) due to *in vivo* defluorination, which could affect its accurate quantification in regions such as the temporal cortex. Research aimed at clinical translation of  $^{18}\text{F}$ -FSL25.1188 to human brain imaging is still at an early stage.

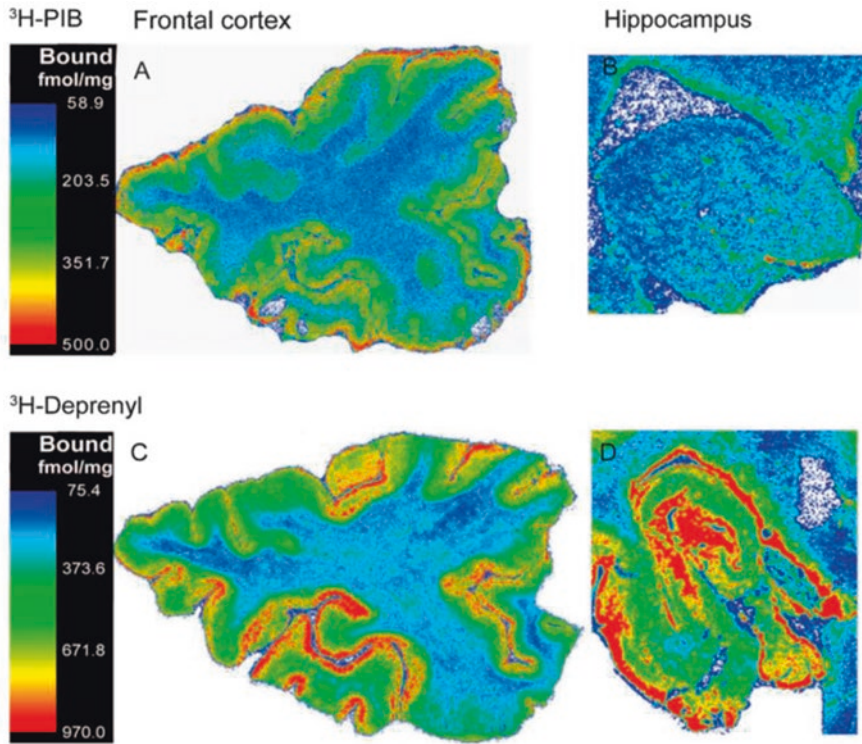
## 15.4 In Vitro Molecular Imaging of MAO-B by Autoradiography

In parallel with the development of novel MAO-B PET tracers, *in vitro* autoradiography studies in postmortem human brain tissue have been performed to test the specificity and binding properties of the tracers (Tong et al. 2013). More detailed *in vitro* autoradiography investigations are also performed to compare the regional and laminar distributions of MAO-B PET tracers vs. other tracers specific to amyloid- $\beta$  or tau in adjacent brain sections, often in combination with immunohistochemical measures to gain greater understanding of the types of neuropathological features associated with tracer binding. Despite the fact that *in vitro* autoradiography is an active field of research, *antemortem* vs. *postmortem* comparisons of MAO-B PET tracers are still rare.

Autoradiography studies of MAO-B began in the 1980s using  $^{11}\text{C}$ -labelled L-deprenyl in whole-hemisphere frozen sections from postmortem healthy human brain tissue. This tracer showed a pattern of high binding in the basal ganglia, thalamus, and hippocampus and moderate binding in the cerebral cortex; non-specific binding was low, amounting to  $\sim 20\%$  of the total binding (Jossan et al. 1989). Quantitative autoradiography using either  $^3\text{H}$ -L-deprenyl (Jossan et al. 1991) or the reversible MAO-B tracer  $^3\text{H}$ -lazabemide (Saura et al. 1994) showed higher binding in AD compared to control cases.  $^3\text{H}$ -L-deprenyl was elevated in spinal cord, motor cortex, and white matter in ALS compared to control cases (Jossan et al. 1994).

Frequently, autoradiography studies are performed in combination with immunohistochemical staining in adjacent sections. A positive correlation was observed between  $^3\text{H}$ -L-deprenyl binding and GFAP staining in ALS patients (Ekblom et al. 1993, 1994; Jossan et al. 1994), in support of the astroglial localization of MAO-B. Similarly,  $^3\text{H}$ -lazabemide binding was correlated with plaque-associated astrogliosis as measured by GFAP staining in AD brain tissue (Saura et al. 1994). Also using  $^3\text{H}$ -lazabemide autoradiography, Saura et al. (Saura et al. 1997) reported an overall positive association between MAO-B and age in control brains, which followed a biphasic trajectory characterized by a negligible change in MAO-B up to age 50, and a significant increase in MAO-B after 50 years, likely reflecting age-associated neuroinflammation.

These early autoradiography findings, together with a recently renewed interest in neuroinflammation as a therapeutic target in neurodegenerative diseases, have motivated further autoradiography studies using  $^{11}\text{C}$ -L-deprenyl combined with GFAP immunohistochemical staining in AD brain tissue.  $^{11}\text{C}$ -L-deprenyl binding showed the highest intensity in correlation with GFAP staining at early Braak stages of AD compared to that in more advanced disease stages, suggesting that astrogliosis plays an early role in disease progression (Gulyás et al. 2011). Subsequently, multitracer autoradiography studies (Lemoine et al. 2017; Marutle et al. 2013) were designed to investigate MAO-B in relation to other relevant pathological features in the AD brain including amyloid- $\beta$  and tau deposition. Both studies revealed similar laminar patterns of  $^3\text{H}$ -L-deprenyl binding in AD, which differed from the laminar distribution of fibrillar amyloid- $\beta$  deposition as measured by  $^3\text{H}$ -PiB (Lemoine et al. 2017; Marutle et al. 2013). In the frontal cortex,  $^3\text{H}$ -L-deprenyl binding was highest



**Fig. 15.3** In vitro autoradiography studies in Alzheimer's disease postmortem brain tissue using  $^3\text{H}$ -PiB in frontal cortex (a) and hippocampus (b), compared to  $^3\text{H}$ -L-deprenyl in frontal cortex (c) and hippocampus (d). Figure adapted from Marutle et al. (2013), under the terms of the Creative Commons CC BY license

in superficial layers, while  $^3\text{H}$ -PiB binding was elevated in all layers; in contrast, the hippocampus showed low  $^3\text{H}$ -PiB binding to fibrillar amyloid- $\beta$  but intense  $^3\text{H}$ -L-deprenyl binding in all layers, as illustrated in Fig. 15.3 (Marutle et al. 2013). GFAP staining confirmed abundant reactive astrocytes surrounding amyloid- $\beta$  plaques in correlation with  $^3\text{H}$ -L-deprenyl binding in superficial layers of the frontal cortex (Marutle et al. 2013). The laminar pattern of  $^3\text{H}$ -L-deprenyl was more closely associated with tau deposition as measured by  $^3\text{H}$ -THK5117 than it was to  $^3\text{H}$ -PiB (Lemoine et al. 2017), suggesting a close association between astrocytosis and tau pathology (Lemoine et al. 2017). Of note, early-generation tau tracers such as  $^3\text{H}$ -THK5117 have limitations due a certain amount of off-target binding to MAO-B most prominently in the basal ganglia (Lemoine et al. 2018). Therefore, the correlations between  $^3\text{H}$ -L-deprenyl and  $^3\text{H}$ -THK5117 PET tracers in these subcortical regions should be taken with a degree of caution (Lemoine et al. 2017; Murugan et al. 2019). Overall, there was a close association between  $^3\text{H}$ -L-deprenyl binding and GFAP in AD brain indicating a good level of correspondence between MAO-B and activated astrocytes; adjacent paraffin sections showed similar staining patterns for the tau marker AT8 and GFAP, adding support to the observed association between tau deposits and activated astrocytes (Lemoine et al. 2017).

There are also some challenges and limitations in the autoradiography studies of MAO-B. As pointed out in early studies (Fowler et al. 1987), the procedures for the isolation of MAO-B from its native environment within a tissue that are necessary for performing *in vitro* studies may change some properties of the enzyme, so differences in MAO-B activity may be observed when comparing *in vivo* and *in vitro* analyses. For example, while elevated *in vivo*  $^{11}\text{C}$ -deuterium-L-deprenyl binding was observed in a transgenic mouse model of AD compared to age-matched controls, no differences were observed by *in vitro* autoradiography with  $^3\text{H}$ -L-deprenyl (Rodriguez-Vieitez et al. 2015), which may point to limitations of the *in vitro* imaging technique in mouse brains. Finally, while there are some isolated case studies (Engler et al. 2012), no systematic comparisons between antemortem and postmortem MAO-B activity in human brain have been performed; these types of studies would be informative to validate the *in vivo* findings against *in vitro* data in the same individuals and to unravel the biological mechanisms and pathological changes underlying the *in vivo* PET signal.

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## 15.5 Applications of In Vivo PET Imaging of MAO-B in Health and Disease

### 15.5.1 Aging

An *in vivo* PET study with  $^{11}\text{C}$ -L-deprenyl- $\text{D}_2$  in healthy human brain (21 participants, age range 23–86 years) using arterial plasma sampling and a 2-tissue-compartment irreversible kinetic model showed an increase in  $^{11}\text{C}$ -L-deprenyl- $\text{D}_2$  binding ( $\lambda k_3$ ) with age of about 7% per decade in all brain regions except the cingulate gyrus, likely due to age-related neuroinflammation (Fowler et al. 1997). In contrast to the age-associated increase in MAO-B, subjects showed the expected decline in cerebral blood flow with age, as measured by the blood-to-brain influx constant  $K_1$ , as well as in glucose metabolism in 15 participants for whom both  $^{11}\text{C}$ -L-deprenyl- $\text{D}_2$  and  $^{18}\text{F}$ -FDG PET were available. Of note, the age-related increase in MAO-B observed *in vivo* was smaller than that previously reported by *in vitro* biochemical methods to determine MAO-B protein concentrations in brain tissues, thus pointing to some discrepancies between *in vivo* and *in vitro* findings.

### 15.5.2 Effects of Smoking

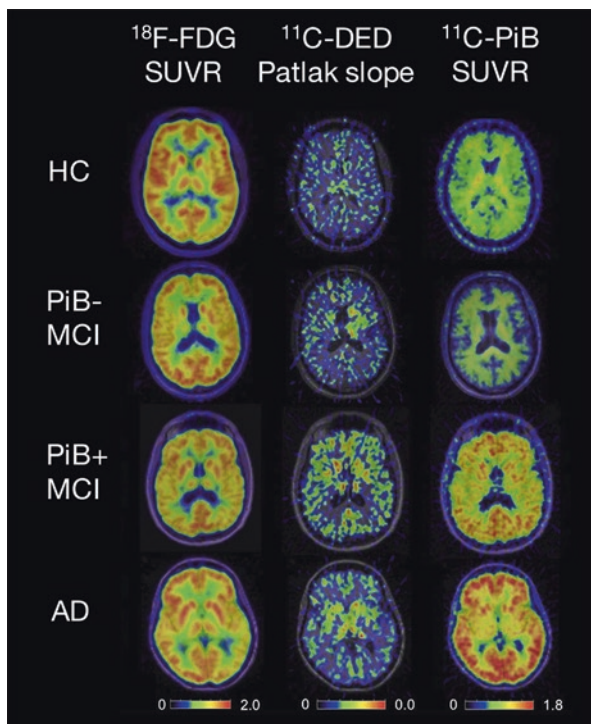
Previous studies have reported that smokers have reduced platelet MAO-B levels (Fowler et al. 2003). While the reasons for the observed reduction in MAO-B levels were unknown, it was hypothesized as due to either slower rates of MAO-B synthesis in smokers compared to non-smokers or to MAO-B inhibition caused by substances present in the tobacco, without excluding the possibility that low MAO-B individuals may be more vulnerable to smoking. PET studies with  $^{11}\text{C}$ -L-deprenyl- $\text{D}_2$  revealed that smokers had up to 40% reduction in MAO-B levels in the brain compared to non-smokers or former smokers (Fowler et al. 1996), while blood-to-brain flow ( $K_1$ ) and glucose metabolism ( $^{18}\text{F}$ -FDG) in smokers were not different from those in non-smokers.

### 15.5.3 Neuroinflammation in Neurodegenerative Diseases

Among the few PET tracers available to quantify astrocytosis *in vivo* in human brain, the most commonly used has been  $^{11}\text{C}$ -deuterium-L-deprenyl, as highlighted in recent reviews (Albrecht et al. 2016; Boche et al. 2019; Carter et al. 2019b; Narayanaswami et al. 2018; Varrone and Nordberg 2015). Elevated  $^{11}\text{C}$ -deuterium-L-deprenyl PET binding has been reported in a range of neurodegenerative diseases including AD (Carter et al. 2012; Rodriguez-Vieitez et al. 2016a), ALS (Johansson et al. 2007), and Creutzfeldt-Jakob disease (CJD) (Engler et al. 2003, 2012), as well as and in the affected hemisphere of epileptic patients (Bergström et al. 1998). A multitracer PET study in patients with probable CJD (Engler et al. 2003) revealed both high  $^{11}\text{C}$ -deuterium-L-deprenyl and low  $^{18}\text{F}$ -FDG uptake in several cortical regions, indicating the presence of astrocytosis and neuronal dysfunction.

Using a multitracer PET imaging study design including  $^{11}\text{C}$ -deuterium-L-deprenyl,  $^{11}\text{C}$ -PiB and  $^{18}\text{F}$ -FDG in sporadic AD patients (Rodriguez-Vieitez and Nordberg 2018), significantly increased  $^{11}\text{C}$ -deuterium-L-deprenyl binding was observed in prodromal stages of AD in comparison to healthy controls or to patients with AD dementia (Carter et al. 2012), as illustrated in Fig. 15.4. In addition, the 1–4 min early phase of  $^{11}\text{C}$ -deuterium-L-deprenyl standardized uptake value ratio (SUVR) was used as a surrogate marker of brain perfusion, being highly correlated

**Fig. 15.4** Representative *in vivo* multitracer PET scans of  $^{11}\text{C}$ -deuterium-L-deprenyl ( $^{11}\text{C}$ -DED), in comparison with  $^{18}\text{F}$ -FDG and  $^{11}\text{C}$ -PiB. AD Alzheimer's disease, HC healthy controls, MCI mild cognitive impairment, PiB+ (-) Pittsburgh compound B positive (negative), SUVR standardized uptake value ratio. Figure adapted from Rodriguez-Vieitez et al. (2016a), with permission © SNMMI



with glucose metabolism ( $^{18}\text{F}$ -FDG) (Rodriguez-Vieitez et al. 2016a); a limitation of this study was however the lack of an independent measure of brain perfusion such as  $^{15}\text{O}$ -water PET. The same multitracer PET paradigm investigated in familial autosomal dominant AD, in which mutation carriers develop symptomatic AD at a rather predictable age of onset, revealed that the most prominent elevation in  $^{11}\text{C}$ -deuterium-L-deprenyl binding occurs at the presymptomatic stage, up to two decades prior to symptom onset (Schöll et al. 2015). The longitudinal follow-up of this cohort showed increasing amyloid- $\beta$  plaque deposition ( $^{11}\text{C}$ -PiB) together with declining levels of both astrocytosis ( $^{11}\text{C}$ -deuterium-L-deprenyl) and glucose metabolism ( $^{18}\text{F}$ -FDG) in mutation carriers along disease progression (Carter et al. 2019a; Rodriguez-Vieitez et al. 2016b).  $^{11}\text{C}$ -deuterium-L-deprenyl was also found to be elevated from early disease stages in two different transgenic mouse models of AD (Olsen et al. 2018; Rodriguez-Vieitez et al. 2015); in both studies the astrocytic marker GFAP was observed to peak at late stages and not being well correlated to  $^{11}\text{C}$ -deuterium-L-deprenyl, suggesting that the two markers measure different aspects or stages of astrocytosis. Overall, these studies showing elevation of  $^{11}\text{C}$ -deuterium-L-deprenyl in neurodegenerative diseases suggest that the modulation of astrocytosis is a promising early therapeutic target, but further research is needed (Schott and Fox 2016).

More recently, multimodal MRI-PET imaging studies have allowed interrogating the relationships between structural and molecular changes in the brain. Molecular biology studies have shown that astrocytes and microglia interact in neurodegenerative diseases and that their activation is accompanied by morphological remodelling such as early hypertrophy of astrocytes followed by atrophy and astrodegeneration, while microglia undergo morphological changes toward ameboid-like forms (Arranz and De Strooper 2019; Boche et al. 2019; De Strooper and Karran 2016). An emerging field of research proposes using MRI sequences such as diffusion-weighted imaging (DWI) toward investigating microstructural brain changes that may be sensitive to glial morphological changes. Using a DWI approach to assess cortical microstructural changes in the gray matter, a negative association was observed between cortical mean diffusivity and  $^{11}\text{C}$ -deuterium-L-deprenyl PET in autosomal dominant AD, and a positive association with cortical thickness, suggesting that astrocytosis may be linked to macrostructural and microstructural changes in the gray matter (Vilaplana et al. 2020). Further developments using longitudinal, multimodal imaging will contribute to characterizing the complex molecular and morphological changes associated with neuroinflammation in different brain diseases.

### 15.5.4 Neuroinflammation in Other Neurological Diseases

PET imaging of MAO-B can be used to investigate neuroinflammatory aspects in other neurological diseases. While neuroinflammation is believed to play an important role in diseases characterized by chronic pain, its underlying molecular mechanisms are not well-known. An *in vivo* PET study combining  $^{11}\text{C}$ -deuterium-L-deprenyl

and TSPO (translocator protein) tracer  $^{11}\text{C}$ -PBR28, a marker of microgliosis, sought to investigate neuroinflammatory mechanisms in fibromyalgia, a chronic disorder characterized by musculoskeletal pain, fatigue, and cognitive difficulties, which is still poorly understood (Albrecht et al. 2019). Fibromyalgia patients had elevated fronto-parietal  $^{11}\text{C}$ -PBR28 binding, but no difference in  $^{11}\text{C}$ -deuterium-L-deprenyl compared to a control group, therefore suggesting that microgliosis is the main mechanism driving neuroinflammation in this disease. These types of studies will be useful to clarify whether astrocytes and/or microglia are major players in disease initiation or progression and to design therapies aiming at the modulation of the relevant glial cell types and neuroinflammatory phenotypes in the disease of interest.

### 15.5.5 Depression

The monoamine hypothesis of depression proposes that the underlying pathophysiological mechanisms for depression are the deficits in monoamine neurotransmitters including dopamine, norepinephrine and serotonin, which are involved in arousal, mood, motivation, and reward (Pitsillou et al. 2020). Individuals with depression have high levels of MAO-A and MAO-B in the brain. While existing therapeutic strategies have aimed mostly at inhibiting MAO-A, there is a recent interest in exploring also MAO-B as a therapeutic target in depression. The  $^{11}\text{C}$ -SL25.1188 tracer was recently translated for in vivo human PET studies to investigate depression (Moriguchi et al. 2019). In this study, the MAO-B distribution volume ( $V_T$ ) was higher in a group of 20 patients with major depressive disorder compared to an age-matched group of controls, most prominently in the prefrontal cortices, which was also positively correlated with duration of illness; other cortical regions and the thalamus also had high MAO-B expression in patients. However, more PET imaging studies with tracers for MAO-B or other relevant targets would be needed to better characterize the neuroinflammatory underpinnings of depression.

### 15.5.6 Evaluation of Clinical Trials of MAO-B Inhibitors

MAO-B inhibition has been investigated as a potential therapeutic approach in neurodegenerative diseases including AD and PD (Schapira et al. 2006; Thomas 2000; Youdim et al. 2006). In the context of clinical trials, PET imaging of MAO-B is a promising tool to evaluate whether MAO-B inhibitors cross the blood-brain barrier, how they distribute in the brain, and for optimization of drug dosing and to quantify trial outcomes. In clinical trials, PET scans have been performed using  $^{11}\text{C}$ -labelled tracers, in which a scan can be repeated in the same individual before and after administration of the drug. Early studies used  $^{11}\text{C}$ -L-deprenyl PET to find the dose of reversible MAO-B inhibitor lazabemide (Ro 19-6327) that achieved >90% inhibition of MAO-B activity in normal subjects (Bench et al. 1991). This study also showed that brain MAO-B activity was correlated with platelet MAO-B

peripherally, suggesting the potential utility of platelet MAO-B activity as a marker to monitor trial outcomes. A study using serial  $^{11}\text{C}$ -L-deprenyl PET scans in the same individuals investigated the dynamic trajectory of MAO-B after withdrawal from treatment with MAO-B irreversible inhibitor L-deprenyl until recovery to baseline MAO-B levels (Fowler et al. 1994). The half-life of recovery to baseline levels of MAO-B was determined to be 40 days, and because the binding of L-deprenyl to MAO-B involves covalent modification of the enzyme, the 40-day half-life for recovery of brain MAO-B activity can be interpreted as a measure of how long it takes for the brain to synthesize new MAO-B molecules. From a therapeutic point of view, this slow turnover of MAO-B also indicates that a reduced dose of L-deprenyl could have a good therapeutic effect, which would avoid excessive drug use and potential associated side effects. A similar PET imaging study with  $^{11}\text{C}$ -L-deprenyl was used to evaluate the MAO-B inhibitory effect of irreversible inhibitor rasagiline in the brain, and this study also showed a gradual recovery of brain MAO-B activity after withdrawal from drug treatment (Freedman et al. 2005).

Irreversible MAO-B inhibitor drugs have shown certain limitations, such as that in some cases the effects were short-term. Therefore, reversible MAO-B inhibitors have been explored as potential treatment for PD and more recently AD (Borroni et al. 2017). A PET study with  $^{11}\text{C}$ -L-deprenyl- $\text{D}_2$  was used to find the dose of reversible MAO-B inhibitor EVT-301 that resulted in near complete occupancy of brain MAO-B (Hirvonen et al. 2009). More recently, a PET study with  $^{11}\text{C}$ -L-deprenyl- $\text{D}_2$  was applied to optimize the dose of reversible inhibitor sembragiline (also known as EVT-302) leading to the maximum inhibition of brain MAO-B (Sturm et al. 2017), as shown in Fig. 15.5. Novel reversible MAO-B inhibitors are being further investigated as potential treatments for AD (Park et al. 2019).

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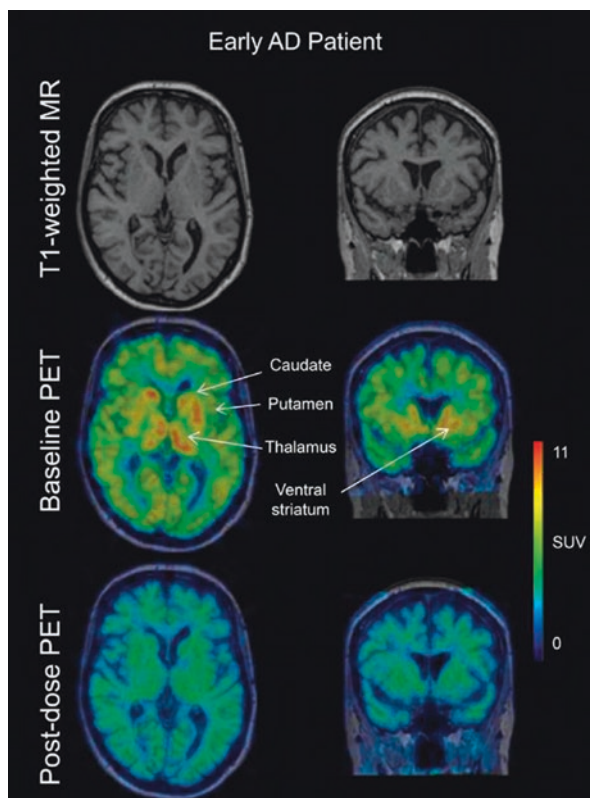
## 15.6 Conclusions and Future Prospects

PET studies of brain MAO-B in healthy volunteers and in patients with different brain disorders have evolved over the past ~30 years since the first MAO-B specific radiotracers were developed and investigated in preclinical and clinical studies. Different factors such as age, smoking, or depressive symptoms have been found to affect the levels of MAO-B in the brain. While the available radiotracers reviewed in this chapter have shown many properties that make them suitable for MAO-B quantification, radiochemists continue to develop novel radiotracers with improved properties, especially focusing on reversible radiotracers and more recently fluorine-18-labelled compounds for more widespread clinical application in centers without on-site cyclotron.

Many scientific questions remain to be investigated using *in vivo* and *in vitro* studies. Translational *in vitro* studies combining autoradiography and immunohistochemical analyses are critically needed to determine exactly what the MAO-B tracers are binding to and their relationship to other pathological markers such as amyloid- $\beta$  and tau. In particular, there is a paucity of antemortem/postmortem



**Fig. 15.5** Illustration of MAO-B inhibition in the brain following treatment with 1 mg sebragiline once a day for 2 weeks, as visualized using  $^{11}\text{C}$ -L-deprenyl- $\text{D}_2$  PET before treatment (baseline PET) and after treatment (post-dose PET). *SUV* standardized uptake value. Reprinted from Sturm et al. (2017), under the terms of the Creative Commons CC BY license



investigations, which would be highly informative to further validate the *in vivo* findings and understand what the MAO-B PET signal means biologically. Multimodal and multitracer MRI-PET imaging approaches have provided unique information on the comparative regional and temporal patterns of *in vivo* brain astrocytosis, amyloid- $\beta$  deposition, glucose metabolism, and brain structure. There are still very few *in vivo* longitudinal studies with MAO-B PET tracers, or to investigate the relationships between astrocytosis and microgliosis by either imaging or using glial markers in the plasma or CSF (e.g., GFAP as a biofluid marker of astrocytosis). Given the heterogeneity of the astrocytic response, there is a need to develop novel PET radiotracers specifically targeting the multiple biological changes associated with astrocytosis. Along these lines, future translational MRI-PET imaging investigations in combination with fluid biomarkers will be needed to advance the understanding of glial cells as therapeutic targets, with a view toward clinical applications.

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# Attempts to Image MRP1 Function in the Blood-Brain Barrier Using the Metabolite Extrusion Method

# 16

Toshimitsu Okamura, Tatsuya Kikuchi,  
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## Abstract

Multidrug resistance-associated protein 1 (MRP1) is a member of the adenosine triphosphate-binding cassette superfamily of transporters and plays an important role in limiting the permeation of xenobiotics and exporting endogenous sub-

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stances across the blood-brain and blood-cerebrospinal fluid barriers and across brain parenchymal cells in the central nervous system. In addition to multidrug resistance, changes in activity or expression of cerebral MRP1 occur in several brain diseases. Therefore, noninvasive and quantitative assessment of MRP1 activity in the brain is valuable for investigating the changes in MRP1 related to brain diseases, toward an understanding of the underlying molecular mechanisms. In this chapter, we will focus on 6-bromo-7- $^{[11}\text{C}]$ methylpurine for imaging MRP1 activity, as well as the concept of the metabolite extrusion method (MEM) for measuring efflux transporter activity. Several positron emission tomography (PET) and single-photon emission computed tomography (SPECT) imaging studies of MRPs activity are also reviewed. Finally, we discuss our recent findings regarding the contribution of Mrp1 in brain parenchymal cells to tracer efflux and the sensitivity of 6-bromo-7- $^{[11}\text{C}]$ methylpurine for measuring MRP1 activity.

## Abbreviations

$^{[18}\text{F}]$ FEP-4MA	<i>N</i> - $^{[18}\text{F}]$ Fluoroethylpiperidin-4ylmethyl acetate
15 <i>R</i> - $^{[11}\text{C}]$ TIC-Me	(15 <i>R</i> )-16- <i>m</i> - $^{[11}\text{C}]$ tolyl-17,18,19,20-tetranorisocarbacyclin methyl ester
$^{99\text{m}}\text{Tc}$ -ECD	<i>N,N'</i> -1,2-ethylenediylbis- <i>L</i> -cysteine diethyl ester
$^{99\text{m}}\text{Tc}$ -HIDA	$^{99\text{m}}\text{Tc}$ - <i>N</i> -(2,6-dimethylphenylcarbonylmethyl)-iminodiacetic acid
$^{99\text{m}}\text{Tc}$ -MIBI	$^{99\text{m}}\text{Tc}$ -hexakis-2-methoxyisobutylisonitrile
ABC	Adenosine triphosphate-binding cassette
BBB	Blood-brain barrier
BCECs	Brain capillary endothelial cells
BEI	Brain efflux index
GSH	Glutathione
GSTs	Glutathione <i>S</i> -transferases
ICI	Intracerebral injection
ID	Injected dose
LT	Leukotriene
MDR	Multidrug resistance
MEM	Metabolite extrusion method
MRP	Multidrug resistance-associated protein
OAT3	Organic anion transporter 3
PET	Positron emission tomography
SPECT	Single-photon emission computed tomography
WT	Wild type



## 16.1 Introduction

Multidrug resistance-associated protein 1 (MRP1), also denoted as ABCC1, is a member of the adenosine triphosphate-binding cassette (ABC) superfamily of transporters and transports a wide range of compounds, including xenobiotics and physiological substances, from cells in an energy-dependent manner (Bakos and Homolya 2007; Cole et al. 1992). The significance of MRP1 in human health and disease has been accepted, and the preclinical and clinical interest has increased, primarily because of its extraordinary transport properties. In the central nervous system, MRP1 is expressed at the blood-brain barrier (BBB) and blood-cerebrospinal fluid barrier (Dallas et al. 2006). In addition to multidrug resistance (MDR) (Loscher and Potschka 2005; Leslie et al. 2005), changes in activity or expression of cerebral MRP1 occur in several brain diseases (Qosa et al. 2015; Krohn et al. 2015; Krohn et al. 2011). In the adult and the developing human brain, MRP1 is also expressed in nonendothelial cells such as astrocytes and neurons (Bernstein et al. 2014; Daood et al. 2008).

Furthermore, a recent meta-analysis suggested that MRP1 is overexpressed in neurons and astrocytes of intractable epilepsy patients (Sun et al. 2016). Moreover, MRP1 expression was increased in samples from traumatic brain injury patients compared with controls, while cells with the morphological appearance of neurons or astrocytes are MRP1-immunoreactive (Willyerd et al. 2016). Therefore, noninvasive and quantitative assessment of MRP1 activity in the brain is valuable for investigating changes in MRP1 related to brain diseases, toward an understanding of the underlying molecular mechanisms.

Here, we will focus on 6-bromo-7- $^{[11C]}$ methylpurine for imaging of MRP1 activity and the concept of the metabolite extrusion method (MEM) for measuring the activity of an efflux transporter. Several studies which imaged the activity of MRPs, including MRP1, MRP2, and MRP4, using PET and SPECT are also reviewed. Finally, we discuss our recent findings suggesting that Mrp1 in brain parenchymal cells, but not at the BBB, may contribute to the efflux of the glutathione (GSH) conjugate of 6-bromo-7- $^{[11C]}$ methylpurine from the brain and the sensitivity of 6-bromo-7- $^{[11C]}$ methylpurine for measuring MRP1 activity.

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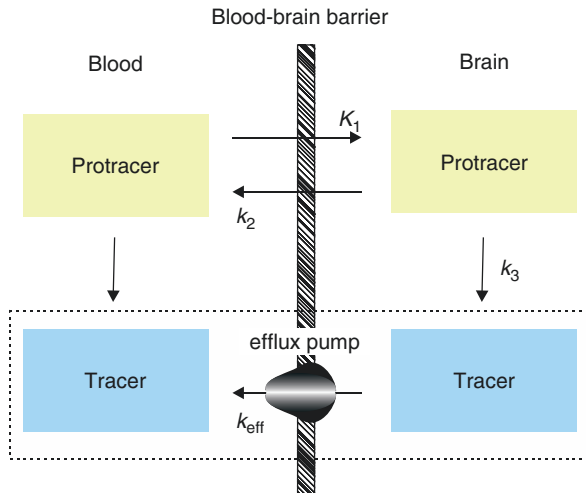
## 16.2 Metabolite Extrusion Method (MEM)

### 16.2.1 Concept of the MEM

The blood-brain barrier (BBB) consists of a monolayer of brain capillary endothelial cells (BCECs) and limits the entry of large or hydrophilic substances into the brain, except for nutrients such as amino acids and glucose (Pardridge 1998; Bradbury 1993; Reese and Karnovsky 1967). For assessment of efflux transporter activity, it would thus be challenging to deliver radiolabeled hydrophilic substrates into the brain via intravenous administration. One of the methods to overcome this problem is intracerebral injection (Cserr and Berman 1978) or the brain efflux index

(BEI) (Kakee et al. 1996; Kusuhara et al. 2003). In the BEI, a radiolabeled substrate is microinjected into the brain, and elimination of the substrate from the brain is measured over time. The efflux rate can then be determined from the slope of a semilogarithmic plot of the percentage of the substrate remaining in the brain against time. Apart from the disadvantage of invasiveness, the BEI provides an ideal situation where the decrease in brain radioactivity depends on the clearance of a single radioactive substrate.

The MEM is a method for the noninvasive measurement of efflux transporter activity (not limited to MRP1) and provides the same situation as the BEI without intracerebral injection (Kikuchi et al. 2016; Okamura et al. 2010; Okamura et al. 2009a; Okamura et al. 2007). In this chapter, we use the term “protracer” to denote the radiolabeled lipophilic compound for the delivery of a radiolabeled hydrophilic substrate for an efflux transporter into the brain, while we use the term “tracer” to denote a radiolabeled hydrophilic substrate resulting from the brain metabolism of the protracer, where the tracer provides information regarding the activity of an efflux transporter. To clearly distinguish an MEM tracer from a non-MEM tracer such as [ $^{11}\text{C}$ ]verapamil or [ $^{11}\text{C}$ ]loperamide, we also use the term “probe” to denote a non-MEM radiolabeled compound for imaging an efflux transporter. The schematic explanation for the MEM is shown in Fig. 16.1. A protracer in the blood enters the brain through the BBB after intravenous administration. The protracer in the brain diffuses back into the blood and also becomes converted into a tracer that



**Fig. 16.1** Diagrammatic representation of the metabolite extrusion method (MEM). The terms of “protracer” and “tracer” are defined in Sect. 16.2.1. A protracer readily enters the brain after intravenous administration and undergoes rapid conversion into a tracer. After the protracer has been completely converted into the tracer, the time course of radioactivity in the brain can be considered one-tissue compartment model inside the dotted box.  $K_1$  and  $k_2$  represent the transport rate constants of the protracer across the BBB, while  $k_3$  denotes the conversion rate of the protracer into the tracer, and  $k_{\text{eff}}$  shows the efflux rate of the tracer from the brain to the blood

is then eliminated by an efflux transporter of interest. The decreased brain radioactivity after the disappearance of the protracer can be considered the elimination of the tracer from the brain. Thus, the efflux rate ( $k_{\text{eff}}$ ) of the tracer can be noninvasively determined from the time course of brain radioactivity.

In the MEM, the conversion rate of a protracer to a tracer (hydrophilic substrate) is crucial for measuring the activity of an efflux transporter of interest. If the conversion rate is slow, the complete disappearance of the protracer is delayed, resulting in the low concentration of the tracer in the brain. These issues may compromise the accurate measurement of the efflux rate.

### 16.2.2 Design of a Protracer and Tracer for Imaging MRP1 Activity

To obtain an accurate measurement of  $k_{\text{eff}}$  by the MEM, a protracer should meet the following criteria: (1) high extraction to the brain tissue from the blood and (2) rapid conversion into a tracer ( $k_2 \ll k_3$ ). Moreover, an easily modified protracer platform is desirable for introducing a lipophilic group, radiolabeling, and adjusting the conversion rate for application to different species including humans. In addition, if the tracer satisfies the kinetic condition of  $k_{\text{eff}} \ll (k_2 + k_3)$  (see below), the MEM provides a monoexponential curve of the tracer after a short lag time, thereby allowing the simple measurement of  $k_{\text{eff}}$ .

In the two-tissue compartment model, the measured brain concentration ( $Cb$ ) is equal to the integral of the arterial plasma input curve (input function) convolved with a term called the tissue impulse response function ( $R$ ):

$$Cb(t) = \text{Input Function} \otimes R(t)$$

In the MEM, however, an efflux rate is determined from the time-activity curve following the conversion of a protracer to a tracer, and the input function can thus be considered a constant value. As described in a previous review (Okamura et al. 2010), the response function ( $R$ ) with respect to time ( $t$ ) is given by:

$$R(t) = Ae^{-(k_2+k_3)t} + Be^{-k_{\text{eff}}t}$$

$$A = \frac{k_2 - k_{\text{eff}}}{k_2 + k_3 - k_{\text{eff}}}, \quad B = \frac{k_3}{k_2 + k_3 - k_{\text{eff}}}$$

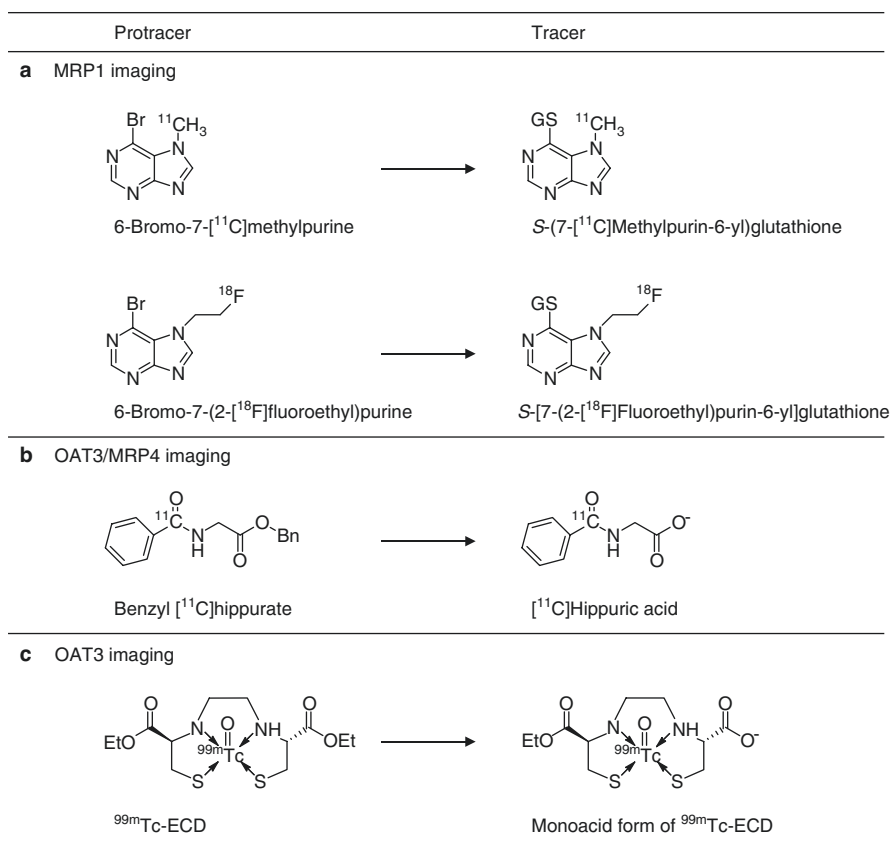
where  $k_2$  and  $k_3$  are the rate parameters describing the diffusional transport of a protracer through the BBB and the conversion of a protracer to a tracer, respectively, and  $k_{\text{eff}}$  (an index of the activity of an efflux transporter) is the parameter describing the rate of the tracer clearance from the brain by an efflux transporter. The condition  $k_{\text{eff}} \ll (k_2 + k_3)$  is required to obtain a monoexponential curve of a tracer after a short lag time.

Based on these requirements, we describe the design of a protracer and tracer for imaging of MRP1 activity. The substrates of MRP1 comprise many lipophilic compounds, conjugated organic anions, and nonconjugated organic

anions, and typical substrates include GSH, glucuronate, and sulfate conjugates (Bakos and Homolya 2007). Initially, we selected GSH conjugates as a candidate tracer, although glucuronide or sulfate conjugates may be candidates. According to the requirements mentioned above, a protracer should be efficiently converted to a tracer (a GSH conjugate). Since 6-chloropurine is converted to *S*-(6-purinyl)glutathione by hepatic and renal glutathione *S*-transferases (GSTs) (Hwang and Elfarra 1993), 6-chloropurine was selected as the basic skeleton of the protracer. Then, 6-chloropurine was methylated with [<sup>14</sup>C]methyl iodide, providing the two candidate protracers, 6-chloro-9-[<sup>14</sup>C]methylpurine and 6-chloro-7-[<sup>14</sup>C]methylpurine. The reaction rate (in vitro  $k_3$ ) of 6-chloro-7-[<sup>14</sup>C]methylpurine with GSH in rat brain homogenates was much higher than that of 6-chloro-9-[<sup>14</sup>C]methylpurine (Okamura et al. 2007). The difference in the reaction rate between these candidate protracers unambiguously accounted for a fraction of each protracer remaining in the in vivo brain. 6-Chloro-9-[<sup>14</sup>C]methylpurine did not satisfy the requirements as a protracer, whereas 6-chloro-7-[<sup>14</sup>C]methylpurine was a promising protracer. The GSH conjugate of 6-chloro-7-methylpurine, *S*-(7-methylpurin-6-yl)glutathione, was shown to be a substrate for MRP1 in vitro (Okamura et al. 2007). Replacement of chlorine by bromine in the 6-chloro-7-[<sup>14</sup>C]methylpurine increased the reaction rate with GSH in mouse brain homogenates (Okamura et al. 2009b). We thus evaluated 6-bromo-7-methylpurine as a protracer for imaging of MRP1 activity in the brain (Okamura et al. 2009c). The structure of 6-bromo-7-[<sup>11</sup>C]methylpurine is shown in Fig. 16.2.

### 16.2.3 Imaging of MRP1 Activity in the Brain Using 6-Bromo-7-[<sup>11</sup>C]Methylpurine

The protracer 6-bromo-7-[<sup>11</sup>C]methylpurine showed high uptake in the brain of wild-type (WT) and *Mrp1*<sup>-/-</sup> mice 1.5 min after intravenous administration (approximately 6% injected dose/g) and then a rapid decrease in the brain radioactivity. After intravenous administration, 6-bromo-7-[<sup>14</sup>C]methylpurine was converted to its GSH conjugate in the brains of WT and *Mrp1*<sup>-/-</sup> mice by 15 min, and therefore, the washout rate of the brain radioactivity from 15 to 60 min after the injection of 6-bromo-7-[<sup>11</sup>C]methylpurine reflects the efflux rate of *S*-(7-[<sup>11</sup>C]methylpurin-6-yl)glutathione from the brain. In addition, the efflux rate was approximately tenfold lower in *Mrp1*<sup>-/-</sup> (0.15 h<sup>-1</sup> or 0.0025 min<sup>-1</sup>) than WT mice (1.4 h<sup>-1</sup> or 0.023 min<sup>-1</sup>) (Okamura et al. 2009c). Moreover, the influx of *S*-(7-[<sup>11</sup>C]methylpurin-6-yl)glutathione from the blood to the brain in WT and *Mrp1*<sup>-/-</sup> mice was extremely low for 60 min after intravenous injection, indicating that this tracer in the blood cannot cross the BBB (Okamura et al. 2020). Thus, 6-bromo-7-[<sup>11</sup>C]methylpurine allows noninvasive and quantitative measurement of MRP1 activity in the brain. However, recent findings showed that multiple efflux transporters contribute to the efflux of the GSH conjugate from the mouse brain. We will address this in Sect. 16.4.



**Fig. 16.2** Structures of MEM tracers. **(a)** The efflux rate of *S*-(7-[<sup>11</sup>C]methylpurin-6-yl)glutathione from the brain after intravenous administration of 6-bromo-7-[<sup>11</sup>C]methylpurine was reduced to nearly zero in *Mrp1*<sup>-/-</sup> mice compared with WT mice, and thus the protracer allows noninvasive and quantitative measurement of *Mrp1* activity in the brain. However, recent studies show that the efflux process of *S*-(7-[<sup>11</sup>C]methylpurin-6-yl)glutathione from the brain via *Mrp1* would not be the rate-limiting step in mice and that *Mrp4* and *Oat3* contribute to the efflux of the tracer (see Sect. 16.4). The <sup>18</sup>F-labeled derivative may be transported from the brain by the same mechanisms as the <sup>11</sup>C-labeled tracer. GS denotes a glutathionyl group. **(b)** Benzyl [<sup>11</sup>C]hippurate is a potential protracer for imaging the activities of *Oat3* in the brains and *Mrp4* in the hearts of mice. **(c)** <sup>99m</sup>Tc-ECD can be used as a protracer for imaging of *Oat3* activity in the mouse brain, but not in primates, including humans

### 16.2.4 MRP1 Imaging in Peripheral Tissues Using 6-Bromo-7-[<sup>11</sup>C]Methylpurine

MRP1 is ubiquitously expressed in normal tissues, including the brain, lung, heart, and kidney (Bakos and Homolya 2007; Flens et al. 1996). The change of MRP1 expression is associated with certain lung diseases (van der Deen et al. 2006), and the noninvasive measurement of pulmonary MRP1 activity would thus help

elucidate pathological conditions and diagnoses of lung diseases. We examined the feasibility of imaging pulmonary MRP1 activity using 6-bromo-7- $^{11}\text{C}$ methylpurine (Okamura et al. 2013). The radioactivity in the lungs of WT and *Mrp1* $^{-/-}$  mice reached a maximum level immediately after the intravenous administration of 6-bromo-7- $^{11}\text{C}$ methylpurine. Thereafter, the radioactivity in the lungs of WT mice was rapidly reduced, whereas in *Mrp1* $^{-/-}$  mice, the radioactivity was mostly retained. In addition, radioactive compounds in the lungs of *Mrp1* $^{-/-}$  mice 5 min after the intravenous administration of 6-bromo-7- $^{11}\text{C}$ methylpurine almost entirely consisted of *S*-(7-methylpurin-6-yl)glutathione (MRP1 substrate), although 27% of the radioactivity in the lungs of WT mice was present as an unknown metabolite. Thus, this protracer showed a kinetic behavior according to the MEM model: 6-bromo-7- $^{11}\text{C}$ methylpurine diffuses into the lung tissue after administration and undergoes conversion into the tracer, its GSH conjugate, which is then specifically effluxed by Mrp1. This finding suggests that 6-bromo-7- $^{11}\text{C}$ methylpurine can be used to measure MRP1 activity in the lungs (Okamura et al. 2013).

Zoufal et al. have recently investigated whether the activity of MRPs in excretory organs can be measured with 6-bromo-7- $^{11}\text{C}$ methylpurine (Zoufal et al. 2019). The elimination rate constant of 6-bromo-7- $^{11}\text{C}$ methylpurine-derived radioactivity was significantly decreased in the kidneys of *Mrp1* $^{-/-}$  mice compared with WT mice. Loss of *Mrp4* $^{-/-}$  also significantly decreased the elimination rate constant in the kidney. These results suggest that 6-bromo-7- $^{11}\text{C}$ methylpurine may be useful for assessing the activity of MRPs in the kidneys. However, further studies are required to identify the MRP subtypes involved in the renal excretion.

MRP1 is highly expressed in the heart (Flens et al. 1996), and it has been shown that MRP1 polymorphisms are related to anthracycline-induced cardiotoxicity in cancer patients (Semsei et al. 2012; Wojnowski et al. 2005). To our knowledge, thus far, no studies are available regarding PET tracers or probes for the measurement of MRP1 activity in the heart. However, based on previous reports, we would like to suggest the possibility of MRP1 imaging in the heart using 6-bromo-7- $^{11}\text{C}$ methylpurine. Zoufal et al. investigated whether the activity of MRPs in excretory organs can be measured with 6-bromo-7- $^{11}\text{C}$ methylpurine (Zoufal et al. 2019). In their study, a region of interest was drawn around the left ventricle of the heart of mice to obtain image-derived blood concentration-time curves. Similar to the kinetics in the lungs, the radioactivity clearance rate in the blood of *Mrp1* $^{-/-}$  mice, which was determined by regression analysis from 17.5 to 80 min, was significantly decreased compared with that of WT mice (Zoufal et al. 2019). In contrast to the image-derived concentration-time curves in the blood, we examined the time course of radioactivity in the blood collected from the stump following decapitation (Okamura et al. 2013). Consequently, the radioactivity in the blood of *Mrp1* $^{-/-}$  mice was 1.4% injected dose (ID)/g at 1 min, 1.1% ID/g at 15 min, and 0.84% ID/g at 60 min, while that of WT mice was 5.8% ID/g at 1 min, 1.1% ID/g at 15 min, and 0.27% ID/g at 60 min. The time course of radioactivity in the blood was inconsistent between the collected blood and the image-derived blood. Given that MRP1 is highly expressed in the heart (Flens et al. 1996), we believe that the elimination rate of the radioactivity in the image-derived blood of *Mrp1* $^{-/-}$  mice would be strongly affected by the

loss of Mrp1 in the heart tissue, hinting a possibility of MRP1 imaging in the heart using 6-bromo-7-[ $^{11}\text{C}$ ]methylpurine. However, further studies in a large animal model are needed.

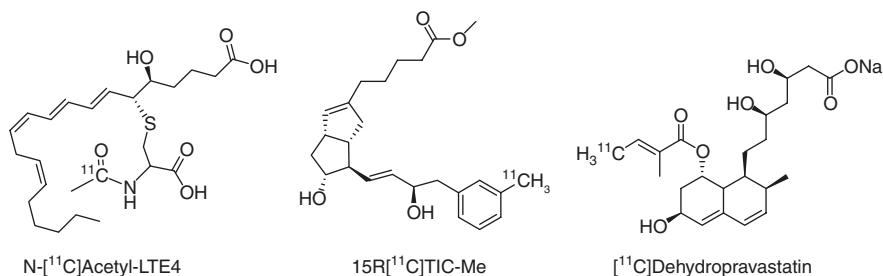
### 16.3 Other Potential Tracers or Probes for the Imaging of MRP1, MRP2, and MRP4

Except for 6-bromo-7-[ $^{11}\text{C}$ ]methylpurine, few reports are available regarding PET protracters or probes for imaging of MRP1 activity. Since  $^{18}\text{F}$  has a longer half-life than  $^{11}\text{C}$  (110 vs. 20 min), a  $^{18}\text{F}$ -labeled derivative of 6-bromo-7-[ $^{11}\text{C}$ ]methylpurine has advantages over 6-bromo-7-[ $^{11}\text{C}$ ]methylpurine such as better image quality and applications in species that show lower rates of GSH conjugation. Thus, a series of  $^{18}\text{F}$ -labeled 6-halopurines were synthesized and evaluated as a protracter for imaging of MRP1 activity in the brain. As a consequence, 6-bromo-7-(2-[ $^{18}\text{F}$ ]fluoroethyl) purine (Fig. 16.2) was suggested to be a promising protracter for noninvasive measurement of MRP1 activity in the brain (Galante et al. 2014).

The SPECT probes  $^{99\text{m}}\text{Tc}$ -hexakis-2-methoxyisobutylisonitrile ( $^{99\text{m}}\text{Tc}$ -MIBI) and  $^{99\text{m}}\text{Tc}$ -tetrofosmin, which function as myocardial perfusion imaging agents (Higley et al. 1993; Wackers et al. 1989), have been suggested to serve as substrates for MRP1 (Li et al. 2001; Hendrikse et al. 1998). Therefore, the SPECT probes may be used for visualization of MRP1 activity in a tissue of interest. For  $^{99\text{m}}\text{Tc}$ -MIBI, the elimination rate from the lungs corresponded to the alteration of MRP1 expression: the rate of inhaled  $^{99\text{m}}\text{Tc}$ -MIBI from the lungs in smokers with high levels of MRP1 expression was slower compared with that in nonsmokers with low levels of MRP1 expression (Mohan et al. 2019; Mohan et al. 2016). However, the use of  $^{99\text{m}}\text{Tc}$ -MIBI for MRP1 imaging in the brain would be difficult because the positive charge of  $^{99\text{m}}\text{Tc}$ -MIBI is unfavorable for the ingress into the brain. It should also be noted that  $^{99\text{m}}\text{Tc}$ -MIBI and  $^{99\text{m}}\text{Tc}$ -tetrofosmin are substrates for the ABC transporter P-glycoprotein (Ballinger et al. 1996; Piwnica-Worms et al. 1993).

A series of flavone derivatives were synthesized and evaluated as probes to visualize MRP1 (Mavel et al. 2006). In vitro experiments using Pgp- or MRP1-overexpressing cell lines showed these compounds to be selective for MRP1-expressing cells and enhanced the sensitivity of MRP1-mediated MDR cell lines to doxorubicin toxicity, suggesting that halogen-substituted flavone derivatives are potential PET and SPECT probes for imaging MRP1-mediated MDR phenomena. However, to date, the radiolabeling for PET or SPECT imaging or the in vivo evaluation of these candidates has not been reported.

MRP2 mainly mediates the hepatobiliary transport of many drugs and endogenous conjugated metabolites, and MRP2 deficiency leads to Dubin-Johnson syndrome, an inherited disorder characterized by conjugated hyperbilirubinemia (Nies and Keppler 2007). Compared with MRP1, there are relatively many studies focusing on the imaging of MRP2 in the liver. Hendrikse et al. examined the substrate specificities of  $^{99\text{m}}\text{Tc}$ -*N*-(2,6-dimethylphenylcarbamoylemethyl)-iminodiacetic acid ( $^{99\text{m}}\text{Tc}$ -HIDA), a hepatobiliary agent (Wistow et al. 1977), for MRP1, MRP2, and



**Fig. 16.3** Structures of potential PET probes for MRP2 imaging

P-glycoprotein and examined the kinetics of <sup>99m</sup>Tc-HIDA in normal and Mrp2-deficient GY/TR<sup>-</sup> rats (Hendrikse et al. 2004). The results showed that <sup>99m</sup>Tc-HIDA was transported by MRP1 and MRP2, but not by P-glycoprotein, and the elimination rate for radioactivity from the liver of control rats (half-life: 7 min) was faster than that of GY/TR<sup>-</sup> rats (half-life: 40 min). Thus, <sup>99m</sup>Tc-HIDA might be useful for assessing the in vivo activity of MRP2 in the liver.

The structures of the potential PET probes for imaging MRP2 are displayed in Fig. 16.3. It has been shown that the inflammatory cytokine leukotriene C<sub>4</sub> (LTC<sub>4</sub>) and its metabolites, LTD<sub>4</sub> and LTE<sub>4</sub>, are substrates for both MRP1 and MRP2 (Ito et al. 2001; Leier et al. 1994). The leukotriene derivatives labeled with a positron emitter could thus be a potential probe for assessing MRP1 or MRP2 activity. Furthermore, *N*-[<sup>11</sup>C]acetyl-LTE<sub>4</sub> was synthesized, and its kinetic behavior was investigated in normal rats, in cholestatic rats, and in Mrp2-deficient GY/TR<sup>-</sup> rats using PET (Hendrikse 2000; Guhlmann et al. 1995). After injection of *N*-[<sup>11</sup>C]acetyl-LTE<sub>4</sub>, a fast clearance from the blood and a transient accumulation in the liver were observed in normal rats, whereas *N*-[<sup>11</sup>C]acetyl-LTE<sub>4</sub> showed a delayed washout from the blood in cholestatic and GY/TR<sup>-</sup> rats. Moreover, there was a delayed decrease in hepatic *N*-[<sup>11</sup>C]acetyl-LTE<sub>4</sub> and negligible amounts of radioactivity in the intestine in GY/TR<sup>-</sup> rats, compared with normal rats. The findings imply that *N*-[<sup>11</sup>C]acetyl-LTE<sub>4</sub> might enable the noninvasive assessment of Mrp2-mediated transport in the liver.

The probe (15*R*)-16-*m*-[<sup>11</sup>C]tolyl-17,18,19,20-tetranorisocarbacyclin methyl ester (15*R*-[<sup>11</sup>C]TIC-Me) was converted in the blood into its acid form, 15*R*-[<sup>11</sup>C]TIC, which was, in turn, converted to at least three metabolites (M1, M2, and M3) (Takashima et al. 2010). An in vitro transport assay using Mrp2-expressing membrane vesicles suggested that the major metabolites (M2 in the plasma and M3 in the bile) were good substrates for Mrp2. Moreover, the intrinsic canalicular efflux clearance of M2 and M3 in Mrp2-deficient rats was reduced compared with controls. Thus, the authors of this study conclude that 15*R*-[<sup>11</sup>C]TIC-Me might be useful for in vivo analyses of Mrp2-mediated hepatobiliary transport.

A recent study suggested that [<sup>11</sup>C]dehydropravastatin can assess the hepatobiliary transport by MRP2 and organic anion-transporting polypeptides in rats and



humans (Kaneko et al. 2018; Shingaki et al. 2013). Rifampicin-treated and MRP2-defective rats decreased the canalicular efflux clearance of [ $^{14}\text{C}$ ]dehydropravastatin (Shingaki et al. 2013). Similarly, the decrease in the canalicular efflux clearance in rifampicin-treated subjects was observed compared with that in control subjects, and the canalicular efflux clearance in a patient with Dubin-Johnson syndrome, caused by hereditary MRP2 deficiency, was also lower than that in control subjects (Kaneko et al. 2018).

MRP4 is thought to play a crucial role in homeostasis since this protein contributes to the transport of several important endogenous substances such as cAMP, cGMP, ADP, prostaglandins, and leukotrienes (Wen et al. 2015), whose alteration is probably responsible for pathophysiological processes. In addition, MRP4 controls cAMP levels in cardiac myocytes by extruding cAMP (Sassi et al. 2012). Noninvasive measurement of MRP4 activity may thus be useful for the understanding of the relationship between MRP4 and certain diseases. To our knowledge, there is only one study that investigated the feasibility of imaging MRP4 activity in vivo using PET (Kikuchi et al. 2016). Benzyl [ $^{14}\text{C}$ ]hippurate is a potential protracer for estimating the activities of MRP4 in the mouse heart and organic anion transporter 3 (OAT3) in the brain (see also Sect. 16.5), and the structure of the protracer is shown in Fig. 16.2. Benzyl [ $^{14}\text{C}$ ]hippurate is converted into [ $^{14}\text{C}$ ]hippuric acid, which is thought to be a substrate of MRP4 (Mutsaers et al. 2011), in the heart. The efflux rate of the [ $^{14}\text{C}$ ]hippuric acid formed in the heart of *Mrp4*<sup>-/-</sup> mice (0.043 min<sup>-1</sup>) after administration of benzyl [ $^{14}\text{C}$ ]hippurate was approximately 60% lower than that of control mice (0.11 min<sup>-1</sup>). The protracer benzyl [ $^{14}\text{C}$ ]hippurate might thus be used to assess MRP4 activity in the mouse heart, although further studies are needed for evaluating MRP4 specificity for [ $^{14}\text{C}$ ]hippuric acid elimination from the heart (Kikuchi et al. 2016).

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## 16.4 Feasibility of MRP1 Imaging in the Blood-Brain Barrier

### 16.4.1 Type of Protracer

As shown in Fig. 16.1, the MEM is described by the simple compartment model, and thus the activity of an efflux transporter at the BBB appears to be assessed from the brain kinetics of a tracer. This would be true for a protracer that can be metabolized to a tracer only in BCECs because a single efflux transporter at the luminal side of BCECs contributes to the efflux of the tracer to the blood. However, the brain has different kinds of cells such as BCECs, neurons, and glial cells, and thus, a protracer that is metabolized in these cells would not always provide information of the activity of an efflux transporter at the BBB. Unfortunately, a protracer that can be metabolized only in BCECs has not been developed yet.

Since 6-bromo-7-[ $^{14}\text{C}$ ]methylpurine can easily enter the brain because of its hydrophobicity, the protracer should be able to diffuse into not only BCECs but also brain parenchymal cells, including neurons and nonneuronal cells. Moreover, GSTs, which catalyze GSH conjugation, are found in neurons, glial cells, and

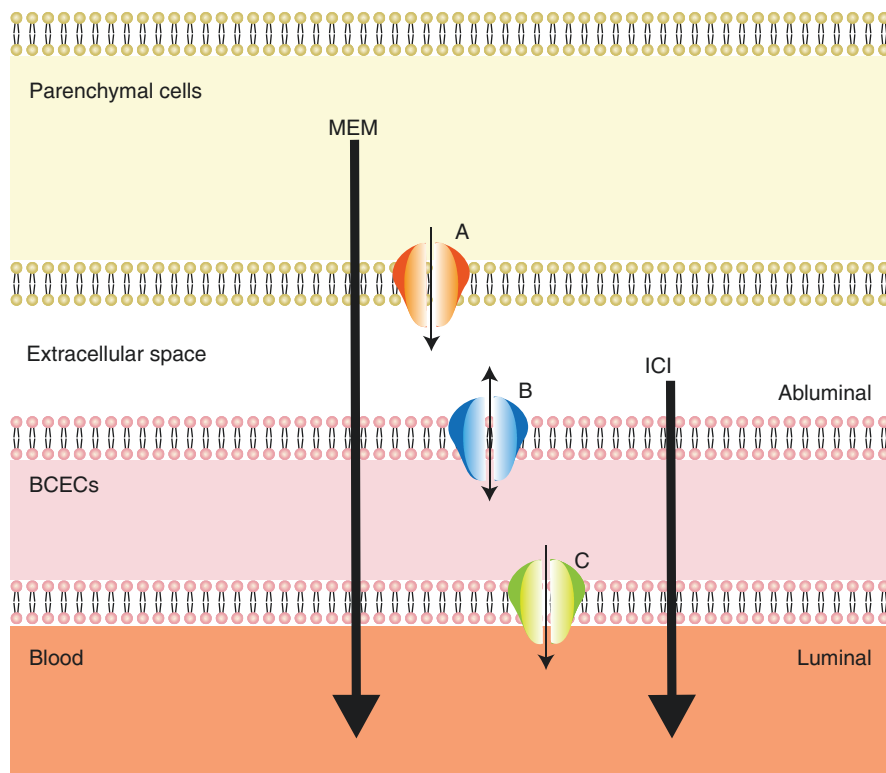
endothelial cells in the rodent brain (Beiswanger et al. 1995; Johnson et al. 1993; Cammer et al. 1989). Therefore, *S*-(7-[<sup>11</sup>C]methylpurin-6-yl)glutathione would be produced within brain parenchymal cells as well as BCECs. The tracer, *S*-(7-[<sup>11</sup>C]methylpurin-6-yl)glutathione, formed in brain parenchymal cells, must penetrate three membranes to be eliminated from the brain to the blood. In other words, at least three transporters, an efflux transporter at the luminal side (blood side), an uptake transporter at the abluminal side (brain side) of BCECs, and an efflux transporter in brain parenchymal cells, are required for the efflux of *S*-(7-[<sup>11</sup>C]methylpurin-6-yl)glutathione from brain parenchymal cells to the blood through the BBB (BCECs).

#### 16.4.2 MRP1 at the Blood-Brain Barrier Versus Brain Parenchymal Cells

In the central nervous system, MRP1 is expressed not only at the BBB and blood-cerebrospinal fluid barrier (Dallas et al. 2006) but also in nonendothelial cells such as astrocytes and neurons in the adult and developing human brain (Bernstein et al. 2014; Daood et al. 2008). As described above, 6-bromo-7-[<sup>11</sup>C]methylpurine allows noninvasive and quantitative measurement of MRP1 activity in the brain. However, the protracer cannot discriminate between the washout of *S*-(7-[<sup>11</sup>C]methylpurin-6-yl)glutathione from brain parenchymal cells and through the BBB (Fig. 16.4); we thus investigated the involvement of MRP1 at the BBB in the tracer efflux using the intracerebral injection method, which is similar to the BEI for analyzing efflux at the BBB (Kakee et al. 1996). In our recent study, no significant differences in the efflux rate of *S*-(7-[<sup>11</sup>C]methylpurin-6-yl)glutathione was observed between WT and *Mrp1*<sup>-/-</sup> mice after intracerebral injection (Okamura et al. 2020). This result shows that intracerebrally injected *S*-(7-[<sup>11</sup>C]methylpurin-6-yl)glutathione was cleared from the brain, presumably through the BBB, but the tracer efflux was not mediated by *Mrp1*. By contrast, the tracer produced in the brain following administration of 6-bromo-7-[<sup>11</sup>C]methylpurine was almost completely trapped in the brain of *Mrp1*<sup>-/-</sup> mice (Okamura et al. 2020). Therefore, *Mrp1* in brain parenchymal cells, but not the BBB, may contribute to the extrusion of *S*-(7-[<sup>11</sup>C]methylpurin-6-yl)glutathione from the brain to the blood.

#### 16.4.3 Involvement of MRP4 and OAT3 in the Efflux of *S*-(7-[<sup>11</sup>C]Methylpurin-6-yl)Glutathione

In addition to *Mrp1*, *Mrp4* and *Oat3* contribute to the efflux of *S*-(7-[<sup>11</sup>C]methylpurin-6-yl)glutathione produced in the brain after intravenous administration of 6-bromo-7-[<sup>11</sup>C]methylpurine (Okamura et al. 2020). In mice, *Mrp4* and *Oat3* are localized at the luminal and abluminal membrane of BCECs, respectively (Ohtsuki et al. 2004; Leggas et al. 2004). It has also been reported



**Fig. 16.4** Putative mechanism for the efflux of *S*-(7-[ $^{11}\text{C}$ ]methylpurin-6-yl)glutathione from the brain. In the MEM, *S*-(7-[ $^{11}\text{C}$ ]methylpurin-6-yl)glutathione would be formed within glial cells, neurons, and BCECs. *S*-(7-[ $^{11}\text{C}$ ]methylpurin-6-yl)glutathione produced within parenchymal cells is therefore effluxed to the extracellular space by transporter A and then is taken up into BCECs by transporter B. Finally, the tracer is extruded into the blood by transporter C at the luminal side of BCECs. A recent study using the MEM and the intracerebral injection (ICI) method, which is similar to the BEI for analyzing efflux at the BBB, suggests that transporters A, B, and C contributing to the tracer efflux from the mouse brain are Mrp1, Oat3, and Mrp4, respectively. Note that this figure shows only three transporters; however, we cannot exclude the possibility that other transporters mediate the transport of *S*-(7-[ $^{11}\text{C}$ ]methylpurin-6-yl)glutathione from the brain

that the interplay between Oat3 and Mrp4 at the BBB may allow vectorial transport of prostaglandins, fatty acids derived from arachidonic acid metabolism, from the brain interstitial fluid to the blood across BCECs (Tachikawa et al. 2014); that is, Oat3 takes up prostaglandins from the interstitial fluid into BCECs, and subsequently, Mrp4 eliminates them from BCECs into the blood. Based on this observation, *S*-(7-[ $^{11}\text{C}$ ]methylpurin-6-yl)glutathione in the extracellular space may be incorporated into BCECs by Oat3 and then effluxed from the cells to the blood by Mrp4 to be cleared from the brain. Nonetheless, we cannot rule out the possibility that other pumps are involved in the transport of the tracer into and out of BCECs.

#### 16.4.4 The Efflux Process Via Mrp1 Is Not the Rate-Limiting Step in the Mouse Brain

6-Bromo-7- $^{11}\text{C}$ methylpurine allows the assessment of a transporter either at the luminal side or the abluminal side of the BBB, or in brain parenchymal cells, probably Mrp4, Oat3, or Mrp1. The determined efflux rate of *S*-(7- $^{11}\text{C}$ methylpurin-6-yl) glutathione would reflect the slowest step of the efflux processes, i.e., the rate-limiting step. Unfortunately, no methods are available for determining the rate-limiting step of the efflux process in the brain. However, in control mice, we found that the efflux rate of intracerebrally injected *S*-(7- $^{11}\text{C}$ methylpurin-6-yl)glutathione was comparable to that of the tracer formed in the brain after intravenous injection of 6-bromo-7- $^{11}\text{C}$ methylpurine (Okamura et al. 2020). This finding shows that the efflux process of the tracer via Mrp1 from brain parenchymal cells to the extracellular space would not be the rate-limiting step, but either the uptake or efflux process at the BBB, partly via Oat3 and Mrp4, is the rate-limiting step. As a result, it is difficult to assess Mrp1 activity in the mouse brain using 6-bromo-7- $^{11}\text{C}$ methylpurine, unless Mrp1 activity markedly changes, such as in gene knockout animals. After the protracer injection, there was only a moderate reduction in the washout rate of brain radioactivity in heterozygous *Mrp1*<sup>+/-</sup> mice, which are expected to have a 50% reduction in Mrp1 expression, compared with WT mice (Zoufal et al. 2019). This finding implies that the transport system is very high capacity, e.g., possesses a high ratio of maximal transport rate to the Michaelis-Menten constant.

6-Bromo-7- $^{11}\text{C}$ methylpurine was shown to possess limited sensitivity for measuring moderate changes in MRP1 activity in the mouse brain (Zoufal et al. 2019); however, the same research group has further investigated whether MRP1 activity in the brain differs between a beta-amyloidosis mouse model (APP/PS1–21) and control mice using the protracer (Zoufal et al. 2020). They partially inhibited MRP1 in the brain with MK571 (300 mg/kg, i.p.), an MRP1 inhibitor, to reduce the transport capacity of MRP1, thereby increasing the sensitivity of 6-bromo-7- $^{11}\text{C}$ methylpurine to detect small changes in MRP1 activity. They found a significant positive correlation between the efflux rate and MRP1 levels in the brain. However, the sensitivity was still poor: when values were roughly read from the figure, Mrp1 levels were increased from 0.5 to 5.5, whereas the elimination rate was slightly increased from 0.9 h<sup>-1</sup> to 1.1 h<sup>-1</sup>, i.e., from 0.015 min<sup>-1</sup> to 0.018 min<sup>-1</sup> (Zoufal et al. 2020). Zoufal et al. speculated that astrocytes were involved in the increases in the efflux parameter based on their strong expression of MRP1 and GSTs and their uniform distribution through the brain. Furthermore, they posited that 6-bromo-7- $^{11}\text{C}$ methylpurine may be used in future studies as a marker of astrocyte function (Zoufal et al. 2020). Considering our findings regarding the rate-limiting step (Okamura et al. 2020) and recent reports regarding the limited sensitivity (Zoufal et al. 2019; Zoufal et al. 2020), however, we posit that 6-bromo-7- $^{11}\text{C}$ methylpurine cannot quantitatively assess Mrp1 activity in either astrocytes or the brains of mice, even if astrocytes among brain parenchymal cells mainly contribute to the efflux of the GSH conjugate from the brain. This question deserves further studies.

## 16.5 Application of the MEM to Other Efflux Transporters

The MEM would apply to noninvasive and quantitative measurement of the activity of other efflux transporters. We have recently developed a protracer, benzyl [ $^{11}\text{C}$ ]hippurate, for measuring the activity of OAT3 in the mouse brain (Kikuchi et al. 2016). The protracer showed moderate brain uptake 0.5 min after intravenous injection and became converted into [ $^{11}\text{C}$ ]hippuric acid, which is thought of as a substrate of MRP4 and OAT3 (Mutsaers et al. 2011; Deguchi et al. 2006), after 5-min postinjection. Moreover, the efflux rate of [ $^{11}\text{C}$ ]hippuric acid in the brain of *Oat3*<sup>-/-</sup> mice was approximately 20% of that of the control mice, indicating that the efflux rate primarily reflected the activity of OAT3 in the brain. Interestingly, MRP4 did not contribute to the elimination of [ $^{11}\text{C}$ ]hippuric acid from the brain although MRP4 is expressed in BCECs (Leggas et al. 2004) and [ $^{11}\text{C}$ ]hippuric acid was retained in the heart of *Mrp4*<sup>-/-</sup> mice (Kikuchi et al. 2016). Given the localization of OAT3 in BCECs, [ $^{11}\text{C}$ ]hippuric acid formed in the mouse brain would be cleared by other transporters (excluding MRP4) after [ $^{11}\text{C}$ ]hippuric acid had been taken up into BCECs mainly via OAT3.

Technetium-99m complex with *N,N'*-1,2-ethylenediylbis-L-cysteine diethyl ester ( $^{99\text{m}}\text{Tc}$ -ECD), whose structure is shown in Fig. 16.2, has been clinically used for brain perfusion imaging (Kapucu et al. 2009). Injected,  $^{99\text{m}}\text{Tc}$ -ECD enters the brain and becomes metabolized into a monoacid in the brain (Walovitch et al. 1994; Walovitch et al. 1991). In mice, *Oat3* mediates the efflux of the  $^{99\text{m}}\text{Tc}$ -ECD metabolite from the brain to the blood (Kikuchi et al. 2014), and thus  $^{99\text{m}}\text{Tc}$ -ECD is a potential protracer for the assessment of *Oat3* activity in the mouse brain. Indeed,  $^{99\text{m}}\text{Tc}$ -ECD was used to analyze the efflux transport function of *Oat3* at the BBB (Kuwahara et al. 2018). However, we should note that  $^{99\text{m}}\text{Tc}$ -ECD as an OAT3 protracer would apply to rodents but not primates because OAT3 is poorly expressed in BCEC in primates, including humans, but highly expressed in mice (Deo et al. 2013). In primates, radioactivity remained trapped in the brain long after the administration of  $^{99\text{m}}\text{Tc}$ -ECD (Walovitch et al. 1994; Walovitch et al. 1991), which is necessary for imaging of cerebral blood flow.

The protracer 6- $^{125}\text{I}$ iodo-9-pentylpurine was designed for the assessment of the efflux transport of iodide from the brain (Okamura et al. 2009a). The protracer readily entered the brain and showed the almost complete conversion to [ $^{125}\text{I}$ ]<sup>-</sup> by 10 min after intravenous administration into mice, whereas the influx of [ $^{125}\text{I}$ ]<sup>-</sup> from the blood to the brain was extremely low for 20 min after intravenous administration. In addition, dose-dependent inhibition of [ $^{125}\text{I}$ ]<sup>-</sup> efflux from the brain was observed in mice pretreated with perchlorate, an inhibitor of iodide transport. Thus, the protracer labeled with  $^{124}\text{I}$  for PET may be able to assess the I<sup>-</sup> efflux from the brain noninvasively, although further studies are needed to identify a carrier protein involved in the efflux of iodide.

*N*- $^{18}\text{F}$ Fluoroethylpiperidin-4-ylmethyl acetate ( $^{18}\text{F}$ ]FEP-4MA) was developed for measuring the activity of acetylcholinesterase in the brain and was shown to be hydrolyzed by this enzyme to its metabolite *N*- $^{18}\text{F}$ fluoroethyl-4-piperidinemethanol (Kikuchi et al. 2010; Kikuchi et al. 2005). In contrast to  $^{18}\text{F}$ ]FEP-4MA, the influx

of the hydrophilic alcohol metabolite from the blood to the brain was negligible after intravenous administration, and the metabolite generated in the brain after [ $^{18}\text{F}$ ]FEP-4MA administration might thus be retained. However, the alcohol metabolite formed in the brain was slowly washed out from the brain (Kikuchi et al. 2010), suggesting that certain transporters may mediate the efflux of the alcohol metabolite from the brain. Thus, [ $^{18}\text{F}$ ]FEP-4MA may be a potential protracer for imaging of the efflux systems, although the carriers that export the alcohol metabolite are currently unknown.

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## 16.6 Conclusions and Perspectives

Our studies using the MEM with 6-bromo-7- $^{[11}\text{C}]$ methylpurine and the intracerebral injection method with *S*-(7- $^{[11}\text{C}]$ methylpurin-6-yl)glutathione in transgenic mice revealed that Mrp1, Oat3, and Mrp4 mediate tracer efflux from the brain. Mrp1 may contribute to the efflux of *S*-(7- $^{[11}\text{C}]$ methylpurin-6-yl)glutathione from brain parenchymal cells, while the tracer in the extracellular space of the brain is likely cleared in part by transport across the BBB by the combination of Oat3 and Mrp4. Although 6-bromo-7- $^{[11}\text{C}]$ methylpurine would not allow the imaging of Mrp1 activity in the mouse brain, it should be noted that the protracer is not always unusable for evaluating MRP1 activity in human brain (parenchymal cells) because there may be species difference in the rate-limiting step of the efflux process via MRP1 between humans and mice. Despite reports suggesting the pathophysiological importance of MRP1 in brain parenchymal cells as described in the Introduction section, past studies have exclusively focused on the *in vivo* role or function of MRP1 at the blood-brain or blood-cerebrospinal fluid interfaces (Lee et al. 2004; Sugiyama et al. 2003; Cisternino et al. 2003; Wijnholds et al. 2000). Methods for noninvasively assessing MRP1 activity in humans have not been developed yet, and hence 6-bromo-7- $^{[11}\text{C}]$ methylpurine is expected to be a potential protracer for the assessment in human brain parenchymal cells, although further clinical studies are needed.

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# Neuroinflammation: From Target Selection to Preclinical and Clinical Studies

# 17

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## Abstract

Inflammation is a highly dynamic and complex adaptive process to preserve and restore tissue homeostasis in neurological disorders and often serves as a prognostic marker for disease outcome. The underlying cellular and factorial heterogeneity represents an opportunity in the development of disease-modifying therapies.

Molecular imaging of neuroinflammation (NI) may support the characterization of key aspects of the dynamic interplay of various inducers, sensors, transducers, and effectors of the multifactorial inflammatory response *in vivo* in animal models and patients. The characterization of the NI response by molecular imaging will (i) support early diagnosis and disease follow-up, (ii) guide (stereotactic) biopsy sampling, (iii) highlight the dynamic changes during disease pathogenesis in a noninvasive manner, (iv) help monitoring existing therapies, (v) support the development of novel NI-modifying therapies, and (vi) aid stratification of patients, according to their individual NI profile.

This book chapter will review the basic principles of NI, recent developments and applications of novel molecular imaging targets, key considerations for the selection and development of imaging targets, as well as examples of successful clinical translation of NI imaging.

## 17.1 Introduction

Inflammation is a highly dynamic and complex process combining local and systemic reactions of multiple cell types, chemical signals, and signaling pathways.

Depending on the inflammatory stimulus and the unique tissue- and disease-specific microenvironment, inflammation can support repair and regeneration processes and, on the opposite, worsen tissue injury (e.g., allergy, chronic infections, neurodegeneration).

The type (mechanical, chemical, infectious, tumoral) of inflammatory stimuli determines the degree and nature of the inflammatory response. However, neuroinflammation (NI) contains unique features, i.e., (i) the CNS has efficient natural

protection from mechanical aggressions by the skull and from biological and chemical aggressions by the blood-brain barrier (BBB); (ii) the CNS has limited regenerative capacities; and (iii) any local impairment of its spatial organization can lead to a definite loss of function.

During NI, resident microglia and infiltrating peripheral immune cells induce a neuroinflammatory response with the overall goal to limit the extent of the disease, to clear tissue damage, and to support repair and regeneration. However, in several cases NI may contribute to the disease process itself, depending on the pathological mechanism.

To improve our understanding of the heterogeneity and dynamics of the disease-specific neuroinflammatory response, including the variety of cells and factors induced and the subsequent changes in tissue organization, exploration methods must take “time” and “space” into account. Molecular, anatomical, and functional *in vivo* imaging methods are essential for (i) early diagnosis and disease follow-up, (ii) investigating dynamic changes during NI noninvasively, (iii) guiding (stereotactic) biopsy sampling, (iv) monitoring existing therapies, (v) supporting the development of novel NI-modifying therapies, and (vi) stratification of patients, according to their specific cellular and molecular composition of NI in correlation to their therapy response. Dedicated imaging paradigms thus have a major impact on patient care (Villa et al. 2018; Zinnhardt et al. 2018). Moreover, *in vivo* imaging technologies are of importance to decipher the molecular mechanisms and functional consequences of neuroinflammatory processes at various disease stages.

It will be important, however, to cross-correlate imaging findings of NI with functional parameters and to integrate imaging data with detailed immunophenotyping by, e.g., flow cytometry and omics data for clinical decision-making. This will allow to gain a holistic understanding of the patients’ individual NI profile and to understand the dynamic interplay of NI and the molecular mechanisms inducing cellular damage.

This book chapter will review the process of target selection and tracer development, as well as current NI imaging strategies in experimental and clinical application.

We describe and discuss the current status and possible future developments of (i) molecular targets for NI, (ii) imaging methods and technologies, and (iii) a variety of neurological diseases. This review will complement other book chapters focusing on dedicated molecular targets of NI.

The focus will be on particular requirements for target selection, molecular mechanisms, and possible targets for imaging in NI, assuming that the reader is familiar with the basic principles of the major imaging technologies, such as optical [OI], radionuclide (positron emission tomography [PET] and single photon emission computed tomography [SPECT]), and magnetic resonance imaging [MRI], as their different strengths and weaknesses have already been extensively commented upon, i.e., a high sensitivity and high spatial resolution for OI; high sensitivity and possibility for true signal quantification for radionuclide imaging; and high spatial resolution and physiological and biochemical imaging for MRI.

This book chapter is divided into three parts. The basic principles of NI are briefly summarized and potential targets for molecular imaging are reviewed in the first part.

In the second part, particular requirements as well as decision points for molecular imaging targets and tracer development are discussed. The third part highlights the application of the concepts of the second part in clinical and experimental molecular imaging of NI.

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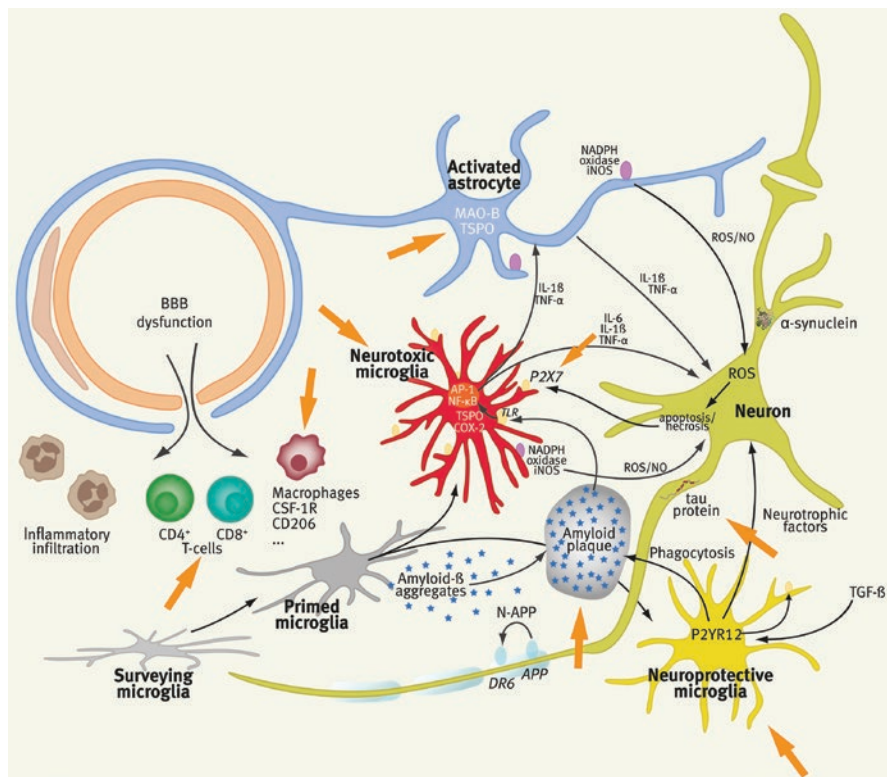
## 17.2 Basic Principles of Neuroinflammation

The term “inflammation” describes a multicellular and multifactorial process, characterized by (i) changes in local vasculature (increased blood flow and vascular permeability), (ii) activation of resident immune cells, (iii) infiltration of peripheral immune cells (neutrophils, macrophages, lymphocytes), and (iv) cytokine production (Graeber et al. 2011). During NI, a certain stimulus (environmental, toxic, hypoxic, viral, bacterial, mechanical, protein deposits, etc.) leads to an immune activation in the CNS, potentially contributing to tissue damage, loss of neurons and dysfunction, and on the other hand neuroregeneration and tissue repair (Graeber et al. 2011).

The underlying mechanisms include:

1. Activation of damage-associated molecular patterns (DAMPs, e.g., high-mobility group box 1 [HMGB1] proteins, heat-shock proteins [HSPs], histones, oxidized lipids, ATP, amyloid- $\beta$ ) by tissue injury or of pathogen-associated molecular patterns (PAMPs) by exogenous pathogens
2. Activation of Toll-like receptors (TLR) of various subtypes and of the receptor for advanced glycation end products (RAGE) with subsequent release of adhesion molecules, cytokines, and chemokines directing activation and targeted migration of effector cells (microglia, macrophages, lymphocytes)
3. Activation and recruitment of microglial cells
4. Infiltration of macrophages and T-lymphocytes of various subtypes (CD4+, CD8+, Th1, Th2, Th17, regulatory T cells)
5. Alteration of blood-brain barrier (BBB) integrity
6. Activation and recruitment of astrocytes
7. Release or expression of a variety of signaling factors, including neuron-derived neuropeptides, membrane proteins (e.g., fractalkine, cannabinoid receptors, major histocompatibility complex [MHC] molecules), semaphorins, and lectins which all take part in the regulation of NI (Jacobs et al. 2012; Stephenson et al. 2018)

In summary, the heterogeneous nature of NI and the pleiotropic functions of different immune cells during the course of NI render NI as attractive source of targets for therapeutic modulation in the treatment of neurological diseases. Molecular imaging of NI has greatly enhanced our understanding of the dynamics of immune



**Fig. 17.1** Overview of selected imaging targets within the neurovascular unit. Abbreviations: *MAO-B* monoamine oxidase B, *TSPO* translocator protein, *IL* interleukin, *TNF- $\alpha$*  tumor necrosis factor alpha, *ROS* reactive oxygen species, *TGF- $\beta$*  transforming growth factor beta, *APP* amyloid precursor protein, *CSF-1R* colony-stimulating factor 1 receptor, *TLR* Toll-like receptor, *BBB* blood-brain barrier (modified from Jacobs et al. 2012 and Perani et al. 2019)

cells and factors in neurological diseases. The following paragraphs will highlight examples of imaging targets in NI (Fig. 17.1).

## 17.3 Potential Imaging Targets

### 17.3.1 DAMPs

Damage-associated molecular patterns (DAMPs) represent a heterogeneous group of nuclear or cytosolic proteins released in response to a pathological stimulus, including the high-mobility group protein B1 (HMGB1), S100 proteins, heat-shock proteins (HSPs), chromogranin A, and chromogranin A $\beta$ . In addition, nucleic acids and nucleotide derivatives, such as mitochondrial DNA (mt-DNA), DNA, and adenosine triphosphate (ATP), may serve as DAMPs (Ardura-Fabregat et al. 2017).

An excellent example of *in vivo* molecular imaging of DAMPs has been the direct visualization of amyloid- $\beta$  (the DAMP of AD) using radiolabeled thioflavine derivatives (e.g., Pittsburgh compound B; [ $^{11}\text{C}$ ]PIB). Next-generation tracers include [ $^{18}\text{F}$ ]florbetapir, [ $^{18}\text{F}$ ]florbetaben, and [ $^{18}\text{F}$ ]flutemetamol, which are currently approved by the Food and Drug Administration and the European Medicines Agency (for review, see Perani et al. 2019).

Another DAMP of interest in NI is the appearance of hyperphosphorylated tau protein. Several classes of tau tracers have been described, including the [ $^{18}\text{F}$ ]THK family, [ $^{18}\text{F}$ ]flortaucipir (formerly known as [ $^{18}\text{F}$ ]AV1451 and [ $^{18}\text{F}$ ]T807), and [ $^{11}\text{C}$ ]PBB3 (for review, see Perani et al. 2019). Recently, tracers with improved characteristics and hence improved selectivity and specificity were introduced (Honer et al. 2017; Kuwabara et al. 2018; Wong et al. 2018). Other DAMP-targeting tracers for neurodegenerative diseases are reviewed elsewhere (Perani et al. 2019).

### 17.3.2 Chemokines

Chemokines and their receptors comprise approximately 50 peptides and 20 receptors in humans, with homologs, orthologs, and related peptides in other vertebrate species. In NI, chemokines and their receptors fulfill multiple roles. They are involved in guiding movement of (immuno-) cells (chemoattraction), modulating the chemokine environment, activation of glial cells, and coordination of complex tissue responses (Ransohoff 2009).

Among the most intensively studied chemokines in AD are CX3C chemokine ligand 1 (CX3CL1, fractalkine) and chemokine ligand 2 (CCL2).

CX3CL1 and CX3CR1 may have a bimodal function in AD, depending on the underlying pathology. CX3CL1 and CX3CR1 play a beneficial role in the context of tau pathology while exerting detrimental functions in amyloid pathology (Ardura-Fabregat et al. 2017). No PET ligands were described for imaging the CX3CL1 and CX3CR1 system. In contrast, the CCL2-CCR2 pathway seems to be positively associated with AD disease severity. Recently, a PET probe for CCR2-directed inflammation was successfully introduced (Liu et al. 2017). However, validation in the context of NI is still missing.

Another chemokine of interest is C-X-C motif chemokine receptor 4 (CXCR4) involved in the regulation of leukocyte trafficking (Jacobson and Weiss 2013). CXCR4 can be visualized by PET using the probe [ $^{68}\text{Ga}$ ]pentixafor (Buck et al. 2017).

### 17.3.3 Microglia/Macrophages

Microglia play an important role in tissue homeostasis and response to pathological stimuli (Hickman et al. 2018). Their responsiveness to injury suggests microglia as diagnostic markers of disease onset or progression. Microglia and their counterparts outside of the CNS, macrophages, represent an interesting target for imaging NI.



Various cell surface and mitochondrial receptors expressed in microglia/macrophage cells are involved in the regulation and function of microglia and macrophages, and some of these receptors were used for the development of ligands for imaging. One of the most studied targets is the 18 kDa translocator protein (TSPO, formerly called peripheral benzodiazepine receptor or PBR), which is upregulated in response to neuroinflammatory stimuli. TSPO will be further discussed later in this chapter as well as in another book chapter (Chap. 33). Besides TSPO, new targets for microglia/macrophage imaging are evolving, as TSPO PET in clinical applications is limited by (i) low target density in neurodegeneration, (ii) lack of selectivity for microglia (astrocytes, peripheral macrophages, endothelial cells), (iii) varying binding affinity due to a genetic polymorphism, and (iv) lack of selectivity for various microglia subtypes (Tronel et al. 2017).

New microglia targets include receptor-based ligands (purinergic receptors P2X7 and P2Y<sub>12</sub>, colony-stimulating factor receptor 1 (CSF-1R), cannabinoid receptors ( $\alpha 7$  and  $\alpha 4\beta 2$ ), nicotinic acetylcholine receptors, adenosine 2A receptor, folate receptor  $\beta$ ) and enzyme-targeting ligands (cyclooxygenase, nitric oxide synthase, matrix metalloproteinase,  $\beta$ -glucuronidase, and enzymes of the kynurenine pathway) (Wagner et al. 2007; de Vries et al. 2008; Martín et al. 2015; Tronel et al. 2017; Colás et al. 2018; Janssen et al. 2018; Villa et al. 2018; Horti et al. 2019).

Macrophages can be imaged, e.g., with the radiolabeled macrophage mannose receptor (MMR, CD206)-targeting [68Ga]NOTA-anti-MMR nanobodies (Movahedi et al. 2012; Varasteh et al. 2019).

### 17.3.4 Lymphocytes

Lymphocytes are important mediators of the NI response in neurological disorders but fulfill also physiological functions, including memory consolidation, hippocampal long-term potentiation, and neurogenesis, and they also contribute to psychological stress resilience. In response to NI, lymphocytes infiltrate into the CNS where they enhance and perpetuate neuroinflammatory processes, e.g., in neurodegeneration (González et al. 2014). The vast heterogeneity of NI-associated lymphocytes makes lymphocytes an attractive but also challenging imaging target.

Two important subsets of T cells, CD4<sup>+</sup> and CD8<sup>+</sup>, can be investigated using single-domain antibody fragments (VHHs) and cys-diabodies (Tavaré et al. 2015; Rashidian et al. 2017). Likewise, first approaches for imaging B cells using <sup>64</sup>Cu-rituximab were described (James et al. 2017). The use of these new imaging biomarkers in the context of NI remains to be elucidated. In the context of lymphocyte-targeting immunotherapies, immuno-PET will inform on the presence and localization of the therapeutic target (Zettlitz et al. 2019). Likewise, direct visualization of immune checkpoints may support the identification of cellular target expression, heterogeneity of inflammatory lesions, as well as therapy effects (Niemeijer et al. 2018).

### 17.3.5 Astrocytes

Astrocytes are a heterogeneous and pleomorphic population of glial cells in the brain. Astrocytes are in direct contact with cerebral blood vessels and neurons, forming the neurovascular unit (NVU). Next to physiological functions as regulators of blood flow, neuronal development and activity, energy metabolism, and circadian rhythm, astrocytes differentially respond to NI stimuli. Depending on the underlying stimulus, astrocytes fulfill beneficial and detrimental functions in NI (González et al. 2014; Boche et al. 2019).

Current astrocyte imaging techniques are limited to  $^{11}\text{C}$ -deuterium-L-deprenyl ( $^{11}\text{C}$ -DED) binding to MAO-B.  $^{11}\text{C}$ -DED may be used to image early stages of AD (Boche et al. 2019). Recently, imidazoline-2 binding site (I2BS)-targeting tracers are gaining attention for the assessment of astrocytes (Boche et al. 2019).

To noninvasively investigate the (over-) expression, activation, and/or increase/decrease of factors, gene reporters and cells in response to NI by molecular imaging, targets of interest, and their respective tracers have to fulfill certain requirements, as described below (for review, see Elsinga 2002, Patel and Gibson 2008, and Serdons et al. 2009).

### 17.3.6 Target Considerations

Before developing and designing novel tracers for the investigation of NI, certain criteria should be considered:

1. Unmet clinical need/specific biological question: The decision on a target for tracer development must be based on recent targets in drug development, a currently unmet medical need and/or a biological question, which could not be answered without noninvasive imaging.
2. Level of expression and maximum concentration of target binding sites ( $B_{\max}$ ): Ideally, the target of interest should be expressed in sufficient amounts to be detected by a tracer. As the level of affinity ( $K_d$ ) of many radiotracers is usually in the pico-/nanomolar range, the expression level of the target should be at least a factor of 4 higher than the  $K_d$  ( $B_{\max}/K_d > 4-10$ ). Targets with low expression levels require a higher radiotracer affinity, in order to show specific binding in vivo (Zhang and Villalobos 2017).
3. Cellular expression: The cellular sources of a certain marker of interest are often not fully characterized. This is of particular importance in tracer development, as this may lead to difficulties in the interpretation of future imaging results. An ideal target is only expressed by a certain cell type at a certain disease state. However, the plasticity of cells and factors and the dynamics of NI make it difficult to achieve this goal. Dedicated immuno-phenotyping and cross-correlation of preclinical and clinical tissue samples may inform on the cellular substrates of the target of interest.

4. **Biological function:** In line with the characterization of the cellular sources of the target of interest, the precise function of the target should ideally be determined in advance. However, the particular function of the target may change during the course of NI.
5. **Target localization:** The localization and biological function of the target determines the molecular structure of the lead radio tracer. Most current tracers are small molecule tracers targeting cellular receptors, transporters, enzymes, as well as intra- and extracellular proteins and protein aggregates (Van de Bittner et al. 2014). However, in the era of immunotherapies, radiolabeled biologicals including antibodies, antibody fragments, and nanobodies are increasingly investigated as radio tracers with longer circulation times and biological half-life.

### 17.3.7 Tracer Considerations

Next to considerations for target selection, the design of NI tracers needs to take the following points into consideration (for review, see Pike 2009, Serdons et al. 2009, Zhang and Villalobos 2017, and Vermeulen et al. 2019):

1. **Blood-brain barrier (BBB) penetration:** To reach their target in the brain, novel NI tracers need to be able to cross the BBB while avoiding to be actively transported out of the brain by, e.g., P-glycoprotein (P-gp) efflux pumps and/or multidrug resistance-associated proteins (Pike 2009). Additionally, tracers need to be able to cross the cell membrane for intracellular targets.
2. **Structure attribute:** Structure of the lead compound allows for high specific activity labeling with short-lived isotopes, including F-18 or C-11.
3. **Affinity:** The affinity for the target should be ideally in the nano- to picomolar range.
4. **Specificity/selectivity:** The tracer should have a high specificity and selectivity for the target to avoid off-target tracer binding to, for example, plasma proteins.
5. **Binding characteristics:** The binding characteristics of the lead structure (e.g., covalent vs. non-covalent, internalization, metabolic trapping, etc.) are of importance for the quantification of the imaging signal.
6. **Metabolic stability:** The biological half-life and the bioavailability of a tracer are determined by the chemical structure of the lead compound and the tracer environment. Nevertheless, most of the tracers are rapidly metabolized by metabolizing enzymes in blood and tissues. It remains important that the resulting radiometabolites are polar and less lipophilic than the parent tracer for rapid washout from the brain, as radiometabolites are indistinguishable from the parent tracer fraction in PET. Ideally, the parent tracer should not be metabolized at all, allowing for determination of true quantitative measures (e.g., regional values of volumes of distribution ( $V_T$ ) or binding potentials (BP) of the target) (Pike 2009).

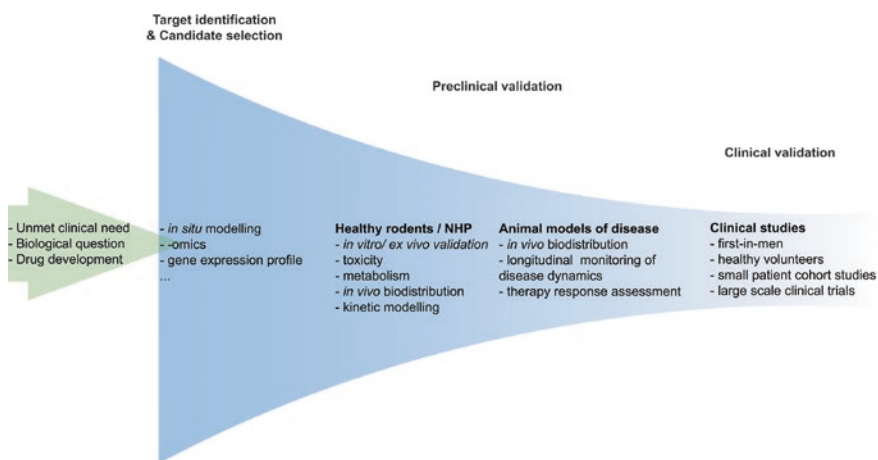
Additionally, the biological half-life should be shorter than for a therapeutic drug, as a fast clearance of the tracer from the blood pool will improve the target-to-background ratio (Serdons et al. 2009).

Next to the biological half-life, the half-life of the chosen radioisotope determines the duration of possible biological investigations. For example, isotopes like C-11 with a half-life of about 20 min do not allow repeated imaging sessions, whereas isotopes with longer half-life (e.g., Zr-89,  $T_{1/2}$ , 78.4 h) allow repeated imaging sessions with a single injection of tracer. However, isotopes with a longer half-life also bear the risk of longer exposure and higher organ doses for the patient (Van Dongen et al. 2015). For quantification purposes it is essential that no brain-permeable radioactive metabolites occur (Pike 2009).

## 17.4 Preclinical Imaging and Clinical Imaging in Neurological Diseases

After target selection, lead definition, and initial radiotracer synthesis, tracer candidates undergo further *ex vivo* and *in vivo* investigations before entering clinical trials.

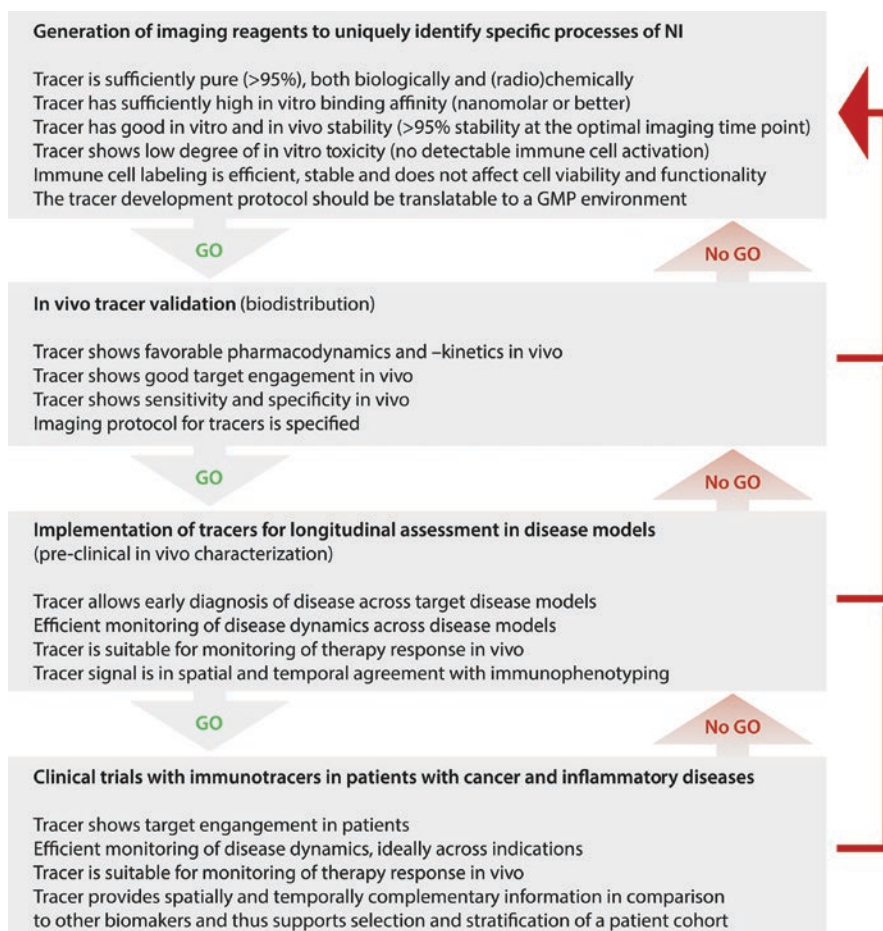
First, tracer candidates are screened for stability and *ex vivo* binding affinities (e.g., cellular binding and autoradiography, competition assays). The most promising tracer candidates are then transferred into small animals (mice and rats), including healthy animals (biodistribution, pharmacokinetics, toxicity, kinetic modeling) and disease models of interest. The candidate tracers are then further tested in larger animals, e.g., nonhuman primates (NHP) before entering first-in-man studies in healthy volunteers (dosimetry) and small patient cohorts (Serdons et al. 2009; Zhang and Villalobos 2017). The general workflow of tracer development is summarized in Fig. 17.2.



**Fig. 17.2** Simplified workflow of tracer development

Along the development of novel imaging biomarkers, important go/no-go decision points have to be taken into consideration. For example, in the initial phase of target selection, tracers are required that have sufficient biological and radiochemical purities (>95%), as well as favorable affinity and in vivo stability (>95%). If one of these points is not met, the target/lead structure should be reconsidered. In accordance, tracers in first in vivo validation not showing favorable pharmacodynamics and pharmacokinetics, target engagement, and sensitivity have to be returned for radiochemical modification. Potential decision points are summarized in Fig. 17.3.

The following paragraphs will highlight examples of NI tracers and their transition from ex vivo to *first-in-patient* applications.



**Fig. 17.3** Overview of potential go/no-go decision criteria in the development of imaging biomarkers

## 17.5 From Ex Vivo Validation to First-in-Man Studies: The Story of [<sup>18</sup>F]DPA-714

Upregulation of the 18 kDa translocator protein (TSPO, formerly called peripheral benzodiazepine receptor or PBR) is a hallmark of neuroinflammation. TSPO is located in the outer mitochondrial membrane and involved in the transport of cholesterol into mitochondria. Under physiological conditions, TSPO is expressed by ependymal cells, olfactory bulb neurons, and endothelial cells, but its global expression is low (Rupprecht et al. 2010). In response to neuroinflammatory stimuli, TSPO becomes rapidly and extensively upregulated. However, despite more than 30 years of research, the precise functions of TSPO remain to be elucidated. Initially, TSPO tracers were developed to image microglial activation, highlighting the importance of characterization of cellular sources by immuno-phenotyping. It has to be pointed out that the cellular sources may be depending on the disease context, disease severity, and disease stage (Lavissee et al. 2012; Zinnhardt et al. 2015, 2018; Wiesmann et al. 2017; Villa et al. 2018).

Nevertheless, TSPO meets most of the target criteria for tracer development as described above.

Following the process of target selection, several TSPO tracers were developed. The first-generation TSPO tracer (*R*)-[<sup>11</sup>C]PK11195 is among them. However, the application of (*R*)-[<sup>11</sup>C]PK11195 was severely limited by poor signal-to-noise ratios, as well as the short half-life of the <sup>11</sup>C-label. Second-generation tracers with improved bioavailability, higher specificity, and favorable signal-to-noise ratios were developed, including [<sup>18</sup>F]DPA-714 (Jacobs et al. 2012). Using the example of [<sup>18</sup>F]DPA-714, the following paragraphs will exemplarily highlight the transition from ex vivo to in vivo applications, including major decision points for transition to the next phase.

The synthesis and first characterization of [<sup>18</sup>F]DPA-714 were first described by James et al. (2008). The in vitro binding affinity of the lead molecule DPA-714 for its target TSPO was determined in isolated mitochondria from rat kidneys and C6 glioma cells. Partition coefficients (Log *D*) were determined by HPLC. In vivo validation was performed in rats after QA-induced excitotoxic damage. For validation of the in vivo biodistribution findings, several brain regions and peripheral organs were harvested and measured ex vivo in a gamma counter. After obtaining encouraging imaging results, pre-dosing with cold DPA-714, DPA-713, and PK11195, respectively, was performed. Pre-dosing led to substantial blocking of the signal, indicating specificity of the tracer.

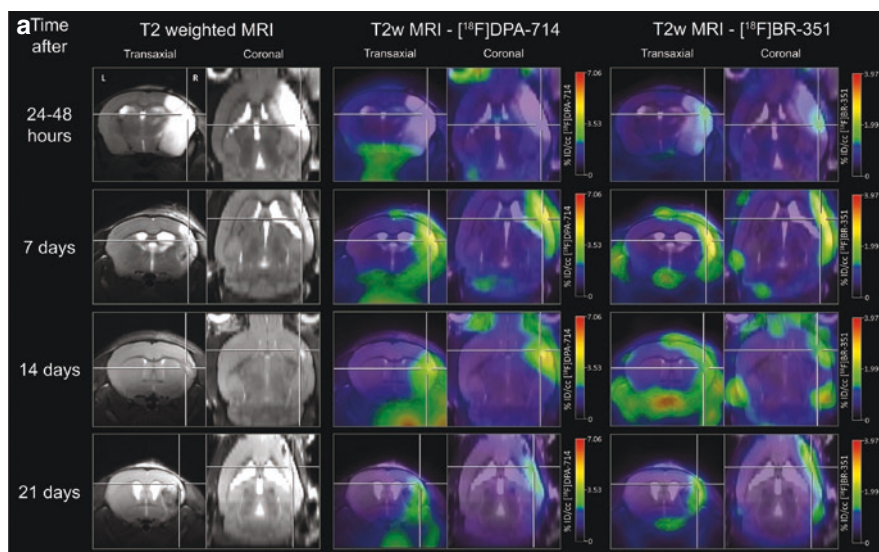
Biodistribution studies in healthy rats were not performed; instead the biodistribution and specificity of [<sup>18</sup>F]DPA-714 were investigated in a more clinically relevant nonhuman primate model after pretreatment with PK11195 5 min before radioligand injection, as well as with a displacement study, where non-labeled DPA-714 was administered 20 min after the injection of [<sup>18</sup>F]DPA-714 (James et al. 2008).

Despite an initial validation and proof of specificity, several points remained to be addressed for the ultimate goal of translating [<sup>18</sup>F]DPA-714 into clinical

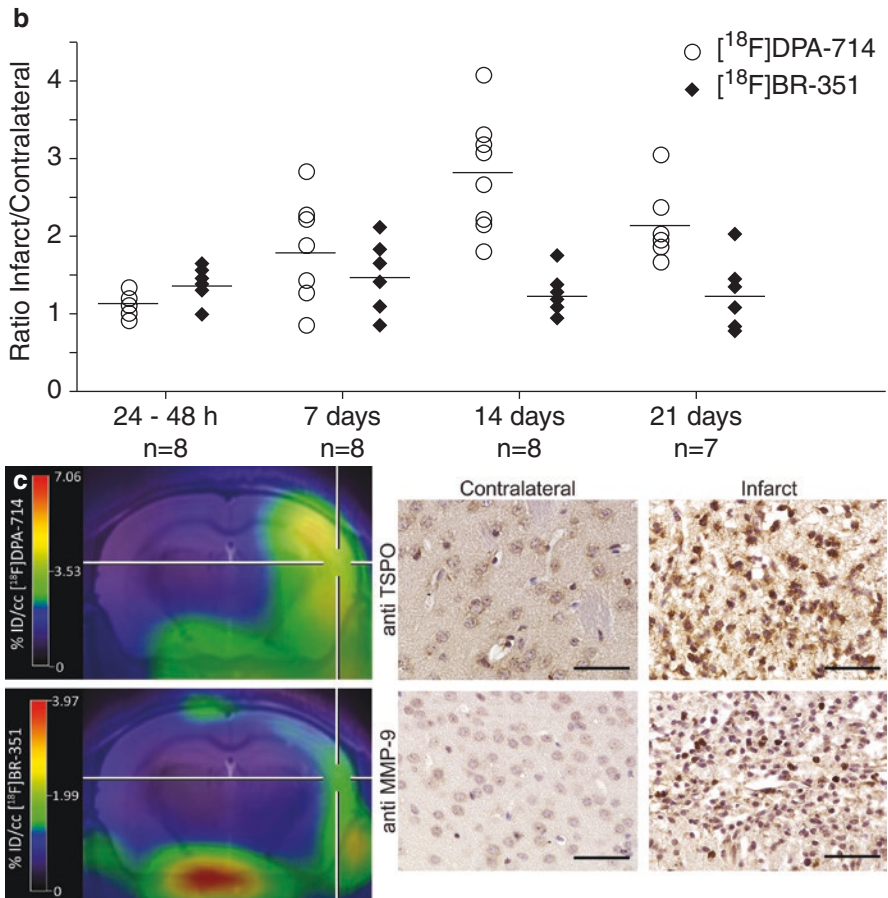
application, including biodistribution in healthy and murine models of disease, histological validation, cross-correlation with other (imaging-) biomarkers, longitudinal studies for the investigation of disease dynamics across various indications, investigations of therapy-induced changes, first-in-man and clinical studies, as well as quantitative modeling of [ $^{18}\text{F}$ ]DPA-714 PET signals.

Consequently, the biodistribution of [ $^{18}\text{F}$ ]DPA-714 was further investigated in mice (Vicidomini et al. 2015) and healthy NHP (Lavissee et al. 2015b). In addition, indispensable for the pharmacodynamic assessment of a lead tracer, the plasma and brain metabolism of [ $^{18}\text{F}$ ]DPA-714 was determined ex vivo and in vivo (Peyronneau et al. 2013).

The in vivo characteristics of [ $^{18}\text{F}$ ]DPA-714 were further analyzed at single time points and longitudinally in various rodent models of NI, including stroke (Fig. 17.4) (Martín et al. 2010; Zinnhardt et al. 2015), glioma (Fig. 17.5) (Tang et al. 2012; Winkeler et al. 2012; Zinnhardt et al. 2017; Pigeon et al. 2019), models of multiple sclerosis (Fig. 17.6) (Abourbeh et al. 2012; Zinnhardt et al. 2019), Alzheimer's disease (AD) (Sérierie et al. 2015; Takkinen et al. 2017; Chaney et al. 2019), Parkinson's disease (Fig. 17.7) (Fricke et al. 2016), ALS (Gargiulo et al. 2016), traumatic brain injury (Wang et al. 2014a; Israel et al. 2016), epilepsy (Harhausen et al. 2013; Nguyen et al. 2018), subarachnoid hemorrhage (Lee et al. 2015), experimental BBB opening (Sinharay et al. 2019), depression and anxiety (Wang et al.



**Fig. 17.4** (a) Example of a longitudinal study combining two imaging biomarkers of NI [ $^{18}\text{F}$ ]DPA-714 (TSPO) and [ $^{18}\text{F}$ ]BR-351 (MMPs) in the postischemic phase after stroke. (b) TSPO PET signals peaked 14 days after stroke, whereas MMP PET preceded peaking 1 week after stroke. (c) Histological validation of target expression. The combination of imaging biomarkers and in vivo PET supported the characterization of the reactive postischemic inflammation (modified from Zinnhardt et al. 2015)



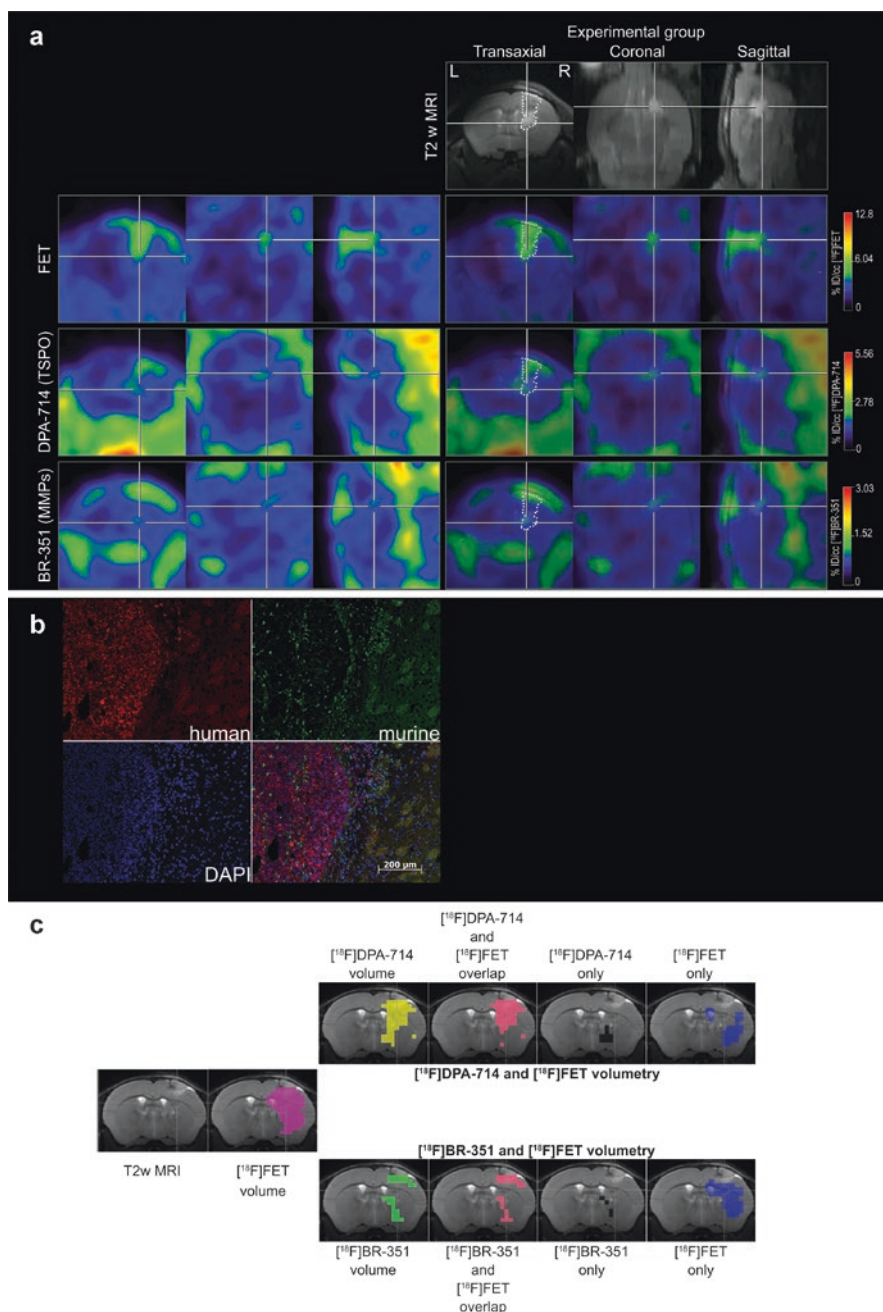
**Fig. 17.4** (continued)

2018), alcohol exposure (Saba et al. 2018), morphine exposure (Auvity et al. 2016, 2017), HIV-associated NI (Lee et al. 2015), and Zika virus-associated NI (Lee et al. 2015).

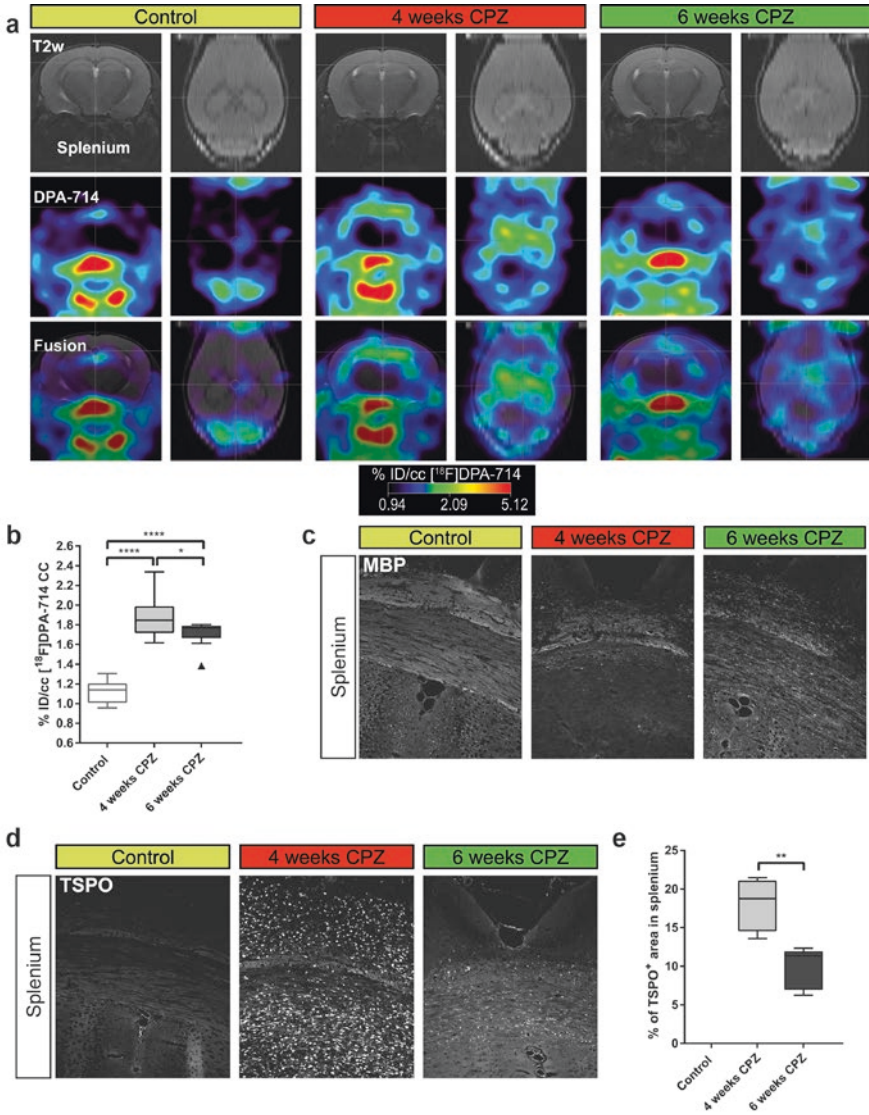
The suitability of  $[^{18}\text{F}]\text{DPA-714}$  as imaging biomarker to detect therapy-induced changes of disease-modifying therapies was also successfully investigated in various animal models of NI (Martin et al. 2011; Awde et al. 2013; Wang et al. 2014b; Domercq et al. 2016; Wiesmann et al. 2017; Colás et al. 2018; Tan et al. 2018; Zinnhardt et al. 2018).

Importantly, head-to-head comparisons with, for example, first-generation TSPO ligands were performed to understand the relative improvement of tracer

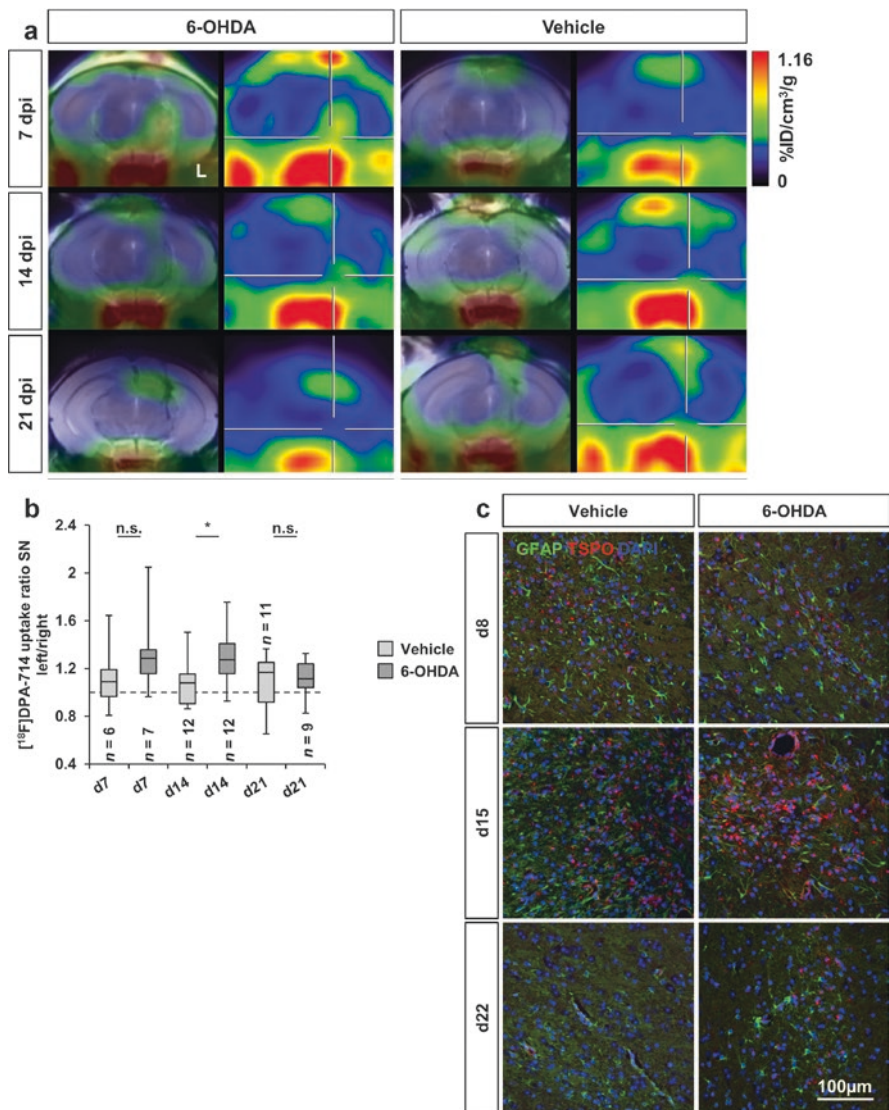




**Fig. 17.5** (a) The combination of the imaging biomarkers [<sup>18</sup>F]FET, [<sup>18</sup>F]DPA-714, and [<sup>18</sup>F]BR-351 reveals complementary information of the glioma immune microenvironment. (b) Histological confirmation of TSPO expression in human glioma cell lines and infiltrating murine immune cells. (c) A thresholding-based volumetric analysis reveals distinct areas of tracer uptake, potentially representing areas of tumor infiltration and immune escape (modified from Zinnhardt et al. 2017)



**Fig. 17.6** (a) [<sup>18</sup>F]DPA-714 (TSPO) imaging of two distinct disease stages (demyelination at week 4 vs. remyelination at week 6) in the cuprizone model of multiple sclerosis. (b) Increased uptake of [<sup>18</sup>F]DPA-714 was observed during demyelination, whereas uptake levels decreased at remyelination. (c) Immunofluorescence for myelin basic protein (MBP) confirmed time-dependent demyelination and remyelination in the cuprizone model. (d, e) Immunofluorescence confirmed increased TSPO expression in the splenium of the corpus callosum at demyelination (week 4). At remyelination (week 6), TSPO levels decreased again (modified from Zinnhardt et al. 2019)



**Fig. 17.7** (a, b) Intranigral 6-OHDA-induced degeneration of the nigrostriatal system is accompanied by increased TSPO levels, measured by [<sup>18</sup>F]DPA-714 in the substantia nigra. (c) Histological markers validate the presence of micro- and astrogliosis in the substantia nigra (modified from Fricke et al. 2016)

characteristics, including an improved higher signal-to-noise ratio (Chauveau et al. 2009; Boutin et al. 2013; Sridharan et al. 2017).

Through histological assessment, most of the studies focusing on NI identified myeloid cells as the major source of TSPO expression (Rupprecht et al. 2010; Jacobs et al. 2012; Chaney et al. 2019). However, the general concept of [ $^{18}\text{F}$ ]DPA-714 as microglia tracer was challenged by Lavissee et al. in Lavissee et al. 2012, highlighting astrocytes as additional source of TSPO expression (Lavissee et al. 2012). Although the expression of TSPO in astrocytes seems to be disease context-dependent and mostly associated with chronic disease stages (Wiesmann et al. 2017; Zinnhardt et al. 2019), these findings again highlight the importance of a detailed characterization of the cellular sources of the target of interest in radiotracer development.

After successful characterization in animal models, another important step toward clinical application was marked with the description of an automated synthesis of [ $^{18}\text{F}$ ]DPA-714 required for manufacturing qualification for clinical production (Kuhnast et al. 2012). Consequently, healthy volunteers were imaged with [ $^{18}\text{F}$ ]DPA-714 including arterial and venous blood sampling to understand the bio-distribution characteristics, in vivo stability, and dosimetry of [ $^{18}\text{F}$ ]DPA-714 in humans (Arlicot et al. 2012). [ $^{18}\text{F}$ ]DPA-714 was described with excellent in vivo stability, biodistribution, and acceptable effective dose estimation, paving the way for small cohort studies in patients with neurological diseases, including stroke (Ribeiro et al. 2014), ALS (Corcia et al. 2012), and AD (Hamelin et al. 2016, 2018).

In summary, the development of [ $^{18}\text{F}$ ]DPA-714 as NI tracer represents an excellent example of structured radiotracer development leading to successful clinical applications.

However, the application of [ $^{18}\text{F}$ ]DPA-714 may be complicated by (i) the lack of selectivity of TSPO to distinguish different immune cell subsets, in particular microglia, (ii) the differential binding affinities within the human population (Owen et al. 2010, 2012, 2015), (iii) the intracellular localization of the target, and (iv) challenges in the exact quantification of [ $^{18}\text{F}$ ]DPA-714 signals, due to a lack of a TSPO devoid reference region within the brain (Golla et al. 2015, 2016; Lavissee et al. 2015a; García-Lorenzo et al. 2018; Wimberley et al. 2018).

The challenge to image microglia has led to the identification and investigation of additional targets for imaging NI. In an integrated approach within the EU FP7 framework “Imaging Inflammation in Neurodegenerative diseases,” Villa et al. aimed to identify new translational markers of the anti-inflammatory/protective state of microglia for the development of novel PET tracers. Using bioinformatics, genes encoding candidate targets were selected based on the following criteria:

1. The level of expression of the candidate mRNA in the anti-inflammatory state had to be at least twofold higher compared to the resting/surveillance state and to the pro-inflammatory activation state.
2. The genes had to be similarly regulated in rodents and humans.
3. The genes had to be preferentially expressed in microglia.
4. Cell surface expression rather than intracellular expression.

5. Higher expression in microglia versus macrophages (or other neural cell types).
6. Available literature on small molecule compounds binding the encoded proteins (to be used for the development of a PET tracer).

The resulting target candidates were further investigated in human primary macrophage and microglia cultures. Polarization with recombinant IL-4 and LPS is allowed to select markers for anti- and pro-inflammatory microglia, respectively. These candidates were further investigated in microglia isolated from murine brains after intracerebral injection of IL-4 and LPS. The results suggested P2ry12/P2RY12 as a promising translational target for PET imaging of the anti-inflammatory activation state of microglia, due to its high specificity for microglial cells versus other brain cells and macrophages and the high and consistent induction in mouse models of disease *in vivo* and in human microglia *in vitro*. Consequently, P2RY12 was chosen for the development of novel PET tracers for application in preclinical and clinical *in vivo* imaging (Villa et al. 2018).

Together, this study highlights the potential of a rationale-based approach for the identification of new molecular targets for imaging NI and may serve as blueprint for the identification of novel biomarkers in NI.

Currently, several other targets, including the role of purinergic receptors for imaging NI, are under investigation (Jacobs et al. 2012; Janssen et al. 2018; Perani et al. 2019). These will be reviewed elsewhere in this book (e.g., Chap. 26).

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## 17.6 Conclusions

CNS diseases represent a pathophysiologically heterogeneous group of diseases, where neuroinflammation is the common link between cellular stress response, synapse dysfunction, and neuronal damage. Targeting the NI response represents an exciting strategy in drug discovery and patient management to protect neuronal function. However, noninvasive tools to assess the NI response for integration into the drug discovery process and clinical decision-making are still scarce.

Assessing the NI response by molecular imaging remains an important denominator to understand the underlying pathophysiological changes in neurological diseases. In the era of targeted immunotherapies, the detailed characterization of the immuno-environment in neurological diseases is of great fundamental and clinical interest. Novel tracers will provide quantifiable endpoints for the development and early assessment of therapeutic efficacy in various neurological diseases and may support decision-making in drug development. It will be important to cross-correlate imaging findings with in-depth immuno-phenotyping as well as benchmark novel tracer developments against existing NI and clinical standard tracers. The applicability of novel NI tracers in clinical studies will also be strongly dependent on the availability of improved methodology, including the level of standardization, harmonization across centers, and available quantification tools.

Additionally, the application of a single tracer will likely not be sufficient to image the complexity of NI, suggesting that multi-tracer and multi-modality

combinations may be the method of choice to cope with and disentangle the complexity of neurological diseases. To be successful, an integrated, multidisciplinary approach is necessary to facilitate the transition of novel tracers targeting the NI response into clinical application.

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# Preclinical and Clinical Aspects of Nicotinic Acetylcholine Receptor Imaging

# 18

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## Abstract

Innovations in radiochemistry and pharmacology are opening new vistas for studies of nicotinic acetylcholine receptors (nAChRs) in human brain by positron emission tomography (PET) and by single-photon emission computed tomography (SPECT). In parallel, instrumentation optimized for molecular imaging in rodents facilitates preclinical studies in models of human diseases with perturbed nAChR signalling, notably Alzheimer's disease and other neurodegenerative conditions, schizophrenia and other neuropsychiatric disorders, substance abuse and traumatic brain injury. The nAChRs are ligand-gated ion channels composed of five subunits forming a central pore for cation flux. The most abundant nAChRs in the central nervous system are heteropentamers (designated  $\alpha 4\beta 2$ ), followed by the  $\alpha 7$  homopentamer. We present a systematic review of published findings with the various nAChR ligands using imaging techniques in vivo, emphasizing preclinical models and human studies. Molecular PET imaging of the  $\alpha 4\beta 2$  nAChR subtype with the antagonist 2-[ $^{18}\text{F}$ ]fluoro-A-85380 is hampered by the long acquisition times. Newer agents such as (-)-[ $^{18}\text{F}$ ]flubatine, [ $^{18}\text{F}$ ]XTRA or [ $^{18}\text{F}$ ]nifene permit quantitation of  $\alpha 4\beta 2$  receptors with PET recordings lasting 90 min or less and without the toxicity risk of earlier epibatidine derivatives. The early PET studies of  $\alpha 7$  nAChRs suffered from low pharmacological specificity, further hampered by low natural abundance of the receptor. However, several good  $\alpha 7$  nAChR ligands such as [ $^{18}\text{F}$ ]ASEM and [ $^{18}\text{F}$ ]DBT10 have emerged in the past few years. There are still no ligands selective for  $\alpha 6$ -containing nAChRs, despite their importance for nicotine-induced dopamine release in striatum. Selective  $\alpha 3\beta 4$  nAChR radioligands are under development but remain untested in clinical studies of depression and addiction. Several nAChR ligands find use for pharmacological occupancy studies, and competition from endogenous acetylcholine reduces  $\alpha 4\beta 2$  binding site availability, a property that enables monitoring by PET of acetylcholine release in living brain.

## 18.1 Introduction

Nicotine is an addictive drug named after the French diplomat Jean Nicot, who introduced cultivation of the tobacco plant (*Nicotiana tabacum*) in Europe. As early as 1828, nicotine was characterized as the major pharmacologically active substance of this solanaceous herb (Posselt and Reimann 1828). Elucidation of its chemical structure followed 60 years later (Pinner and Wolffenstein 1891; Pinner 1893), and soon thereafter Amé Pictet obtained the first successful synthesis (Pictet 1903). The stimulation of sympathetic ganglia by nicotine, first observed by John Newport Langley in Cambridge (Langley 1901), led to the concept that receptors mediate drug actions (Langley 1905; Bennett 2000) and ultimately to the modern concept of molecular neurotransmission.

Nicotine and related alkaloids are present in other *Solanaceae*, such as tomato, potato, peppers and eggplant, and also in tea leaves (Schep et al. 2009), such that the mean daily dietary nicotine intake is about 1.4  $\mu\text{g}$  per day (Siegmund et al. 1999). Nicotine absorption in the gut is rapid, and it is highly permeable to the blood-brain barrier (Oldendorf et al. 1979; Allen and Lockman 2003). Indeed, inhaled nicotine enters the brain within seconds (Rose et al. 2010), which may account for the perniciousness of smoking as a vehicle for nicotine self-administration. Within the brain, nicotine binds with high affinity to heteromeric (mainly the  $\alpha 4\beta 2$  subtype) and homomeric (mainly the  $\alpha 7$  subtype) nicotinic acetylcholine receptors (nAChRs) (Changeux 2010; Bouzat and Sine 2018), which are the focus of this chapter. Signalling by the endogenous agonist acetylcholine via cerebral nAChRs is critically involved in attention, vigilance and cognition, as well as locomotion and reward mechanisms (Changeux 2010; Graef et al. 2011). Activation of nAChRs stimulates dopamine release in the basal ganglia, especially in the ventral striatum (Pradhan et al. 2002; Cumming et al. 2003), which almost certainly underlies the addictive potential of nicotine and tobacco smoking (Hogg et al. 2003). Furthermore, nAChRs play a major role in brain development (Hruska et al. 2009; Ross et al. 2010). Recently allosteric modulators of nAChRs are undergoing intensive consideration for drug development (Chatzidaki and Millar 2015; Wang and Lindstrom 2018).

Nicotinic receptors, in particular the  $\alpha 7$  subtype, are expressed by many classes of neurons and indeed by virtually all cell types of the brain, including astrocytes (Sharma and Vijayaraghavan 2001), microglia (De Simone et al. 2005; Suzuki et al. 2006), oligodendrocyte precursor cells (Sharma and Vijayaraghavan 2002) and endothelial cells (Hawkins et al. 2005). Perhaps consistent with this protean cellular distribution, there is evidence for dysfunction of nAChRs in diverse human neurological and psychiatric diseases (Jasinska et al. 2014; Bertrand et al. 2015; Dineley et al. 2015), which motivates the present search for optimal molecular imaging agents.

The nAChRs belong to the Cys-loop superfamily of pentameric ligand-gated ion channels, which also includes the serotonin 5-HT<sub>3</sub>,  $\gamma$ -aminobutyric acid (GABA<sub>A</sub> and GABA<sub>C</sub>) and glycine receptors, as has been reviewed in detail (Paterson and Nordberg 2000; Taly et al. 2009; Plested 2016). Functional nAChRs consist of

pentamers of homologous or heterologous subunits forming a central cation channel permeable for  $\text{Na}^+$ ,  $\text{K}^+$  and  $\text{Ca}^{2+}$  ions. There are at least 17 genes encoding the following subunits:  $\alpha 1$ – $10$ ,  $\beta 1$ – $4$ ,  $\delta$ ,  $\epsilon$  and  $\gamma$  (Karlin 2002), all of which occur in mammals except for the avian  $\alpha 8$  subunit. Each monomer possesses an extracellular N-terminal domain, four transmembrane helices and a small cytoplasmic region (Karlin 2002). Also, the 1.76 Å resolution X-ray structure of a nAChR homologue, the prokaryotic acetylcholine-binding protein (AChBP), has been reported (Brejc et al. 2001; Bourne et al. 2005; Hibbs et al. 2009), and a refined model of the membrane-associated nAChR from Torpedo electric organ based on 4 Å resolution electron microscopy data has been presented (Unwin 2005). An X-ray crystal structure is now available for the human  $\alpha 4\beta 2$  subtype, bound to nicotine (Giastas et al. 2018).

The consensus nomenclature for nAChRs is based on the predominant subunit composition of the receptor (Alexander et al. 2017). Although most functional receptors are heteromers, the  $\alpha 7$ – $10$  subunits form functional homomers *in vivo*. Most neuronal nAChRs contain  $\alpha$ - and  $\beta$ -subunits only ( $\delta$ ,  $\epsilon$  and  $\gamma$  are expressed in peripheral tissues). Of the various possible  $\alpha\beta$  permutations, nAChRs with six of the  $\alpha$ -subunits (2–7) and three of the  $\beta$ -subunits (2–4) have been identified in mammalian brain, with heteromeric  $\alpha 4\beta 2$  nAChRs predominating (Gotti et al. 2006). The next most abundant cerebral nAChR is the homomeric  $\alpha 7$  subtype, which is functionally distinct from the heteromeric nAChRs due to its lower affinity for the agonists acetylcholine and nicotine and higher affinity for  $\alpha$ -bungarotoxin, an antagonistic derived from snake venom. Indeed, radiolabelled  $\alpha$ -bungarotoxin serves admirably for selective  $\alpha 7$  autoradiography *in vitro* (Clarke et al. 1984). The  $\alpha 7$  nAChRs show relatively fast activation and have the highest permeability to  $\text{Ca}^{2+}$  of all nAChR subtypes, whereas the  $\alpha 4\beta 2$  hetero-oligomer is characterized by a high affinity for ACh and slow desensitization (Changeux 2010). A recent study indicates that heteromeric  $\alpha 7\beta 2$  nAChRs are also naturally present in the brain and are functionally distinct from the  $\alpha 7$  nAChR (Wu et al. 2016).

In general, activation of nAChRs requires cooperativity between subunits, with the agonist binding sites being located at subunit interfaces (Taylor et al. 1994). Activation of nAChR heteromers requires binding of two ACh molecules at orthosteric binding sites, which are formed within the hydrophilic extracellular domain from three peptide domains on the  $\alpha$ -subunit and three domains on the adjacent subunits ( $\beta$  or other) (Kalamida et al. 2007). Homomers have five acetylcholine binding sites, one between each  $\alpha$ - $\alpha$ -subunit interface (Millar and Harkness 2008).

As noted above, the nAChRs can shift between functionally distinct conformational states. Four such states have been identified: resting (R), activated (A) with rapid opening within 1 ms and low affinity ( $\mu\text{M}$  to  $\text{mM}$ ) for agonists, and two desensitized, which are closed channel states refractory to opening for intervals lasting ms (I, insensitive) or minutes (D, desensitized), although still possessing high affinity (pM to nM) for receptor agonists (Decker et al. 2000; Auerbach 2015). Thus, binding of ligands either at the orthosteric site or any of several allosteric sites alters the functional state of nAChRs by favouring particular conformational states, consequently modifying the equilibrium between the four states of the receptor (Taly

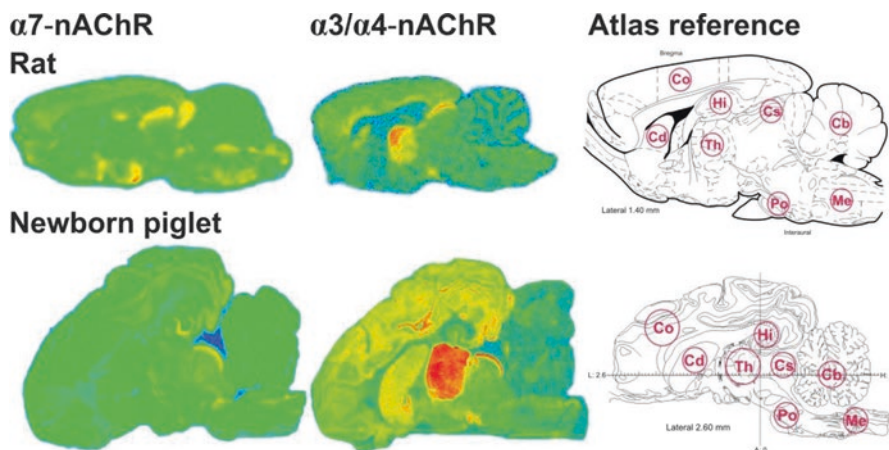


et al. 2009). Activation of nAChRs increases cation influx. This has a spectrum of consequences, including (i) immediate effects, such as neurotransmitter release; (ii) short-term effects, such as receptor desensitization and recovery; and (iii) long-lasting adaptive effects, such as neuroprotection or brain plasticity via altered gene expression (Radcliffe and Dani 1998; Leonard 2003; Shen and Yakel 2009). The particular pathways of intracellular signalling evoked by activation of nAChRs are complex and cell-specific (Frazier et al. 1998; Schilström et al. 2000; Berg and Conroy 2002).

Neuronal nAChRs in the brain are localized at post-, pre-, peri- and extrasynaptic sites of cholinergic or other neurones, affording multiple ways in which to modulate brain function. Prolonged exposure of nAChRs to agonists (e.g. nicotine) results in upregulation of functional receptors (i.e. an increase in the number of [<sup>3</sup>H]nicotine binding sites) rather than the downregulation typically seen with G-protein-coupled metabotropic receptors. This upregulation is thought to occur via altered nAChR turnover, resulting in increased insertion of receptors into the cell membrane, or decreased removal via altered endocytotic trafficking or degradation rates (Peng et al. 1994; Darsow et al. 2005). The predominant nAChRs in the brain,  $\alpha 4\beta 2$  and  $\alpha 7$ , seem particularly sensitive to this form of post-translational regulation. In analogy to dopamine receptors (Cumming 2011), multiple affinity states of nAChRs are differentially sensitive to agonist and antagonist ligands. Exposure to acetylcholine, other agonists and pharmacological chaperones can alter the affinity states of the nAChRs receptor to influence receptor translocation and modulate the cell surface expression of nAChR subtypes (Darsow et al. 2005; Kumari et al. 2008; Govind et al. 2009; Lester et al. 2009; Crespi et al. 2018).

The total concentration of nAChRs in membrane homogenates from rodent brain is between 8 and 15 pmol/g tissue (Wang et al. 2011), and in human brain up to 10 pmol/g tissue (Shimohama et al. 1985; Marutle et al. 1998), which is similar to the abundance of dopamine receptors in striatum (Cumming 2011). The predominant heteromeric  $\alpha 4\beta 2$  nAChRs receptors account for some 80% of the total abundance of brain nAChRs (Wang et al. 2011). For the less abundant  $\alpha 7$  nAChRs, a  $B_{\max}$  of 5 pmol/g tissue has been described in mice (Whiteaker et al. 1999), while levels of [<sup>125</sup>I] $\alpha$ -bungarotoxin binding in the temporal cortex were only 1 pmol/g tissue for human and monkey (Gotti et al. 2006). Notably, these studies are based on tissue homogenates. However, the native environment of nAChRs is important for accurate quantitation of nAChRs in the brain (Wang et al. 2011); a very similar phenomenon has been described for other receptors, e.g. dopamine (Cumming 2011) and opioid receptors (Quelch et al. 2014), suggesting that many receptors may be lost during the preparation of membranes or that not all receptors are externalized in the plasma membrane. Therefore, to predict the potential of nAChR radioligands for molecular imaging, binding to nAChRs should in general be tested by autoradiographic analysis of sections from frozen, intact brain.

The relatively few quantitative autoradiographic studies consistently report higher nAChR densities than homogenate-based studies. Autoradiographic studies with ligands for heteromeric  $\alpha 4\beta 2$  nAChRs characteristically reveal particularly intense binding in the thalamus (Fig. 18.1). Using [<sup>18</sup>F]FNEP, the  $B_{\max}$  for



**Fig. 18.1** Representative autoradiograms of rat (*upper row*) and neonate pig brains (*lower row*) obtained with an  $\alpha 7$  nAChR ligand ( $[^{125}\text{I}]\alpha$ -bungarotoxin) and the semi-selective  $\alpha 4\beta 2$  ligand  $[^3\text{H}]\text{epibatidine}$ . For anatomic reference, the right-hand side of the figure shows comparable sagittal planes modified from the rat brain atlas (Paxinos and Watson 1998) and the pig brain atlas (Felix et al. 1999). Structures marked in the references atlases are *Co* cerebral cortex, *Cd* caudate nucleus, *Cs* superior colliculus, *Cb* cerebellum, *Hi* hippocampus, *Th* thalamus, *Po* pons, *Me* medulla oblongata

heteromeric nAChRs in human thalamus autoradiograms was 20 pmol/g tissue (Gatley et al. 1998), which is comparable to the density of binding sites of 2- $[^{18}\text{F}]\text{fluoro-A-85380}$  in porcine thalamus, i.e. 46 pmol/g tissue (Deuther-Conrad et al. 2006). In quantitative receptor autoradiographic studies with the  $\alpha 7$  selective ligands  $[^3\text{H}]\text{MLA}$  and  $[^{125}\text{I}]\alpha$ -bungarotoxin, densities were as high as 40 pmol/g tissue in mouse brain (Whiteaker et al. 1999), whereas other studies with  $[^{125}\text{I}]\alpha$ -bungarotoxin have indicated densities of 4–20 pmol/g tissue in rat and 2–10 pmol/g tissue in neonate pig (Hoffmeister et al. 2011). Across species, autoradiographic studies have revealed a characteristic distribution pattern for  $\alpha 7$  nAChRs (Fig. 18.1), which is rather diffuse, but with focally high density in the hippocampus, the colliculi and the hypothalamus (except in newborn piglet), with moderate radiotracer binding in thalamus and low expression in the cerebellum (Breese et al. 1997; Whiteaker et al. 1999; Hoffmeister et al. 2011). The regional distribution of the  $\alpha 7$  nAChR ligand  $[^3\text{H}]\text{AZ11637326}$  in rat brain by autoradiography *ex vivo* (in which the tracer had been administered while the animal was alive) was consistent with autoradiography findings *in vitro* (Maier et al. 2011).

Molecular brain imaging of nAChRs in general refers to the use of radiolabelled receptor ligands, although optical imaging has been used to investigate the cholinergic system (Prakash and Frostig 2005). Furthermore, clinical imaging of  $\alpha 4\beta 2$  nAChR in Alzheimer's disease is reviewed elsewhere (see Sabri et al. in PET and SPECT in Neurology). Therefore, the current review is focussed primarily, albeit not exclusively, on preclinical aspects of PET and SPECT brain imaging of the

nAChR in vivo, along with a presentation on instrumentation and radiotracer development. This review is updated and extended from an earlier chapter on the same topic (Brust et al. 2014).

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## 18.2 Advances in Animal PET and SPECT Technology

As noted above, studies of nAChRs abundance in vitro and ex vivo use selective radioligands in conjunction with quantitative autoradiography. However, molecular imaging with positron emission tomography (PET) or single-photon emission computed tomography (SPECT) enables the detection of neuroreceptors in the living brain. Contrary to autoradiography in vitro, imaging procedures in vivo allow for longitudinal studies in individual animals, thereby reducing intersubject variability and allowing intervention or challenge studies. Animal PET studies using clinical scanners, most having a spatial resolution of approximately 5 mm, are barely adequate for resolution of structures within mouse brain, which measures only 10 mm along its longest axis.

Although dedicated small-animal PET systems such as the Focus series (Siemens), microPET P4 (Concorde Microsystems) or ClearPET (Raytest) enable microPET imaging in rodents and small non-human primates with high resolution and sensitivity, these instruments are no longer commercially available. However, current efforts aim to improve the performance of stand-alone small-animal PET scanners by using detector systems approaching the physical limits of spatial resolution (circa 1 mm) with optimized reconstruction algorithms (Yang et al. 2016). Other approaches are to improve the spatial resolution of commercially available systems by optimizing the arrangement of detectors (Bolwin et al. 2017), or applying new detector material (Abbaszadeh and Levin 2017). Commercial stand-alone PET systems designed for small-animal imaging include the beta-CUBE (Molecubes) or VECTor (MILabs). Contemporary small-animal PET scanners increasingly combine the PET and MRI imaging modalities in one instrument. Sequential PET and MR measurements can be realized with instruments such as the nanoScan PET/MRI systems (Mediso). The PET CLIP-ON system (MR Solutions). The PET/MR 3T system (Bruker) or the PET INSERT system (MR Solutions) give simultaneous multimodal acquisitions.

Despite spatial resolution approaching 1 mm, images from small-animal PET and SPECT instruments suffer from a lack of anatomic information. In common practice, the emission images are registered to digitized brain atlases, based on histology or magnetic resonance imaging (MRI) atlases for rodent brain (Jupp et al. 2007; Rominger et al. 2010). Contemporary multimodal imaging systems combine small-animal PET with SPECT, X-ray computed tomography (CT) and/or MRI. PET-CT presents a great advantage for brain studies in that a high-resolution structural brain image in perfect registration with the PET image is obtained for each individual animal, without resorting to some standard atlas. In addition, the CT scan serves to correct the PET images for attenuation by tissue, thus providing

absolute quantitation of radioactivity concentrations in the brain without requiring an additional time-consuming transmission scan. MRI offers better tissue contrast than CT, but the combining of PET and MRI instrumentation initially presented a greater technical challenge.

Initially, the instrumentation for simultaneous small-animal imaging entailed the development of PET inserts, with special efforts made by research teams in academia and industry to improve the PET detector technology with regard to MRI compatibility. An early prototype PET/MRI scanner developed at the University of Cambridge (Lucas et al. 2007) was followed by the commercial PET/MRI system for rodents developed by Mediso; its 1 tesla permanent magnet limits MRI applications but offers great flexibility for animal PET studies. PET inserts using silicone-based photoelectron multipliers are compatible with magnetic fields (Wehner et al. 2015), and a recently developed PET insert can operate within the bore of a 7 T magnet (Thiessen et al. 2016). Likewise, a SPECT camera that can be placed within an MRI magnet has recently been developed for small-animal studies (Meier et al. 2011). Whereas PET attenuation correction is usually obtained by CT scanning, MR-based attenuation correction has been demonstrated for PET and SPECT imaging using clinical scanners (Marshall et al. 2011) and has recently successfully been proven for small-animal studies (Kranz et al. 2014, 2016; Sattler et al. 2014). Despite considerable progress, simultaneous PET/MRI scanners have lower detection sensitivity in comparison with dedicated PET scanners (Cabello and Ziegler 2018; Hallen et al. 2018). However, in recent years the focus has shifted towards the development of fully integrated scanners (Ko et al. 2016; Parl et al. 2019).

Most animal PET studies are confounded by the need to use anaesthesia, which can profoundly alter radiotracer pharmacokinetics. Monkeys are trained to tolerate head fixation during PET recordings lasting as long as 30 min (Sandiego et al. 2013). The “rat conscious animal PET” (ratCAP) with a head-mounted PET detector was developed at Brookhaven for imaging in awake, behaving rats (Schulz et al. 2011). While this technology has been slow to mature, there is a recent report of FDG PET recordings in awake, behaving chickens and rats (Gold et al. 2018).

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### 18.3 PET and SPECT Radioligands Targeting nAChR

Many autoradiographic studies of nAChRs in human post-mortem brain specimens have employed rather non-selective agonist ligands such as [<sup>3</sup>H]acetylcholine, [<sup>3</sup>H]nicotine, [<sup>3</sup>H/<sup>125</sup>I]epibatidine or [<sup>3</sup>H]cytisine, or alternately the antagonist ligand [<sup>3</sup>H/<sup>125</sup>I]α-bungarotoxin (Paterson and Nordberg 2000). These studies suggested the existence of at least three native receptor subtypes, i.e. α-bungarotoxin binding at homomeric α7 nAChRs and acetylcholine/nicotine binding mainly at heteromeric α4β2 nAChRs, plus a relatively small population of heteromeric receptors containing the α3 subunit. Given the predominance of α4β2 and the α7 subtypes in the brain as documented above, they have presented the main targets for molecular imaging of nAChRs. The available information on the distribution, density and functional role of other subtypes in the brain is relatively sparse (Gotti et al. 2006; Sharma and

Vijayaraghavan 2008; Zoli et al. 2015). Furthermore, the current emphasis on  $\alpha 4\beta 2$  and the  $\alpha 7$  subtypes as targets for pharmaceutical development may account for the paucity of selective high-affinity drugs for the other subtypes (Gündisch and Eibl 2011) due to a kind of circularity.

### 18.3.1 Radioligands for $\alpha 4\beta 2$ nAChRs

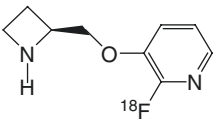
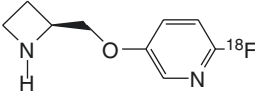
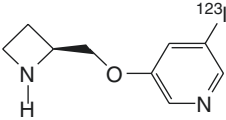
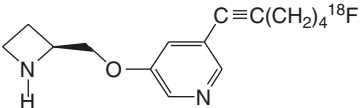
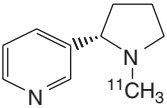
Reviews summarizing the history of ligand development for the nAChRs (Sihver et al. 2000a, b; Ding and Fowler 2005; Horti and Villemagne 2006; Horti et al. 2010) have helped galvanize research groups to generate new lead compounds for molecular imaging and therapeutics. Selectivity and high affinity (nM) in vitro are necessary but not sufficient for successful visualization of nAChRs. In general, PET and SPECT tracers, if they are to be successful, must fulfil a multitude of additional criteria. Chief among these are (1) low non-specific binding and absence of brain-penetrating radiometabolites, (2) rapid clearance from non-specific brain regions and plasma to reduce background in the target tissue, (3) high membrane permeability, (4) high permeability and low efflux at the blood-brain barrier (BBB) and (5) attainment of equilibrium binding with a tolerable time interval. Settling upon optimal tracers is often a matter of fierce competition.

The vast majority of novel tracers for  $\alpha 4\beta 2$  nAChRs derive from three compounds: nicotine, epibatidine and 3-pyridyl ether. Seven of these tracers have so far been used successfully to image  $\alpha 4\beta 2$  receptors in human brain: *S*-[ $^{11}\text{C}$ ]nicotine (Nordberg 1993); the two halogen-substituted derivatives of A-85380, namely, 2-[ $^{18}\text{F}$ ]fluoro-A-85380 (Kimes et al. 2003) and 6-[ $^{18}\text{F}$ ]fluoro-A-85380 (Ding et al. 2000a; Horti et al. 2000); the two epibatidine derivatives (–)-[ $^{18}\text{F}$ ]flubatine and (+)-[ $^{18}\text{F}$ ]flubatine (Sabri et al. 2018; Tiepolt et al. 2018); [ $^{18}\text{F}$ ]nifene (Betthausen et al. 2017); and [ $^{18}\text{F}$ ]XTRA (Coughlin et al. 2018c). Receptor-ligand interactions frequently entail stereoselective features (Smith and Jakobsen 2007), as has been formally demonstrated for the case of nAChRs in human PET studies with the two stereoisomers of [ $^{11}\text{C}$ ]nicotine (Nordberg et al. 1991, 1992), and also several of the radioligands discussed below. A summary of the compounds that have been investigated for imaging of  $\alpha 4\beta 2$  nAChRs with PET and SPECT, their associated references, binding affinities and results of biodistribution ex vivo and PET imaging studies is given in Table 18.1.

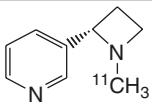
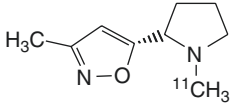
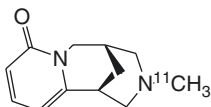
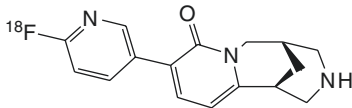
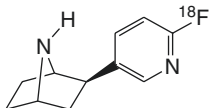
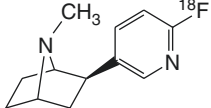
#### 18.3.1.1 Nicotine Derivatives

The tracer *S*-[ $^{11}\text{C}$ ]nicotine, one of the very first positron-emitting receptor ligands (Maziere et al. 1976), was initially developed to investigate the biodistribution of nicotine in the context of tobacco addiction and for investigation of diabetes insipidus and for insecticide research (Soloway 1976; Gündisch 2000). With the advent of PET, *S*-[ $^{11}\text{C}$ ]nicotine was tested for imaging nAChRs in human brain (Nordberg 1993). However, co-administration of unlabelled nicotine failed to displace much of the radioligand, indicating that the PET signal did not sensitively reveal specific binding to  $\alpha 4\beta 2$  nAChRs (Nyböck et al. 1994); cerebral *S*-[ $^{11}\text{C}$ ]nicotine uptake

**Table 18.1** Molecular structures, IUPAC names, binding affinities and results of biodistribution ex vivo and PET imaging studies in vivo of the cerebral binding of  $\alpha 4\beta 2$  nAChR-selective radioligands

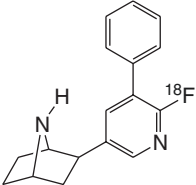
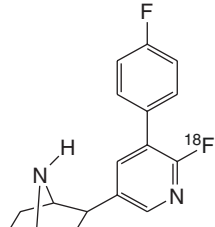
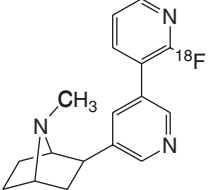
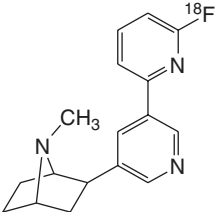
	Main findings
 <p>2-<sup>[18F]</sup>fluoro-A-85380 ((S)-3-(azetidin-2-ylmethoxy)-2-<sup>[18F]</sup>fluoropyridine)</p>	<p>Thalamic uptake of radioactivity in rat and baboon peaked at 60 min. In humans, cerebral uptake pattern was consistent with the known distribution of <math>\alpha 4\beta 2</math> nAChRs. The total distribution volume was significantly higher in smokers than in non-smokers, except in the thalamus. Radioactivity in the brain reached steady state by 6 h. <math>K_i = 0.061</math> nM (Dollé et al. 1999; Kimes et al. 2003, 2008; Mukhin et al. 2008)</p>
 <p>6-<sup>[18F]</sup>fluoro-A-85380 ((S)-3-(azetidin-2-ylmethoxy)-6-<sup>[18F]</sup>fluoropyridine)</p>	<p>In baboon dynamic PET, faster peak uptake and clearance as well as higher thalamus-to-cerebellum ratios than obtained for 2-<sup>[18F]</sup>fluoro-A-85380. <math>K_i = 0.025</math> nM (Ding et al. 2000a, Horti et al. 2000)</p>
 <p>5-<sup>[123/125I]</sup>iodo-A-85380 (((S)-3-(azetidin-2-ylmethoxy)-5-<sup>[123/125I]</sup>iodopyridine)</p>	<p>In rhesus dynamic PET, regional distribution in brain consistent with the known nAChR distribution pattern. Relatively slow kinetics, with maximal binding ratios at more than 4 h. In baboon, significant displacement of radioactivity from cerebellum by cytosine, indicating this region inappropriate as reference region. <math>K_d</math> (rat) = 10 pM, <math>K_d</math> (human) = 12 pM (Chefer et al. 1998; Fujita et al. 2000)</p>
 <p><sup>[18F]</sup>ZW-104 ((S)-3-(azetidin-2-ylmethoxy)-5-(6-<sup>[18F]</sup>fluorohex-1-yn-1-yl)pyridine)</p>	<p>In baboon dynamic PET, rather slow kinetics in thalamus. High affinity towards multiple <math>\beta 2</math>-containing nAChR subtypes. <math>K_i = 0.21</math> nM (Valette et al. 2009)</p>
 <p>S-<sup>[11C]</sup>nicotine ((S)-3-(1-<sup>[11C]</sup>methylpyrrolidin-2-yl)pyridine)</p>	<p>In human and rhesus monkey dynamic PET studies, binding was less selective for nAChR subtypes than in the case of <sup>[11C]</sup>MPA. <math>K_i = 10</math> nM (Sihver et al. 1999b)</p>

**Table 18.1** (continued)

	Main findings
 <p>[<sup>11</sup>C]MPA (<i>S</i>)-3-(1-[<sup>11</sup>C]methylazetididin-2-yl)pyridine)</p>	Specific binding in rhesus monkey proven by nicotine displacement in dynamic PET studies, but specific binding was found to be rather low. $K_d = 0.011$ nM (Sihver et al. 1999b)
 <p>[<sup>11</sup>C]ABT418 (<i>S</i>)-3-(1-[<sup>11</sup>C]methyl-5-(1-methylpyrrolidin-2-yl)isoxazole)</p>	In rhesus dynamic PET, low uptake and rapid washout, with no evidence for displacement by unlabelled ABT-418. Increased uptake following <i>S</i> (-)-nicotine pretreatment. $K_i = 3$ nM. (Armeric et al. 1994; Valette et al. 1997; Sihver et al. 1999b)
 <p><i>N</i>-[<sup>11</sup>C]methylcytisine ([<sup>11</sup>C]caulophylline) ((1<i>R</i>,5<i>S</i>)-3-[<sup>11</sup>C]methyl-2,3,4,5-tetrahydro-1,5-methanopyrido[1,2-<i>d</i>][1,4]diazepin-7(<i>1H</i>)-one)</p>	No evidence of specific binding in rhesus monkey brain by dynamic PET. $K_i = 5.7$ nM (Valette et al. 1997; Imming et al. 2001)
 <p>[<sup>18</sup>F]FPpCYT (1<i>R</i>,5<i>S</i>)-8-(6-[<sup>18</sup>F]fluoropyridin-3-yl)-2,3,4,5-tetrahydro-1,5-methanopyrido[1,2-<i>d</i>][1,4]diazepin-7(<i>1H</i>)-one)</p>	In rat biodistribution study, low (0.3% ID/g) and uniform brain uptake, with little evidence of specific binding. $K_i = 24$ nM (Roger et al. 2003)
 <p>[<sup>18</sup>F]NFEP (1<i>R</i>,2<i>R</i>,4<i>S</i>)-2-(6-[<sup>18</sup>F]fluoropyridin-3-yl)-7-azabicyclo[2.2.1]heptane)</p>	High brain uptake by mouse biodistribution and baboon PET studies and clear indication of specific binding. Applications in humans limited by high toxicity. $K_{app} = 0.02$ nM (Ding et al. 1996, 1999; Liang et al. 1997; Villemagne et al. 1997; Dolci et al. 1999)
 <p>[<sup>18</sup>F]<i>N</i>-methyl-NFEP (1<i>R</i>,2<i>R</i>,4<i>S</i>)-2-(6-[<sup>18</sup>F]fluoropyridin-3-yl)-7-methyl-7-azabicyclo[2.2.1]heptane)</p>	In baboon dynamic PET, higher peak uptake in all brain regions than for [ <sup>18</sup> F]NFEP. Despite milder toxicity than [ <sup>18</sup> F]NFEP, evidence against safe use in humans. $K_d = 0.028$ nM (Ding et al. 1999)

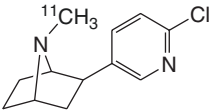
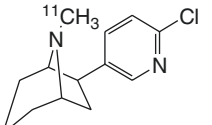
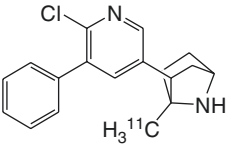
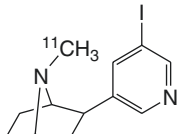
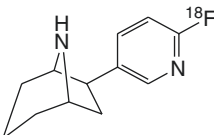
(continued)

**Table 18.1** (continued)

	Main findings
 <p><b>[<sup>18</sup>F]FphEP</b>        ((1<i>R</i>,2<i>R</i>,4<i>S</i>)-2-(6-[<sup>18</sup>F]fluoro-5-phenylpyridin-3-yl)-7-azabicyclo[2.2.1]heptane)</p>	<p>In baboon dynamic PET; more favourable kinetics than 2-[<sup>18</sup>F]fluoro-A-85380. Peak uptake at 20 min, but no evidence for displacement by nicotine challenge. <math>K_d = 0.66</math> nM (Roger et al. 2006; Valette et al. 2007)</p>
 <p><b>[<sup>18</sup>F]F<sub>2</sub>PhEP</b>        ((1<i>R</i>,2<i>R</i>,4<i>S</i>)-2-(6-[<sup>18</sup>F]fluoro-5-(4-fluorophen-1-yl)pyridin-3-yl)-7-azabicyclo[2.2.1]heptane)</p>	<p>Dynamic PET in baboon did not indicate reduction in brain distribution volume following pretreatment with nicotine. <math>K_i = 0.029</math> nM (Valette et al. 2007)</p>
 <p><b>[<sup>18</sup>F]XTRA, JHU86428</b>        ((1<i>R</i>,2<i>R</i>,4<i>S</i>)-2-(2'-[<sup>18</sup>F]fluoro-[3,3'-bipyridin]-5-yl)-7-methyl-7-azabicyclo[2.2.1]heptane)</p>	<p>Dynamic PET showed peak activity in baboon thalamus at 75 min after bolus, requiring several hours for steady-state measurement. <math>K_i = 0.058</math> nM (Horti and Wong 2009; Coughlin et al. 2018c)</p>
 <p><b>[<sup>18</sup>F]AZAN, JHU87522</b>        ((1<i>R</i>,2<i>R</i>,4<i>S</i>)-2-(6-[<sup>18</sup>F]fluoro-[2,3'-bipyridin]-5'-yl)-7-methyl-7-azabicyclo[2.2.1]heptane)</p>	<p>Baboon dynamic PET showed rapid brain kinetics, favourable metabolic profile and high <math>BP_{ND}</math>, reliably measured with 90 min scans. However, part of in vivo binding could be related to other nAChRs with <math>\beta</math>-subunits. <math>K_i = 0.26</math> nM (Gao et al. 2008b, Kuwabara et al. 2012)</p>

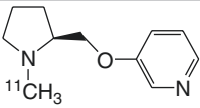
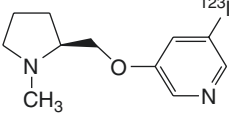
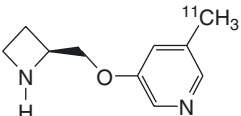
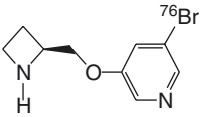
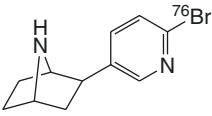
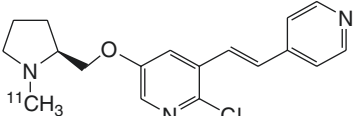


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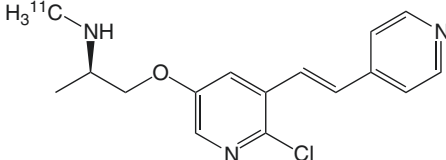
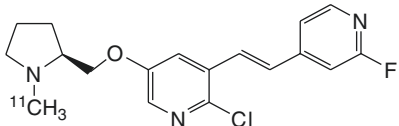
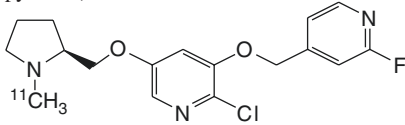
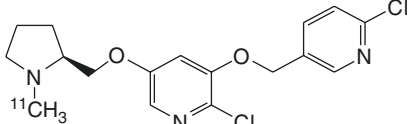
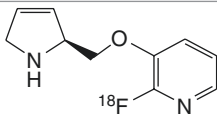
	Main findings
 <p><i>N</i>-[<sup>11</sup>C]methyl-epibatidine ((1<i>R</i>,2<i>R</i>,4<i>S</i>)-2-(6-chloropyridin-3-yl)-7-[<sup>11</sup>C]methyl-7-azabicyclo[2.2.1]heptane)</p>	Different kinetics of brain uptake and washout seen for the stereoisomers in rats and mice. Dynamic PET showed high enrichment of radioactivity in the thalamus of the pig, but steady-state not attained during 60 min scans. Applications in humans limited by high toxicity. $K_i = 0.027$ nM (Patt et al. 1999; Spang et al. 2000)
 <p><i>N</i>-[<sup>11</sup>C]methyl-homoepibatidine ((1<i>S</i>,5<i>R</i>,6<i>R</i>)-6-(6-chloropyridin-3-yl)-8-[<sup>11</sup>C]methyl-8-azabicyclo[3.2.1]octane)</p>	In pig dynamic PET, the (–)-enantiomer showed a regional distribution high accumulation in the thalamus consistent with representative for the $\alpha 4\beta 2$ nAChR, which could be displaced cytosine. Distribution of the (+)-enantiomer non-specific. Applications in humans limited by high toxicity. $K_i = 0.13$ nM (Malpass et al. 2001; Patt et al. 2001)
 <p>2-(6-chloro-5-phenylpyridin-3-yl)-7-[<sup>11</sup>C]methyl-7-azabicyclo[2.2.1]heptane</p>	Rat biodistribution study showed high displaceable binding. Dynamic baboon PET showed rapid peak in thalamus but increasing ratio relative to cerebellum over at least 2 h. $K_i = 0.032$ nM (Huang et al. 2004, 2005)
 <p>[<sup>11</sup>C]NMI-EPB ((1<i>R</i>,2<i>R</i>,4<i>S</i>)-2-(5-iodopyridin-3-yl)-7-[<sup>11</sup>C]methyl-7-azabicyclo[2.2.1]heptane)</p>	Baboon dynamic PET showed high uptake and displaceable thalamic binding. Higher uptake and faster kinetics than 2-[ <sup>18</sup> F]fluoro-A-85380; however, (–)-enantiomer did not reach steady state within 90 min post-injection. $K_i = 0.068$ nM (Ding et al. 2006; Gao et al. 2008a)
 <p>[<sup>18</sup>F]flubatine (((1<i>R</i>,2<i>R</i>,4<i>S</i>)-2-(6-[<sup>18</sup>F]fluoropyridin-3-yl)-7-azabicyclo[2.2.1]heptane))</p>	Higher uptake of radioactivity in mouse than for 2-[ <sup>18</sup> F]fluoro-A-85380. Binding equilibrium of the (–)-enantiomer was reached significantly earlier (~60 min p.i.) than that of the (+)-enantiomer. $K_i(+)$ = 0.064 nM, $K_i(-)$ = 0.112 nM (Brust et al. 2008; Deuther-Conrad et al. 2008)

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**Table 18.1** (continued)

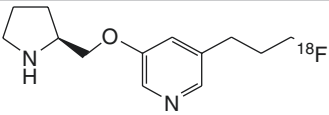
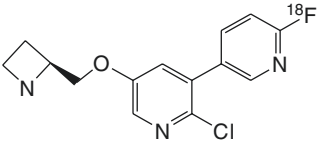
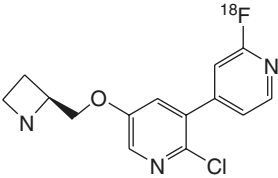
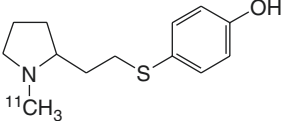
	Main findings
 <p><sup>11</sup>C]A-84543 ((S)-3-((1-[<sup>11</sup>C]methylpyrrolidin-2-yl)methoxy)pyridine)</p>	<p>Mouse biodistribution showed high brain uptake and a distribution consistent with the density of <math>\alpha 4\beta 2</math> nAChRs. <math>K_i = 0.38</math> nM (Kassiou et al. 1998)</p>
 <p>5-[<sup>123</sup>I]iodo-A-84543 ((S)-3-((1-methylpyrrolidin-2-yl)methoxy)-5-[<sup>123</sup>I]iodopyridine)</p>	<p>Nearly homogeneous distribution in mouse brain, likewise seen in baboon dynamic SPECT. <math>K_i = 0.016</math> nM (Fan et al. 2001; Henderson et al. 2004)</p>
 <p>5-[<sup>11</sup>C]methyl-A-85380, 5-MA ((S)-3-((azetidin-2-yl)methoxy)-5-[<sup>11</sup>C]methylpyridine)</p>	<p>Mouse biodistribution consistent with <math>\alpha 4\beta 2</math> nAChRs. Rhesus monkey dynamic PET showed high non-specific binding. <math>K_i = 0.27</math> nM (Iida et al. 2004)</p>
 <p>[<sup>76</sup>Br]BAP ((S)-3-((azetidin-2-yl)methoxy)-5-[<sup>76</sup>Br]bromopyridine)</p>	<p>In comparison with rat biodistribution, baboon dynamic PET showed higher non-specific binding with nicotine or cystine displacement. <math>K_i = 0.023</math> nM (Sihver et al. 1999a)</p>
 <p>[<sup>76</sup>Br]BrPH ((1R,2R,4S)-2-(6-[<sup>76</sup>Br]bromopyridin-3-yl)-7-azabicyclo[2.2.1]heptane)</p>	<p>Rat biodistribution and baboon dynamic PET showed binding to nAChRs but without subtype selectivity. <math>K_d = 0.008</math> nM (Kassiou et al. 2002)</p>
 <p>[<sup>11</sup>C]Me-p-PVC ((S,E)-2-chloro-5-((1-[<sup>11</sup>C]methylpyrrolidin-2-yl)methoxy)-3-(2-(pyridin-4-yl)vinyl)pyridine)</p>	<p>Rapid accumulation in mice ex vivo, and rhesus PET studies, thus quantifiable with 2 h recordings, but <math>BP_{ND}</math> slightly lower than for 2-[<sup>18</sup>F]fluoro-A-85380. <math>K_i = 0.028</math> nM (Brown et al. 2004)</p>

**Table 18.1** (continued)

	Main findings
 <p><math>[^{11}\text{C}]p\text{-PVP-MEMA}</math>, (<i>R,E</i>)-1-(6-chloro-5-(2-(pyridin-4-yl)vinyl)pyridin-3-yloxy)-<i>N</i>-<math>[^{11}\text{C}]</math>methylpropan-2-amine.</p>	<p>In dynamic PET studies in baboon, <math>[^{11}\text{C}]p\text{-PVP-MEMA}</math> entered the brain rapidly with a peak concentration of 0.5% ID/mL 2 min p.i. but with a low thalamus/cerebellum ratio of about 1 at 1 h p.i., possibly due to low metabolic stability of the parent compound. <math>K_i = 0.077</math> nM (Dolle et al. 2008)</p>
 <p><math>[^{11}\text{C}]</math>JHU85270 (<i>S,E</i>)-2-chloro-3-(2-(2-fluoropyridin-4-yl)vinyl)-5-((1-<math>[^{11}\text{C}]</math>methylpyrrolidin-2-yl)methoxy)pyridine)</p>	<p>In baboon dynamic PET, thalamic <math>\text{BP}_{\text{ND}}</math> lower than that of 2-<math>[^{18}\text{F}]</math>fluoro-A-85380 due to rapid metabolism to brain-penetrating metabolites. <math>K_i(\text{JHU85270}) = 0.09</math> nM, <math>K_i(\text{JHU85208}) = 0.05</math> nM, <math>K_i(\text{JHU85157}) = 0.02</math> nM (Gao et al. 2009)</p>
 <p><math>[^{11}\text{C}]</math>JHU85208 (<i>S</i>)-2-chloro-3-((2-fluoropyridin-4-yl)methoxy)-5-((1-<math>[^{11}\text{C}]</math>methylpyrrolidin-2-yl)methoxy)pyridine)</p>	
 <p><math>[^{11}\text{C}]</math>JHU85157 (<i>S,E</i>)-2-chloro-3-(2-(6-chloropyridin-3-yl)vinyl)-5-((1-<math>[^{11}\text{C}]</math>methylpyrrolidin-2-yl)methoxy)pyridine)</p>	
 <p><math>[^{18}\text{F}]</math>nifene (<i>S</i>)-3-((2,5-dihydro-1<i>H</i>-pyrrol-2-yl)methoxy)-2-<math>[^{18}\text{F}]</math>fluoropyridine)</p>	<p>In rhesus dynamic PET, fast kinetics with peak thalamic binding at less than 10 min, thalamus to cerebellum ratio of ~2. <math>K_i = 0.5</math> nM (Pichika et al. 2006)</p>

(continued)

**Table 18.1** (continued)

	Main findings
 <p>[<sup>18</sup>F]nifrolidine (<i>S</i>)-3-((2,5-dihydro-1<i>H</i>-pyrrol-2-yl)methoxy)-5-(3-[<sup>18</sup>F]fluoroprop-1-yl)pyridine)</p>	In rhesus dynamic PET, good labelling of thalamus, with slightly faster kinetics than for 2-[ <sup>18</sup> F]fluoro-A-85380, with maximal binding at 70 min, and plateau thalamus-to-cerebellum ratio of 1.7 at 2 h. $K_i = 0.8$ nM (Chattopadhyay et al. 2005)
 <p>[<sup>18</sup>F]NIDA52189 (<i>S</i>)-2-chloro-6'-[<sup>18</sup>F]fluoro-5-((1-methylazetidin-2-yl)methoxy)-3,3'-bipyridine)</p>	In rhesus dynamic PET, distribution consistent with $\alpha 4\beta 2$ nAChRs, but $BP_{ND}$ 2.5 times higher than for 2-[ <sup>18</sup> F]fluoro-A-85380. $K_d = 0.005$ nM (Zhang et al. 2004)
 <p>[<sup>18</sup>F]NIDA522131 (<i>S</i>)-2-chloro-2'-[<sup>18</sup>F]fluoro-5-((1-methylazetidin-2-yl)methoxy)-3,4'-bipyridine)</p>	In rhesus dynamic PET, $BP_{ND}$ in thalamus 3–4-fold greater than with 2-[ <sup>18</sup> F]fluoro-A-85380 but at least 8 h recordings required for stable estimation. $K_d = 0.005$ nM (Chefer et al. 2008)
 <p>[<sup>11</sup>C]SIB-1553A (4-((2-(1-methylpyrrolidin-2-yl)ethyl)thio)phenol)</p>	In rat biodistribution study, 0.5% ID/g at 10 min and 0.25% ID/g at 30 min, but no evidence of specific binding discernible against high background. SIB-1553A was more potent than nicotine on $\alpha 2\beta 4$ ( $EC_{50} = 0.59$ $\mu$ M, (nicotine) = 1.95 $\mu$ M) and $\alpha 3\beta 4$ ( $EC_{50} = 1.10$ $\mu$ M, (nicotine) = 7.50 $\mu$ M), but no effect on $\alpha 4\beta 2$ and $\alpha 3\beta 2$ receptors was observed potentially explaining the absence of specific binding in brain (Sobrio et al. 2008)

proved mainly to be determined by blood flow, rather than local abundance of nAChRs in vivo (Gündisch 2000), this in keeping with its considerable lipophilicity. Indeed, labelled nicotine serves admirably as a cerebral blood flow tracer.

### 18.3.1.2 Cytisine Derivatives

The nAChR agonist [<sup>3</sup>H]cytisine is a useful radioligand for characterization of  $\alpha 4\beta 2$  nAChRs in vitro. However, the cytisine derivatives [<sup>11</sup>C]ABT-418 and *N*-[<sup>11</sup>C]methylcytisine (Valette et al. 1997) failed in vivo due to their low cerebral uptake and

rapid washout. Another derivative, [ $^{11}\text{C}$ ]MPA, showed high-affinity binding to  $\alpha 4\beta 2$  nAChRs ( $\sim 10$ – $100$ -fold higher than [ $^{11}\text{C}$ ]ABT-418 and *S*-[ $^{11}\text{C}$ ]nicotine) in membranes from rat forebrain (Sihver et al. 1998). Furthermore [ $^{11}\text{C}$ ]MPA showed rapid uptake into monkey brain, with similar permeability as [ $^{11}\text{C}$ ]ABT-418 and *S*-[ $^{11}\text{C}$ ]nicotine. Pre-administration of unlabelled *S*-nicotine (0.02 mg/kg) decreased the peak uptake of [ $^{11}\text{C}$ ]MPA in monkey brain by about 20%, indicating the presence of some specific binding to nAChRs (Sihver et al. 1999b), but no further studies have been published with this ligand. [ $^{18}\text{F}$ ]fluoropyridinylcytisine was developed as another candidate radioligand for  $\alpha 4\beta 2$  nAChR imaging. However, its distribution in living rat brain did not match the regional distribution of nAChRs, and blocking studies with nicotine failed to demonstrate specific binding of this tracer (Roger et al. 2003).

### 18.3.1.3 Epibatidine Derivatives

Epibatidine, an alkaloid from the skin of the Ecuadoran poison arrow frog *Epipedobates anthonyi*, has long been known for its high-affinity binding to heteromeric nAChRs (Daly 1998). It has considerable toxicity due to its potent activation of many different neuronal nAChR subtypes (Avalos et al. 2002), notably the  $\alpha 3\beta 4$  nAChR (Tomizawa et al. 2001; Avalos et al. 2002). Nonetheless, radiolabelled epibatidine derivatives have found limited use in human PET studies (Bohnen and Frey 2007). Theoretically, subtype-specific analogues of epibatidine might have favourable tracer properties with lesser toxicity (Avalos et al. 2002). Although [ $^{18}\text{F}$ ]NFEP ([ $^{18}\text{F}$ ]FPH, norchlorofluorepibatidine) and [ $^{18}\text{F}$ ]N-Me-NFEP showed good brain uptake and signal-to-background ratios in mouse and baboon brain (Ding et al. 1996; Dolci et al. 1999), their toxicity, even when prepared at high specific activity, was too high for use in man (Horti et al. 1997; Molina et al. 1997; Villemagne et al. 1997; Ding et al. 1999). However, Horti and co-workers successfully synthesized epibatidine derivatives with lesser toxicity (Horti et al. 1998a).

Fluorine-18-labelled FPhEP (Roger et al. 2006), a functional nAChR antagonist with much reduced toxicity, had faster brain kinetics in baboon than did 2-[ $^{18}\text{F}$ ]fluoro-A-85380 (discussed below), whereas its fluorophenyl analogue [ $^{18}\text{F}$ ]F<sub>2</sub>PhEP had higher specific binding (Valette et al. 2007). Its binding on PET was quantified as *binding potential* ( $BP_{ND}$ ), which is proportional to the ratio  $B_{\max}/K_d$ . However, nicotine failed to displace either radioligand, indicating low specific binding. Patt and co-workers developed *N*-[ $^{11}\text{C}$ ]methylepibatidine and *N*-[ $^{11}\text{C}$ ]methylhomoepibatidine (Patt et al. 1999, 2001) and compared the uptake and binding of *N*-[ $^{11}\text{C}$ ]methyl-epibatidine enantiomers in the brain of living mouse, rat and pig. Whereas the (–)-enantiomer showed slower uptake and gradual accumulation in the brain, the (+)-enantiomer had very rapid uptake and washout, indicating distinct binding mechanisms. Because of its better kinetics and higher selectivity, *N*-[ $^{11}\text{C}$ ]methyl-(–)-epibatidine was investigated in pigs by PET, showing high brain uptake. However, steady-state binding in the highest binding regions (thalamus) was not attained within 1 h recordings, which is a disadvantage for quantitation of carbon-11-labelled tracers, due to the 20 min physical half-life. Furthermore there was high toxicity, precluding its use in humans (Patt et al. 1999). The analogue

*N*-[<sup>11</sup>C]methylhomo-epibatidine had a better toxicity profile; as with *N*-[<sup>11</sup>C]methyl-epibatidine, the (–)-enantiomer of *N*-[<sup>11</sup>C]methylhomo-epibatidine showed high uptake in pig brain, while the (+)-enantiomer was rapidly washed out. Although findings of *N*-[<sup>11</sup>C]methyl(–)-homoepibatidine binding in pig brain suggested suitability for PET imaging, its toxicity in mice and rats was comparable to that of *N*-methyl-epibatidine, again precluding its safe use in humans (Patt et al. 2001).

Further development of epibatidine derivatives has focussed on maintaining or improving kinetic profiles while reducing toxicity. One such compound, 2-(6-chloro-5-phenylpyridin-3-yl)-7-[<sup>11</sup>C]methyl-7-aza-bicyclo[2.2.1] heptane, showed a thalamus/cerebellum ratio of 4.2 at 90 min after injection, indicating high specific binding in rat brain, which was displaceable by nicotine treatment (1 mg/kg). A preliminary PET study of this tracer in a baboon revealed fast brain uptake and high thalamic binding consistent with  $\alpha 4\beta 2$  nAChR distribution. However binding equilibrium was not reached within 2 h, which is the absolute limit for PET recordings with <sup>11</sup>C-labelled radioligands (Huang et al. 2004).

Mu et al. synthesized another series of labelled epibatidine and homoepibatidine analogues (Mu et al. 2006). Of these, the 8-[<sup>11</sup>C]methyl-8-aza-bicyclo[3.2.1]octane derivative had a double bond conjugated with the pyridine nucleus, thus restricting free rotation of the pyridine ring. It showed high affinity (2 nM) in vitro and at least 100-fold selectivity for  $\alpha 4\beta 4$  over  $\alpha 7$  nAChRs. Furthermore, its toxicity was 50-fold lower than for epibatidine. Although promising for PET studies, there are no further reports on its evaluation. The antagonistic epibatidine derivative ( $\pm$ )-[<sup>11</sup>C]NMI-EPB (Ding et al. 2006) had 2.5-fold higher uptake in the baboon brain than did the 3-pyridyl ether 2-[<sup>18</sup>F]fluoro-A-85380, which is discussed in more detail below (Ding et al. 2006). Surprisingly, separation of the ( $\pm$ )-[<sup>11</sup>C]NMI-EPB enantiomers revealed that (+)-[<sup>11</sup>C]NMI-EPB had fast kinetics and but low affinity, whereas (–)-[<sup>11</sup>C]NMI-EPB appeared to be suitable for imaging but had a slow kinetics (Gao et al. 2007a).

Two further sets of epibatidine analogues (Gao et al. 2007a, 2008b) were subsequently developed by the group from Johns Hopkins University, including (–)-[<sup>18</sup>F]JHU87522 (now termed [<sup>18</sup>F]AZAN) (Gao et al. 2008b; Horti et al. 2010), which had promising properties with respect to brain uptake, kinetics, metabolic stability and low toxicity. Results of [<sup>18</sup>F]AZAN toxicology and human radiation dosimetry have been reported (Horti et al. 2010). PET studies in baboons confirmed that [<sup>18</sup>F]AZAN rapidly enters brain, attaining steady state within 90 min after injection (Kuwabara et al. 2012). Furthermore, blocking experiments with cytosine showed [<sup>18</sup>F]AZAN to bind specifically to  $\beta 2$ -containing (predominantly  $\alpha 4\beta 2$ ) nAChRs, supporting its suitability for nicotinic drug evaluation. Based on its exceptionally high affinity in vitro and improved lipophilicity over 2-[<sup>18</sup>F]FA, another epibatidine analogue, (–)-[<sup>18</sup>F]JHU86428 ([<sup>18</sup>F]XTRA), was proposed as a potential tracer for the less abundant extrathalamic  $\alpha 4\beta 2$  nAChRs (Gao et al. 2008b, Horti et al. 2010). An improved radiosynthesis methodology has been reported (Gao et al. 2010). Following successful imaging in non-human primates (Kuwabara et al. 2017), [<sup>18</sup>F]XTRA proved fit for quantitation of  $\alpha 4\beta 2$  nAChRs in hippocampus of healthy humans and showed an age-dependent decline (Coughlin et al. 2018c).

Toxicity of epibatidine analogues arises from high affinity for the ganglionic  $\alpha 3\beta 4$  receptors. This has motivated the search for derivatives with higher  $\alpha 4\beta 2$  selectivity. To this end,  $^{18}\text{F}$ -labelled stereoisomers of the chloro-fluoro-substituted homoepipatidine analogue, flubatine (previously called NCFHEB), have been synthesized. The flubatine enantiomers both bind with subnanomolar affinity to membranes from rat thalamus or HEK293 cells expressing the human  $\alpha 4\beta 2$  nAChR (Deuther-Conrad et al. 2004), with twofold higher affinity for the (+)-enantiomer. Previous work shows that fluoro- and norchloro-analogues of epibatidine have selectivity for  $\beta 2$ -containing receptors (Avalos et al. 2002). Indeed, the affinity of both flubatine enantiomers for  $\alpha 4\beta 2$  nAChRs was comparable to that of epibatidine, but affinity to ganglionic  $\alpha 3\beta 4$  nAChRs was 20–60-fold lower (Deuther-Conrad et al. 2004). The increased subtype selectivity of flubatine seemingly results in lesser pharmacological side effects compared to epibatidine; injection of 25  $\mu\text{g}/\text{kg}$  (+)-flubatine or (–)-flubatine to awake mice was without important pharmacological effects (Deuther-Conrad et al. 2008). The doses encountered in a human PET study with [ $^{18}\text{F}$ ]flubatine was 1000-fold lower (Vaupel et al. 2005), which entails a considerable margin of safety. In addition to its selectivity for  $\alpha 4\beta 2$ , (–)-flubatine and (+)-flubatine also had considerably better selectivity for  $\alpha 4\beta 2$  over  $\alpha 7$  receptors than did (–)-epibatidine (Deuther-Conrad et al. 2004).

*N*-methyl and *N*-ethyl derivatives of flubatine have been synthesized but displayed lower target affinities and were consequently not considered for radiolabelling (Deuther-Conrad et al. 2004). Similar distribution patterns for (+)-[ $^{18}\text{F}$ ]flubatine and (–)-[ $^{18}\text{F}$ ]flubatine were observed in mice, rat and porcine brain (Brust et al. 2008; Deuther-Conrad et al. 2008; Sabri et al. 2008). Allen et al. provided evidence that nicotine analogues are transported into the brain via the blood-brain barrier (BBB) choline transporter (Allen et al. 2003); this mechanism may also be involved in the brain uptake of epibatidine and homoepipatidine derivatives (Deuther-Conrad et al. 2008). Indeed, addition of flubatine to the incubation medium inhibited with an  $\text{IC}_{50}$  of  $370 \pm 90 \mu\text{M}$  the uptake of [ $^3\text{H}$ ]choline in immortalized rat brain endothelial cells, which are known to express the blood-brain barrier (BBB) choline transporter (Deuther-Conrad et al. 2008). This result is comparable to the  $K_i$  of  $65 \mu\text{M}$  obtained for hemicholinium-3 in the same experimental system (Friedrich et al. 2001). Furthermore, *in vivo* experiments in rats have confirmed the postulated interaction of flubatine with the BBB choline transporter; 50  $\mu\text{M}$  flubatine reduced the transport rate of [ $^3\text{H}$ ]choline by 21%, whereas equimolar epibatidine resulted in a ~ 40% reduction (Deuther-Conrad et al. 2008). The stronger interaction of epibatidine is consistent with its higher uptake in mouse brain (London et al. 1995) compared to (+)-[ $^{18}\text{F}$ ]flubatine or (–)-[ $^{18}\text{F}$ ]flubatine.

PET studies in young pigs were performed to compare the brain uptake and kinetics of (+)-[ $^{18}\text{F}$ ]flubatine and (–)-[ $^{18}\text{F}$ ]flubatine with that of 2-[ $^{18}\text{F}$ ]fluoro-A-85380, a 3-pyridyl ether discussed in detail below. The brain uptake of both enantiomers proved to be two- to threefold higher than that of 2-[ $^{18}\text{F}$ ]fluoro-A-85380. The binding equilibrium of (–)-[ $^{18}\text{F}$ ]flubatine was reached significantly earlier (~ 60 min p.i.) than that of the (+)-enantiomer (Brust et al. 2008), consistent with its lesser affinity *in vitro*. The specific binding of (–)-[ $^{18}\text{F}$ ]flubatine in porcine brain

was comparable to that of 2-[<sup>18</sup>F]fluoro-A-85380, but (+)-[<sup>18</sup>F]flubatine displayed about twofold higher specific binding. Thus, both [<sup>18</sup>F]flubatine enantiomers may present advantages over 2-[<sup>18</sup>F]fluoro-A-85380 for application in human PET studies, especially as pertains to the time to equilibrium binding.

The metabolites of (+)-[<sup>18</sup>F]flubatine in pig urine and plasma have been determined by HPLC-MS (Ludwig et al. 2018). Displaceable binding of (–)-[<sup>18</sup>F]flubatine in human cerebellum upon smoking (Bhatt et al. 2018) raises a red flag for its quantitation through reference tissue methods, which assume absence of specific binding in the reference region. Nonetheless, (–)-[<sup>18</sup>F]flubatine PET studies in non-human primate showed complete displacement with nicotine, revealing a non-displaceable distribution volume ( $V_{ND}$ ) of 6 mg/ml, and  $BP_{ND}$  of 4 in the thalamus, 1 in the frontal cortex and putamen and only 0.1 in the cerebellum (Bois et al. 2015), consistent with an earlier semi-quantitative analysis in non-human primate (Hockley et al. 2013). The extensive displaceable binding in human cerebellum might reflect a species difference.

#### 18.3.1.4 3-Pyridyl Ethers

First developed as experimental treatments for Alzheimer's disease, the 3-pyridyl ethers were identified as promising radioligands for nAChRs imaging (Gündisch 2000; Horti and Villemagne 2006; Horti et al. 2010). The 3-pyridyl ethers are equipotent to epibatidine at the mainly  $\alpha 4\beta 2$  nAChR [<sup>3</sup>H]cytisine binding sites in the brain but are 100-fold less potent than epibatidine as agonists at  $\alpha 3\beta 4$  nAChRs (Abreo et al. 1996). As noted above, this predicts larger dose safety margins with minimal cardiovascular or other toxic side effects. The prototype compound A-85380 has similar binding affinities at recombinant  $\alpha 2\beta 2$ ,  $\alpha 3\beta 2$  and  $\alpha 4\beta 2$  nAChRs in vitro (Xiao and Kellar 2004). In addition, the iodinated derivative 5-[<sup>125</sup>I]iodo-A-85380 binds with high affinity to  $\alpha 6\beta 2\beta 3$  nAChRs in monkey and rat striatum (Kulak et al. 2002). Therefore, 3-pyridyl ethers are properly regarded as  $\beta 2$ -selective compounds (Jensen et al. 2005; Lai et al. 2005), also considering their lack of affinity for the  $\alpha 7$  nAChR (Sullivan et al. 1996).

As noted above, 2-[<sup>18</sup>F]fluoro-A-85380 and 6-[<sup>18</sup>F]fluoro-A-85380 (Horti et al. 1998b, 2000) showed early promise for PET imaging, having less toxicity than epibatidine analogues. Of the two, 6-[<sup>18</sup>F]fluoro-A-85380 (Scheffel et al. 2000) had superior kinetics, characterized by earlier peak and faster clearance from the brain (Ding et al. 2004), and better target-to-background ratios than were obtained with 2-[<sup>18</sup>F]fluoro-A-85380 in a comparative study in baboon (Ding et al. 2000a). Although 6-[<sup>18</sup>F]fluoro-A-85380 has nonetheless not yet found wide use, both derivatives proved successful in human brain imaging (see Sabri et al. in PET and SPECT in Neurology). The SPECT analogue 5-[<sup>123</sup>I]iodo-A-85380 was tested in non-human primates (Chefer et al. 1998), and has also served for nicotine challenge studies in human smokers (Esterlis et al. 2010a), as described below. Although suffering from slow kinetics, 5-[<sup>123</sup>I]iodo-A-85380 was sensitive to competition from endogenous acetylcholine (Fujita et al. 2003), a property also to be discussed in some detail below. The methyl-substituted derivative 5-[<sup>123</sup>I]iodo-A-84543 (Henderson et al. 2004) displayed faster kinetics, but homogenous uptake of



radioactivity in baboon brain, in contrast to the spatially heterogeneous pattern of specific binding for the analogue [ $^{11}\text{C}$ ]A-84543 seen in mouse brain (Kassiou et al. 1998). PET studies in pigs with [ $^{11}\text{C}$ ]A-186253, a structurally similar tracer, showed little displacement by cytosine, indicating excessive non-specific binding (Itier et al. 2004).

The comparably low brain uptake of 2- $^{18}\text{F}$ fluoro-A-85380 was similar to that previously found with 5- $^{11}\text{C}$ methyl-A-85380 (2.2% ID/g brain tissue at 30 min p.i. (Iida et al. 2004)). This may be related to their rather high polarity (Zhang et al. 2004), which is a critical parameter for brain radiotracer uptake (Waterhouse 2003). Nevertheless 2- $^{18}\text{F}$ fluoro-A-85380 was successfully used to image nAChRs in non-human primates (Chefer et al. 1999, 2003; Valette et al. 1999; Le Foll et al. 2007) and in rat microPET studies (Vaupel et al. 2007). An optimized radiosynthesis of 2- $^{18}\text{F}$ fluoro-A-85380 and improved analytical techniques (Mitkovski et al. 2005; Schmaljohann et al. 2005; Kimes et al. 2008) have facilitated its use in human PET studies (Ellis et al. 2009b) (Lotfipour et al. 2012b). Despite the disadvantage of its slow binding kinetics, requiring interrupted or continuous PET recordings lasting at least 6 h (Gallezot et al. 2005; Horti and Villemagne 2006), 2- $^{18}\text{F}$ fluoro-A-85380 remains the most frequently utilized ligand in human PET studies (see Sabri et al. in PET and SPECT in Neurology); despite its limitations, it remains for the present the standard against which other nAChR PET ligands have been compared.

A series of 5-substituted-6-halogeno derivatives of A-85380 have potentially improved lipophilicity and affinity (Zhang et al. 2004). PET studies in rhesus monkey with two such  $^{18}\text{F}$ -labelled derivatives showed higher lipophilicity than for 2- $^{18}\text{F}$ fluoro-A-85380, resulting in enhanced target-to-background ratios. Imaging studies with another 5-substituted A-85380 derivative, [ $^{11}\text{C}$ ]5-MA (Iida et al. 2004), demonstrated lower total brain uptake and lower target-to-background ratios than for 2- or 6- $^{18}\text{F}$ fluoro-A-85380. A further analogue,  $^{18}\text{F}$ ZW-104 has had initial testing (Kozikowski et al. 2005; Valette et al. 2009; Saba et al. 2010). In baboon PET studies,  $^{18}\text{F}$ ZW-104 showed regional radioactivity distribution resembling that of 2- $^{18}\text{F}$ fluoro-A-85380 and some superior properties, including higher accumulation in the brain, earlier peak uptake in the thalamus and faster washout kinetics. However, it also displayed considerable affinity for  $\alpha 3\beta 2$  and  $\alpha 2\beta 2$  receptors in vitro (Valette et al. 2009) and rather high non-displaceable (by nicotine) uptake in the striatum, a region with comparably low density of nAChRs.

Many further derivatives have been tested in rodent and non-human primate PET studies: [ $^{76}\text{Br}$ ]BAP (Sihver et al. 1999a); [ $^{76}\text{Br}$ ]BrPH (Kassiou et al. 2002); [ $^{11}\text{C}$ ]Me-p-PVC (Brown et al. 2004) and its analogues [ $^{11}\text{C}$ ]JHU85208, [ $^{11}\text{C}$ ]JHU85157 and [ $^{11}\text{C}$ ]JHU85270 (Gao et al. 2007b, 2009);  $^{18}\text{F}$  nifene (Pichika et al. 2006; Easwaramoorthy et al. 2007); and two carbon-11- and fluorine-18-labelled isotopomers of one pyridine-derived ligand. While some of these ligands had better kinetics than 2- $^{18}\text{F}$ fluoro-A-85380, their low  $BP_{ND}$ , high non-specific binding or non-selectivity discouraged further development for human imaging (Easwaramoorthy et al. 2007). Despite this, specific binding of  $^{18}\text{F}$  nifene was later attributed entirely to  $\alpha 4\beta 2$  nAChRs based on studies in  $\beta 2$ -knockout mice (Bieszczad

et al. 2012), and subsequent human studies demonstrated the safety and test-retest reliability of [ $^{18}\text{F}$ ]nifene (Betthausen et al. 2017; Lao et al. 2017). This was further supported by a study showing no significant decline in [ $^{18}\text{F}$ ]nifene binding over five decades of healthy human ageing (Mukherjee et al. 2018). Despite the partial selectivity of the tracer for  $\alpha 4\beta 2$  nAChRs, there was low binding in human habenula (which contains  $\alpha 3\beta 2$  nAChRs) and in the red nucleus (which contains abundant  $\alpha 2\beta 2$  nAChRs).

PET imaging of  $\alpha 4\beta 2$  nAChRs using the pyridyl ether analogue [ $^{18}\text{F}$ ]nifrolidine has been tested in non-human primate (Chattopadhyay et al. 2005). Although having favourable kinetics, the thalamus-to-cerebellum ratio was lower than that of other  $\alpha 4\beta 2$ -targeting pyridyl ether analogues (Chattopadhyay et al. 2005), attaining a value of 4 at several hours after administration in non-human primate and showing considerable displaceability by nicotine (Pichika et al. 2013). Similarly, there have been no follow-up reports on the nifrolidine homologue [ $^{18}\text{F}$ ]nifzetidine (Mukherjee et al. 2004), except for a report showing a continuous increase of the thalamus/cerebellum ratio up to 3 h after administration to non-human primates (Pichika et al. 2011). Such slow kinetics is unfavourable for quantitation.

The series of pyridyl ether-based compounds [ $^{18}\text{F}$ ]NIDA52189, [ $^{18}\text{F}$ ]NIDA522131 and [ $^{18}\text{F}$ ]NIDA52289 have been synthesized and evaluated by PET in rhesus monkeys (Horti and Villemagne 2006), especially intended for imaging of the relatively sparse extrathalamic  $\alpha 4\beta 2$  nAChRs. Among these, [ $^{18}\text{F}$ ]NIDA52189 (Zhang et al. 2004) and [ $^{18}\text{F}$ ]NIDA522131 (Chefer et al. 2008) were deemed superior to 2- $^{18}\text{F}$ ]FA with respect to extrathalamic binding but suffered from slow kinetics in vivo.

#### 18.3.1.5 Non-Epiatidine-and-Non-A-85380-Related Compounds

Carbon-11-labelled Me-*p*-PVC (Brown et al. 2002) and *p*-PVP-MEMA, which were selected from the class of (4-pyridinyl)vinylpyridines developed by Abbott Laboratories, possessed picomolar affinity towards  $\alpha 4\beta 2$  receptors. Nevertheless, [ $^{11}\text{C}$ ]Me-*p*-PVC had low  $BP_{ND}$  in thalamus of non-human primate (Brown et al. 2004), and [ $^{11}\text{C}$ ]p-PVP-MEMA had a low target-to-background ratio in a preliminary PET study (Dollé et al. 2008).

### 18.3.2 Imaging of Heteromeric $\beta 4$ -Containing nAChR Subtype

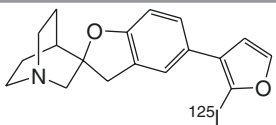
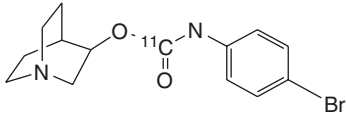
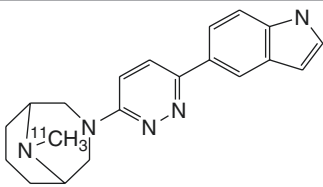
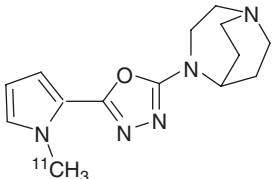
The putative  $\beta 4$ -selective agonist [ $^{11}\text{C}$ ]SIB-1553A was assessed by biodistribution and ex vivo brain autoradiography in rats (Sobrio et al. 2008). Its low specific binding did not encourage further development, and there was no attempt made to separate and individually investigate the enantiomers of this racemic radioligand.

### 18.3.3 Radioligands for $\alpha 7$ nAChRs

Efforts to develop a radiopharmaceutical for PET imaging of  $\alpha 7$  nAChR have met with growing success in the past few years. Structurally diverse classes of

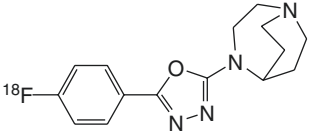
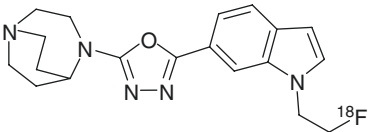
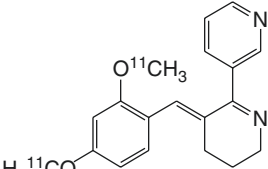
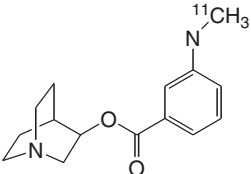
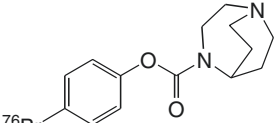
compounds meet the steric and electronic requirements of this binding site, as reviewed recently (Brust et al. 2012; Mo et al. 2014). As noted above, the cerebral expression of  $\alpha 7$  nAChR is comparatively low, constituting perhaps one quarter of the density for  $\alpha 4\beta 2$  receptors (Spurden et al. 1997; Whiteaker et al. 1999; Hellström-Lindahl and Court 2000). Because of this low natural abundance of  $\alpha 7$  sites, high affinity is particularly important for an effective PET tracer. Binding properties of various  $\alpha 7$  tracers tested to date are summarized in Table 18.2.

**Table 18.2** Results of biodistribution ex vivo and PET imaging studies in vivo of the cerebral binding of  $\alpha 7$  nAChR selective radioligands

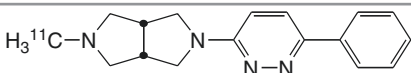
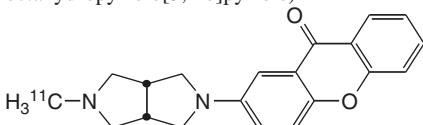
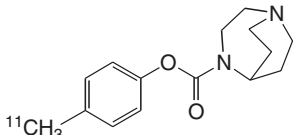
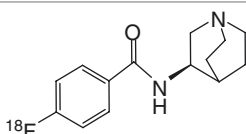
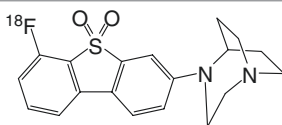
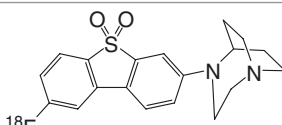
	Main findings
 <p><b>[<sup>125</sup>I]4</b>            ((1'S,2R,4'S)-5-(2-[<sup>125</sup>I]iodofuran-3-yl)-3H-4'-azaspiro[benzofuran-2,2'-bicyclo[2.2.2]octane))</p>	Biodistribution in CD1 mice showed very limited uptake in brain, and no evidence of displaceable binding. $K_i = 0.33$ nM (Pomper et al. 2005)
 <p><b>[<sup>11</sup>C]1</b>            ((1S,3R,4S)-quinuclidin-3-yl (4-bromophenyl) [<sup>11</sup>C]carbamate)</p>	Biodistribution study in rats showed no regionally selective or specific binding. Affinity has not been reported (Dollé et al. 2001)
 <p><b>[<sup>11</sup>C]NS12857</b>            ((1R,5S)-3-(6-(1H-indol-5-yl)pyridazin-3-yl)-9-[<sup>11</sup>C]methyl-3,9-diazabicyclo[3.3.1]nonane)</p>	High uptake in pig brain by dynamic PET but lack of displacement in vivo. $K_i = 0.51$ nM (Lehel et al. 2009)
 <p><b>[<sup>11</sup>C]NS14492</b>            (2-(1,4-diazabicyclo[3.2.2]nonan-4-yl)-5-(1-[<sup>11</sup>C]methyl-1H-pyrrol-2-yl)-1,3,4-oxadiazole)</p>	Dose-dependent decline in cerebral binding after receptor blockade in pigs. $K_i = 2.2$ nM (Ettrup et al. 2011)

(continued)

**Table 18.2** (continued)

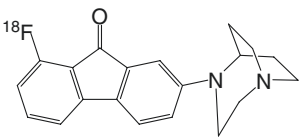
	Main findings
 <p><sup>18</sup>F]NS10743 (2-(1,4-diazabicyclo[3.2.2]nonan-4-yl)-5-(4-<sup>18</sup>F]fluorophenyl)-1,3,4-oxadiazole)</p>	High uptake in the pig brain by dynamic PET, with clear evidence of displaceable binding in regions of relatively high tracer accumulation. $K_i = 9.27$ nM (Deuther-Conrad et al. 2011)
 <p><sup>18</sup>F]NS14490 (((1<i>R</i>,5<i>S</i>)-3-(6-(1-(2-[<sup>18</sup>F]fluoroethyl)indol-5-yl)pyridazin-3-yl)-9-methyl-3,9-diazabicyclo[3.3.1]nonane))</p>	Moderate uptake in the pig brain by dynamic PET, with clear evidence of displaceable binding in brain and cerebral vasculature. $K_i = 2.5$ nM (Rötering et al. 2014)
 <p>H<sub>3</sub><sup>11</sup>CO 2/4-methoxy-[<sup>11</sup>C]GTS21 (<i>E</i>)-3-(2,4-dimethoxybenzylidene)-3,4,5,6-tetrahydro-2,3'-bipyridine)</p>	In baboon dynamic PET, very high initial uptake followed by rapid clearance; relatively little evidence for specific $\alpha 7$ nAChR binding. Brain penetrating radiometabolites detected in plasma. $K_i = 211$ nM (Kim et al. 2007b)
 <p>[<sup>11</sup>C]MeQAA (1<i>S</i>,3<i>R</i>,4<i>S</i>)-quinuclidin-3-yl 3-([<sup>11</sup>C]methylamino)benzoate</p>	In rhesus monkey dynamic PET, the <i>R</i> -enantiomer had high cerebral uptake and distribution consistent with $\alpha 7$ nAChRs. $K_i = 41$ nM (Ogawa et al. 2010)
 <p><sup>76</sup>Br [<sup>76</sup>Br]SSR180711 (4-[<sup>76</sup>Br]bromophenyl 1,4-diazabicyclo[3.2.2]nonane-4-carboxylate)</p>	In rhesus monkey dynamic PET, substantial and heterogeneous brain accumulation. Binding globally reduced to level seen in cerebellum by pretreatment with the $\alpha 7$ nAChR agonist SSR180711. Biton et al. (2007) reported $K_i$ values of $22 \pm 4$ and $14 \pm 1$ nM in rat and human, respectively (Hashimoto et al. 2008)

**Table 18.2** (continued)

	Main findings
 <p><math>H_3^{11}C</math>-N [<math>^{11}C</math>]A-582941 (2-[<math>^{11}C</math>]methyl-5-(6-phenylpyridazin-3-yl) octahydropyrrolo[3,4-<i>c</i>]pyrrole)</p>  <p><math>H_3^{11}C</math>-N [<math>^{11}C</math>]A-844606 (2-(5-methylhexahydroindolizino[1,2-a] pyrrol-2(1<i>H</i>)-yl)-9<i>H</i>-xanthen-9-one)</p>	<p>In rhesus monkey dynamic PET studies, evidence for regional distribution consistent with <math>\alpha 7</math> nAChR expression. <math>K_i</math> (A-582941, rat) = 10.8 nM, <math>K_i</math> (A-582941, human) = 17 nM, <math>IC_{50}</math> (A-844606, rat) = 11 nM (Toyohara et al. 2010)</p>
 <p><math>^{11}CH_3</math> [<math>^{11}C</math>]CHIBA-1001 (4-[<math>^{11}C</math>]methylphenyl)-1,4-diazabicyclo[3.2.2] nonane-4-carboxylate)</p>	<p>In a PET study of one healthy human, evidence for preferential binding in the hippocampus, cortex and basal ganglia, with slow washout, and least binding in cerebellum. <math>K_i</math> = 35 nM (Toyohara et al. 2009)</p>
 <p><math>^{18}F</math> [<math>^{18}F</math>]4 (4-[<math>^{18}F</math>]fluoro-<i>N</i>-((1<i>S</i>,3<i>R</i>,4<i>S</i>)-quinuclidin-3-yl) benzamide)</p>	<p>Low brain uptake in rats, homogenous distribution. <math>K_i</math> = 14 nM (Pin et al. 2014)</p>
 <p><math>^{18}F</math> [<math>^{18}F</math>]ASEM (7-(1,4-diazabicyclo[3.2.2]nonan-4-yl)-4- [<math>^{18}F</math>]fluorodibenzo[<i>b,d</i>]thiophene 5,5-dioxide)</p>	<p>The radiotracer readily entered the baboon brain and specifically labelled <math>\alpha 7</math>-nAChR. SSR180711 blocked the binding in the baboon brain in a dose-dependent manner. <math>K_i</math> = 0.84 nM (Horti et al. 2014; Teodoro et al. 2015)</p>
 <p><math>^{18}F</math> [<math>^{18}F</math>]DBT10 (7-(1,4-diazabicyclo[3.2.2]nonan-4-yl)-2- [<math>^{18}F</math>]fluorodibenzo[<i>b,d</i>]thiophene 5,5-dioxide)</p>	<p>Uptake of the radiotracer in monkey brain occurred rapidly, reaching SUV values of 2.9–3.7 within 30 min. There was dose-dependent blockade by ASEM throughout the brain. <math>K_i</math> = 0.60 nM (Hillmer et al. 2016b)</p>

(continued)

**Table 18.2** (continued)

	Main findings
 <p>[<sup>18</sup>F]FLN28 (7-(1,4-diazabicyclo[3.2.2]nonan-4-yl)-1- [<sup>18</sup>F]fluoro-9H-fluoren-9-one)</p>	<p>A fluoren-9-one derived <math>\alpha 7</math> nAChR selective PET ligand (Teodoro et al. 2018) with high brain uptake in mice and rats. <math>K_i = 2.98</math> nM (Wang et al. 2018)</p>

### 18.3.3.1 Quinuclidine-Based Ligands

The lead compound of a series of azabicyclo carbamate  $\alpha 7$  receptor agonists developed by Astra Laboratories has been labelled with carbon-11 and evaluated in rats (Dollé et al. 2001). Despite having relatively good brain uptake, no regionally selective or specific binding could be seen. Another series of potential  $\alpha 7$ -selective imaging agents based on the quinuclidine moiety has been labelled with carbon-11 and iodine-125 (Pomper et al. 2005). Target selectivities of these compounds were modest, and the most affine compounds had significant binding to the 5-HT<sub>3</sub> receptor, a structural homologue of  $\alpha 7$  nAChR (Zwart et al. 2004).

### 18.3.3.2 GTS-21

Other potential  $\alpha 7$  ligands originate from benzylidene anabasein compounds such as GTS-21 (de Fiebre et al. 1995; Meyer et al. 1998). Indeed, GTS-21 (3-(2,4-dimethoxybenzylidene)-anabasein) showed early promise as a  $\alpha 7$  nAChR agonist medication for improving cognition in patients with schizophrenia (Freedman et al. 2008; Tregellas et al. 2011). It has been labelled with iodine-123 (Zhang et al. 2001) and carbon-11 (Kim et al. 2007b). Consistent with the relatively low affinity and specificity of GTS-21 for  $\alpha 7$  nAChRs, the distribution and kinetics of 5- [<sup>123</sup>I]GTS-21 and 2- [<sup>11</sup>C]GTS-21 in the brain of living baboon and mice were dominated by non-specific binding (Kim et al. 2007a).

### 18.3.3.3 Diazabicyclononane Derivatives

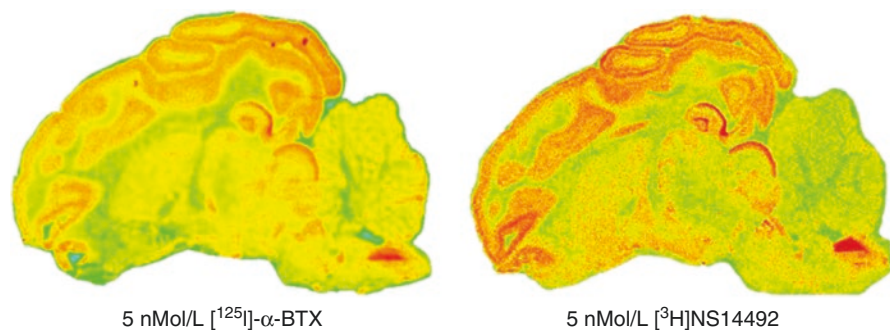
The 1,4-diazabicyclo-[3.2.2]nonane skeleton (Bunnelle et al. 2004) was identified as a motif for  $\alpha 7$  nAChR ligands, and the two novel diazabicyclononane-derived PET ligands [<sup>76</sup>Br]SSR180711 and [<sup>11</sup>C]CHIBA-1001 were evaluated by PET in conscious non-human primates, with testing in a model of schizophrenia (Hashimoto et al. 2008). Of the two tracers, [<sup>11</sup>C]CHIBA-1001 demonstrated superior accumulation in the brain, revealing a heterogeneous regional distribution consistent with the localization of  $\alpha 7$  in the brain; specific binding was blocked by selective  $\alpha 7$  but not  $\alpha 4\beta 2$  agonists revealing some potential for measuring occupancy by pharmaceuticals at  $\alpha 7$  nAChRs. A first clinical PET study confirmed the suitability of [<sup>11</sup>C]CHIBA-1001, although the regional binding differences were small in the human brain (Toyohara et al. 2009). Notably,  $\alpha 7$ -specific binding could not be demonstrated in vitro: there was no displacement of 30 nM [<sup>3</sup>H]CHIBA binding from rat

brain membranes by 1  $\mu\text{M}$   $\alpha$ -bungarotoxin (Tanibuchi et al. 2010). Another rodent study reported low in vitro binding affinity of [ $^3\text{H}$ ]CHIBA and poor in vivo selectivity to  $\alpha 7$  nAChRs in rodent brain (Ding et al. 2012). Furthermore, differences in the regional distribution of the binding sites of CHIBA-1001 relative to [ $^{125}\text{I}$ ] $\alpha$ -bungarotoxin were evident in monkey and human brain samples (Tanibuchi et al. 2010).

The 1,4-diazabicyclo[3.2.2]nonane derivatives developed by NeuroSearch emerged as promising PET ligands for imaging of cerebral  $\alpha 7$  nAChR (Peters et al. 2007). Two carbon-11 radioligands were developed, [ $^{11}\text{C}$ ]NS12857 (Lehel et al. 2009) and [ $^{11}\text{C}$ ]NS14492 (Ettrup et al. 2011), along with the fluorine-18 compounds [ $^{18}\text{F}$ ]NS10743 (Deuther-Conrad et al. 2009) and [ $^{18}\text{F}$ ]NS14490 (Rötering et al. 2014). The cerebral uptake of these three 1,4-diazabicyclo[3.2.2]nonane derivatives exceeded that of [ $^{11}\text{C}$ ]CHIBA-1001. Although the uptake of [ $^{11}\text{C}$ ]NS12857 was not displaced by  $\alpha 7$  nAChR-selective compounds, specific binding was clearly evident for [ $^{11}\text{C}$ ]NS14492, [ $^{18}\text{F}$ ]NS14490 and [ $^{18}\text{F}$ ]NS10743, the ligands with higher target affinity (Brust et al. 2012).

NS14492 has been labelled with tritium to allow in vitro autoradiographic studies on  $\alpha 7$  nAChR assessing  $\alpha 7$  nAChR density in the porcine brain and by extension other species (Magnussen et al. 2015). Figure 18.2 shows the autoradiographic comparison of a saturating concentration of [ $^{125}\text{I}$ ] $\alpha$ -bungarotoxin and [ $^3\text{H}$ ]NS14492 in whole brain sections from newborn piglets. Brain region-specific binding pattern of [ $^3\text{H}$ ]NS14492 is very similar to that of [ $^{125}\text{I}$ ] $\alpha$ -bungarotoxin, with the non-specific binding being considerably lower. Furthermore, the use of [ $^3\text{H}$ ]NS14492 provides a higher resolution, which is especially noticeable in the laminar layers of the neocortex. [ $^3\text{H}$ ]NS14492 therefore is an interesting alternative to [ $^{125}\text{I}$ ] $\alpha$ -bungarotoxin when assessing  $\alpha 7$  nAChR density in the porcine brain and by extension other species.

The general suitability of the diazabicyclononane derivatives for PET imaging of  $\alpha 7$  nAChRs was shown by preclinical PET studies in pigs (Lehel et al. 2009; Deuther-Conrad et al. 2011; Ettrup et al. 2011), although the magnitude of  $BP_{ND}$ , about 0.5, is rather low for a useful PET tracer. In view of the low natural abundance



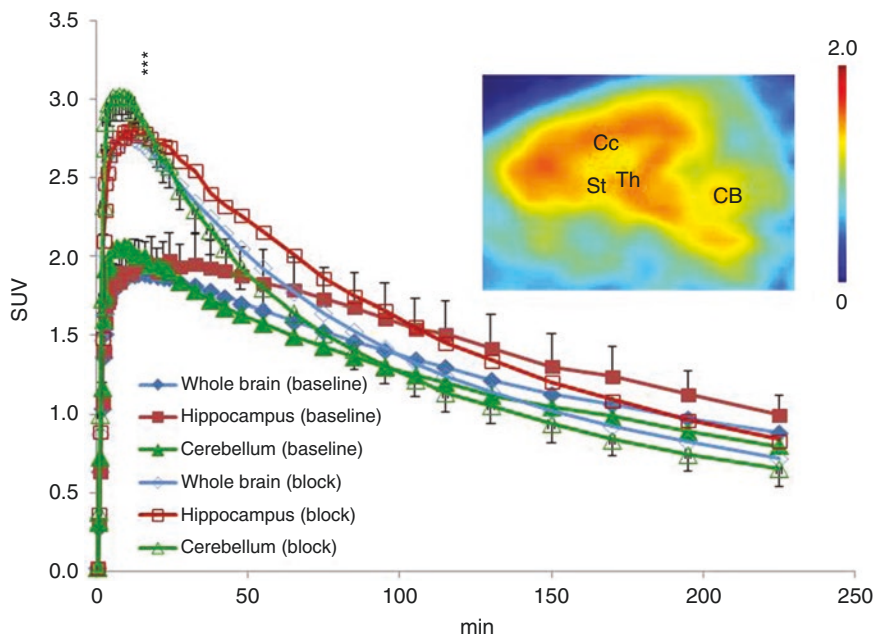
**Fig. 18.2** Autoradiographic comparison of [ $^{125}\text{I}$ ] $\alpha$ -bungarotoxin and [ $^3\text{H}$ ]NS14492 binding to brain sections from newborn piglets. For anatomic reference see Fig. 18.1

of the  $\alpha 7$  nAChR in the brain, a substantial increase in  $\alpha 7$  affinity of PET radiotracers may be required for sensitive quantitation (Brust et al. 2012); target affinities of [ $^{11}\text{C}$ ]CHIBA-1001 ( $K_i \sim 35$  nM) (Hashimoto et al. 2008; Toyohara et al. 2009) and [ $^{18}\text{F}$ ]NS10743 ( $K_i \sim 10$  nM) (Deuther-Conrad et al. 2009) do not predict adequate specific signal in vivo, given the low  $B_{\text{max}}$  (Koepppe 2001). NS14490, a novel diazabicyclononane derivative, with a  $K_i$  of 3 nM may be more promising in this regard (Brust and Deuther-Conrad 2012). The distribution of [ $^{18}\text{F}$ ]NS14490 binding in mouse brain autoradiograms correlated with the known pattern of  $\alpha 7$  nAChR expression and was displaced with the  $\alpha 7$  nAChR ligand methyllycaconitine (Brust and Deuther-Conrad 2012).

With tilorone, an amphiphilic molecule possessing high interferon-inducing potential, a novel  $\alpha 7$  nAChR pharmacophore has been identified (Briggs et al. 2008; Schrimpf et al. 2012), leading to a new series of diazabicyclononane-substituted dibenzothiophene derivatives for PET imaging developed independently by two groups (Gao et al. 2013; Scheunemann et al. 2014). The 2-fluoro dibenzothiophene sulfone derivative (DBT10) and 4-fluoro dibenzothiophene sulfone derivative (ASEM) have been identified as potential  $\alpha 7$  nAChR imaging agents, as shall be discussed in more detail below. In addition, a small library of further tilorone-based derivatives was synthesized to explore further the impact of the isomeric effect and effects of different cationic centres on the ligand features. However, an isomer, substituted with fluorine and the cationic centre in the same benzo ring, did not achieve criteria for further  $^{18}\text{F}$ -labelling. Increased flexibility of the tertiary amine of the cationic centre 9-methyl-3,9-diazabicyclo[3.3.1]nonane and 3-methyl-3,8-diazabicyclo[3.2.1]octane resulted in a remarkable loss of binding affinity (Teodoro et al. 2015). Most recently, a series of novel fluoren-9-one-based diazabicyclononane derivatives has been developed, showing low nM affinity towards the  $\alpha 7$  nAChR and >1000-fold selectivity over the  $\alpha 4\beta 2$  nAChR (Teodoro et al. 2018; Wang et al. 2018). Two derivatives were radiolabelled leading to the corresponding carbonyl bioisosteres of [ $^{18}\text{F}$ ]DBT10 and [ $^{18}\text{F}$ ]ASEM. The carbonyl derivative of [ $^{18}\text{F}$ ]DBT10 exhibited high initial brain uptake (12% ID/g at 15 min post-injection) and displaceable binding (Wang et al. 2018).

Both [ $^{18}\text{F}$ ]DBT10 and [ $^{18}\text{F}$ ]ASEM have been investigated as PET tracers in non-human primates, showing a favourable kinetic profile for quantitation of the  $\alpha 7$  nAChR in living brain (Horti et al. 2014; Hillmer et al. 2016b). Recently, a direct comparison in rhesus monkey of [ $^{18}\text{F}$ ]DBT10 and [ $^{18}\text{F}$ ]ASEM indicated very similar pharmacokinetics (Hillmer et al. 2017). A blocking study with [ $^{18}\text{F}$ ]DBT10 in pigs revealed a 75% decrease of the binding potential  $BP_{\text{ND}}$  after treatment with 3 mg•kg $^{-1}$  i.v. of the  $\alpha 7$  nAChR partial agonist NS6740 (Fig. 18.3) (Teodoro et al. 2015). The total distribution volume ( $V_T$ ; ml g $^{-1}$ ) of [ $^{18}\text{F}$ ]ASEM has been quantified in non-smoking healthy volunteers over a broad range of ages, showing a positive correlation between [ $^{18}\text{F}$ ]ASEM  $V_T$  and age in various brain regions of interest, with  $V_T$  increasing from 20 to 30 ml g $^{-1}$  (Coughlin et al. 2018b). Occupancy by the experimental drug DMXB-A at central  $\alpha 7$  nAChRs could be estimated from [ $^{18}\text{F}$ ]ASEM binding changes in the brain of healthy volunteers (Wong et al. 2018).





**Fig. 18.3** Specific uptake values (SUV) of [ $^{18}\text{F}$ ]DBT10 in porcine brain under baseline and blocking ( $3\text{ mg}\cdot\text{kg}^{-1}$  NS6740 i.v.) conditions

Given this success and in expectation of future human applications, an automated cGMP-compliant radiosynthesis of [ $^{18}\text{F}$ ]DBT10 was established, and toxicity and radiation dosimetry studies were performed. The single-dose toxicity in rats (No-Observed-Effect-Level =  $620\text{ }\mu\text{g}\cdot\text{kg}^{-1}$ ) and the effective dose estimated from mouse and pig studies ( $12.7$  and  $13.7\text{ }\mu\text{Sv}/\text{MBq}$ , resp.) indicated the safe use of [ $^{18}\text{F}$ ]DBT10 in human PET studies ((Kranz et al. 2014; Teodoro et al. 2015).

#### 18.3.3.4 [ $^{11}\text{C}$ ]A-582941 and [ $^{11}\text{C}$ ]A-844606

A new series of octahydropyrrolo[3,4-c]pyrrole derivatives was described by Abbott Laboratories, two of which were selected for labelling with carbon-11 (Toyohara et al. 2010). Whereas no regional heterogeneity or displaceable binding was evident in mouse brain ex vivo, pretreatment with an  $\alpha 7$ -specific agonist decreased the total distribution volumes of both tracers in conscious monkey PET studies, indicative of a  $\text{BP}_{\text{ND}}$  close to 0.5, as with the NeuroSearch (NS) compounds cited above.

#### 18.3.3.5 [ $^{125}\text{I}$ ]I-TSA

A diazabicyclooctane-derived PET ligand with high affinity and selectivity has been radiolabelled and evaluated in mice (Ogawa et al. 2006). Despite a subnanomolar affinity, the high non-specific binding made [ $^{125}\text{I}$ ]I-TSA inadequate for imaging of brain  $\alpha 7$  receptors.

### 18.3.3.6 *R*-[<sup>11</sup>C]MeQAA

The two enantiomers of [<sup>11</sup>C]MeQAA, an azabicyclooctylester-derived compound, were evaluated in mice and conscious monkey PET studies (Ogawa et al. 2009). Although (*R*)-[<sup>11</sup>C]MeQAA showed target-specific accumulation, the in vivo selectivity was insufficient due to binding to the serotonin 5HT<sub>3</sub>-R. Nonetheless, the tracer was used more recently in a multitracer study of aged monkeys (Nishiyama et al. 2015). The hippocampal binding of (*R*)-[<sup>11</sup>C]MeQAA correlated inversely with binding of a marker for mitochondrial complex I and positively with the binding of a marker for beta-amyloid deposition in the aged animals. The authors interpreted their results to indicate a significant (adaptive?) upregulation of  $\alpha 7$  nAChRs in metabolically compromised and degenerating brain tissue.

### 18.3.4 Radioligands for $\alpha 3\beta 4$ nAChRs

Recently, a series of quinuclidine *anti*-1,2,3-triazole derivatives was synthesized with the aim of developing an <sup>18</sup>F-labelled radioligand for imaging the  $\alpha 3\beta 4$  nAChR in the brain. This subtype attracts interest because of its involvement in drug addiction and depression pathways (Rahman et al. 2015). In contrast to  $\alpha 4\beta 2$  and  $\alpha 7$ , the  $\alpha 3\beta 4$  subtype expression is mainly in the autonomic ganglia (hence the toxicity of epibatidine) but also in some specific brain regions and neuronal subpopulations. These regions notably include the medial habenula, nucleus interpeduncularis, dorsal medulla, pineal gland and retina (Gotti et al. 2009). Binding studies in vitro revealed that stereochemistry at the C3 position of the quinuclidine scaffold plays an important role in the nAChR subtype selectivity (Sarasamkan et al. 2016). Whereas the (*R*)-enantiomers are selective to  $\alpha 7$  over  $\alpha 4\beta 2$  (by factors of 44–225) and to a smaller degree over  $\alpha 3\beta 4$  (3–33), their (*S*)-counterparts prefer  $\alpha 3\beta 4$  over  $\alpha 4\beta 2$  (62–237) as well as over  $\alpha 7$  (5–294). Two potent compounds (*S*)-T1 and (*S*)-T2 were identified that bind selectively to  $\alpha 3\beta 4$  nAChR over  $\alpha 7$  nAChR. The compound (*S*)-T1 was chosen for radiolabelling and first preclinical evaluation (Sarasamkan et al. 2017). The brain uptake and the brain-to-blood ratio of (*S*)-[<sup>18</sup>F]T1 in mice at 30 min post-injection were 2.02 (SUV) and 6.1, respectively. According to an ex vivo analysis, the tracer remained intact (>99%) in the brain. Only one major radiometabolite was detected in plasma and urine samples. In vitro autoradiography on pig brain slices revealed binding of (*S*)-[<sup>18</sup>F]T1 to brain regions associated with the expression of  $\alpha 3\beta 4$  nAChRs, which could be reduced by the  $\alpha 3\beta 4$  nAChR selective drug AT-1001. These findings suggest (*S*)-[<sup>18</sup>F]T1 as a promising tool for non-invasive PET imaging of  $\alpha 3\beta 4$  nAChRs in the brain.

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## 18.4 nAChR Imaging of Neurodegenerative Diseases

Reductions in cortical nAChR binding have been found in patients with diverse forms of neurodegeneration, including Alzheimer's disease, Parkinson's disease, Lewy body disease, progressive supranuclear palsy and Down's syndrome (Perry

et al. 1986; Picciotto and Zoli 2002) (see also Sabri et al. in PET and SPECT in Neurology). However, there have been few preclinical PET and SPECT studies of nAChRs in animal models of neurodegenerative disease. This is attributable to two considerations: First, models in transgenic mice have only recently become available for some of these diseases, while remaining lacking for others, and second, the spatial resolution of small-animal PET and SPECT instruments has until recently been inadequate for regional analysis of neuroreceptors in rodent brain, as noted in the instrumentation section above. Therefore, the majority of such investigations use autoradiography *in vitro*. Furthermore, the occurrence of species differences is a hindrance to the interpretation of preclinical imaging studies (Pauly et al. 1989; Quik et al. 2000; Han et al. 2003). Despite these limitations, molecular imaging is emerging as a powerful tool for investigating pathophysiological changes in animal models of neurodegenerative diseases, especially when conducted in conjunction with techniques such as *in vivo* microdialysis, electrophysiology and histopathology (Higuchi et al. 2012).

#### 18.4.1 Alzheimer's Disease

A link between cognitive performance and  $\alpha 4\beta 2$  nAChR expression in the forebrain of healthy rats has been demonstrated in a PET study using the ligand [ $^{18}\text{F}$ ]nifene (Bieszczad et al. 2012), as confirmed by autoradiography *ex vivo* and *in vitro*. The three imaging methods showed the same rank order of specific binding by brain region. We anticipate that [ $^{18}\text{F}$ ]nifene PET should allow tracking of dynamic changes in nAChRs during learning acquisition and memory consolidation, in rodents and also in large-brained animals. In contrast to the case for humans (Zanardi et al. 2002), normal ageing does not seem to reduce nAChR density in rat brain (Picciotto and Zoli 2002; Schliebs and Arendt 2011).

The hallmark histopathological features of Alzheimer's disease are extraneuronal amyloid plaques composed of aggregated amyloid- $\beta$  peptides ( $\text{A}\beta$ ), and intraneuronal neurofibrillary tangles, which are composed largely of hyperphosphorylated forms of tau, a microtubule-associated protein (Thal and Braak 2005). Molecular imaging studies in animal models have primarily targeted fibrillar protein assemblies such as  $\beta$ -amyloid and tau depositions, neuroinflammatory processes and cerebral glucose metabolism (Higuchi et al. 2012). However, these features are closely related to cholinergic hypofunction found in Alzheimer's disease and relevant animal models (Schliebs and Arendt 2011).

Cognitive impairment in Alzheimer's disease is at least partially associated with loss of cortical nAChRs, which may arise due to toxicity of soluble  $\beta$ -amyloid (Zanardi et al. 2002; Schliebs 2005; Schliebs and Arendt 2011). Nicotine treatment in a transgenic mouse model (3xTg-AD) can mediate increased tau phosphorylation and decrease  $\beta$ -amyloid load (Rubio et al. 2006). Impaired cholinergic neurotransmission has been described in the brain of Tg2576 mice, which express the Swedish mutation of human  $\beta$ -amyloid precursor protein (Apelt et al. 2002). Enzyme activities for acetylcholine synthesis (choline acetyltransferase) and degradation

(acetylcholine esterase; AChE) did not differ between transgenic mice and non-transgenic littermates. However, a reduction of high-affinity choline uptake and  $M_1$ -muscarinic receptor density was observed. Autoradiography with [ $^3\text{H}$ ]cytisine revealed a significant 20% loss of  $\alpha 4$ -containing nAChRs in cingulate and parietal cortices of these animals at an age of 17 months. However, there was no change in the number of basal forebrain cholinergic neurons in the transgenic mice, compared to age-matched wild-type animals.

Evidence for an involvement of  $\alpha 7$  nAChR in Alzheimer's disease was first presented three decades ago (Davies and Feisullin 1981). More recently, a very high-affinity binding of (soluble)  $\beta$ -amyloid to  $\alpha 7$  nAChRs has been described in vitro (Wang et al. 2000), supporting the hypothesis that  $\beta$ -amyloid at very low concentrations may initiate neuronal degeneration via an  $\alpha 7$  nAChR-mediated inflammatory process (Bencherif and Lippiello 2010). A 20% reduction in  $\alpha 7$  nAChRs labelled with [ $^{125}\text{I}$ ] $\alpha$ -bungarotoxin was evident in the hippocampus, retrosplenial and parietal cortices and thalamus of 3xTg-AD mice at 6 months of age. There was a significant correlation between intraneuronal  $\beta$ -amyloid and reduced  $\alpha 7$  nAChR binding in the same mouse model (Oddo et al. 2005). Whereas chronic nicotine administration did not alter  $\alpha 7$  nAChR levels in these mice, there was an increase in  $\alpha 4\beta 2$  nAChRs labelled with [ $^{125}\text{I}$ ]epibatidine (Oddo et al. 2005). In contrast to the earlier report, no alteration in  $\alpha 7$  nAChR binding was noted in a subsequent study of the triple transgenic 3xTg-AD mice, which closely emulate several features of natural Alzheimer's disease (Hedberg et al. 2010); this unexpected negative finding was attributed to unknown environmental and/or genetic factors.

The abundance of  $\alpha 7$  nAChRs was determined using nanogold-conjugated  $\alpha$ -bungarotoxin in the APP<sub>(SWE)</sub> mouse model of Alzheimer's disease (Jones et al. 2004). Interestingly, the  $\alpha 7$  nAChR binding increased in the transgenic animals until 9 months of age but had declined at 12 months, most notably in areas of gliosis associated with  $\beta$ -amyloid plaques (Jones et al. 2006). Also [ $^{125}\text{I}$ ] $\alpha$ -bungarotoxin binding decreased in APP<sub>(SWE)</sub> mice between 9 months of age and 16 months (Hellström-Lindh et al. 2004).

With the establishment of reliable cholinergic PET tracers, a number of molecular imaging studies of Alzheimer's disease have appeared in recent years. The first phase of this research consisted of studies of the  $\alpha 4\beta 2$  subtype of nAChR with 2- $^{18}\text{F}$ fluoro-A-85380 (2-FA). One such quantitative PET study of 14 early AD patients showed no difference in  $V_T$  compared to healthy controls, nor was there any relationship between tracer uptake and cognitive scores (Ellis et al. 2008). The same research group did not detect any reduction in binding with healthy ageing (Ellis et al. 2009a), and did not find any effect of precognitive treatment with galantamine on 2-FA binding in Alzheimer's disease patients (Ellis et al. 2009b). Another study with 2-FA using a white matter reference tissue calculation of binding potential ( $\text{BP}_{\text{ND}}$ ) found a 40% reduction in cerebral cortex of a group of nine patients with moderate Alzheimer's disease, but no such change in the thalamus (Kendziorra et al. 2011). Similar findings in a group of mild cognitive impairment (MCI) patients suggested that the loss of  $\alpha 4\beta 2$  sites was already complete at the prodromal state of the disease. Others reported a 20–40% decline in 2-FA  $V_T/f_p$  (i.e. the distribution

volume corrected for the plasma free fraction) in the thalamus, caudate, hippocampus anterior cingulate cortex and insula in a group of 24 Alzheimer's disease patients compared to a healthy control group (Sultzer et al. 2017). Similarly, a dual tracer study with 2-FA and the beta-amyloid tracer [ $^{11}\text{C}$ ]PiB showed an inverse relationship between the binding ratio of  $\alpha_4\beta_2$  tracer and the amyloid  $BP_{ND}$  in the medial frontal cortex and basal forebrain of Alzheimer's disease patients (Okada et al. 2013). While the preponderance of such studies indicates a widespread loss of  $\alpha_4\beta_2$  binding sites in Alzheimer's disease, there is a certain lack of congruence about its spatial extent and magnitude. This may reflect the various endpoints used for quantitation and the very long interval of 4–6 h required for attainment of equilibrium binding with 2-FA. Another PET study with (–)-[ $^{18}\text{F}$ ]flubatine PET in patients with early Alzheimer's disease showed the reductions in  $\alpha_4\beta_2$  nAChR mainly in target regions of the basal forebrain-cortical and septohippocampal cholinergic projections. The same study showed relationship between lower  $\alpha_4\beta_2$  nAChR availability and impairment of distinct cognitive domains, notably episodic memory and executive function/working memory (Sabri et al. 2018). Further details, in particular related to the  $\alpha_4\beta_2$  radioligands (–)-[ $^{18}\text{F}$ ]flubatine and (+)-[ $^{18}\text{F}$ ]flubatine, are reviewed elsewhere (Sabri et al. 2014).

Very recently, uptake of the  $\alpha_7$  nAChR ligand [ $^{11}\text{C}$ ]-(*R*)-MeQAA was measured in groups of 20 Alzheimer's disease patients and 10 healthy age-matched controls. In the nucleus basalis magnocellularis and medial prefrontal cortex, the  $\alpha_7$  nAChR binding correlated positively with individual beta-amyloid PET results. Furthermore, the  $\alpha_7$  nAChR binding correlated positively with memory and executive function (Nakaizumi et al. 2018). This stands in agreement with findings in aged monkeys cited above, where [ $^{11}\text{C}$ ]-(*R*)-MeQAA binding, ostensibly to  $\alpha_7$  NACHR, was associated with increased beta-amyloid binding (Nishiyama et al. 2015). Similar increases in Alzheimer's disease mice were attributed to microglial activation (Matsumura et al. 2015) at sites of amyloid deposition. At time of writing, the jury is out whether this might represented a compensatory and salutogenic reaction, or another aspect of pathology.

[ $^{18}\text{F}$ ]fluoroethoxybenzovesamicol (FEOBV) is a novel PET radiotracer, which binds selectively to the vesicular acetylcholine transporter in terminals of basal forebrain neurons. SUVR analysis of this tracer showed widespread decreases in tracer binding in a group of Alzheimer's disease patients (Aghourian et al. 2017). Together with findings reported above, this suggests pre- and postsynaptic elements of basal-forebrain cholinergic pathways are compromised in Alzheimer's disease.

## 18.4.2 Movement Disorders

Parkinson's disease is, after Alzheimer's disease, the second most common neurodegenerative disorder. The hallmark neuropathology of Parkinson's disease is selective degeneration of midbrain dopaminergic neurons of the substantia nigra pars compacta (SNpc) and the presence of intra-cytoplasmic inclusions (Lewy bodies) consisting of aggregated  $\alpha$ -synuclein (Spillantini et al. 1997) in surviving dopamine

neurons. Neurotoxins such as MPTP and 6-hydroxydopamine have been used in animal models emulating the nigrostriatal degeneration of Parkinson's disease (Quik 2004). Lesions of the nigrostriatal pathway in rats reveal a population of [ $^3\text{H}$ ]nicotine binding sites on dopamine terminals, where they are positioned to influence dopamine release (Clarke and Pert 1985). More recently, the stimulation of striatal dopamine release by nicotine has been linked specifically to  $\alpha 6\beta 2\beta 3$  and  $\alpha 6\alpha 4\beta 2\beta 3$  nAChRs, which predominate in the basal ganglia (Quik et al. 2011). Quantitative analysis with the  $\alpha 6\beta 2$  nAChR subtype ligand [ $^{125}\text{I}$ ] $\alpha$ -conotoxin MII in conjunction with plasma membrane dopamine transporter measurements in MPTP-lesioned mice shows an association with presynaptic nigrostriatal terminals (Quik et al. 2003). In that study, much smaller reductions in the binding of [ $^{125}\text{I}$ ]epibatidine (multiple sites) and 5-[ $^{125}\text{I}$ ]iodo-A-85380 ( $\beta 2$ -sites) were noted after dopamine lesioning, while no change was detected in  $\alpha 7$  nAChRs measured with [ $^{125}\text{I}$ ] $\alpha$ -bungarotoxin binding after nigrostriatal lesions. Displacement of [ $^{125}\text{I}$ ] $\alpha$ -conotoxin MII binding with the analogue E11A was biphasic, allowing resolution of the  $\alpha 6\beta 2\beta 3$  and  $\alpha 6\alpha 4\beta 2\beta 3$  components (Bordia et al. 2007); autoradiographic studies in MPTP-treated mouse and non-human primates, as well as in material from idiopathic Parkinson's disease patients, revealed the  $\alpha 6\alpha 4\beta 2\beta 3$  nAChR subtype population to be selectively vulnerable to nigrostriatal damage. Chronic oral nicotine administration was able to protect nicotinic receptors and dopaminergic markers in MPTP-treated monkeys (Bordia et al. 2006; Quik et al. 2006). Treatment of rats with a selective  $\alpha 7$  nAChR agonist protected against nigrostriatal degeneration in the 6-OHDA model of parkinsonism, an effect that was linked to attenuated microglial activation (Serriere et al. 2015).

In a 2-[ $^{18}\text{F}$ ]fluoro-A-85380 PET study, the density of  $\alpha 4\beta 2$  nAChRs was slightly reduced in the basal ganglia of non-smoking Parkinson's disease patients (Kas et al. 2009). Another 2-[ $^{18}\text{F}$ ]fluoro-A-85380 PET study of Parkinson's disease patients reported a widespread reduction of  $\alpha 4\beta 2$  nAChR availability in cortical and subcortical regions, which correlated with the severity of mild cognitive or depressive symptoms (Meyer et al. 2009). There have been no  $\alpha 7$  nAChR studies in Parkinson's disease, despite the preclinical evidence suggesting a role for these receptors in protecting against nigrostriatal degeneration. There is a current lack of agents for molecular imaging of the particular  $\alpha 6$ -containing nAChRs subtypes present in the basal ganglia, which may be of great relevance to Parkinson's disease given their involvement in dopamine release.

Huntington's disease is an autosomal dominant hereditary disorder proceeding to severe cognitive impairment and motor symptoms, notably hyperkinetic involuntary movements (chorea) (Roos 2010). A transgenic rat model of HD, which carries a truncated huntingtin cDNA fragment with 51 CAG repeats under control of the native rat huntingtin promoter, has been developed (von Horsten et al. 2003). Early investigations of nicotinic receptors in *post-mortem* brain from Huntington's disease patients did not reveal any significant changes (Perry et al. 1987; Whitehouse and Kellar 1987). However, autoradiographic assessment of 2-year-old transgenic rats revealed significant increase of nAChR in various regions in heterozygous but not homozygous animals (Bauer et al. 2005). There have not yet been any PET studies of nAChR in patients with Huntington's disease.

## 18.5 Epilepsy

Some forms of epilepsy have recently been associated with alterations of  $\alpha 4$  nAChR subtype expression (Raggenbass and Bertrand 2002), and there is experimental evidence that the  $\alpha 7$  nAChR may play a role in epileptogenesis (Dobelis et al. 2003). The autosomal dominant nocturnal frontal lobe epilepsy (ADNFLE) can be caused by mutations in the neuronal nicotinic acetylcholine receptor (nAChR) subunit genes *CHRNA4* and *CHRNA2* (Steinlein et al. 1995; Phillips et al. 2001). Relative to age-matched non-smoking subjects, there was a 10–20% increase in the binding of 2-[ $^{18}\text{F}$ ]fluoro-A-85380 to  $\alpha 4\beta 2$  receptors in the brain of patients with ADNFLE (Picard et al. 2006); knock-in mice bearing a culprit mutant  $\alpha 4$  gene have been prepared (Lipovsek et al. 2008), but have not been investigated by receptor autoradiography or PET. A recent clinical PET study of patients with idiopathic generalized epilepsy using the  $\alpha 4\beta 2$  nAChR ligand [ $^{18}\text{F}$ ]A-85380 showed focal 25% increases in the binding ratio within the anterior cingulate cortex (ACC), which could distinguish individual patients from healthy controls (Garibotto et al. 2019).

## 18.6 nAChR Imaging of Stroke and Neuroinflammation

Stroke is the leading cause of adult disability in the United States and Europe and the second leading cause of death worldwide. Stroke is characterized by a loss of brain functions due to rapid disturbances in cerebral blood supply, either as reduced blood flow by thrombosis or embolism (ischemic stroke) or bleeding (haemorrhagic stroke). Hyperacute mechanisms of stroke-related brain tissue damage, such as excitotoxicity, can be discriminated from delayed factors such as inflammation and apoptosis. All cellular components of the so-called neurovascular unit, which includes neurons, astrocytes and endothelial cells, express nAChRs (Paulson et al. 2010). Insofar as long-term tobacco smoking is a risk factor for ischemic stroke (Hawkins et al. 2002), it may be relevant that nAChRs were altered in a *post-mortem* study of smokers;  $\alpha 4$  expression was increased in neurons and dendritic processes, and  $\alpha 7$  expression was decreased in hippocampal neurons and astrocytes (Teaktong et al. 2004). In hypertensive stroke-prone rats, cortical  $\alpha 7$  nAChRs are reduced, without concomitant changes in the  $\alpha 4\beta 2$  nAChRs (Ferrari et al. 1999). Activation of nAChRs by nicotine promotes endothelial cell proliferation (Villablanca 1998) and leucocyte migration (Yong et al. 1997), which together may increase thrombotic risk. However, nAChR agonism was neuroprotective against excitotoxicity *in vitro*, an effect mediated by growth factors (Belluardo et al. 1998) and also by inactivation of the toxins (O'Neill et al. 2002). These effects were blocked by the  $\alpha 7$  nAChR antagonist  $\alpha$ -bungarotoxin (Donnelly-Roberts et al. 1996). Interestingly, nicotine increased oedema/infarct size in a rodent stroke model (Paulson et al. 2010). A particular contribution of  $\alpha 7$  nAChRs to stroke-related excitotoxicity might reflect their high  $\text{Ca}^{2+}$  permeability, especially under depolarizing conditions. Treatment with the acetylcholinesterase inhibitor methanesulfonyl fluoride attenuated stroke-induced learning and memory deficits in rats (Borlongan et al. 2005).

These rather discordant findings of effects of nAChR agonism in stroke models may reflect the different contributions of excitotoxicity and neuroinflammatory processes. The  $\alpha 7$  nAChRs expressed on microglia (Shytle et al. 2004a) seem particularly poised to mediate inflammatory responses. Nicotinic agonism at these sites suppressed inflammation by decreasing TNF- $\alpha$  production, while nicotine antagonists had the opposite effect (Shytle et al. 2004a). In addition to central mechanisms, cholinergic signalling in the “cholinergic anti-inflammatory pathway” involving the vagus nerve may suppress the release of pro-inflammatory cytokines and influence migration of T cells from the periphery to brain areas affected by stroke or multiple sclerosis (Borovikova et al. 2000) by a mechanism sensitive to  $\alpha$ -bungarotoxin (Pavlov et al. 2003).

Despite this extensive background, there have been very few molecular imaging studies of nAChRs in relevant neuroinflammation models. We have undertaken PET studies to assess  $\alpha 7$  nAChR alterations in the sheep stroke model (Boltze et al. 2008). Here, we measured [ $^{18}\text{F}$ ]DBT10 binding at different time points after permanent medial cerebral artery occlusion (pMCAO) by dynamic imaging using a clinical hybrid PET/MRI system. We found increased tracer uptake in the stroke-border zone 14 days after pMCAO. In these areas, microglia activation and macrophage infiltration were histologically confirmed. Ongoing studies aim to establish better the time course and histological correlates of the  $\alpha 7$  nAChR changes in our stroke model. A recent dual tracer study monitored longitudinal changes in 2-[ $^{18}\text{F}$ ]fluoro-A85380 and [ $^{11}\text{C}$ ]PK11195 binding during a month after middle cerebral artery occlusion (MCAO) in rats. In the ischemic territory, both ligands showed progressive binding increase from day 3 to 7 post-injury, followed by a progressive decrease (Martin et al. 2015). *Post-mortem* analysis linked the changes to increased  $\alpha 4\beta 2$  nAChR and TSPO expression on microglia and macrophages.

In PET studies in a rat stroke model with transient MCAO occlusion, the PET signal from the  $\alpha 7$  nAChR ligand [ $^{11}\text{C}$ ]NS14492 increased around the core of the infarct, consistent with activation of microglia and astrocytes following the injury (Colas et al. 2018). In that study, treatment with the  $\alpha 7$  agonist PHA 568487 1 week after the stroke lowered TSPO binding (suggesting a rescue from microgliosis) while diminishing the ultimate infarct volume. Thus, the  $\alpha 7$  nAChR is a promising target for disease-altering interventions against stroke.

Atherosclerosis is a kind of chronic inflammatory condition that brings a high risk of cardiovascular events. Especially the “vulnerable” atherosclerotic plaques have a high risk of rupture, which is predictable from their avidity on FDG PET, as an index of macrophage infiltration into the vessel wall. As such,  $\alpha 7$  nAChR PET presents an unexplored channel for investigating inflammatory changes in atherosclerosis (Boswijk et al. 2017). This follows also for a wide range of chronic inflammatory conditions such as rheumatoid arthritis, Crohn’s disease, etc.

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## 18.7 nAChR Imaging of Traumatic Brain Injury

Traumatic brain injury is a permanent or temporary impairment of brain functions with an associated diminished or altered state of consciousness caused by external mechanical force transferring kinetic energy to the brain tissue. In developed



countries, TBI is the most important cause of death and disability in young adults, in both civilian and military contexts (Olesen and Leonardi 2003) (Dewan et al. 2018). Indeed, TBI causes some 30–50% of all injury-related deaths (de Ramirez et al. 2012; Kamal et al. 2016). Apart from the distinct acute injuries, TBI is a continuous disease process (Masel and DeWitt 2010), with survivors often suffering from persistent or permanent physical and cognitive impairments (Fleminger and Ponsford 2005) occurring as long-term sequelae of the initial injury. This profile imposes a considerable socio-economic burden (Humphreys et al. 2013), which is further exacerbated by the failure of clinical trials aiming to improve outcomes (Loane and Faden 2010; Gruenbaum et al. 2016).

### 18.7.1 Animal Models of TBI

Experimental animal models of TBI provide a solid body of evidence for specific cholinergic alterations. A plethora of different TBI models has been developed since the late 1940s. There is a distinction to be made between closed head injuries and open-head models, whereby in the former model the underlying cortical tissue is damaged by impact on the intact *dura mater*. Most animal studies of cholinergic responses to TBI make use of fluid-percussion injuries (FPI), weight drop and controlled cortical impact (CCI) (Xiong et al. 2013) (O'Connor et al. 2011). This can be performed either through a fluidic wave, as in the FPI models (Thompson et al. 2005), rigid impactors such as weight drop (Marmarou et al. 1994) or CCI models (Lighthall 1988). Unfortunately, there is little data on cholinergic alterations provoked by the more clinically relevant (repeated) closed-head or concussive models. The CCI model, which was first developed in small animals (Lighthall 1988) and later used in pigs (Duhaime et al. 2000; Alessandri et al. 2003; Manley et al. 2006), entails a piston strike, which results in more focal brain injury than is afforded by the FPI model.

Early evidence of the involvement of the cholinergic system in TBI was obtained in dogs and cats, in which increased acetylcholine concentration was noted in cerebrospinal fluid up to 48 h post-injury (Bornstein 1946). In subsequent years, studies of TBI implicate diverse aspects of cholinergic neurotransmissions, including muscarinic and nicotinic receptors, enzymatic pathways and vesicular transporters (Hayes et al. 1992; Arciniegas 2011; Kelso and Oestreich 2012; Shin and Dixon 2015). In an autoradiographic study of rats with TBI, there were widespread and substantial bilateral reductions in cortical and hippocampal  $\alpha 7$  nAChRs labelled with [ $^{125}$ I] $\alpha$ -bungarotoxin, in contrast to lesser, more focal and ipsilateral effects on heteromeric nAChRs labelled with [ $^3$ H]epibatidine (Verbois et al. 2000). We have seen a similar decline in  $\alpha 7$  binding in the brain of rats and neonate piglets following FP injuries (Donat et al. 2008, 2010a, b; Hoffmeister et al. 2011). In a saturation binding study, the reductions were linked to declines in  $B_{\max}$ , rather than affinity changes (Verbois et al. 2002), an issue that must always be considered in autoradiographic or PET studies performed with a single ligand concentration. Peak effects occurred at 2 days post-injury but persisted for as long as 2 weeks. Brain regions not directly subjected to mechanical damage, such as the thalamus, were found to

exhibit reduced density of  $\alpha 7$  and  $\alpha 3/4$  nAChR, indicating retrograde and anterograde changes.

Most interestingly, brain areas showing reduced nAChR binding after TBI also showed delayed neuroinflammation, as indicated by increased binding of a tracer for the 18 kDa translocator protein, a marker of microglial activation (Donat et al. 2016), suggesting that nAChRs regulate neuroinflammatory responses (Egea et al. 2015). Indeed, in a model of blast-induced TBI, midbrain structures showed lowered gene transcripts of muscarinic and  $\alpha 7$  nAChR, accompanied by increased gene transcripts of pro-inflammatory markers (Valiyaveetil et al. 2013). Treatments with agonists and positive allosteric modulators of  $\alpha 7$  nAChR attenuated microglia activation after TBI while rescuing blood-brain barrier permeability increases and reducing motor deficits (Gatson et al. 2015; Dash et al. 2016). This is consistent with anti-inflammatory effects  $\alpha 7$  nAChR activation on microglia and potentially endothelial cells (Kimura, Dohgu et al. 2018), as indicated before (Cortes et al. 2017; Zhang et al. 2017).

Cognitive impairment was demonstrated in the rat FPI model (Scheff et al. 1997), suggesting that the  $\alpha 7$  nAChR reductions may contribute to these cognitive deficits caused by brain trauma. Also in the FPI paradigm, prolonged treatment with nicotine partially attenuated the cognitive deficits seen in the Morris water maze performance and reduced the magnitude and spatial extent of the  $\alpha 7$  changes (Verbois et al. 2003a, b). Indeed, nicotine has been effective in a number of other brain lesion models (Visanji et al. 2006; Huang et al. 2009; Zafonte et al. 2009; Quik et al. 2010), even though human patients who are smokers do not show improved outcome after TBI (Ostberg and Tenovuo 2014), because of smoking-associated health impairments such as vascular diseases. Furthermore, investigations in  $\alpha 7$  knockout mice failed to show much effect of  $\alpha 7$  receptor expression on lesion volume or microglia activation (Kelso et al. 2006), which could be confounded by compensatory mechanisms (Smith et al. 2014). Nonetheless, dietary choline supplements partially rescued the spatial memory deficits and  $\alpha 7$  receptor deficits after CCI (Guseva et al. 2008; Guseva et al. 2013).

### 18.7.2 Human TBI Studies

Neuroinflammation, a process mediated primarily by glia cells, i.e. microglia and astrocytes, is not just a hallmark pathology of TBI, but also a potential avenue of intervention (Morganti-Kossmann et al. 2019). Activated microglia, which are the resident macrophages of the brain, can assume many different roles post-injury, as characterized by diverse morphologies and molecular expression patterns. The dichotomy of pro-inflammatory M1 microglial phenotype versus anti-inflammatory M2 phenotype (Donat et al. 2017) may be an oversimplification (Ransohoff 2016), but it should not be assumed that all microglia are equal. A similar model of functions and phenotypes has been recently reported for astrocytes (Liddelov et al. 2017).

Glial activation can persist for many years after TBI (Ramlackhansingh et al. 2011). TBI is a risk factor for neurodegenerative diseases including chronic

traumatic encephalopathy (McKee et al. 2016) and dementias, including Alzheimer's disease (AD) (LoBue et al. 2018). The link between TBI and dementia may be mediated either by tau pathology or by altered cholinergic neurotransmission particularly involving hetero- and homomeric nAChR, as proposed for AD in the 1980s (Shimohama et al. 1986), in association with the cholinergic hypothesis of AD (Bohnen et al. 2018; Hampel et al. 2018). Historically, nAChR changes were linked to cognitive symptoms (Wallace and Bertrand 2013), as most  $\alpha 7$  and  $\alpha 4/3\beta 2/4$  nAChR are expressed on cortical neurons. However, cholinergic enzymes and receptors are also implicated in central and peripheral inflammatory response (Fujii et al. 2017), pointing to the above-mentioned cholinergic anti-inflammatory pathway (Martelli et al. 2014). Acetylcholine receptors may be essential mediators of this pathway and are likely involved in the neuroinflammatory response after TBI (Ren et al. 2017). Indeed, microglia express  $\alpha 7$  nAChR (Shytle et al. 2004b), as do astrocytes (Shen and Yakel 2012), where they seem to be directly involved in regulating glia activation. Reduced  $\alpha 7$  nAChR expression or antagonist exposure can promote the pro-inflammatory phenotype, while treatment with agonists may shift microglia towards an anti-inflammatory phenotype (Cortes et al. 2017; Zhang et al. 2017).

Several studies have investigated key players of the cholinergic neurotransmission following TBI. However, only very few studies investigated human patients or tissue samples. In a human *post-mortem* study, performed after traumatic brain injury, reduced choline acetyl transferase activity and synaptophysin immunoreactivity in cerebral cortex were found, indicating damage of the cholinergic innervation, but nAChR binding determined by either [ $^3\text{H}$ ]nicotine or [ $^{125}\text{I}$ ] $\alpha$ -bungarotoxin was unchanged (Murdoch et al. 1998), suggesting mainly presynaptic mechanisms. *Post-mortem* cortical tissue from patients acquired between 1 and 300 h after injury showed a reduction in choline acetyltransferase activity, but no change in muscarinic (M1 and M2 receptors) or nicotinic receptors, determined by either [ $^3\text{H}$ ]nicotine or [ $^{125}\text{I}$ ] $\alpha$ -bungarotoxin (Dewar and Graham 1996; Murdoch et al. 1998). Reduced synaptophysin immunoreactivity was seen as an indicator of mainly presynaptic pathomechanisms, even though a follow-up study showed extensive damage in basal forebrain cholinergic neurons (Murdoch et al. 2002). Interestingly, diffusion tensor imaging indicated axonal injury in the same areas (Hong et al. 2012). More importantly, recent studies of CTE showed a gradual emergence of pretangle pathology and oligomeric tau accumulation in cholinergic neurons of the basal forebrain, which might directly relate to cholinergic impairments and ultimately to cognitive dysfunction and immune dysregulation (Mufson et al. 2016). Laser capture microdissection and gene profiling of these neurons showed, among other findings, a reduction in CHRN2 transcripts, encoding the nAChR  $\beta 2$  subunit (Mufson et al. 2018). Another potential factor in TBI is autoimmune consequences (Raad et al. 2014). One study reported a significant increase in blood autoantibodies to regions of the  $\alpha 7$  subunit in children with TBI, which correlated to severity (Sorokina et al. 2012).

In vivo molecular imaging with nAChR subtype-specific tracers is still lacking in clinical TBI. However, PET imaging with [ $^{11}\text{C}$ ]MP4A in chronic TBI patients

indicated loss of AChE activity (Östberg et al. 2011; Ostberg et al. 2018). Treatment of post-traumatic cognitive deficits using cholinergic compounds, e.g. acetylcholinesterase inhibitors, has been attempted in several studies of small patient groups. However, most of these exploratory studies did not meet their endpoints justifying larger randomized placebo-controlled studies, and systematic reviews therefore concluded only weak evidence of efficacy (Wheaton et al. 2011; Bengtsson and Godbolt 2016).

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## 18.8 nAChR Imaging of Addiction and Psychiatric Disorders

### 18.8.1 Physiological Effects of Nicotine in the Context of Addiction

Whereas acetylcholine is rapidly inactivated by AChE, nicotine exerts prolonged agonism due to its metabolic stability. This property accounts for the well-known phenomenon of desensitization of nicotinic receptors by nicotine (Govind et al. 2012), in which inactivated receptors concentrate in the plasma membrane due first to agonist-evoked conformational changes, which is followed by decreased degradation of subunits (Govind et al. 2012). In the case of  $\alpha 4\beta 2$  nAChRs, upregulation is mediated by activation of protein kinase C, which results in phosphorylation of the  $\alpha 4$  subunit (Wecker et al. 2010). This mechanism is supported by studies in rats and 3xTd-AD mice showing upregulation of  $\alpha 4\beta 2$  nAChR after chronic nicotine administration (Flores et al. 1992; Oddo et al. 2005). Chronic nicotine treatment persistently increased the binding of the  $\alpha 4\beta 2$  ligand 5-[ $^{123}\text{I}$ ]iodo-A-85380 by approximately 50% in the brain of baboons (Kassiou et al. 2001). Similarly, a 2-[ $^{18}\text{F}$ ]fluoro-A-85380 PET study in chronic smokers abstinent for at least 24 h revealed persistent upregulation of  $\alpha 4\beta 2$  nAChR throughout the brain (Mukhin et al. 2008); these findings stress the requirement for strict control of exposure to nicotine in the design of clinical and preclinical imaging studies of nAChRs. Thus, exposure of rats to tobacco smoke not only induced nicotine dependence but also increased the  $\alpha 7$  nAChR density in the CA2/3 area (+ 25%) and the stratum oriens (+ 18%) of the hippocampus (Small et al. 2010).

Since the nAChRs are ligand-gated cation channels, their activation facilitates depolarization and enhances the release of dopamine and other neurotransmitters. Consistent with the increased energy demands associated with depolarization, acute challenge with nicotine increases the cerebral metabolic rate for glucose (CMRglc) as measured by [ $^{14}\text{C}$ ]-deoxyglucose autoradiography; these effects are most notable in the thalamus and other rat brain regions in which nicotinic receptors are most abundant (London et al. 1988; Marenco et al. 2000). Despite the phenomenon of upregulation/desensitization of nAChRs, stimulation of CMRglc in rat brain was still evident after reinstatement of nicotine following a period of withdrawal (Schröck and Kuschinsky 1991). Effects of nicotine on CMRglc are poorly documented in human brain, but one PET study showed that nicotine-evoked stimulation of 2-[ $^{18}\text{F}$ ]fluorodeoxyglucose uptake in non-smokers was dependent on the hostility

trait, as was also seen in smoking subjects treated with a higher dose of nicotine (Fallon et al. 2004). The latter finding seems consistent with desensitization of the cerebrometabolic response to nicotine challenge in the smoking group. Smoking provoked a delayed increase in the cerebral consumption of oxygen in the brain of acutely withdrawn smokers, suggesting a global effect of nicotine or other constituents of tobacco smoke on mitochondrial respiration as distinct from glycolysis (Vafaei et al. 2015).

Baseline levels of 2-[<sup>18</sup>F]fluoro-A-85380 binding in the brainstem of healthy squirrel monkeys predicted their motivation to subsequently self-administer nicotine, as indicated by number of bar presses (Le Foll et al. 2009), suggesting that nAChRs mediate a trait vulnerability specifically for nicotine abuse. It is unclear whether this association with nicotine addiction generalizes to other addictive drugs. However, activation of nAChRs on dopamine neurons and terminals is central to the reinforcing and addictive properties of nicotine. Stimulation of striatal dopamine release by nicotine has recently been linked to  $\alpha 6\beta 2\beta 3$  and  $\alpha 6\alpha 4\beta 2\beta 3$  nAChRs, which predominate in the basal ganglia (Quik et al. 2011). Activation of these receptors increases the firing rate and augments phasic bursting of midbrain dopamine neurons (De Biasi and Dani 2011). The  $\alpha 7$  nAChRs in the ventral tegmental area may mediate nicotine's stimulatory effect on mesolimbocortical dopaminergic function and consequently its reinforcing and dependence-producing properties (Nomikos et al. 2000).

In an early molecular imaging study of nAChRs, binding of the SPECT tracer 5-[<sup>123</sup>I]-A-85380 at  $\alpha 4\beta 2$  sites was increased by one third in the cortex and striatum of smokers with confirmed abstinence for 1 week (Staley et al. 2006). This study recapitulated in human smokers the well-known upregulation and inactivation of nAChRs following repeated nicotine exposure seen in experimental animals. The same group showed normalization of nAChR availability after 6 weeks of abstinence (Cosgrove et al. 2009). The competition paradigm affords the possibility of detecting occupancy by exogenous nicotine or other drugs at nAChR in living individuals. Here, comparison of molecular imaging results at baseline and after drug challenge reveals the percentage of receptors occupied by the drug. In one such 5-[<sup>123</sup>I]-A-85380 SPECT study, smoking to satiety (about two cigarettes) induced a 67% reduction in their chosen endpoint,  $V_T/f_p$  (Esterlis et al. 2010a). Occupancy after use of a nicotine inhaler (56%) was significantly lower than after smoking cigarettes to satiety (Esterlis et al. 2010b).

These nAChR occupancies may be underestimated due to the slow kinetics of the SPECT tracer relative to the pharmacodynamics of nicotine. Corresponding PET studies with 2-[<sup>18</sup>F]fluoro-A-85380 showed 79% occupancy after smoking a low-nicotine cigarette and 26% after smoking a denicotinized cigarette (Brody et al. 2009). In another 2-[<sup>18</sup>F]fluoro-A-85380 PET study, a few puffs on a normal cigarette induced 50% occupancy persisting 3 h later, whereas an entire cigarette provoked 88% occupancy (Brody et al. 2006). Even exposure to second-hand smoke led to 20% occupancy (Brody et al. 2011). A single low dose (0.5 mg) of varenicline provoked complete saturation of 2-[<sup>18</sup>F]fluoro-A-85380 binding sites in human brain (Lotfipour et al. 2012a). Thus, varenicline is apt to evoke full agonism at  $\alpha 7$

nAChRs, with partial agonism at other sites, i.e.  $\alpha 4\beta 2$ , as well as  $\alpha 3\beta 4$ , and  $\alpha 6\beta 2$  subtypes.

Altered dopamine release can be detected in PET studies with [ $^{11}\text{C}$ ]raclopride and other benzamide antagonists of  $\text{D}_{2/3}$  receptors, wherein ligand binding is reduced by competition from endogenous dopamine (Laruelle 2000; Cumming et al. 2003). Whereas powerful psychostimulants such as amphetamine can evoke 30% decreases in striatal [ $^{11}\text{C}$ ]raclopride binding, challenge with nicotine reduced this binding by only 5–10% in striatum of anaesthetized pigs, most notably in the ventral striatum (Cumming et al. 2003). There were effects of similar magnitude in the brain of non-human primates (Marenco et al. 2004), although others saw no such effects in PET scans of awake monkeys (Tsukada et al. 2002), suggesting a confounding effect of general anaesthesia. Similarly, intranasal administration of nicotine did not greatly reduce striatal [ $^{11}\text{C}$ ]raclopride binding in healthy habitual smokers, although baseline binding correlated with scores of nicotine dependence, and there was a positive relationship between binding changes and individual reports of pleasant subjective experience (Montgomery et al. 2007), as likewise seen in another [ $^{11}\text{C}$ ]raclopride PET study of smokers (Barrett et al. 2004). In general, preclinical studies with non-contingent pharmacological challenge may not replicate the motivational aspects of self-administered nicotine.

Dopamine synthesis capacity in living striatum can be measured by PET studies with DOPA decarboxylase substrates such as [ $^{11}\text{C}$ ]DOPA or 6- $^{18}\text{F}$ ]fluoro-DOPA (FDOPA). Whereas acute nicotine treatment had no effect on [ $^{11}\text{C}$ ]DOPA utilization in striatum of awake monkeys, this utilization was reduced by one third after overnight abstinence in monkey habituated to nicotine and normalized rapidly following reinstatement of nicotine treatment (Domino et al. 2009). On the other hand, in an isolated clinical PET finding, FDOPA utilization was 20–30% higher in striatum of human smokers (Salokangas et al. 2000). Another study showed a 20% reduction in the striatal utilization of FDOPA in a larger group of dependent smokers (Rademacher et al. 2016), which may have greater face validity as an adaptive response to chronic nicotine exposure.

## 18.8.2 Alcohol Dependence

There is very high comorbidity between nicotine and alcohol dependence, and considerable overlap between their neural mechanisms of addiction (Larsson and Engel 2004). Indeed, a possible role for nAChR in alcohol addiction has been proposed (Meyerhoff et al. 2006). Altered nACh receptor density *in vitro* has been observed in rats in response to ethanol using a combination of [ $^3\text{H}$ ]nicotine, [ $^3\text{H}$ ]MLA and [ $^{125}\text{I}$ ]alpha-bungarotoxin, but the direction of change was inconsistent and region-specific (Yoshida et al. 1982; Booker and Collins 1997; Robles and Sabria 2008). *In vivo*, binding of the SPECT radioligand [ $^{123}\text{I}$ ]5-IA-85380 in rhesus monkey was unaltered following chronic alcohol self-administration, although decreases in cortical and thalamic binding were observed following abstinence (Cosgrove et al. 2010). More recently, using [ $^{18}\text{F}$ ]nifene in a PET study in non-human primates,

reductions in nACh receptor density were observed following chronic alcohol exposure (Hillmer et al. 2014). Further validation work is required in human subjects to resolve these inconsistencies.

### 18.8.3 Schizophrenia and Depression

Nicotine addiction is notoriously prevalent among patients with schizophrenia (Lohr and Flynn 1992), a chronic psychiatric disease characterized by behavioural changes, which are difficult to model in experimental animals. The DSM-IV criteria for schizophrenia consist of positive symptoms (such as agitation, paranoia and hallucinations), negative symptoms (including emotional blunting, avolition and social withdrawal) and specific cognitive and psychomotor deficits (Goldman-Rakic and Selemon 1997). An increasing appreciation of the importance of cognitive deterioration has motivated a search for treatments to improve processing speed, attention/vigilance and working memory (Nuechterlein et al. 2004; Heinrichs 2005). This has emerged in the context of a hypothesis implicating gene-mediated dysfunction of  $\alpha 7$  nAChRs (Freedman et al. 1997; Stephens et al. 2009; Dome et al. 2010) in the cognitive impairments of schizophrenia (Nomikos et al. 2000). Indeed, in one *post-mortem* study, the hippocampal [ $^{125}$ I] $\alpha$ -bungarotoxin binding was decreased in a schizophrenic patient population consisting mainly of smokers (Freedman et al. 1995). Furthermore there were significant differences in  $\alpha 7$  nAChR (*CHRNA7*) expression at both mRNA and protein levels between smokers and non-smokers with schizophrenia (Mexal et al. 2010).

Nicotinic receptors and treatments have been investigated in a number of murine models emulating some behaviour aspects of schizophrenia. An auditory gating defect in dilute brown non-Agouti (DBA/2) mice is rectified by treatment with an  $\alpha 4\beta 2$  agonist (Wildeboer and Stevens 2008), but other nAChR subtypes such as  $\alpha 7$  may also mediate sensory gating in these mice (Radek et al. 2006). Mice lacking the plasma membrane dopamine transporter show hyperactivity and cognitive deficits, and are hypersensitive to the locomotor stimulant effect of nicotine, which models schizophrenia. Treatment of these mice with nicotine improved their performance in spatial and cued learning tasks (Weiss et al. 2007a), whereas autoradiography showed a small decline in the  $\beta 2$  subunit, a large decrease in  $\beta 6$  and substantially increased  $\alpha 7$  nAChR expression (Weiss et al. 2007b). Deficiency of the microtubule-stabilizing protein STOP results in impaired hippocampal plasticity and behavioural hypersensitivity to psychostimulants, likewise associated with a decrease in  $\beta 6$ , and a substantial increase in  $\alpha 7$  nAChRs (Bouvrais-Veret et al. 2007), suggesting a common nicotinic pathway in these two models of schizophrenia.

A preliminary study with the  $\alpha 7$ -nAChR ligand [ $^{18}$ F]ASEM showed a moderate reduction of binding in the brain of a group of patients with schizophrenia (Wong et al. 2018). This finding was recapitulated in another study of recent onset psychosis (Coughlin et al. 2018a). Given the association of  $\alpha 7$  nAChR expression with microglia, the resident brain macrophage, it is a matter of interest if there are changes in inflammatory pathways in association with schizophrenia. A recent

meta-analysis of many PET TSPO studies in patients with schizophrenia showed an overall finding of reduced microglial activation (Plaven-Sigra et al. 2018). While this was contrary to general expectation of an active inflammatory process during schizophrenia, it may be consistent with the  $\alpha7$ -nAChR findings.

Availability of  $\alpha4\beta2$  nAChR binding sites was globally reduced in a 5-[<sup>123</sup>I]iodo-A-85380 SPECT study of patients with major depression (Saricicek et al. 2012). Changes in behaviour, and in endocrine, and immune and neurotransmitter systems, modelling symptoms of patients with major depression have been described in rats following bilateral olfactory bulbectomy (Song and Leonard 2005). Such rats present characteristic alterations in cholinergic function (Hozumi et al. 2003). Although degeneration of cholinergic neurons may well underlie the impairments of learning and memory-related behaviour in olfactory bulbectomy rats, densities of nAChRs have not yet been investigated in this model. Interestingly, a recent TSPO PET study showed globally increased microglial expression in cortex of patients with major depression and an association with cognitive deficits (Li et al. 2018). However, there has hitherto been no PET study of  $\alpha7$ -nAChR depressed (non-smoking) patients.

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## 18.9 nAChR Imaging for Measurement of Endogenous Acetylcholine

The competition model for measuring dopamine release by molecular imaging, i.e. the [<sup>11</sup>C]raclopride binding model described above, has delivered substantial insights into the role of dopamine in the pathophysiology of schizophrenia and nicotine addiction. However, attempts to generalize the dopamine competition model to other neurotransmitter systems have met with mixed success (Paterson et al. 2010; Finnema et al. 2015). A molecular imaging assay for fluctuations in extracellular acetylcholine would be a useful tool for evaluating new pharmacologic treatments for Alzheimer's disease and other conditions, and the first demonstration of an acetylcholine sensitive  $\alpha4\beta2$  PET tracer was recently published (Hillmer et al. 2016a). Competition between endogenous acetylcholine and radioligands targeting muscarinic acetylcholine receptors is reported in the literature (Dewey et al. 1993; Sahara et al. 1994; Ma et al. 2004; Eckelman 2006), but the case is better established for nAChR ligands, the details of which are presented here.

Preclinical studies in rodents suggest that treatment with AChE inhibitors may raise acetylcholine levels sufficiently to elicit changes in binding potential with PET. Treatment with AChE inhibitors, an important class of compounds used for alleviation of Alzheimer's disease symptoms, affords a convenient way of enhancing acetylcholine levels in living brain in tests of the competition model. In the first such study, the uptake of 2-[<sup>18</sup>F]fluoro-A-85380 in rat thalamus measured *ex vivo* was 45% reduced by the AChE inhibitor physostigmine (Dollé et al. 1999); decreases of similar magnitude were seen in animals treated with typical  $\alpha4\beta2$  agonist ligands (2-fluoro-A-85380, nicotine, epibatidine and cytisine) but not by  $\alpha7$ - or 5-HT<sub>3</sub>-specific antagonists. Previous microdialysis studies had confirmed that the same



dose of physostigmine (300  $\mu\text{g}/\text{kg}$ ) caused large increases in interstitial acetylcholine (Cuadra et al. 1994), thus substantiating the competition model.

In a non-human primate PET study, Ding et al. investigated whether modulation of acetylcholine levels could modulate [ $^{18}\text{F}$ ]NFEP binding (Ding et al. 2000b). As predicted, binding was 25% lower in the thalamus and striatum following physostigmine administration. Furthermore, treatment with the dopamine  $D_{2/3}$  antagonist raclopride reduced the striatal binding of [ $^{18}\text{F}$ ]NFEP by 22%, and the agonist quinpirole increased striatal binding by 26%, but selective dopamine  $D_1$  drugs had no such effects. These findings were consistent with results of cerebral microdialysis studies showing that dopamine  $D_{2/3}$  receptors tonically inhibit acetylcholine release in the striatum (Damsma et al. 1991; DeBoer et al. 1996).

In a 5- $^{123}\text{I}$ iodo-A-85380 SPECT study in non-human primate, a bolus plus constant infusion paradigm was used to obtain equilibrium binding prior to physostigmine treatment (Fujita et al. 2003). Due to instability of the plasma 5- $^{123}\text{I}$ iodo-A-85380 concentration following physostigmine, the  $\text{BP}_{\text{ND}}$  was not calculated, but there was a 14–17% decrease in thalamic tracer uptake. This decline was independent of the physostigmine dose administered, suggesting a ceiling effect in the competition from endogenous acetylcholine. In a subsequent 2- $^{18}\text{F}$ fluoro-A-85380 study, a more prolonged infusion of physostigmine evoked a 40% reduction in 2- $^{18}\text{F}$ fluoro-A-85380  $V_T$  in putamen and more modest reduction in cerebral cortex (Valette et al. 2005). The effect was dose-dependent, and infusion of galantamine, a weaker AChE inhibitor, did not alter 2- $^{18}\text{F}$ fluoro-A-85380 binding, which again suggests a fairly narrow relationship between interstitial acetylcholine concentration and increased competition.

In PET studies in rats, cortical and thalamic [ $^{18}\text{F}$ ]nifene binding was found to be significantly reduced (3–10%) following physostigmine and galantamine administration (Hillmer et al. 2013) suggesting this ligand may also be sensitive to changes in endogenous acetylcholine. In vitro binding studies with [ $^{18}\text{F}$ ]nifene and [ $^3\text{H}$ ]cytosine showed displacement of both ligands upon addition of an AChE inhibitor to the medium (Easwaramoorthy et al. 2007). This highlights the rapid hydrolysis of endogenous acetylcholine in the presence of native AChE and raises the consideration that inactivation/upregulation of nAChRs may arise from “supernormal” activation during AChE inhibition, or during exposure to nicotine or other long-lived agonists. Breakdown of released acetylcholine is usually complete within a matter of seconds (Bruno et al. 2006). As such, the PET observations described above cannot unambiguously be ascribed to the competition model; some of the decreases in receptor binding after treatment with AChE inhibitors might rather be attributable to transition of the receptors to a low-affinity state or to trafficking mechanisms. This is especially a consideration in studies with ligands with slow kinetics, such as in the above reports.

More recently, (–)- $^{18}\text{F}$ flubatine ((–)- $^{18}\text{F}$ ]NCFHEB) was shown to be sensitive to displacement by endogenous acetylcholine in non-human primates. Treatment with AChE inhibitors donepezil and physostigmine led to dose-dependent, significant reductions (10–34%) in  $V_T$  in bolus and bolus-infusion protocols, respectively (Gallezot et al. 2014). This finding was soon followed by a similar study in human

subjects (Hillmer et al. 2016a), in which a bolus and bolus-infusion protocol with [ $^{18}\text{F}$ ]flubatine revealed small but significant reductions in  $V_T$  in cortical regions following physostigmine administration.

Overall these data suggest that measuring acetylcholine changes using selective  $\alpha 4\beta 2$  nAChR radioligands may be possible and that we should expect future applications of this methodology in clinical populations such as Alzheimer's disease, substance dependence and schizophrenia.

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## 18.10 Conclusion

The development of selective ligands for nAChRs has been challenging due to the diversity of subtypes existing in the brain and due to the unfavourable kinetics and toxicity profiles of some lead compounds. Promising novel tracers targeting  $\alpha 4\beta 2$  nAChRs include [ $^{18}\text{F}$ ]AZAN, [ $^{18}\text{F}$ ]nifene, [ $^{18}\text{F}$ ]XTRA and both the (–) and (+) enantiomers of [ $^{18}\text{F}$ ]flubatine. For the  $\alpha 7$  nAChR, [ $^{18}\text{F}$ ]DBT10 and [ $^{18}\text{F}$ ]ASEM are promising candidates. Despite the importance of  $\alpha 6$  nAChRs in the action of nicotine on dopamine release, selective tracers for this target remain elusive.

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# Development of PET and SPECT Radioligands for In Vivo Imaging of NMDA Receptors

# 19

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## Abstract

*N*-Methyl-d-aspartate receptors (NMDARs) play an important role in the neurotransmission of the central nervous system (CNS). On the other hand, aberrant functioning of the NMDARs has been implicated in various CNS disorders. Positron emission tomography (PET) and single photon emission computed tomography (SPECT) imaging of NMDARs may provide novel insights on CNS functions and various CNS disorders that could lead to the discovery of potential drug therapies. Despite numerous efforts to develop a number of radioligands for NMDARs since the late 1980s, there are no clinically useful in vivo imaging probes. However, recent continuous efforts have provided several promising

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radiotracers for NMDARs, such as 3-benzazepine derivatives targeting GluN2B. In this chapter, we summarize the progress in the development of PET and SPECT imaging agents for NMDARs and discuss the prospects in this field.

## Abbreviations

% ID/g	Percent injected dose per gram of tissue
% ID/mL	Percent injected dose per milliliter of tissue
3MPICA	3-[2-[(3-methoxyphenylamino)carbonyl]ethenyl]-4,6-dichloroindole-2- carboxylic acid
AD	Alzheimer's diseases
BBB	Blood-brain barrier
BP	Binding potential
CBZ	Benzyloxycarbonyl
<i>cis</i> -HPTC	<i>cis</i> -2-Hydroxymethyl- <i>r</i> -1-( <i>N</i> -piperidyl)-1-(2-thienyl)cyclohexane
CNS 1261	<i>N</i> -(1-Naphthyl)- <i>N'</i> -(3-iodophenyl)- <i>N'</i> -methylguanidine
CNS 5161	<i>N</i> -(2-Chloro-5-(methylmercapto)phenyl)- <i>N'</i> -methylguanidine monohydrochloride
CNS PET MPO	Central nervous system PET multiparameter optimization
CNS	Central nervous system
CP-101,606	( <i>1S,2S</i> )-1-(4-Hydroxyphenyl)-2-(4-hydroxy-4-phenylpiperidino)-1-propanol
DAO	<sub>D</sub> -Amino acid oxidase
DCKA	5,7-Dichlorokynurenic acid
DTG	1,3-Di- <i>o</i> -tolylguanidine
EMD-95885	6-3-4-(4-Fluorobenzyl)piperidinopropionyl-3H-benzoxazol-2-one
GE-179	<i>N</i> -(2-Chloro-5-(2-fluoroethylthiophenyl)- <i>N'</i> -(3-thiomethylphenyl)- <i>N'</i> -methylguanidine
GMOM	<i>N</i> -(2-Chloro-5-thiomethylphenyl)- <i>N'</i> -(3-methoxy-phenyl)- <i>N'</i> -methylguanidine
GV150526A	3-(2-((Phenylamino)carbonyl)ethenyl)-4,6-dichloroindol-2-carboxylic acid
HACH242	<i>N</i> -((5-(4-Fluoro-2-methoxyphenyl)pyridin-3-yl)methyl)cyclopentanamine
HON0001	7-Hydroxy-6-methoxy-2-methyl-1-(2-(4-(trifluoromethyl)phenyl)ethyl)-1,2,3,4-tetrahydroisoquinoline hydrochloride
HPLC	High-performance liquid chromatography
Ifenprodil	4-[2-(4-Benzylpiperidin-1-yl)-1-hydroxypropyl]phenol
Ketamine	( <i>R,S</i> )-2-(2-Chlorophenyl)-2-(methylamino)cyclohexanone
Kryptofix 2.2.2	4, 7,13,16,21,24-Hexaoxa-1,10-diazabicyclo[8.8.8]-hexacosane
L-689,560	<i>trans</i> -2-Carboxy-5,7-dichloro-4-phenylaminocarbonylamino-1,2,3,4-tetrahydroquinoline

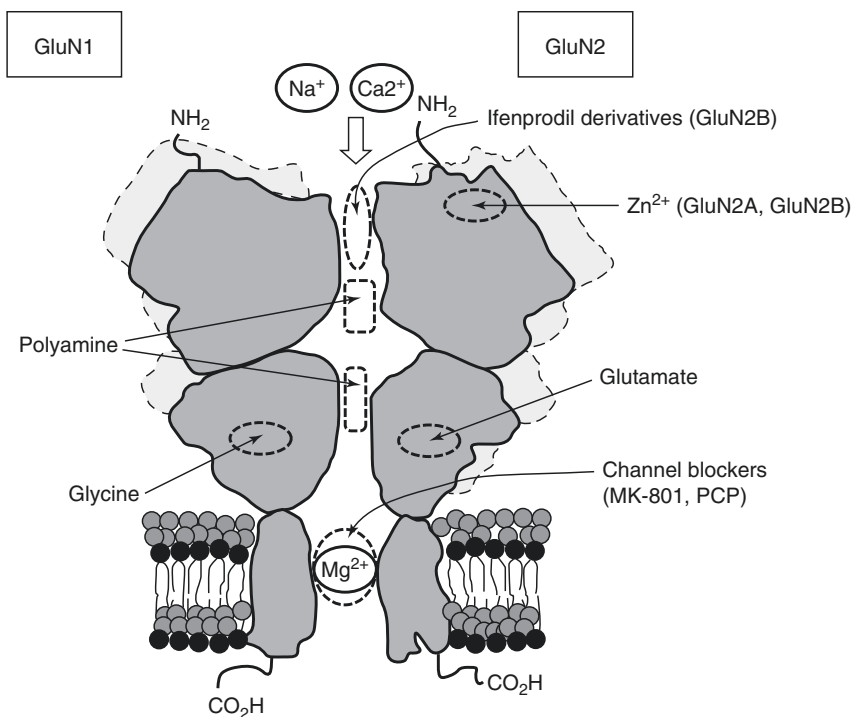
L-703,717	3-[3-(4-Methoxybenzyl)phenyl]-4-hydroxy-7-chloroquinolin-2(1H)-one
MDL-104,653	3-Phenyl-4-hydroxy-7-chloro-quinolin-2(1H)-one
MDL-105519	(Z)-2-Carboxy-4,6-dichloroindole-3-(2'-phenyl-2'-carboxy)-ene
MEM	1-Amino-3-fluoromethyl-5-methyl-adamantane
Memantine	3,5-Dimethyladamantan-1-amine
MK-0657	(3 <i>S</i> ,4 <i>R</i> )-4-Methylbenzyl 3-fluoro-4-((pyrimidin-2-yl-amino)methyl) piperidine-1-carboxylate
MK-801	Dizocilpine maleate {(+)-10,11-dihydro-5-methyl-5 <i>H</i> -dibenzo[ <i>a,d</i> ] cyclohepten-5,10-diylidiammonium maleate}
N2B-05182	((1-(4-Fluoro-3-methylphenyl)-1 <i>H</i> -1,2,3-triazol-4-yl)methoxy)-5-methoxypyrimidine
NCAM	<i>N</i> -[2-( <i>N</i> -(2-Mercaptoethyl)aminoethyl)- <i>N</i> -(2-mercaptoethyl)-3,5-dimethyl acetamide amantadine
NHAM	[1-[ <i>N</i> -[ <i>N</i> -(2-Mercaptoethyl)]- <i>N</i> -[2-(2-mercaptoethyl)amino] amino ethyl] amino-3,5-dimethyladamantane
NMDAR	<i>N</i> -Methyl- <i>d</i> -aspartate receptors
NPS 1506	[3-Fluoro- $\gamma$ -(3-fluorophenyl)- <i>N</i> -methylbenzenepropamine]
PAMQX	( <i>D</i> )-7-Iodo- <i>N</i> -(1-phosphonoethyl)-5-aminomethylquinoline-2,3-dione
PCP	Phencyclidine {1-(1-phenylcyclohexyl) piperidine}
PD	Parkinson's disease
PET	Positron emission tomography
PK-209	(3-(2-Chloro-5-(methylthio)phenyl)-1-(3-(fluoromethoxy)phenyl)-1-methylguanidine)
RGH-896	2-[4-[(4-Fluorophenyl)methyl]piperidin-1-yl]-2-oxo- <i>N</i> -(2-oxo-3 <i>H</i> -1,3-benzoxazol-6-yl)acetamide
Ro 25-6981	[( $\pm$ )-( <i>R</i> *, <i>S</i> *)]- <i>a</i> -(4-Hydroxyphenyl)- <i>b</i> -methyl]-4-(phenylmethyl)-1-piperidine propanol
Ro-647312	[2-(3,4-Dihydro-1 <i>H</i> -isoquinolin-2-yl)-pyridin-4-yl]-imethylamine
SAR	Structure-activity relationship
SPECT	Single photon emission computed tomography
ST3	{( <i>S</i> )-5-[( <i>R</i> )-2-Amino-2-carboxyethyl]-1-[4-(3-fluoropropyl)phenyl]-4,5-dihydro-1 <i>H</i> -pyrazole-3-carboxylic acid
TCP	<i>N</i> -(1-[2-Thienylcyclohexyl]-3,4-piperidine
TLC	Thin-layer chromatography
$V_T$	Total volume of distribution
WMS-1405 (Me-NB1)	7-Methoxy-3-(4-phenylbutyl)-2,3,4,5,-tetrahydro-1 <i>H</i> -3-benzazepin-1-ol

## 19.1 Introduction

*N*-Methyl-d-aspartate receptors (NMDARs) are glutamate-gated ion channels that play a central role in excitatory neurotransmission and are known to be involved in learning, memory, and synaptic plasticity (Riedel et al. 2003; Lau and Zukin 2007; Traynelis et al. 2010). On the other hand, excessive glutamate causes overactivation of NMDARs, which results in an increase of intracellular calcium. Subsequent downstream events, including activation of nitric oxide synthase, calcium-sensitive proteases, and mitochondrial damage, can induce neuronal death. This glutamate excitotoxicity is thought to be implicated in the pathophysiology of various neurological diseases, including epilepsy, ischemia, stroke, and neurodegenerative disorders, such as Parkinson's disease (PD), Alzheimer's disease (AD), and Huntington's disease. Hypofunction of NMDARs has been suggested to be related to the pathology of schizophrenia. Therefore, NMDARs have been considered to be attractive therapeutic targets for these diseases (Kemp and McKernan 2002; Kalia et al. 2008; Lau and Tymianski 2010; Paoletti et al. 2013).

NMDARs are unique ligand-gated ion channels. Activation of the receptors not only requires simultaneous binding of two amino acids, glycine and glutamate, but also removal of magnesium ( $Mg^{2+}$ ) blockade by membrane depolarization. These events lead to the opening of the receptor pore, followed by calcium influx into cells. Consequently,  $Ca^{2+}$ -dependent signal transduction cascades trigger the modulation of many aspects of neuronal function (Dingledine et al. 1999; Traynelis et al. 2010). The functional NMDARs require assembly of two GluN1 subunits (previously named NR1 subunits) together with either two GluN2 (previously named NR2 subunits) or a combination of GluN2 and GluN3 (previously named NR3 subunits) (Laube et al. 1998; Furukawa et al. 2005; Ulbrich and Isacoff 2008; Paoletti et al. 2013). GluN1 have only one gene (eight splice variants) with ubiquitous distribution in the brain. GluN2 are divided into four types (GluN2A–D). These subunits show distinct anatomical distributions in the brain which determine the biophysical and pharmacological properties of the NMDARs (Köhr 2006; Paoletti and Neyton 2007). GluN3 have glycine-binding sites, and glycine alone is sufficient to activate GluN1/GluN3 receptors. However, the function of GluN3-containing NMDARs remains to be fully elucidated (Low and Wee 2010).

The NMDAR's architecture has been reported to be clamshell-like with several binding sites for endogenous and exogenous ligands as shown in Fig. 19.1. While the glutamate-binding site is exhibited on the GluN2, the glycine-binding site is located on the GluN1 or GluN3 (Furukawa and Gouaux 2003; Furukawa et al. 2005; Yao et al. 2008). Zinc ( $Zn^{2+}$ ) has been reported to bind to GluN2A or GluN2B and inhibits the ion channel activity in a non-competitive manner (Rachline et al. 2005; Karakas et al. 2009). Polyamines are allosteric potentiators of GluN2B-containing NMDARs, presumably through recognition of the dimer interface of GluN1 and GluN2B (Mony et al. 2009). A number of exogenous ligands for the several binding sites of NMDARs have been developed as neuroprotective agents. Numerous small compounds have been identified to exhibit binding on the glutamate-binding sites or glycine-binding sites in a competitive manner (Danysz and Parsons 1998; Dingledine et al. 1999). Aliphatic cyclic amines, such as dizocilpine maleate (MK-801) and phencyclidine (PCP), have been reported to inhibit NMDARs strongly in a non-competitive and



**Fig. 19.1** Structure of NMDARs and location of ligand binding site

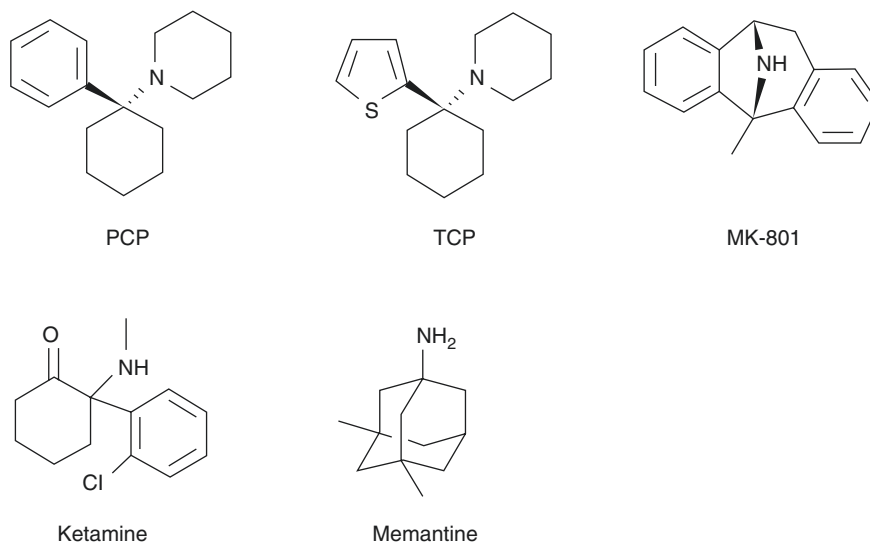
voltage-dependent manner (Anis et al. 1983; Foster and Wong 1987). These compounds are called open channel blockers because the blocking effect on NMDARs is caused by binding to the channel pore region near the Mg<sup>2+</sup> binding site (Kashiwagi et al. 1997). Recently, ifenprodil and related compounds, which are GluN2B-selective antagonists, have been identified to bind at the interface between GluN1 and GluN2B (Karakas et al. 2011).

Positron emission tomography (PET) and single photon emission computed tomography (SPECT) imaging for NMDARs could be powerful tools for obtaining information about the functional mechanism of the NMDA ion channel in the living brain, the pathophysiology of related neurological disorders, and in vivo occupancy of agonists or antagonists of NMDARs. Although a conclusive relationship between NMDAR expression and neurodegenerative disease has yet to be elucidated, several reports suggested that NMDAR expression is deregulated in neurological patients. It is reported that mRNA and protein levels of NMDAR GluN2A and GluN2B were decreased in AD patients (Hynd et al. 2004; Jacob et al. 2007). In addition, [<sup>3</sup>H]Ro 25-6981 (GluN2B-selective ligand as described below) binding was increased in the putamen of PD patients experiencing motor complications (Calon et al. 2003), and protein levels of the GluN1 were decreased in the postmortem left hippocampus in schizophrenic patients (Vrajová et al. 2010). Therefore, nuclear medicine imaging of NMDARs could lead to characterization of disease progression and early diagnosis of such diseases.

Development of imaging agents for NMDARs has been extensively reviewed previously (Sobrio et al. 2010; Fuchigami et al. 2015; Gruber and Ametamey 2017; Fu et al. 2019a). General strategies for the development of radioligands for NMDARs have been based on the structural modification of NMDAR antagonists for application in PET or SPECT imaging. Thus, the target sites for the radioligands could be divided into four classes as follows: (1) PCP-binding site (channel blocker binding site), (2) glutamate-binding site, (3) glycine-binding site, and (4) ifenprodil-binding site (GluN2B negative modulator binding site). Despite enormous efforts, there are still no satisfactory radioligands for the visualization of NMDARs in the living brain. In this chapter, we summarize the history and present status of radioligands for these four sites.

### 19.1.1 Radioligands for PCP-Binding Site

Open channel blockers of NMDARs, such as MK-801 and PCP derivatives (Fig. 19.2), have been reported to bind to NMDARs in a receptor activation-dependent manner (Loo et al. 1986; Foster and Wong 1987; Wong et al. 1988). Thus, in vivo imaging of the PCP-binding site of NMDARs would not only provide quantification of NMDAR density in the living brain but also allow receptor activation under normal and pathophysiological conditions to be monitored. In recent decades, a number of reports have emerged regarding the development of radioligands based on dissociative anesthetics (PCP, MK-801, ketamine, memantine derivatives, diarylguanidines) as shown in Fig. 19.2.



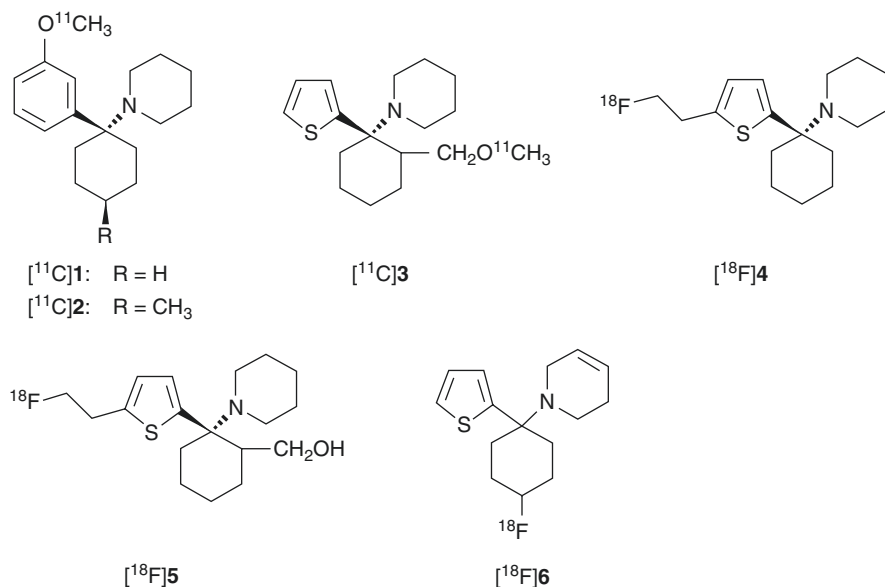
**Fig. 19.2** Chemical structure of open channel blockers of NMDARs

### 19.1.1.1 Dissociative Anesthetic Derivatives

#### PCP and TCP Derivatives

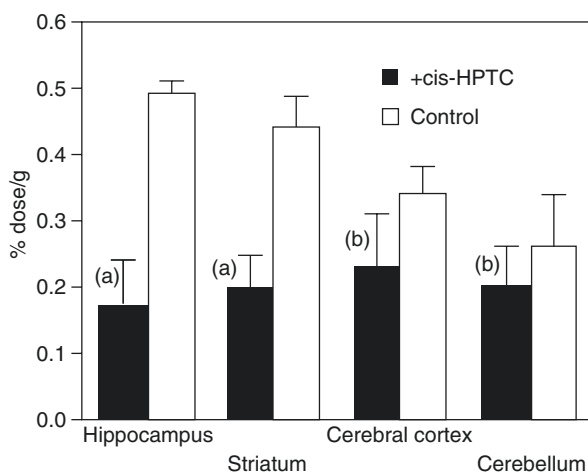
PCP, one of the dissociative anesthetics, has been found to inhibit the actions of NMDARs by blocking the ion channel (Anis et al. 1983). On the other hand, TCP ([2-thienyl]-cyclohexyl-PCP), one of the PCP derivatives, has been reported to show severalfold higher affinity than PCP itself toward the PCP-binding site (Vincent et al. 1979; Tsukiyama et al. 1991).

In structure-activity relationship (SAR) studies of PCP derivatives, the binding affinity of the compounds was expressed as  $K_{0.5}$ , which means the half maximal (50%) inhibitory concentration of [ $^3\text{H}$ ]PCP binding. Methoxy analogs **1** and **2** (Fig. 19.3) showed a higher binding affinity (**1**;  $K_{0.5} = 90$  nM, **2**;  $K_{0.5} = 58$  nM) than PCP itself (PCP;  $K_{0.5} = 250$  nM) (Chaudieu et al. 1989). The  $\text{IC}_{50}$  value of methoxymethyl analog of TCP (**3**, Fig. 19.3) for the [ $^3\text{H}$ ]TCP-binding site was 16 nM, which was slightly weaker than TCP ( $\text{IC}_{50} = 7.9$  nM) but stronger than PCP ( $\text{IC}_{50} = 52$  nM) (Tsukiyama et al. 1991). These compounds were labeled with C-11 and then evaluated as PET radioligands for the PCP-binding site of NMDARs (Haradahira et al. 1998). The [ $^{11}\text{C}$ ]-radiotracers were prepared from hydroxyl precursors reacting with [ $^{11}\text{C}$ ]CH $_3\text{I}$  to reach a specific activity of more than 41 GBq/ $\mu\text{mol}$ . In the biodistribution studies in mice, initial uptake of the TCP analog [ $^{11}\text{C}$ ]**3** was twofold higher (2.6–2.9% ID/g at 1 min) than the PCP analogs [ $^{11}\text{C}$ ]**1** (1.4–1.6% ID/g at 1 min) and [ $^{11}\text{C}$ ]**2** (1.1–1.5% ID/g at 1 min). On the other hand, [ $^{11}\text{C}$ ]**3** showed faster clearance in the brain than [ $^{11}\text{C}$ ]**1** and [ $^{11}\text{C}$ ]**2**. However, there was no significant difference in



**Fig. 19.3** Chemical structure of radiolabeled PCP and TCP derivatives

**Fig. 19.4** Regional brain distribution of radioactivity in rats at 15 min after injection of [ $^{18}\text{F}$ ]4. In the blocking study, *cis*-HPTC (1.7 mmol/kg) was pre-injected 5 min before administration of [ $^{18}\text{F}$ ]4. Student's t-test with respect to the control, (a)  $P < 0.01$ , (b) not significant (Orita et al. 1993)



the accumulated radioactivity level between the NMDAR-rich regions and NMDAR-poor regions. These results indicated that the radioligands had no small binding interaction with brain components other than PCP-binding site. Since no blocking studies using the PCP-binding site antagonists were performed, it is unclear whether these radioligands bound specifically in certain brain regions (Haradahira et al. 1998).

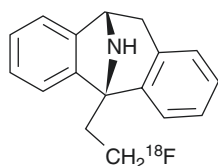
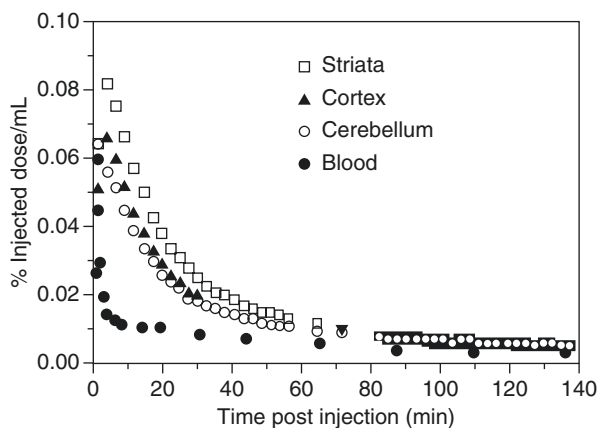
Orita et al. developed [ $^{18}\text{F}$ ]fluoroethyl analog of TCP ([ $^{18}\text{F}$ ]4, Fig. 19.3) with  $\text{IC}_{50}$  values of 61 nM for the PCP-binding site of NMDARs. The [ $^{18}\text{F}$ ]4 showed heterogeneous in vivo distribution that was similar to NMDAR expression. In addition, co-injection of high-affinity ion channel blocker *cis*-HPTC (1.7  $\mu\text{mol/kg}$ ) yielded reduction of regional cerebral distribution of [ $^{18}\text{F}$ ]4 (Fig. 19.4). However, this tracer may be an unsuitable PET radioligand for NMDARs because of the high non-specific binding in brain tissues (Orita et al. 1993).

Another [ $^{18}\text{F}$ ]fluoroethyl analog of TCP ([ $^{18}\text{F}$ ]5, Fig. 19.3) has been synthesized, and biodistribution analysis in rats revealed that this tracer exhibited peak uptake (0.25% ID/g) in the brain at 10 min. However, no regional difference was observed in the brain tissues, and blocking effect of *cis*-HPTC (1.7  $\mu\text{mol/kg}$ ) was not observed for all the tissues investigated. Non-specific brain distribution of [ $^{18}\text{F}$ ]5 could be attributed to the hydrophobic property and low affinity for NMDARs ( $\text{IC}_{50} = 1.5 \mu\text{M}$  vs [ $^3\text{H}$ ]TCP) of these ligands (Shibayama et al. 1996).

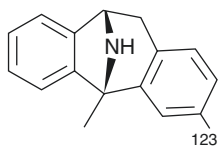
Ouyang et al. have reported a fluorocyclohexyl analog of TCP ([ $^{18}\text{F}$ ]6, Fig. 19.3) with  $\text{IC}_{50}$  value of 97 nM for the PCP-binding site (vs [ $^3\text{H}$ ]MK-801). In PET studies in the rhesus monkey, [ $^{18}\text{F}$ ]6 showed up to 0.08%ID/mL of brain uptake, but the radioactivity was cleared rapidly from all regions of the brain indicative of little or no retention of the radiotracer in the receptor-rich regions (Fig. 19.5) (Ouyang et al. 1996). Because [ $^{18}\text{F}$ ]6 has only moderate affinity and the specificity for NMDARs has not been confirmed, new radioligands with significant improvement of binding affinity and target specificity have to be developed.



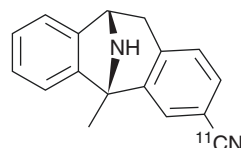
**Fig. 19.5** PET data on the binding of [ $^{18}\text{F}$ ]6 in a rhesus monkey brain. (Ouyang et al. 1996)



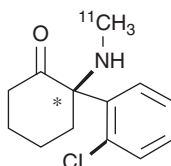
[ $^{18}\text{F}$ ]FMM



[ $^{123}\text{I}$ ]3-iodo-MK-801



[ $^{11}\text{C}$ ]MKC



[ $^{11}\text{C}$ ]Ketamine

**Fig. 19.6** Chemical structure of radiolabeled MK-801 derivatives and [ $^{11}\text{C}$ ]ketamine

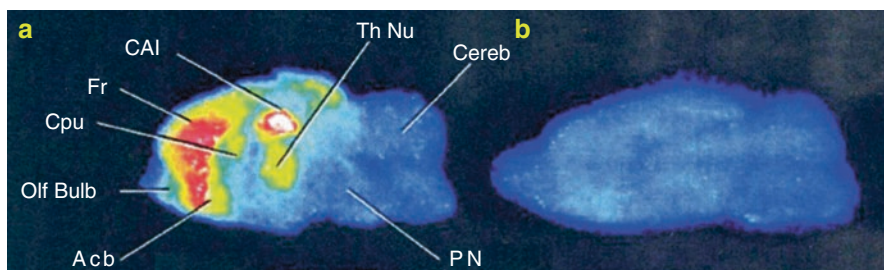
### MK-801 Derivatives

The anticonvulsant MK-801 (Fig. 19.2) has been identified as a potent and selective non-competitive NMDAR antagonist (Wong et al. 1986). Because MK-801 has high binding affinity for the PCP-binding site of NMDARs (Wong et al. 1988) and easy penetrability into the brain (Clineschmidt et al. 1982), MK-801 analogs were expected to be suitable as in vivo imaging probes for NMDARs. Several MK-801 derivatives labeled with  $^{11}\text{C}$ ,  $^{18}\text{F}$ , or  $^{123}\text{I}$  have been synthesized and evaluated as PET or SPECT radioligands for NMDARs.

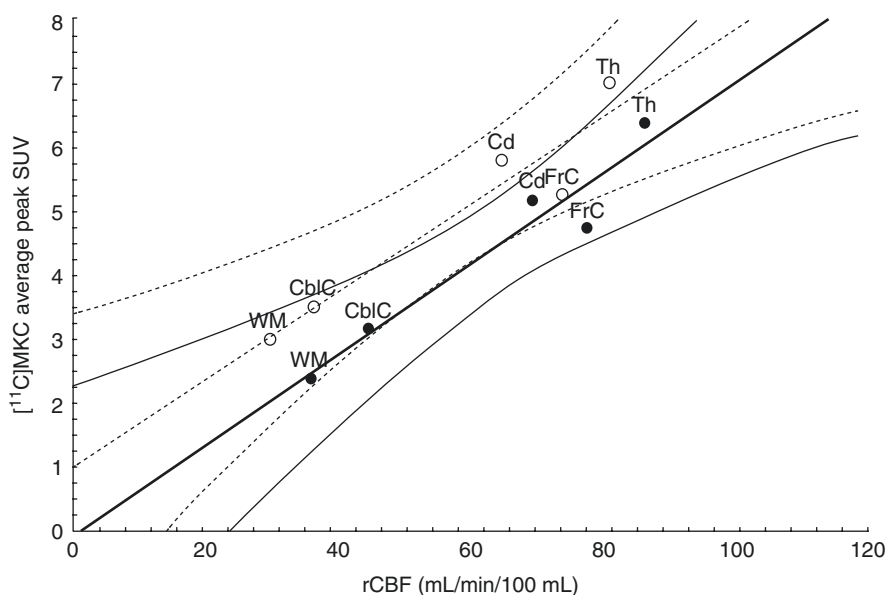
Blin et al. reported the preparation of [ $^{18}\text{F}$ ]fluoromethyl analog of MK-801 ([ $^{18}\text{F}$ ]FMM, Fig. 19.6) and PET studies on baboons (Blin et al. 1991). In the baseline PET studies, initial accumulation into the brain of [ $^{18}\text{F}$ ]FMM was marked with peak uptake values as high as 0.03% ID/mL. However, the ratios of [ $^{18}\text{F}$ ]FMM in the

cerebral cortex and striatum (NMDAR-rich regions) to the cerebellum (NMDAR-poor regions) were only minimal (maximal ratios ranging from 1.1 to 1.6 during the first 20–40 min). In addition, there was no clear reduction in the brain uptake and target to non-target ratios by pretreatment of the open channel blockers (PCP or MK-801). Because binding of [ $^3\text{H}$ ]MK-801 to the PCP-binding site has been enhanced severalfold in the presence of high amount of glutamate (Wong et al. 1988), the authors investigated further using PET studies of [ $^{18}\text{F}$ ]FMM on ischemic baboons with excessive glutamate activation. However, the brain kinetics of [ $^{18}\text{F}$ ]FMM was not significantly different between the ischemic and control groups. It can be concluded that [ $^{18}\text{F}$ ]FMM is an unsuitable radioligand for the PCP-binding site of NMDARs (Blin et al. 1991). Ransom et al. developed the 3- $^{123}\text{I}$ iodo analog of MK-801 ( $^{123}\text{I}$ 3-iodo-MK-801, Fig. 19.6) as a SPECT imaging agent. Over 95% of [ $^{123}\text{I}$ ]3-iodo-MK-801 binding was specific in rat brain homogenates. MK-801 derivatives were shown to bind with high affinity for the [ $^{123}\text{I}$ ]3-iodo-MK-801 binding site (3-iodo-MK-801,  $\text{IC}_{50} = 3.4 \text{ nM}$ ; (+)-MK-801,  $\text{IC}_{50} = 5.6 \text{ nM}$ ) indicating excellent binding affinity of this tracer for the PCP-binding site (Ransom et al. 1990). These positive in vitro results prompted further in vivo imaging studies. Owens et al. reported the clinical SPECT investigation of [ $^{123}\text{I}$ ]3-iodo-MK-801. In normal subjects, [ $^{123}\text{I}$ ]3-iodo-MK-801 showed rapid uptake into the brain, but the tracer has a high non-specific retention in the brain tissues probably due to its high lipophilicity (Owens et al. 1997). Brown et al. reported the clinical SPECT studies of [ $^{123}\text{I}$ ]3-iodo-MK-801 in Alzheimer's diseases (AD) and healthy subjects. There were slight differences in the brain uptake/rCBF (regional cerebral blood flow) ratio and washout of [ $^{123}\text{I}$ ]3-iodo-MK-801 between the AD patients and control groups, which suggested that the binding of this tracer in the brain was increased in AD patients. However, the authors concluded that [ $^{123}\text{I}$ ]3-iodo-MK-801 did not provide accurate measurement of NMDARs since the alteration was not significant, and the brain distribution of this tracer was homogeneous in both AD and control groups (Brown et al. 1997).

Andersson et al. successfully synthesized 3- $^{11}\text{C}$ cyano analog of MK-801 ( $^{11}\text{C}$ ]MKC, Fig. 19.6) with high specific activity (220–600 GBq/ $\mu\text{mol}$ ) from 3-iodo precursor using palladium(0)-promoted [ $^{11}\text{C}$ ]cyanation (Andersson et al. 1998). Saturation assay in rat cortical membranes revealed that [ $^{11}\text{C}$ ]MKC had a single high-affinity binding site with a  $K_d$  of 8.2 nM and a  $B_{\text{max}}$  of 1.6 pmol/mg protein (Sihver et al. 1998). In vitro autoradiography demonstrated that the highest specific binding in rat slices of [ $^{11}\text{C}$ ]MKC was observed in the hippocampus and frontal cortex. Moderate binding was found in the caudate putamen and thalamus. Low-density binding sites were observed in the cerebellum, pons, midbrain, and brainstem (Fig. 19.7). This correlated with the distribution of [ $^3\text{H}$ ]MK-801 (Bowery et al. 1988). The in vivo PET studies demonstrated that [ $^{11}\text{C}$ ]MKC showed a rapid and high uptake in the brain tissues. In addition, the radioactivity level was higher in the frontal cortex than cerebellar cortex, but the brain distribution pattern was well correlated with the rCBF (Fig. 19.8). Neither MK801 (0.1 mg/kg) nor ketamine (2.5 mg/kg) as pretreatment dose caused a significant change in regional brain



**Fig. 19.7** In vitro autoradiography of [ $^{11}\text{C}$ ]MKC in the rat brain slices. Total binding (A) and non-specific binding (B) are shown in sagittal planes. Non-specific binding was measured in the presence of MK-801 (1 mM). Abbreviations: *Acb* accumbens nucleus, *CA1* CA1 of hippocampus, *Cereb* cerebellum, *Cpu* caudate putamen, *Fr* frontal cortex, *Olf bulb* olfactory bulb, *PN* pontine nuclei, *Th Nu* thalamic nuclei. Sihver et al. (1998)



**Fig. 19.8** Typical graph presenting the relation between peak [ $^{11}\text{C}$ ]MKC uptake and rCBF in the same regions in the baseline measurement (open circles and dotted lines) and ketamine-pretreatment measurement (closed circles and continuous lines). Curved lines indicate 95% confidence. Abbreviations: *FrC* frontal cortex, *Cd* right caudate nucleus, *Th* thalamic nuclei, *CblC* cerebellar cortex, *TM* white matter. Sihver et al. (1998)

kinetics. These results could be caused by the very low specific binding fraction of [ $^{11}\text{C}$ ]MKC under in vivo conditions (Sihver et al. 1998). Development of new radioligands with lower lipophilicity and higher NMDAR selectivity should be tried to reduce non-specific binding.

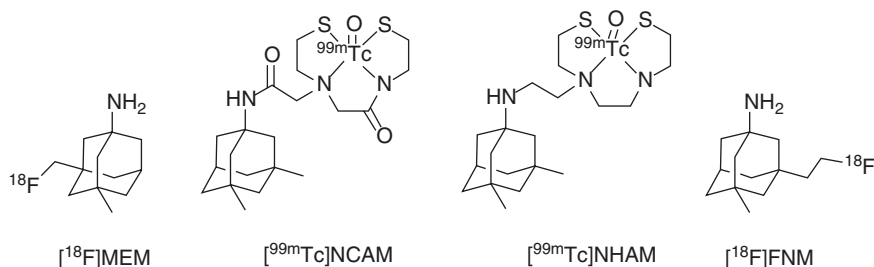
### [<sup>11</sup>C]Ketamine

Ketamine (Fig. 19.2) is known as a PCP-like anesthetic drug which has a weak binding affinity for the PCP-binding site of NMDARs (Lodge and Johnson 1990). In the displacement assay using pig brain homogenates, the  $K_i$  values of (*S*)-ketamine and (*R*)-ketamine for [<sup>3</sup>H]MK-801 binding site of NMDARs were 1.2 and 5.0  $\mu\text{M}$ , respectively (Oye et al. 1992). [<sup>11</sup>C]-labeled (*R,S*)-ketamine ([<sup>11</sup>C]ketamine, Fig. 19.6) and its enantiomers were synthesized, and then in vivo evaluation was performed (Hartvig et al. 1994; Shiue et al. 1997). Although [<sup>11</sup>C]ketamine showed high blood-brain barrier (BBB) permeability (2.80% ID/g at 5 min after injection) in mice brain, unchanged [<sup>11</sup>C]ketamine was only 50% at 5 min. PET studies in a baboon demonstrated that the influx of [<sup>11</sup>C]ketamine and its enantiomers into the brain were high for the first few minutes, but radioactivity then declined rapidly and a similar uptake in the striatum and cerebellum was observed. In addition, these three tracers were also metabolized rapidly in the baboon plasma. Because [<sup>11</sup>C]ketamine and its enantiomers were revealed to be metabolically unstable and had inconsistent binding with NMDAR expression, these tracers failed to act as useful radioligands for in vivo imaging of NMDARs (Shiue et al. 1997).

### Memantine Derivatives

Memantine (Fig. 19.2) is a drug used in the treatment of various neurological and psychiatric disorders, including Parkinson's disease and Alzheimer's disease. Memantine has been identified as a non-competitive open channel blocker of NMDARs with good BBB permeability (Parsons et al. 1993; Lipton 2004). Therefore, several radiolabeled memantine derivatives have been developed as imaging agents for NMDARs.

In 1998, Samnick et al. reported the development of 3-[<sup>18</sup>F]fluoromethyl analog of memantine ([<sup>18</sup>F]MEM, Fig. 19.9) for PET imaging of ion channel site. The MEM showed a slightly weaker inhibitory activity to the current response to NMDARs ( $\text{IC}_{50} = 6.0 \mu\text{M}$ ) than memantine ( $\text{IC}_{50} = 2.9 \mu\text{M}$ ) and PCP ( $\text{IC}_{50} = 1.0 \mu\text{M}$ ). In the biodistribution studies in mice, [<sup>18</sup>F]MEM showed an excellent brain uptake (3.6% ID/g at 60 min), and the accumulation in the frontal cortex and hippocampus was higher than in the cerebellum. In addition, co-injection of MK-801 (0.10 mg/kg) led to 25–36% reduction of radioactivity in NMDAR-rich regions. PET studies



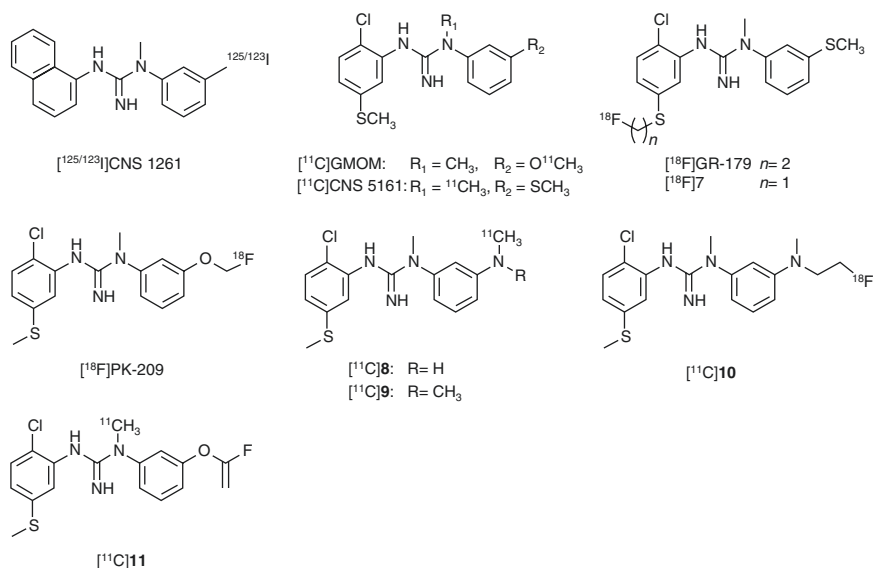
**Fig. 19.9** Chemical structure of radiolabeled memantine derivatives

in a rhesus monkey revealed that [ $^{18}\text{F}$ ]MEM displayed high brain uptake and low clearance of the radiotracer from the brain tissues. Pretreatment with memantine (0.50 and 1 mg/kg) yielded a reduction of uptake of up to 32% from 60 min in the examined brain regions. However, the highest uptake was observed in the cerebellum, which is an NMDAR-poor region. In addition, pretreatment with haloperidol, which is a ligand with high affinity for dopamine D2 and sigma receptors, also caused similar reduction of radioactivity in the brain (Sammick et al. 1998). In clinical PET studies, [ $^{18}\text{F}$ ]MEM showed perfusion-dependent uptake in the brain rather than NMDAR expression (Ametamey et al. 2002). Because of inconsistent binding with NMDAR expression and low specificity for NMDARs, it would appear that [ $^{18}\text{F}$ ]MEM is an unsuitable radioligand (Sammick et al. 1998; Ametamey et al. 2002). However, the different brain distribution pattern of [ $^{18}\text{F}$ ]MEM observed among species was noteworthy. New technetium-labeled memantine derivatives such as [ $^{99\text{m}}\text{Tc}$ ]NCAM and, in 2012, [ $^{99\text{m}}\text{Tc}$ ]NHAM have been reported as novel SPECT imaging probes (Fig. 19.9). The [ $^{99\text{m}}\text{Tc}$ ]NCAM had a single binding site with a  $K_d$  of 701 nM. Furthermore, the *in vitro* binding of [ $^{99\text{m}}\text{Tc}$ ]NCAM was blocked by ketamine and MK-801 in a competitive manner. In the biodistribution studies in mice, [ $^{99\text{m}}\text{Tc}$ ]NCAM showed 1.2% ID/g brain uptake at 5 min, and the target (frontal cortex, hippocampus) to non-target (cerebellum) ratio of radioactivity reached 2.5–3.0 at 60 min. The *in vivo* blocking studies were not so far reported in the literature (Zhou et al. 2012).

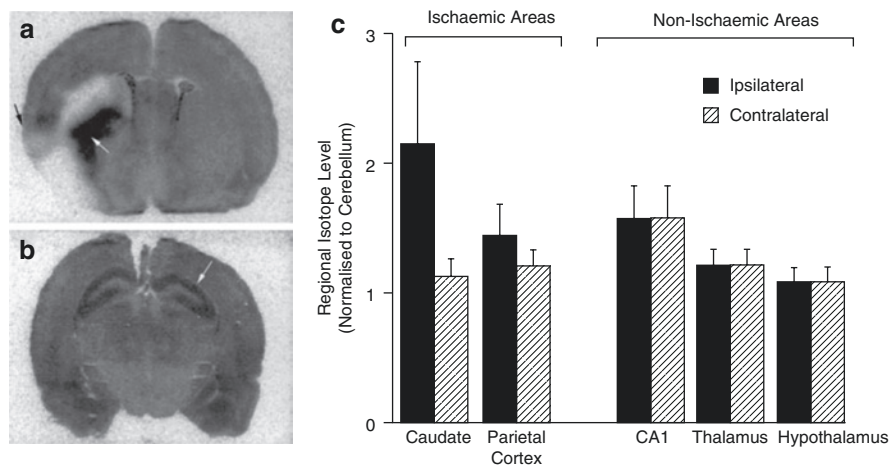
[ $^{18}\text{F}$ ]Fluoroethylnormemantine ([ $^{18}\text{F}$ ]FNM) has been developed as a moderately lipophilic PET tracer ( $\text{Log}D_{7.4} = 1.93$ ). Although [ $^{18}\text{F}$ ]FNM showed a high BBB permeability and good *in vivo* metabolic stability, this tracer exhibited low affinity to NMDAR ( $\text{IC}_{50} = 5 \mu\text{M}$  vs [ $^3\text{H}$ ]TCP binding to rat brain slices), homogeneous *in vivo* brain distribution, and unclear specificity to NMDAR (Salabert et al. 2015, 2018).

### 19.1.1.2 Diarylguanidine Derivatives

Initially reported imaging probes, such as PCP and MK-801 analogs, suffered from high non-specific *in vivo* accumulation, which was probably due to the high lipophilicity and lack of selectivity for NMDARs. Diarylguanidine derivatives were discovered as NMDAR open channel blockers with high potency and selectivity. These compounds showed therapeutic effect in various *v v* of brain disease (Reddy et al. 1994; Hu et al. 1997). Because most diarylguanidines are more hydrophilic than PCP and MK-801, several radiolabeled diarylguanidine analogs were developed for PET or SPECT imaging of NMDARs. In the first study, [ $^{125}\text{I}$ ]CNS 1261 (Fig. 19.10) was developed as a high-affinity imaging agent for NMDARs with a  $K_i$  value of 4.2 nM for the [ $^3\text{H}$ ]MK-801 binding site with moderate lipophilicity ( $\text{Log} D = 2.19$ ). Binding assays to other neurotransmitter receptors demonstrated that CNS 1261 had a weaker binding affinity for the other 41 neurotransmitter receptors. *Ex vivo* autoradiography studies with normal rats showed that the [ $^{125}\text{I}$ ]CNS 1261 uptake in the hippocampus was 2.4–2.9-fold higher than that in the cerebellum, in which the accumulation matched the distribution of NMDARs. In the ischemic rat brain, [ $^{125}\text{I}$ ]CNS 1261 uptake was considerably increased in the neocortex and striatum,

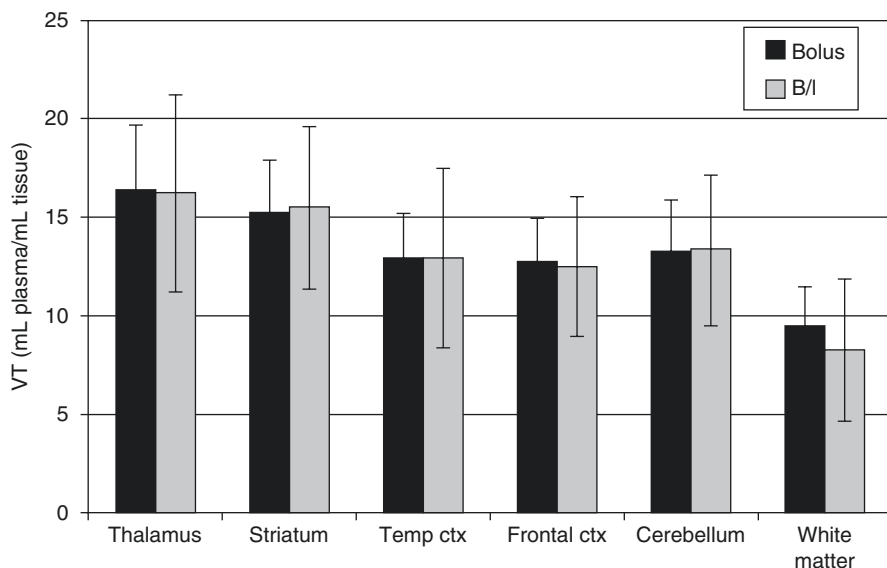


**Fig. 19.10** Chemical structure of radiolabeled diarylguanidine derivatives



**Fig. 19.11** Ex vivo autoradiography of [<sup>125</sup>I]CNS 1261 in coronal sections of rat brain at the level of (a) the caudate nucleus (white arrow) and circumscribed zones of the cerebral cortex (b) and CA1 region of the hippocampus (white arrow). The animals were injected with [<sup>125</sup>I]CNS 1261 15 min after permanent occlusion of the middle cerebral artery (left hemisphere) and sacrificed 120 min after tracer injection. (c) Accumulated radioactivity levels in ROIs to the cerebellum. Owens et al. (2000)

with a twofold higher uptake in the ischemic hemisphere of the caudate nucleus uptake than the same regions on the contralateral non-ischemic hemisphere (Fig. 19.11). This suggested that [<sup>125</sup>I]CNS 1261 could recognize the areas of



**Fig. 19.12** The mean  $V_T$  values ( $\pm$ SD) of [ $^{125}$ I]CNS 1261 for ROIs from healthy volunteers scanned with the bolus protocol ( $n = 7$ ) (dark bars) and with the bolus/infusion protocol ( $n = 6$ ) (light bars). Bressan et al. (2004)

NMDAR activation (Owens et al. 2000). Based on these results, several clinical SPECT studies of [ $^{123}$ I]CNS 1261 were conducted. SPECT imaging of [ $^{123}$ I]CNS 1261 in healthy volunteers was conducted by using both bolus and bolus plus constant infusion paradigms. Total volume of distribution ( $V_T$ ) values were approximately the same in the NMDAR-rich regions (striatum, hippocampus, frontal cortex) and NMDAR-poor region (cerebellum). The  $V_T$  value in the thalamus, which is a region with considerably high expression of NMDARs, was slightly higher than in other regions (Fig. 19.12) (Erlandsson et al. 2003; Bressan et al. 2004). Bolus plus infusion of ketamine in healthy volunteers caused a global reduction in  $V_T$  of [ $^{123}$ I]CNS 1261 compared with placebo in most brain regions, including the putamen, temporal, and pericallosal regions (Stone et al. 2006). Subsequent work by the same group demonstrated that the degree of reduction in the  $V_T$  values of [ $^{123}$ I]CNS 1261 correlated with the negative symptoms of schizophrenia in all brain regions examined (Stone et al. 2008). In addition, they reported that drug-free patients with schizophrenia had reduced binding of [ $^{123}$ I]CNS 1261 in the left hippocampus relative to the whole cortex compared with healthy controls (Pilowsky et al. 2006). In contrast, the same group found that the  $V_T$  values of [ $^{123}$ I]CNS 1261 did not differ between healthy normal volunteers and drug-free or typical antipsychotic-treated schizophrenia patients. On the other hand, clozapine-treated patients showed a significant reduction of  $V_T$  in all brain regions examined, which indicated a decrease of NMDAR activation in this group. However, the regional brain distribution of [ $^{123}$ I]CNS 1261 was almost homogeneous in all groups (Bressan et al. 2005). Although there was no conclusive evidence that [ $^{123}$ I]CNS 1261 could recognize

NMDAR activation, these reports suggested that protein expression levels of NMDARs could be reduced in the brain of schizophrenia patients. Numerous reports indicated the involvement of NMDAR hypofunction in the pathophysiology of positive and negative symptoms of schizophrenia (Kantrowitz and Javitt 2010; Lin et al. 2012). Recently, Knol et al. reported further *in vivo* studies of [ $^{123}$ I]CNS 1261 using rats. This tracer showed higher accumulation in the NMDAR-rich regions such as the hippocampus and frontal cortex than that of NMDAR-poor cerebellum, in which the ratio was maximum (about 1.4) at 2 h post-injection. Pretreatment with the NMDAR co-agonist, d-serine (10 mg/kg), led to no significant change in the hippocampus/cerebellar ratio. On the other hand, treatment with d-serine and MK-801 resulted in a significantly lower hippocampus/cerebellar ratio, which indicated the existence of specific trapping of a certain amount of [ $^{123}$ I]CNS 1261 in the NMDAR-rich regions. However, the authors concluded that the use of [ $^{123}$ I]CNS 1261 would be limited in terms of quantification and detection of small changes in receptor availability because of a considerable amount of non-specific binding of this tracer (Knol et al. 2009).

To provide potential PET radioligands for the PCP-binding site, [ $^{11}$ C]-labeled diarylguanidine derivatives have been synthesized and evaluated. [ $^{11}$ C]GMOM (Fig. 19.10) has been found to exhibit a strong binding affinity for NMDARs ( $K_i = 5.2$  nM). In rats, brain uptake of [ $^{11}$ C]GMOM was high and peaked during the early part of the study. Regional brain distribution at 10 min was 1.23, 0.93, and 0.74% ID/g for the frontal cortex, hippocampus, and cerebellum, respectively, as regards the appearance of contrast in the NMDAR-rich and NMDAR-poor regions. Pretreatment with the open channel blockers GMOM or MK-801 (1 mg/kg) or the NB2B-selective NMDAR modulator Ro 25-6981 led to a moderate and uniform decrease of [ $^{11}$ C]GMOM uptake in brain regions. On the other hand, the NMDAR co-agonist d-serine increased [ $^{11}$ C]GMOM binding in all regions. PET imaging in baboons revealed that [ $^{11}$ C]GMOM exhibited high BBB penetration. However, the brain distribution of [ $^{11}$ C]GMOM was almost homogeneous, and pre-administration of MK801 did not reduce regional  $VT$ 's or regional  $V_T$  ratios. The authors hypothesized that these inconsistent results could be caused by the blocking effect of anesthesia (ketamine and isoflurane) to NMDAR activity. Further PET investigations of [ $^{11}$ C]GMOM in other conditions, such as in a conscious state, have not been reported so far (Waterhouse et al. 2004). In clinical studies, [ $^{11}$ C]GMOM showed high brain uptake and relatively consistent localization with NMDAR expression. Treatment with ketamine led to a significant reduction in the net influx rate constant ( $K_i$ ) in the gray matter. As mentioned above, [ $^{11}$ C]GMOM has been rapidly metabolized, and the brain kinetics of the metabolites should be taken into consideration. Further investigation would be required in a larger number of subjects with test-retest studies (van der Doef et al. 2016). The limitation of such clinical studies is that the available blockers are restricted because of the high toxicity of ion channel blockers with the NMDARs.

Another diarylguanidine derivative, CNS 5161 (Fig. 19.10), has been shown to exhibit high affinity for the PCP-binding site ( $K_i = 1.9$  nM) (Hu et al. 1997; Dumont et al. 2002). Clinical trials using CNS 5161 have been evaluated in the treatment of



neuropathic pain (Walters et al. 2002; Forst et al. 2007). [ $^3\text{H}$ ]CNS 5161 showed a heterogeneous in vivo brain distribution in rats and a cortex/cerebellum ratio of 1.4. Pretreatment with NMDA increased the ratio in the hippocampus/cerebellum to 1.6–1.9, while MK801 diminished this increase that resulted in ratios close to 1. [ $^3\text{H}$ ]CNS 5161 appeared to be unstable in arterial blood, as 65% of the parent compound was metabolized to several more hydrophilic compounds even 20 min after tracer injection (Biegon et al. 2007). The [ $^{11}\text{C}$ ]CNS 5161 was synthesized by [ $^{11}\text{C}$ ]methylation of the desmethyl guanidine precursor with low specific activity (41 GBq/mmol) (Zhao et al. 2006). In human PET studies, [ $^{11}\text{C}$ ]CNS 5161 exhibited the largest uptake in the putamen and thalamus and the lowest uptake in the cerebellum, but relatively lower radioactivity was observed in the hippocampus compared with other gray matter regions. Similar to rats, the metabolism of [ $^{11}\text{C}$ ]CNS 5161 was rapid in human plasma. No blocking studies were reported in the literature (Asselin et al. 2004). Recently, clinical PET studies of [ $^{11}\text{C}$ ]CNS 5161 were carried out to assess the effect of levodopa on striatal and cortical NMDAR activity in patients with Parkinson's disease with and without dyskinesias (Ahmed et al. 2011). The levodopa-treated dyskinetic patients had higher [ $^{11}\text{C}$ ]CNS 5161 uptake in the caudate, putamen, and precentral gyrus compared with patients without dyskinesias. These results are consistent with reports that upregulation of NMDARs has been associated with the development of levodopa-induced dyskinesias in animal models of Parkinsonism (Oh et al. 1998). However, no significant increase in [ $^{11}\text{C}$ ]CNS 5161 uptake was observed in patients with Parkinson's disease when compared with control subjects. In addition, relatively high uptake of [ $^{11}\text{C}$ ]CNS 5161 in the cerebellum was observed, which was inconsistent with the brain distribution of [ $^{11}\text{C}$ ]CNS 5161 in a previous clinical PET study reported by Asselin et al. (Asselin et al. 2004). Further investigations are necessary in order to prove the usefulness of these diarylguanidines as radioligands for measuring NMDAR function by PET or SPECT.

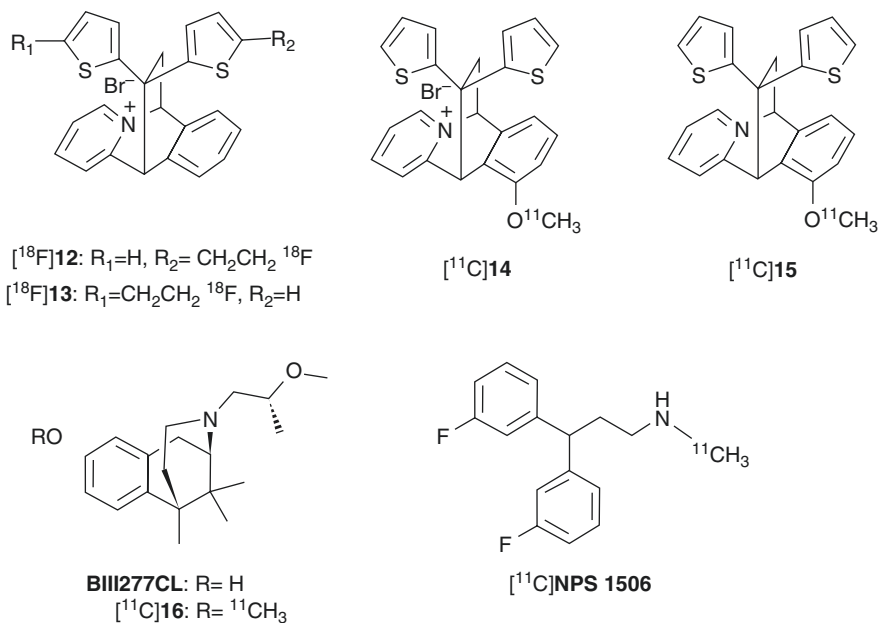
Recently,  $^{18}\text{F}$ -labeled *S*-fluoroalkyl diarylguanidines were synthesized and used for in vitro evaluation. Among them, [ $^{18}\text{F}$ ]GE-179 and [ $^{18}\text{F}$ ]7 (Fig. 19.10) showed high affinity for the PCP-binding site ( $K_d = 2.4$  nM for [ $^{18}\text{F}$ ]GE-179 and  $K_d = 1.4$  nM for [ $^{18}\text{F}$ ]7). Further studies could not be performed on [ $^{18}\text{F}$ ]7 owing to low stability of this ligand. A broad receptor binding assay for 60 receptors demonstrated that [ $^{18}\text{F}$ ]GE-179 had high selectivity for the PCP-binding site of NMDARs (Robins et al. 2010). Although this tracer showed high brain uptake, the  $V_T$  of each region was correlated to cerebral blood flow rather than the levels of NMDAR expression. However, the distribution of the radioactivity of [ $^{18}\text{F}$ ]GE-179 might be explained by subunit composition as the channel open probability of GluN2A is reported to be higher than that of the other subunits. No blocking studies using unlabeled GE-179 or other channel blockers were feasible in this clinical study (McGinnity et al. 2014). In studies by other groups, PET/CT imaging of [ $^{18}\text{F}$ ]GE-179 along with drug competition in rats demonstrated that none of the ion channel blockers (GE-179, MK801, PCP, and ketamine), allosteric inhibitor (ifenprodil), and NMDAR activator (methamphetamine) significantly altered the PET signals in the brain. PET/MR in rhesus macaques exhibited that pretreatment with GE-179 led to no meaningful

reduction in the PET signal. These results indicated that a large part of the [ $^{18}\text{F}$ ]GE-179 signals seems to be non-specific (Schoenberger et al. 2018; Sander et al. 2019). However, it cannot be excluded that anesthesia reduced  $B_{\text{max}}$  of [ $^{18}\text{F}$ ]GE-179 in the brain, which abolished the accessibility of this tracer to the target site (McGinnity et al. 2019). Golla et al. reported [ $^{18}\text{F}$ ]PK-209 (Fig. 19.10) as a high-affinity PET tracer for the ion channel site ( $K_i = 18$  nM vs [ $^3\text{H}$ ]MK-801). In line with the moderate lipophilicity ( $\log D_{7.4} = 1.45$ ), this tracer entered the brain easily. However, over 70% of the parent compound was metabolized, and the distribution was inconsistent with the expression patterns of NMDA receptors (Golla et al. 2015). Klein et al. developed three new meta-substituted GMOM analogs, [ $^{11}\text{C}$ ]8, [ $^{11}\text{C}$ ]9, and [ $^{11}\text{C}$ ]10, with favorable  $K_i$  values for the ion channel of the NMDAR (19.1, 1.35, and 4.81 nM, respectively). Although ex vivo studies of these tracers in awake rats demonstrated good brain uptake, pretreatment of animals with MK-801 increased the brain uptake. Over 38% of the radioactivity in the brain tissue represents metabolites, which may obscure specific binding of these tracers (Klein et al. 2016). In order to overcome the low metabolic stability of guanidine derivatives, Klein et al. developed the fluorovinylxy analog [ $^{11}\text{C}$ ]11 with a high affinity for the ion channel site of NMDAR. However, pretreatment with MK-801 showed no significant inhibitory effect on [ $^{11}\text{C}$ ]11, and only a small improvement in the metabolic stability was achieved (Klein et al. 2017).

### 19.1.1.3 Other Open Channel Blocker Derivatives

The 6,11-ethanobenzo[b]quinolizinium derivatives are highly hydrophilic compounds reported to exhibit high affinity and specificity to the PCP-binding site of the NMDARs (Mallamo et al. 1994). Based on the SAR studies, new fluorine-18 radioligands ([ $^{18}\text{F}$ ]12 and [ $^{18}\text{F}$ ]13) and a carbon-11 ligand ([ $^{11}\text{C}$ ]14) have been developed as novel PET radioligands for the PCP-binding site of NMDARs as shown in Fig. 19.13 (Sasaki et al. 1998; Ishibashi et al. 2000). These compounds showed moderate binding affinity for the PCP-binding site (12,  $\text{IC}_{50} = 47$  nM; 13,  $\text{IC}_{50} = 89$  nM; 14,  $\text{IC}_{50} = 19$  nM vs [ $^3\text{H}$ ]TCP) (Sasaki et al. 1998). In vivo biodistribution studies demonstrated that these three radioligands showed a quite low BBB permeability (0.02–0.05% ID/g at 15 min) presumably because of their cationic property and extremely low lipophilicity (Ishibashi et al. 2000). To improve brain uptake and receptor specificity, the reduced derivative [ $^{11}\text{C}$ ]15 was developed as a prodrug for [ $^{11}\text{C}$ ]14. In the preliminary in vivo studies, cationic [ $^{14}\text{C}$ ]15 was easily metabolized to neutral [ $^{14}\text{C}$ ]14 at 15 min after injection. However, most of [ $^{14}\text{C}$ ]14 was cleared from the brain at 45 min. Although [ $^{11}\text{C}$ ]15 showed a higher BBB penetration (0.7–0.8% ID/g at 15 min) than that of [ $^{14}\text{C}$ ]14, no regional difference was observed between the forebrain regions and cerebellum (Sasaki et al. 2001). Although [ $^{11}\text{C}$ ]15 was not confirmed to be a useful PET ligand for NMDARs, this prodrug strategy can be applied to the development of in vivo imaging agents.

BIII277CL (Fig. 19.13), a potent anticonvulsant and neuroprotective agent, has been reported to have high affinity ( $K_i = 4.5$  nM vs [ $^3\text{H}$ ]MK-801) at the PCP-binding site of NMDARs (Grauert et al. 1997, 1998). Kocic et al. have developed a methyl analog of BIII277CL (16, Fig. 19.13) with substantial affinity for the PCP-binding



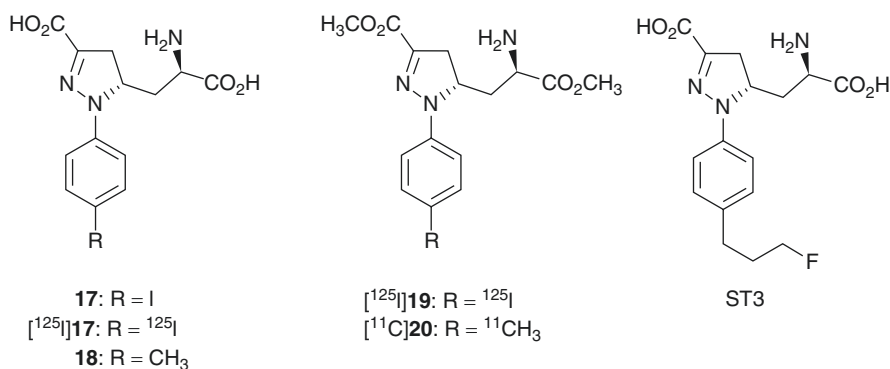
**Fig. 19.13** Chemical structure of other radiolabeled open channel blocker derivatives

site ( $K_i = 49$  nM vs  $[^3\text{H}]\text{TCP}$ ) as a candidate imaging agent for NMDARs. PET studies of  $[^{11}\text{C}]\mathbf{16}$  in pigs revealed homogeneous localization in the cortex and cerebellum. The kinetic analysis using one-tissue compartment model revealed that  $V_T$  of  $[^{11}\text{C}]\mathbf{16}$  was approximately 1 in both the cortex and cerebellum. These results indicated a lack of specific binding of  $[^{11}\text{C}]\mathbf{16}$  for NMDARs (Kokic et al. 2002).

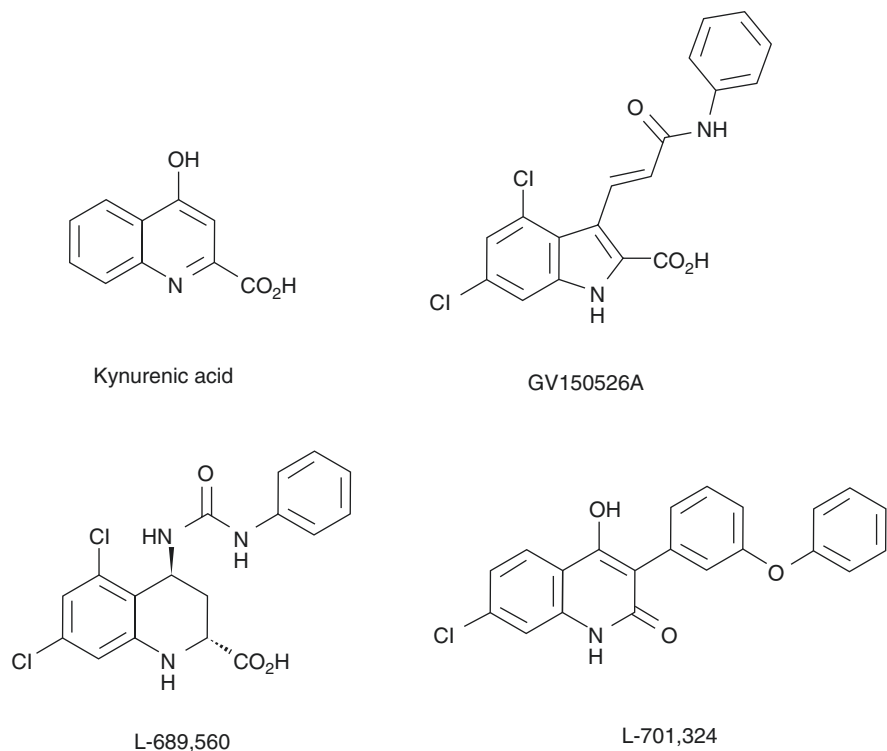
NPS 1506 (Fig. 19.13) has been reported as a non-competitive NMDAR antagonist with moderate affinity ( $\text{IC}_{50} = 664$  nM vs  $[^3\text{H}]\text{MK-801}$ ). NPS 1506 is neuroprotective in a variety of animal models of stroke and head injury with no PCP-like psychotomimetic effects at doses between 1 and 5 mg/kg i.p. in rats (Mueller et al. 1999). Considering such a pharmacological profile, which is different from the previous NMDAR antagonists, NPS 1506 was labeled with carbon-11 and evaluated as a novel PET tracer for the PCP-binding site of NMDARs. Biodistribution of  $[^{11}\text{C}]\text{NPS } \mathbf{1506}$  in mice and rat demonstrated that uptake into the brain was rapid and occurred at high levels. However, the regional brain distribution was fairly uniform and did not clearly reflect the known localization of NMDARs in the rodent brain. In addition, treatment with unlabeled NPS 1506 did not cause any change in uptake in the mouse brain compared with the control group. An activator of NMDARs, 3-nitropropionic acid (20 mg/kg), did not produce any change in the regional uptake in the hippocampus nor the striatum. Therefore,  $[^{11}\text{C}]\text{NPS } \mathbf{1506}$  may be an unsuitable in vivo tracer for NMDARs because of its large non-specific binding fraction and low in vitro binding affinity for NMDARs (Fuchigami et al. 2003).

### 19.1.2 Radioligands for Glutamate-Binding Site

Most competitive antagonists for the glutamate-binding site have an acid moiety that may cause low brain uptake and have demonstrated low selectivity for NMDAR against the other glutamate receptors. Therefore, few radioligands for this site have been published in the literature. Considerable effort has been devoted to developing GluN1/2A-selective NMDA receptor ligands targeting the glutamate-binding site for treatment and imaging of GluN1/2A-related CNS disorders, including schizophrenia, mood disorder, PD, and AD (He et al. 2019). Tamborini et al. reported a radiotracer targeting the competitive glutamate site with the aim of imaging GluN1/2A-containing NMDAR. They discovered compounds **17** and **18** (Fig. 19.14) with  $K_i$  values of 250 and 48.3 nM for GluN1/2A/NMDAR, respectively. Although ex vivo autoradiography of [ $^{125}$ I]**19** (Fig. 19.14), a prodrug of [ $^{125}$ I]**17**, in mice was consistent with the distribution of GluN1/2A subtype receptor mRNA expression, the BBB permeability of this tracer may not be enough to visualize the target. Compound **18** with more GluN1/2A selectivity has been derivatized to prodrug [ $^{11}$ C]**20** (Fig. 19.14); however, ex vivo studies showed still poor brain uptake of [ $^{11}$ C]**20**, and its localization suggested binding to both GluN1/2B and GluN1/2C (Tamborini et al. 2016). Recently, Lind et al. developed ST3 {(*S*)-5-[(*R*)-2-amino-2-carboxyethyl]-1-[4-(3-fluoropropyl)phenyl]-4,5-dihydro-1H-pyrazole-3-carboxylic acid}, which has high affinity for GluN1/2A ( $K_i = 52$  nM) (Fig. 19.14) with a 15-fold selectivity over GluN1/2B (Lind et al. 2017). It is expected that continuous efforts lead to the development of prospective selective radioligands for the glutamate site of GluN1/2A-containing NMDAR.



**Fig. 19.14** Chemical structure of antagonists and radioligands for glutamate-binding site



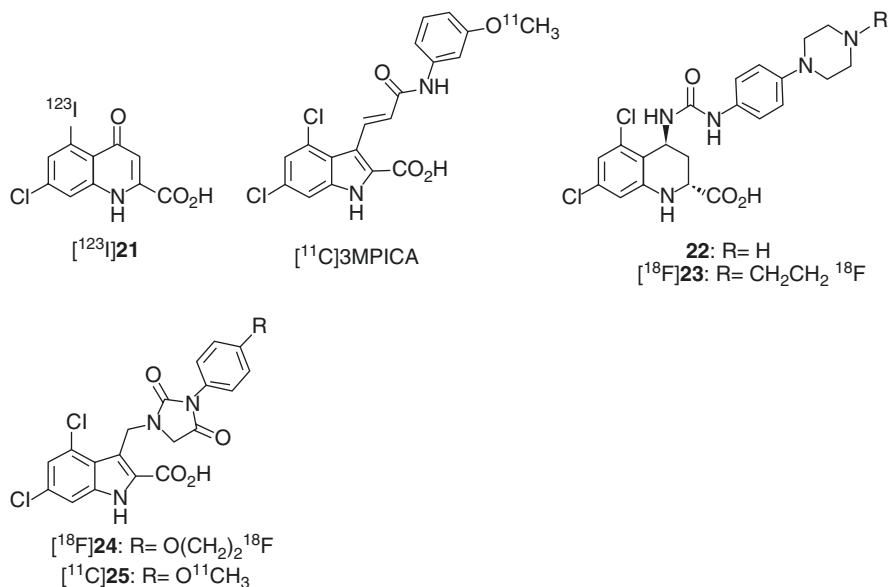
**Fig. 19.15** Chemical structure of glycine-binding site antagonists

### 19.1.3 Radioligands for Glycine-Binding Site

Because glycine and d-serine are essential co-agonists for NMDA ion channel activation, various classes of glycine-binding site antagonists (Fig. 19.15) have been reported as anticonvulsants and neuroprotective agents (Danysz and Parsons 1998). Hypofunction of NMDARs is considered to be associated with the pathophysiology of schizophrenia. Indeed, it was reported that the enhancement of NMDAR activity via the glycine site was efficacious in the treatment of schizophrenia (Millan 2005). Imaging of the glycine site of NMDARs was considered useful for obtaining information about the functional mechanism of the glycine site in the living brain and in the diagnosis of various neurological disorders.

#### 19.1.3.1 Radioligands Based on Cyclic Amino Acids

7-Chloro-5-iodokynurenic acid (**21**, Fig. 19.16) has been identified as a potent ( $IC_{50}$  vs [ $^3H$ ]glycine binding, 32 nM) and selective antagonist for the glycine site of NMDARs (Leeson et al. 1991). [ $^{123}I$ ]**21** was successfully synthesized by halogen exchange reaction of a mixture of 5-bromo-7-chloro- and 7-bromo-5-chlorokynurenic acid at a specific activity of 37 GBq/ $\mu$ mol. In the biodistribution studies in



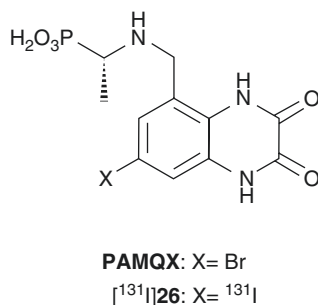
**Fig. 19.16** Chemical structure of radiolabeled cyclic amino acid derivatives

mice, [<sup>123</sup>I]**21** showed fast deiodination (85% of parent compound was changed to <sup>123</sup>I<sup>-</sup>) and low brain uptake (0.61% ID/g at 5 min). Thus, [<sup>123</sup>I]**21** had been demonstrated to be an unsuitable *in vivo* tracer (Dumont and Slegers 1997).

GV150526A (Fig. 19.15) is one of the most potent glycine site antagonists ( $K_i = 3.2$  nM vs [<sup>3</sup>H]glycine) with reasonable lipophilicity (log *P* = 2.24) (Di Fabio et al. 1997). Based on the report, Waterhouse et al. developed a meta-methoxy derivative [<sup>11</sup>C]3MPICA (Fig. 19.16) with high affinity for the glycine site ( $K_i = 4.8$  nM vs [<sup>3</sup>H]MDL 105,519) (Waterhouse et al. 2002a). In the biodistribution studies in rats, regional uptake of [<sup>11</sup>C]3MPICA at 2 min ranged from 0.11% ID/g to 0.18% ID/g. The radioactivity was cleared rapidly from all brain regions examined. Blocking studies using unlabeled compound (1 mg/kg) did not cause a marked reduction in the ratio of the brain tissues/blood. Treatment with warfarin (100 mg/kg) led to reduction in blood radioactivity concentrations, while brain uptake was not changed significantly, which indicated that factors other than binding to serum albumin could prevent BBB penetration of [<sup>11</sup>C]3MPICA (Waterhouse et al. 2002b).

*trans*-5,7-Dichloro-4-substituted-2-carboxytetrahydroquinolines such as **22** (Fig. 19.16) showed high affinities ( $K_i = 6.0$  nM vs [<sup>3</sup>H]5,7-dichlorokynurenic acid) for the glycine site of NMDARs (Leeson et al. 1992). Based on the SAR studies, fluoroethyl derivative [<sup>18</sup>F]**23** (Fig. 19.16) was designed, synthesized, and evaluated as a PET radioligand for glycine site. Component **23** had a relatively low lipophilicity (log *D* = 1.3) and high affinity ( $K_i = 12$  nM vs [<sup>3</sup>H]MDL 105,519) for the glycine-binding site. Biodistribution studies revealed that [<sup>18</sup>F]**23** showed homogeneous low levels of accumulation in all considered brain regions. Because regional differences

**Fig. 19.17** Chemical structure of PAMQX and radioiodinated PAMQX derivative



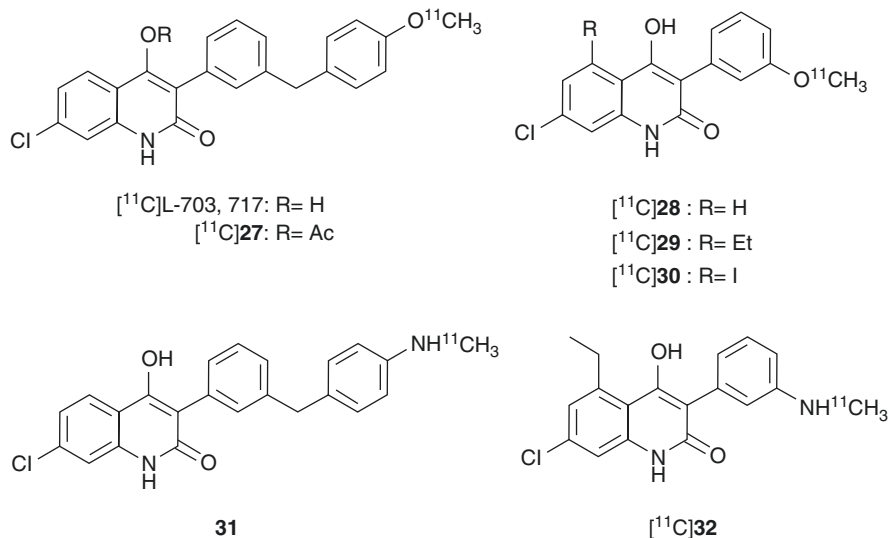
in the brain distribution of [<sup>18</sup>F]23 could not be detected, no blocking experiments were performed (Piel et al. 2003). Recently, hydantoin-substituted indole carboxylic acids were developed as novel PET ligands for the glycine-binding site with a higher lipophilicity than 23. Compounds 24 and 25 showed moderate affinity for the glycine site ( $K_i = 53$  nM for 24 and  $K_i = 31$  for 25 in [<sup>3</sup>H]MDL 105,519 binding assay). The logD values of 17 and 18 were 2.07 and 2.26, respectively. [<sup>18</sup>F]24 and [<sup>11</sup>C]25 were synthesized with poor overall decay-corrected radiochemical yields of 5–7% and 6–9% and specific activities of 24–67 GBq/μmol and 8–26 GBq/μmol, respectively (Bauman et al. 2011). No further in vivo evaluations of these radioligands have been reported so far.

### 19.1.3.2 5-Aminomethylquinoxaline-2,3-Dione Derivative

(D)-7-Bromo-*N*-(1-phosphonoethyl)-5-aminomethylquinoxaline-2,3-dione (PAMQX, Fig. 19.17) has been discovered as a selective and high-affinity glycine site antagonist with an  $IC_{50}$  value of 6 nM against [<sup>3</sup>H]MDL-105519 binding (Auberson et al. 1999). The 7-iodo derivative of PAMQX (26, Fig. 19.17) exhibited comparable affinity ( $IC_{50} = 8$  nM vs [<sup>3</sup>H]MDL-105,519) for the glycine site with PAMQX (Ametamey et al. 2000). Radiosynthesis of [<sup>131</sup>I]26 was achieved by iododestannylation of *N*-CBZ-protected tributylstannyl precursor followed by deprotection of CBZ group. Biodistribution studies of [<sup>123</sup>I]26 in mice revealed an extremely low BBB permeability of [<sup>123</sup>I]26 (0.03% ID/g at 30 min) probably due to its high polarity and low log *D* value ( $\log D_{7,4} = -2.5$ ) (Ametamey et al. 2000).

### 19.1.3.3 4-Hydroxyquinolone Derivatives

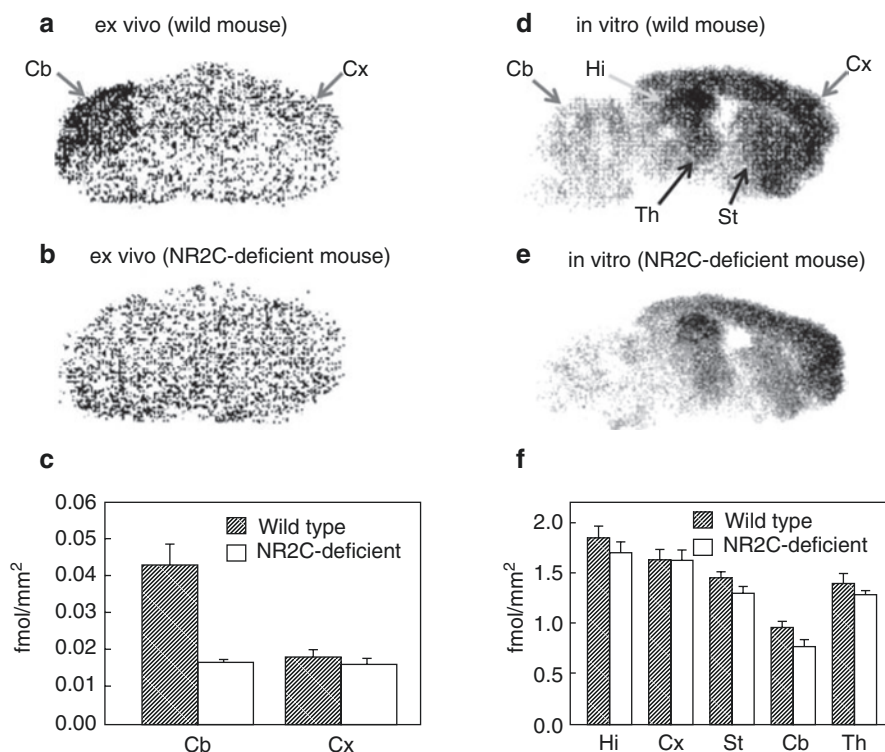
The radioligands consisting of carboxylic acids showed poor brain penetration due to the highly polar nature of the charged carboxylate. The 4-hydroxyquinolones (4-HQs)-devised carboxylic bioisostere had been developed as one of the most potent antagonists for the glycine-binding site of NMDARs. As expected, this series showed in vivo anticonvulsant activity, which suggested an ability to penetrate the BBB (Kulagowski et al. 1994; Rowley et al. 1997). In line with this, Haradahira et al. synthesized and evaluated [<sup>11</sup>C]L-703,717 (Fig. 19.18), one of the most potent glycine site antagonists ( $IC_{50} = 4.5$  nM vs [<sup>3</sup>H]L-689,560), as a PET radioligand for the glycine-binding site. In a biodistribution study in mice, [<sup>11</sup>C]L-703,717 demonstrated a low BBB permeability (0.32–0.36% ID/g at 1 min) and extremely high



**Fig. 19.18** Chemical structure of radiolabeled 4-hydroxyquinolone derivatives

blood concentration presumably due to the high affinity for the warfarin-binding site of the plasma protein. Indeed, warfarin led to a dose-dependent enhancement of the initial brain uptake of  $[^{11}\text{C}]$ L-703,717 that resulted in a fivefold increase in the brain radioactivity at the dose of 200 mg of warfarin. However, the radioactivity in the cerebrum was rapidly decreased (0.2% ID/g) with some retention observed in the cerebellum (0.65% ID/g) at 30 min after injection. This suggested inconsistent binding with known distribution of NMDARs. In addition, a significant blocking effect by non-radioactive L-703,717 (2 mg/kg) was observed only in the cerebellum (Haradahira and Suzuki 1999; Haradahira et al. 2000). To improve brain penetration of  $[^{11}\text{C}]$  L-703,717, the acetyl derivative of L-703,717 ( $[^{11}\text{C}]$ 27, Fig. 19.18) was prepared and evaluated as prodrug of  $[^{11}\text{C}]$ L-703,717. Initial brain uptake of  $[^{11}\text{C}]$ 27 (0.71% ID/g) was twofold higher than that of  $[^{11}\text{C}]$ L-703,717 (0.36% ID/g). Metabolism experiments of  $[^{11}\text{C}]$ 27 in rat brain homogenates confirmed that about 80% of  $[^{11}\text{C}]$ 27 was converted to  $[^{11}\text{C}]$ L-703,717 at 20 min after injection. Ex vivo autoradiography demonstrated that  $[^{11}\text{C}]$ 27 exhibited similar cerebellar localization of radioactivity to  $[^{11}\text{C}]$  L-703,717 (Haradahira et al. 2001). Further in vivo evaluation revealed that the specific cerebellar binding of  $[^{11}\text{C}]$ L-703,717 was abolished in NR2C-deficient mice as shown in Fig. 19.19 (A–C). This result suggested that the in vivo cerebellar localization of  $[^{11}\text{C}]$ L-703,717 could be caused by preferential binding to the NR2C subunit-containing NMDARs predominantly expressed in the cerebellum. However,  $[^{11}\text{C}]$ L-703,717 exhibited high specific localization in the hippocampus and cerebral cortex under in vitro condition as shown in Fig. 19.19 (D–F), indicative of the absence of intrinsic selectivity for the NR2C subunit-containing NMDARs in the cerebellum (Haradahira et al. 2002b). Thus, the authors next hypothesized that endogenous NMDAR co-agonists might play a key role in

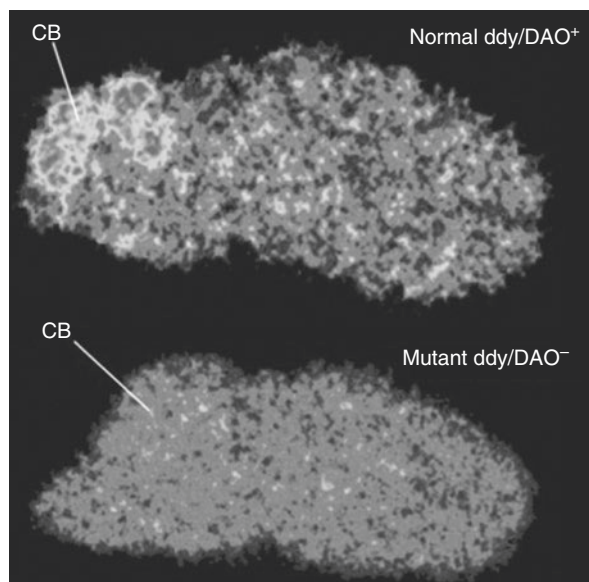




**Fig. 19.19** Ex vivo and in vitro autoradiograms of  $[^{11}\text{C}]$ L-703,717 in sagittal sections of wild-type and NR2C-deficient mice brain. (a, b) Ex vivo images obtained at 20 min after injection of  $[^{11}\text{C}]$ L-703,717 and warfarin (60 mg/kg). (c) Quantified values of the ex vivo binding in the cerebral cortex (Cx) and cerebellum (Cb). (d, e) In vitro images of  $[^{11}\text{C}]$ L-703,717. (f) Quantified values of the in vitro binding in the hippocampus (Hi), cerebral cortex (Cx), striatum (St), cerebellum (Cb), and thalamus (Th). Haradahira et al. (2002b)

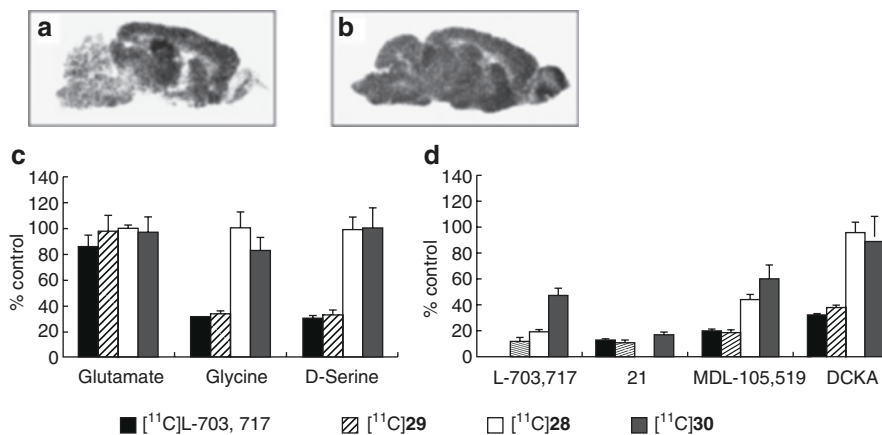
the unusual localization in the cerebellum of  $[^{11}\text{C}]$ L-703,717. Both glycine and d-serine are known as co-agonists of NMDARs and exist at micromolar level in the living brain. Glycine is distributed throughout the brain, while d-serine is localized only in the forebrain regions (Hashimoto et al. 1993; Schell et al. 1997). d-Serine is reported to be nearly undetectable in the cerebellum due to the highest level of d-amino acid oxidase (DAO) activity (Schell et al. 1995). Haradahira et al. demonstrated that the cerebellar localization of  $[^{11}\text{C}]$ L-703,717 was abolished in mutant  $\text{ddY}/\text{DAO}^-$  mice, in which the d-serine level in the cerebellum was markedly increased (Fig. 19.20). Hence, the authors proposed that the low level of  $[^{11}\text{C}]$ L-703,717 binding in forebrain regions could be explained by the strong inhibition of high amount of endogenous d-serine, whereas specific localization in the cerebellum might be due to the lack of d-serine (Haradahira et al. 2003). Considering that  $[^{11}\text{C}]$ L-703,717 has been demonstrated to recognize cerebellar NMDARs to some extent, clinical PET studies of  $[^{11}\text{C}]$ 27 (a prodrug of  $[^{11}\text{C}]$  L-703,717) were

**Fig. 19.20** Ex vivo autoradiogram of [ $^{11}\text{C}$ ]L-703,717 in the brain of normal ddY/DAO $^{+}$  and mutant ddY/DAO $^{-}$  mice. The brain slices (50  $\mu\text{m}$ ) were obtained at 30 min after injection of [ $^{11}\text{C}$ ]L-703,717 and warfarin (60 mg/kg). Haradahira et al. (2003)



performed for the visualization of NMDARs in the cerebellum (Matsumoto et al. 2007). Binding potential (BP) calculated from white matter as a reference region in the cerebellar cortex was twofold higher than in the cerebral cortices (cerebellar cortex, BP = 2.2; cerebral cortices, BP = 1.1). Regional brain distributions were consistent with previous reports of rodents (Haradahira et al. 2001). However, the brain uptake of [ $^{11}\text{C}$ ]27 was too low for the visualization of the cerebellar NMDAs by PET (Matsumoto et al. 2007).

3-Methoxy-MDL-104,653 (28, Fig. 19.18) has been reported as a weak antagonist ( $\text{IC}_{50} = 204 \text{ nM}$  vs [ $^3\text{H}$ ]L-689,560) for the glycine site of NMDARs (Rowley et al. 1997). PET studies on monkeys demonstrated that [ $^{11}\text{C}$ ]28 exhibited higher localization in the cortex regions than in the cerebellum, in which binding was not blocked by glycine site agonists (Haradahira et al. 2002c). It is reported that glycine site agonists and antagonists bind to overlapping but different sites on the NMDARs (Kuryatov et al. 1994; Wafford et al. 1995). Previous studies on [ $^{11}\text{C}$ ]L-703,717 suggested that most radioligands for the glycine site were susceptible to inhibition by endogenous agonists (agonist-sensitive) which resulted in strong inhibition of in vivo binding by agonists (Haradahira et al. 2002b, 2003). On the other hand, [ $^{11}\text{C}$ ]27 might be an atypical glycine site antagonist, in which the binding site overlaps with the glycine antagonist-binding site, but not with the agonist-binding site, namely, agonist-insensitive ligand. Although insufficient BBB permeability and high non-specific binding limited further investigations of [ $^{11}\text{C}$ ]28, structural optimization of [ $^{11}\text{C}$ ]28 in order to increase BBB permeability as well as improved in vivo specificity would produce a useful agonist-insensitive radioligand for the glycine site of NMDARs, especially in the forebrain regions which play crucial roles in various brain functions. On the basis of SAR studies, we developed 5-ethyl



**Fig. 19.21** Typical autoradiogram of in vitro binding of  $[^{11}\text{C}]\mathbf{28}$  (a) and  $[^{11}\text{C}]\mathbf{29}$  (b). Effect of NMDAR agonists (c) and antagonists (d) on in vitro specific  $[^{11}\text{C}]$ 4-hydroxyquinolones binding to the hippocampus. Abbreviations: *DCKA* 5,7-dichlorokynurenic acid. Fuchigami et al. (2008)

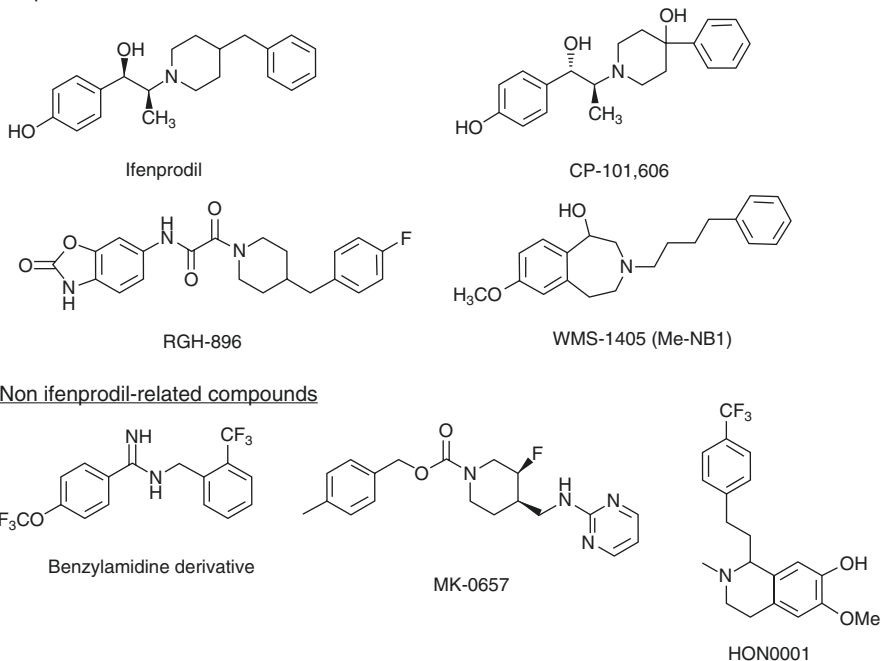
(**29**) and 5-iodine (**30**) derivative of  $[^{11}\text{C}]\mathbf{28}$  (Fig. 19.18) with high affinity for the glycine site ( $K_i = 7.2$  and  $10.3$  nM vs  $[^3\text{H}]\text{MDL-105,519}$ , respectively). In vitro autoradiography demonstrated that  $[^{11}\text{C}]\mathbf{29}$  and  $[^{11}\text{C}]\mathbf{30}$  displayed quite different binding properties.  $[^{11}\text{C}]\mathbf{29}$  showed high accumulation in the forebrain regions but lowest in the cerebellum. The binding was inhibited both by glycine site antagonists and agonists as shown in Fig. 19.21 (a, c, and d) which suggested that  $[^{11}\text{C}]\mathbf{29}$  is an agonist-sensitive ligand. On the other hand,  $[^{11}\text{C}]\mathbf{30}$  exhibited homogeneous accumulation throughout the brain, and the binding was inhibited by glycine site antagonists but not by agonists as shown in Fig. 19.21 (b, c, and d) which suggested that  $[^{11}\text{C}]\mathbf{30}$  is an agonist-insensitive ligand (Fuchigami et al. 2008). These results indicated that 4-HQs could be divided into two groups “agonist-sensitive ligands ( $[^{11}\text{C}]\text{L-703,717}$  and  $[^{11}\text{C}]\mathbf{29}$ )” and “agonist-insensitive ligands ( $[^{11}\text{C}]\mathbf{28}$  and  $[^{11}\text{C}]\mathbf{30}$ ).” It is possible that the agonist-insensitive ligands can bind the glycine antagonist-preferring site, such as homomeric GluN1 or inactivated GluN1/GluN2 site, of the NMDARs. On the other hand, the agonist-sensitive ligand can preferentially bind to the functional and heteromeric NMDARs, which are localized in the forebrain regions rather than in the cerebellum, and therefore is sensitive to agonist binding. Unfortunately,  $[^{11}\text{C}]\mathbf{29}$  and  $[^{11}\text{C}]\mathbf{30}$  did not exhibit a significant increase in brain uptake compared with  $[^{11}\text{C}]\text{L-703,717}$  and  $[^{11}\text{C}]\mathbf{28}$ , although  $[^{11}\text{C}]\mathbf{29}$  and  $[^{11}\text{C}]\mathbf{30}$  had moderate lipophilicity ( $\log P = 2.25$  for **29** and  $1.81$  for **30**). Consistent with previous reports (Haradahira et al. 2002c), agonist-sensitive  $[^{11}\text{C}]\mathbf{29}$  showed higher localization in the cerebellum, whereas agonist-insensitive  $[^{11}\text{C}]\mathbf{30}$  showed homogeneous brain uptake. However, poor brain uptake and high non-specific binding of  $[^{11}\text{C}]\mathbf{29}$  and  $[^{11}\text{C}]\mathbf{30}$  limited further detailed in vivo investigations. Because warfarin increased brain uptake of  $[^{11}\text{C}]\mathbf{29}$  and  $[^{11}\text{C}]\mathbf{30}$ , the binding of these tracers for serum albumin should have considerable effect on in vivo brain uptake (Fuchigami et al. 2008). The 4-HQs are all acidic with pKa values of around 5 or

below (Rowley et al. 1997) and more amenable to binding with plasma proteins than non-acidic compounds, which result in poor BBB permeability. Introduction of the basic amino group into 4-HQs was expected to increase the pK<sub>a</sub> value and reduce unsuitable protein binding, which could increase brain uptake. We synthesized and evaluated several amino 4-HQ derivatives as new PET radioligand candidates for the glycine site. Among these ligands, methylamino derivatives of 4-HQs (**31** and **32**, Fig. 19.18) have been identified as high-affinity glycine site ligands ( $K_i$  values, 11.7 nM for **31** and 11.8 nM for **32**). In vitro autoradiography experiments demonstrated that both [<sup>11</sup>C]**31** and [<sup>11</sup>C]**32** showed higher accumulation in the fore-brain regions than cerebellum, which was consistent with known NMDAR distribution. Furthermore, the in vitro binding of [<sup>11</sup>C]**31** and [<sup>11</sup>C]**32** was strongly inhibited by both glycine agonists and antagonists, which suggests that these tracers are agonist-sensitive glycine site ligands. Although no agonist-insensitive ligands for glycine site were obtained, further in vivo characterization of [<sup>11</sup>C]**31** and [<sup>11</sup>C]**32** was performed. In the biodistribution studies in mice, cerebellum to blood ratio of both [<sup>11</sup>C]**31** and [<sup>11</sup>C]**32** at 30 min was about 0.06 at 30 min after injection, which indicated poor BBB penetration of these radioligands. Although the plasma protein-binding ratio of [<sup>11</sup>C]**32** was much lower than that of methoxy analogs (71% vs 94–98%, respectively), [<sup>11</sup>C]**32** still bound with plasma protein strongly (Fuchigami et al. 2009). Further SAR studies are required for developing favorable PET ligands for the glycine site of NMDARs with improved BBB permeability.

### 19.1.4 Development of Radioligands for Ifenprodil-Binding Site

In the adult brain, GluN2B expression has been reported to be restricted to the fore-brain areas, including the cortex, hippocampus, striatum, thalamus, and olfactory bulb (Akazawa et al. 1994; Monyer et al. 1994). Since GluN2B has been suggested to play a key role in various diseases such as ischemic stroke, Parkinson's disease, Alzheimer's disease, and neuropathic pain, a number of organic compounds that target the GluN2B have been developed for the treatment of such diseases (Gogas 2006; Mony et al. 2009). Ifenprodil was initially discovered as a GluN2B-selective negative modulator of NMDARs ( $K_d = 34$  nM for GluN1a/2B receptors); however, it showed poor selectivity for NMDARs over other receptors such as  $\alpha$ 1-adrenergic and  $\sigma$  receptors (Contreras et al. 1990; Karbon et al. 1990; Williams 1993; Hashimoto and London 1995; Grimwood et al. 2000). Accordingly, several series of GluN2B antagonists with improved selectivity and affinity for GluN2B-containing NMDARs have been developed (Chazot 2004; Gogas 2006; Mony et al. 2009). These compounds are divided into two classes, the "ifenprodil derivatives" and the "non ifenprodil-related compounds" as shown in Fig. 19.22. Ifenprodil derivatives are likely to share the same site with ifenprodil on the interlobe cleft of the GluN2B subunit (Malherbe et al. 2003). However, whether non-ifenprodil-related GluN2B-selective antagonists have a similar binding mode to ifenprodil is unclear.

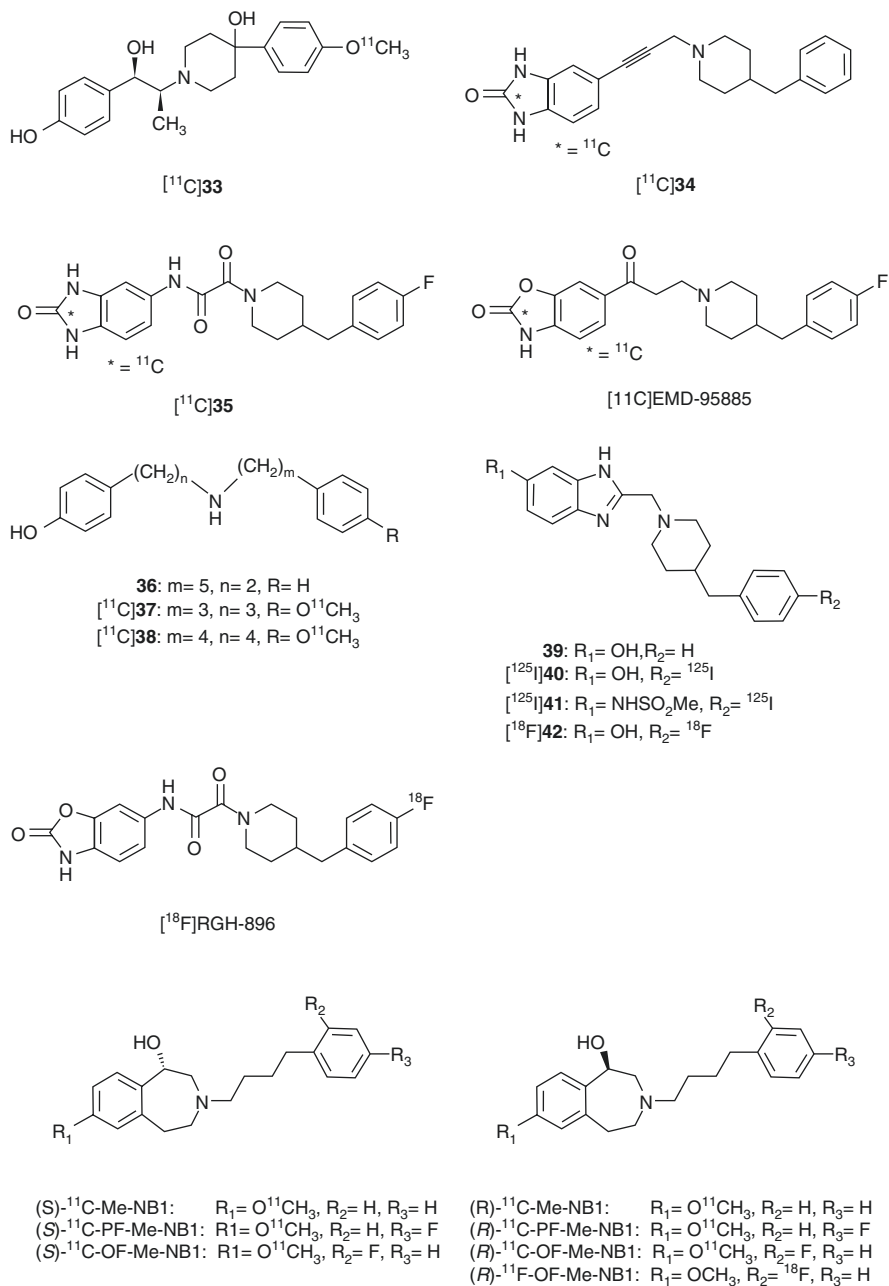
In vivo imaging of GluN2B-containing NMDARs is considered useful for obtaining information about various psychiatric diseases and facilitating drug

Ifenprodil derivatives**Fig. 19.22** Chemical structure of GluN2B-selective NMDAR antagonists

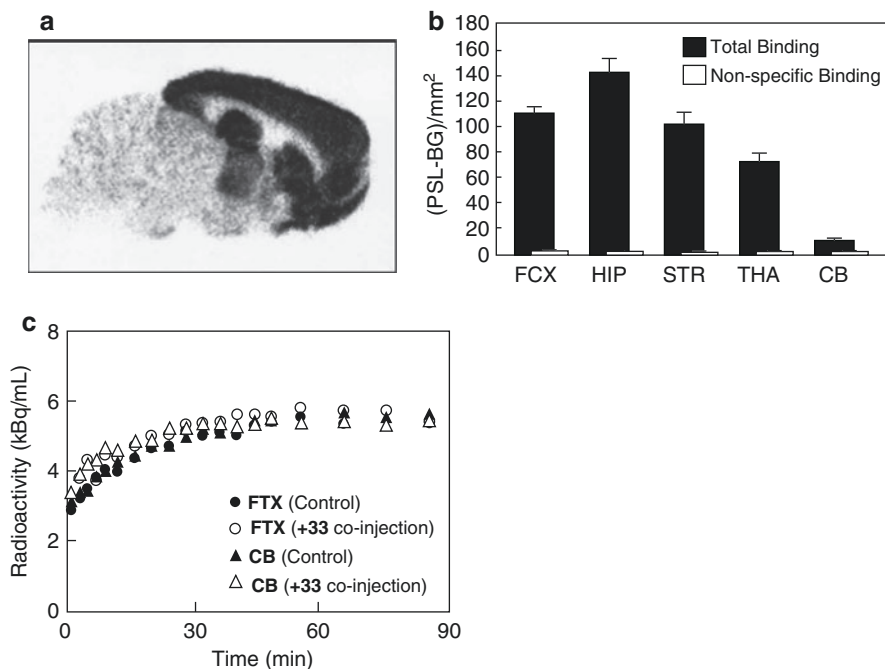
development, as well as in vivo occupancy. Development of PET or SPECT imaging probes for GluN2B has been conducted based on these GluN2B-selective antagonists as described below.

**19.1.4.1 Ifenprodil Derivatives**

It has been shown that CP-101,606 is a potent NMDAR antagonist highly selective for GluN2B-containing receptors with  $K_d$  values of 10 nM (Menniti et al. 1997). A carbon-11-labeled methoxy analog of CP-101,606 ( $[^{11}\text{C}]\mathbf{33}$ , Fig. 19.23) was developed for PET imaging of the GluN2B-containing NMDARs (Haradahira et al. 2002a). In vitro autoradiography in rat brain slices revealed an extremely high localization of  $[^{11}\text{C}]\mathbf{33}$  in the forebrain regions and very low in the cerebellum as shown in Fig. 19.24a. The in vitro distribution pattern of  $[^{11}\text{C}]\mathbf{33}$  matched previously reported  $[^3\text{H}]\text{CP-101,606}$  binding (Menniti et al. 1997) and GluN2B expression (Mori and Mishina 1995). Non-specific binding of  $[^{11}\text{C}]\mathbf{33}$  was found to be less than 5% as shown in Fig. 19.24b. The binding of  $[^{11}\text{C}]\mathbf{33}$  was displaced by CP-101,606 and ifenprodil in a competitive manner. Treatment of spermine and  $\text{Zn}^{2+}$  (1 mM) markedly decreased in vitro binding of  $[^{11}\text{C}]\mathbf{33}$  (74–82%). It is suggested that ifenprodil and  $\text{Zn}^{2+}$  share the binding site on the interlobe cleft of GluN2B (Rachline et al. 2005). These results indicated that this tracer recognizes the ifenprodil-binding site on the GluN2B under in vitro condition. In contrast, in vivo biodistribution



**Fig. 19.23** Chemical structure of radiolabeled ifenprodil derivatives and bis(phenylalkyl)-amines



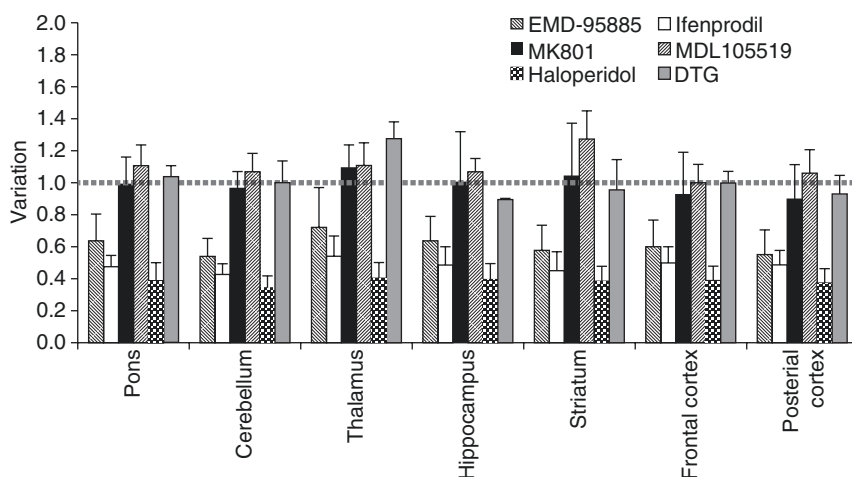
**Fig. 19.24** (a) In vitro autoradiogram of [<sup>11</sup>C]33 in sagittal sections of rat brain. (b) Quantified values of the autoradiogram in the frontal cortex (FCX), hippocampus (HIP), striatum (STR), thalamus (THA), and cerebellum (CB). Non-specific binding was determined in the presence of (+)CP-101,606 (10 μM). (PSL-BG)/mm<sup>2</sup> means photostimulated luminescence (PSL) values per mm<sup>2</sup>. Background PSL values (BG) were subtracted from the total PSL values on ROIs. (c) Time radioactivity curves in the frontal cortex (FCX) and cerebellum (CB) after i.v. injection of [<sup>11</sup>C]33 into monkey. Non-radioactive 33 (2 mg/kg) was injected simultaneously with [<sup>11</sup>C]33 into the same monkey. Haradahira et al. (2002a)

studies on mice and PET studies on monkeys demonstrated that this tracer exhibited a moderate brain uptake, but no apparent specific localization of the radioactivity was observed in any of the brain regions as shown in Fig. 19.24c (Haradahira et al. 2002a).

The 1-(arylkynyl)-4-benzylpiperidine derivative 34 (Fig. 19.23) has been discovered as highly potent (IC<sub>50</sub> = 5.3 nM) GluN2B-selective NMDAR antagonist (Wright et al. 2000). The in vivo brain uptake of [<sup>11</sup>C]34 in rats was fairly low (0.07% ID/g at 30 min), and localization was inconsistent with known GluN2B expression with highest uptake in the cerebellum and lowest in the striata. Pretreatment with ifenprodil (20 mg/kg, i.p.) did not cause a prominent change in brain tissues/plasma ratio of radioactivity, which suggested a high non-specific binding of [<sup>11</sup>C]34 (Roger et al. 2003). Barta-Szalai et al. developed oxamides derived from indole-2-carboxamides as potent GluN2B-selective NMDA receptor antagonists. Among them, compound 35 showed the best affinity in this series with an IC<sub>50</sub> of 5 nM in the [<sup>3</sup>H]Ro-25,6981 binding assay (Barta-Szalai et al. 2004).

Recently, [ $^{11}\text{C}$ ]**35** was synthesized using [ $^{11}\text{C}$ ]phosgene. [ $^{11}\text{C}$ ]**35** was obtained with a radiochemical yield of 30–40% with high specific radioactivity (72–127 GBq/ $\mu\text{mol}$ ) (Labas et al. 2010). Further in vitro and in vivo studies of [ $^{11}\text{C}$ ]**35** have not yet been published. Another 4-benzylpiperidine derivative, EMD-95885 (Fig. 19.23), has been reported to exhibit high affinity ( $\text{IC}_{50} = 3.9 \text{ nM}$ ) and is a selective GluN2B antagonist (Leibrock et al. 1997). [ $^{11}\text{C}$ ] EMD-95885 was prepared by a similar method as used for [ $^{11}\text{C}$ ]**34** using [ $^{11}\text{C}$ ]COCl $_2$  with a specific activity of 37–74 GBq/ $\mu\text{mol}$ . Radiosensitive  $\beta$ -microprobe acquisition and biodistribution studies in rats revealed that [ $^{11}\text{C}$ ]EMD-95885 displayed five- to ninefold higher BBB penetration than [ $^{11}\text{C}$ ]**34**. However, the distribution was almost homogeneous in all brain structures studied. Pretreatment with non-radioactive **34** or ifenprodil led to a considerable reduction in brain uptake of [ $^{11}\text{C}$ ]EMD-95885, whereas other site ligands such as MK-801, MDL-105,519, and DTG (1,3-di-*o*-tolylguanidine) had no blocking effect (Fig. 19.25), suggesting some specific binding in the brain tissues (Roger et al. 2004). It should be taken into consideration that the blocking effects of [ $^{11}\text{C}$ ]EMD-95885 binding were observed not only in GluN2B-rich regions, such as the hippocampus and frontal cortex, but also in GluN2B-poor region, such as the cerebellum. Although treatment with ifenprodil-like compounds resulted in a decrease of in vivo [ $^{11}\text{C}$ ]EMD-95885 binding, this tracer was not a promising PET radioligand for the GluN2B-containing NMDARs.

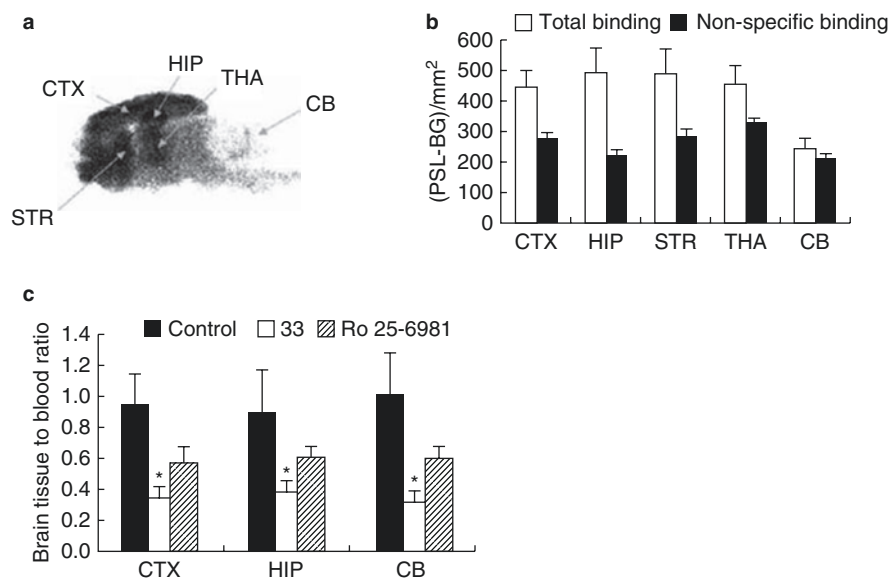
The bis(phenylalkyl)amines were reported to be selective antagonists of GluN1a/2B receptors. The most potent of this series, compound **36** (Fig. 19.23), has an  $\text{IC}_{50}$  value of 8 nM and > 1000-fold selectivity for GluN1a/2B receptors over other subunit-containing receptors (Tamiz et al. 1998). Thus, two  $^{11}\text{C}$ -labeled analogs of bis(phenylalkyl) amine, bis(phenylpropyl)amine ([ $^{11}\text{C}$ ]**37**) and bis(phenylbutyl)amine ([ $^{11}\text{C}$ ]**38**), were



**Fig. 19.25** Effect of various drugs on in vivo localization of [ $^{11}\text{C}$ ]EMD-95885 in rat brain. The y-axis labeled as “Variation” means ratio of the binding in the presence of blocker to the binding in the absence of blocker in each brain region. Roger et al. (2004)



synthesized and evaluated by Sasaki et al. These [ $^{11}\text{C}$ ]-ligands were prepared by *O*-methylation on the corresponding diphenol precursors with [ $^{11}\text{C}$ ]CH $_3$ I with a specific activity of around 75 GBq/ $\mu\text{mol}$ . In the in vitro autoradiography studies, both [ $^{11}\text{C}$ ]**37** and [ $^{11}\text{C}$ ]**38** displayed almost homogeneous distribution throughout the brain. Treatment with non-radioactive **37** and **38** led to only 50% and 40% reduction in the in vitro binding of [ $^{11}\text{C}$ ]**37** and [ $^{11}\text{C}$ ]**38**, respectively. Both spermine and ifenprodil caused 30–40% reduction of these tracers, which indicated that these bis(phenylalkyl) amines shared the binding site with ifenprodil. Further in vivo studies in rodents demonstrated that [ $^{11}\text{C}$ ]**37** exhibited homogeneous accumulation and non-labeled **37** increased the brain uptake. Because of the high non-specific binding both in vitro and in vivo, [ $^{11}\text{C}$ ]**37** and [ $^{11}\text{C}$ ]**38** are unsuitable radiotracers for GluN1a/2B receptors (Sasaki et al. 2004). A new series of benzimidazole derivatives such as compound **39** [ $K_i = 1.5 \text{ nM}$  vs [ $^3\text{H}$ ]{(*E*)-*N* $^1$ -(2-methoxybenzyl) cinnamamide (compound **43** as described below)}] was developed as highly potent GluN2B antagonists (McCauley et al. 2004). Based on the article, we developed 4'-radioiodinated benzimidazoles as SPECT imaging agents for the GluN2B. The compounds **40** and **41** showed high affinity for GluN2B with  $K_i$  values of 7.3 nM and 5.8 nM, respectively (vs [ $^3\text{H}$ ]ifenprodil). In vitro autoradiography revealed that both [ $^{125}\text{I}$ ]**40** and [ $^{125}\text{I}$ ]**41** exhibited high accumulation in the forebrain regions but low in the cerebellum as shown in Fig. 19.26a. These regional distributions of the radioligands were consistent with the expression of the



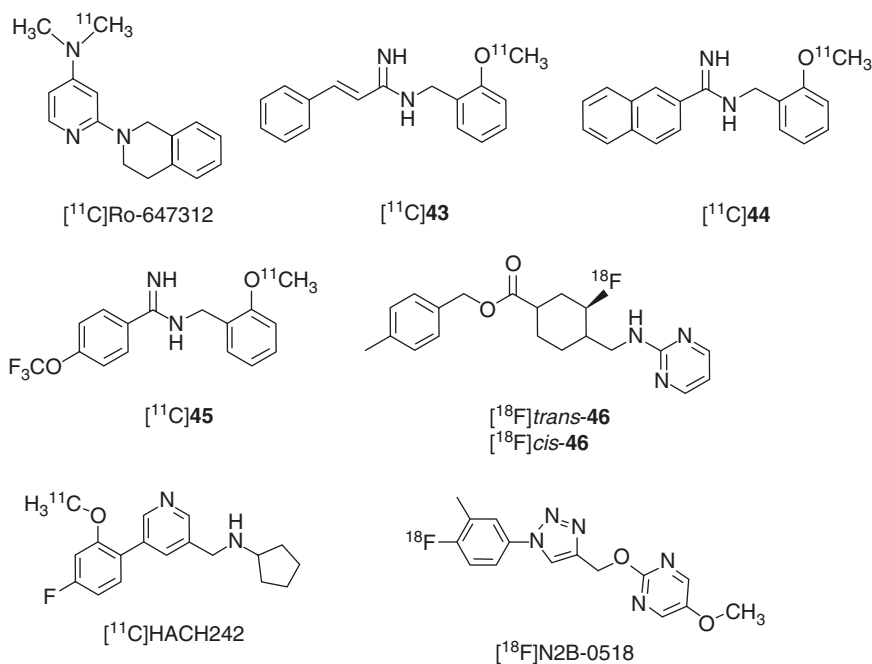
**Fig. 19.26** (a) In vitro autoradiogram of [ $^{125}\text{I}$ ]**40** in sagittal sections of rat brain. (b) Quantified values of the autoradiogram in the cerebral cortex (CTX), hippocampus (HIP), striatum (STR), thalamus (THA), and cerebellum (CB). Non-specific binding was determined in the presence of non-radioactive **40** (10  $\mu\text{M}$ ). (c) Effect of non-radioactive **40** and Ro 25-6981 (GluN2B antagonist) on the brain tissue to blood ratio in mice at 180 min after intravenous injection of [ $^{125}\text{I}$ ]**40**. Drugs were pretreated 30 min before the tracer injection. \* $P < 0.01$  in comparison to the control group. Fuchigami et al. (2010)

GluN2B. The *in vitro* binding of these tracers was blocked by GluN2B antagonist but not by other site ligands, which suggested the high selectivity of [<sup>125</sup>I]**40** and [<sup>125</sup>I]**41** for the GluN2B subunit. However the specific binding of these [<sup>125</sup>I]benzimidazoles was lower than CP-101,606 analog [<sup>11</sup>C]**33** as shown in Fig. 19.26b probably due to high lipophilicity of these [<sup>125</sup>I]ligands. In mice, [<sup>125</sup>I]**40** and [<sup>125</sup>I]**41** showed moderate brain uptake at 5–180 min after administration (0.42–0.56% and 0.44–0.67% ID/g, respectively). The brain-to-blood ratio of [<sup>125</sup>I]**40** led to 34% reduction in the presence of non-radioactive **40** and 59% reduction in the presence of the GluN2B ligand Ro 25-6981 as shown in Fig. 19.26c. Although these radioligands showed poor BBB permeability and inconsistent distribution with GluN2B, [<sup>125</sup>I]**40** may be partially bound to the GluN2B subunit under *in vivo* conditions. Compound **40** may be a potential scaffold for further structural modification in order to develop potential imaging probes for the GluN2B-containing NMDARs (Fuchigami et al. 2010). More recently, high-affinity GluN2B antagonists benzimidazole derivative (**42**, Fig. 19.23) and RGH-896 (Fig. 19.23) were successfully labeled with <sup>18</sup>F by an aromatic nucleophilic radiofluorination followed by a reduction of the para-position carbonyl function (Labas et al. 2011). Despite modification of synthetic conditions, both [<sup>18</sup>F]**42** and [<sup>18</sup>F]RGH-896 were obtained with poor specific activities of 1.7 GBq/μmol and 1.2 GBq/μmol, respectively. *In vivo* brain uptakes of [<sup>18</sup>F]**42** or [<sup>18</sup>F]RGH-896 in rats were quite low at 120 min after injection of radiotracers (0.035% ID/g and 0.054% ID/g, respectively). High accumulation of radioactivity in the bone and further metabolic analysis of [<sup>18</sup>F]fluorobenzyl piperidine fragment implicated that [<sup>18</sup>F]**42** and [<sup>18</sup>F]RGH-896 seemed to be rapidly defluorinated by hydroxylation in the benzyl ring followed by glucuronidation (Labas et al. 2011). It is reported that cleavage and reconstitution of a bond in the piperidine ring of ifenprodil led to 3-benzazepine derivatives as selective GluN1/GluN2B antagonists (Tewes et al. 2010; Szermerski et al. 2018). In particular, WMS-1405 showed the most potent binding affinity ( $K_i = 5.4$  nM) for GluN1/GluN2B NMDA receptors with poor interaction with  $\sigma_1$  and  $\sigma_2$  receptors ( $K_i = 182$  and  $554$  nM, respectively). In 2018, Krämer et al. labeled WMS-1405 with C-11 to yield <sup>11</sup>C-Me-NB1 as a PET tracer of GluN1/GluN2B. *In vitro* autoradiography demonstrated that <sup>11</sup>C-Me-NB1 showed homogeneous distribution throughout the rat brain slices. The binding was blocked by GluN2B antagonists, eliprodil (54 nM and 10 μM) and Ro-25-6981 (100 μM). However, haloperidol (10 μM), an antagonist of D<sub>2</sub>, D<sub>3</sub>, and D<sub>4</sub> receptors and an inverse agonist of  $\sigma_1$  receptor, reduced the binding of <sup>11</sup>C-Me-NB1. *In ex vivo* studies in rat, high brain SUVs were shown in the cortex with high expression of GluN2B ( $3.7 \pm 1.0$ ), whereas a relatively high SUV value was observed in the mid-brain and cerebellum with the GluN2B-poor region ( $3.8 \pm 1.2$  and  $2.8 \pm 0.8$ , respectively). In addition, treatment with eliprodil (2 mg/kg) led to a reduction in <sup>11</sup>C-Me-NB1 binding in all the brain regions. These results indicated that <sup>11</sup>C-Me-NB1 binds not only to GluN2B but also in off-target regions (Krämer et al. 2018). Next, the same group separately synthesized the enantiopure (*R*)- and (*S*)-<sup>11</sup>C-Me-NB1, respectively. Interestingly, these tracers showed quite different properties. *In vitro* autoradiography with rat brain slices demonstrated the heterogeneous brain distribution of (*R*)-<sup>11</sup>C-Me-NB1 with high selectivity for the GluN2B-rich forebrain, whereas (*S*)-<sup>11</sup>C-Me-NB1 displayed a homogeneous accumulation in all brain regions and predominantly

exhibited  $\sigma_1$ -receptor binding. Similar results were observed in the postmortem human brain tissues. In PET imaging with rodents, (*R*)- $^{11}\text{C}$ -Me-NB1 displayed consistent accumulation in the GluN2B-expression pattern. Eliprodil led to the dose-dependent reduction in (*R*)- $^{11}\text{C}$ -Me-NB1 accumulation in the GluN2B-rich regions. These results indicated that (*R*)- $^{11}\text{C}$ -Me-NB1 is a promising PET tracer for imaging the GluN2B subunit of the NMDA receptor (Haider et al. 2019). Introduction of fluoride at the para position of the phenyl moiety in (*R*)-Me-NB1 abolished specific binding for GluN2B, whereas (*R*)-OF-Me-NB1 which introduced fluorine atom at the ortho position maintained binding affinity and specificity to GluN2B as demonstrated both by in vitro and in vivo studies. PET imaging studies of (*R*)- $^{18}\text{F}$ -OF-Me-NB1 in rats showed consistent brain accumulation in the known GluN2B distribution with high specific binding to GluN2B as determined by blocking studies with CP101,606 (Haider et al. 2018). (*R*)- $^{18}\text{F}$ -OF-Me-NB1 may be superior to (*R*)- $^{11}\text{C}$ -Me-NB1 in that  $^{18}\text{F}$  has a longer half-life than  $^{11}\text{C}$  (109.8 vs 20.3 min) and can be easy to handle and evaluate because of a better pharmacokinetic profile. However, further detailed in vivo studies, including clinical studies, would be needed to confirm the feasibility of these PET tracers.

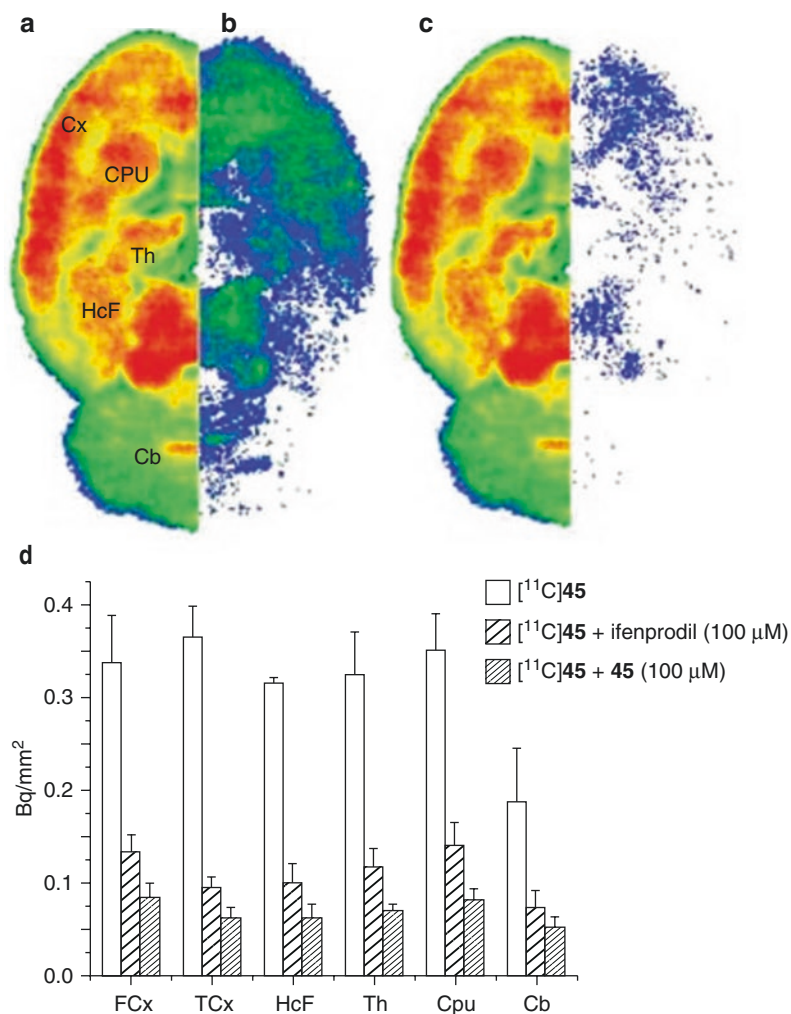
#### 19.1.4.2 Non-ifenprodil-Related GluN2B Antagonist Derivatives

A series of 2-(3,4-dihydro-1H-isoquinolin-2-yl)-pyridines have been identified as a new class of selective antagonists of GluN1/GluN2B-containing NMDARs. Among them, Ro-647312 (Fig. 19.27) displayed high affinity and selectivity for



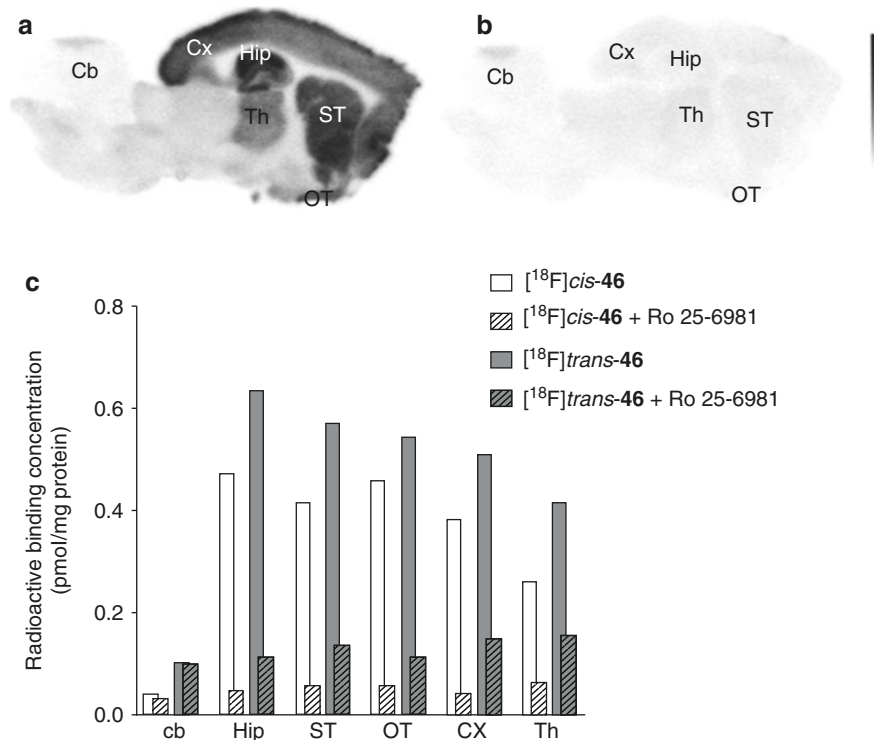
**Fig. 19.27** Chemical structure of radiolabeled non-ifenprodil-related derivatives

GluN2B-containing NMDARs ( $K_i = 8$  nM vs [ $^3$ H]Ro 25-6981) (Büttelmann et al. 2003). The [ $^{11}$ C]Ro-647312 was synthesized using [ $^{11}$ C]methyl triflate from the non-methyl compound with a specific activity of 37–129 GBq/ $\mu$ mol. Biodistribution studies of [ $^{11}$ C]Ro-647312 in rats demonstrated that the distribution of radioactivity was homogeneous in all brain structures and correlated poorly with known distribution of GluN2B (Dollé et al. 2004). Benzylamidines were discovered as novel series of highly potent GluN2B-selective NMDAR antagonists. From this series, (*E*)-styrene derivative (**43**, Fig. 19.27) and 2-naphthyl derivative (**44**, Fig. 19.27) displayed excellent high affinity for GluN2B with  $K_i$  value of 0.7 nM and 1.3 nM (vs [ $^3$ H]ifenprodil), respectively (Claiborne et al. 2003; Curtis et al. 2003). The 4-trifluoromethoxyphenyl derivative (**45**, Fig. 19.27) was identified as a highly potent GluN2B antagonist with  $K_i$  value of 5.7 nM as determined by patch-clamp experiments of GluN1a/GluN2B receptors (Claiborne et al. 2003). In vitro autoradiography studies showed consistent brain localization of [ $^{11}$ C]**45** with distribution of GluN2B. In addition, the in vitro binding of [ $^{11}$ C]**45** was strongly inhibited (over 75%) by non-radioactive **45** and ifenprodil (100  $\mu$ M) (Fig. 19.28), which suggested the high specific binding toward GluN2B-containing NMDARs. [ $^{11}$ C]**43** and [ $^{11}$ C]**44** showed moderate initial brain uptake (about 1.0% ID/g at 5 min), while [ $^{11}$ C]**45** exhibited higher brain uptake with 1.8% ID/g. Metabolite analysis demonstrated that only 31% and 37% of intact [ $^{11}$ C]**43** and [ $^{11}$ C]**44**, respectively, were detected in the brain, whereas 49% of authentic [ $^{11}$ C]**45** remained in the brain at 40 min. Because of metabolic instability, it can be stated that these radiotracers are unsuitable imaging agents for the GluN2B (Arstad et al. 2006). Our independent studies of [ $^{11}$ C]**43** and [ $^{11}$ C]**44** agreed with the study by Arstad et al. The percentage of unchanged radioligands was very low not only in the blood but also in the brain. These data implicated that the metabolite of the radiotracers could enter the brain tissue. Indeed, radio TLC analysis of [ $^{11}$ C]**44** in the whole brain demonstrated that the metabolite deemed as the amide derivative of [ $^{11}$ C]**44**, which could pass easily through BBB, was detected (Fuchigami et al. 2018). MK-0657 (Fig. 19.27) was reported as a highly potent GluN2B antagonist ( $IC_{50} = 3.6$  nM). Clinical trial of MK-0657 has begun for the treatment of neuropathic pain, Parkinson's disease, and major depression (Addy et al. 2009; Mony et al. 2009). Two radiofluorinated diastereoisomers of MK-0657 ([ $^{18}$ F]*trans*-**46** and [ $^{18}$ F]*cis*-**46**) have been developed as PET radioligands for GluN2B-containing NMDARs. Both ligands displayed high affinity for GluN2B with  $K_d$  values ranging from 1.9 to 7.1 nM for [ $^{18}$ F]*trans*-**46** and 5.9–15.2 nM for [ $^{18}$ F]*cis*-**46**, respectively. In vitro autoradiography experiments on rat brain slices showed similar localization of [ $^{18}$ F]*trans*-**46** and [ $^{18}$ F]*cis*-**46** to the known distribution pattern of GluN2B as shown in Fig. 19.29a, b. The in vitro binding of these ligands was strongly inhibited by Ro 25-6981 (10  $\mu$ M), which led to low non-specific binding in all brain regions as shown in Fig. 19.29c (Koudih et al. 2012). These promising in vitro results could lead to further in vivo evaluation of [ $^{18}$ F]*trans*-**46** and [ $^{18}$ F]*cis*-**46** in the future. Christiaans et al. reported a high-affinity PET probe *N*-((5-(4-fluoro-2-[ $^{11}$ C]methoxyphenyl)pyridin-3-yl)methyl) cyclopentanamine ([ $^{11}$ C]HACH242) for the GluN2B subunit ( $K_i = 12.4$  nM vs [ $^3$ H]ifenprodil assay). This probe showed consistent in vitro binding with GluN2B expression in the rat brain sections. However, [ $^{11}$ C]HACH242 revealed inconsistent ex vivo brain



**Fig. 19.28** In vitro autoradiogram of [<sup>11</sup>C]45 in sagittal sections of rat brain, under control conditions (a), in the presence of 100 μM ifenprodil (b) or non-radioactive 49 (c), and quantified values of the autoradiogram under the above three conditions (d). Abbreviations: *Cx* cortex, *FCx* frontal cortex, *TCx* temporal cortex, *Cpu* caudate putamen, *Th* thalamus, *HcF* hippocampal formation, *Cb* cerebellum. Arstad et al. (2006)

distribution with the expression level of the GluN2B subunit presumably due to non-specific binding to the  $\sigma_1$  receptors (Christiaans et al. 2014). PET imaging of [<sup>11</sup>C]HACH242 in the primate brain demonstrated a high brain uptake but homogeneous distribution. No significant reduction in the brain [<sup>11</sup>C]HACH242 SUV values was observed after intravenous infusion of radioprodil. Metabolism experiments indicated that only 26% of the parent fraction of [<sup>11</sup>C]HACH242 remained till 20 min post-injection (van der Aart et al. 2019).



**Fig. 19.29** In vitro autoradiogram of [ $^{18}\text{F}$ ]cis-46 in sagittal sections of rat brain, under control conditions (a), in the presence of 10  $\mu\text{M}$  Ro 25-6981 (b). Radioligands concentration in brain structure after in vitro binding of [ $^{18}\text{F}$ ]cis-46 or [ $^{18}\text{F}$ ]trans-46 alone or co-incubation with Ro 25-6981 (10  $\mu\text{M}$ ) on rat brain sections (c). Abbreviations: Cb cerebellum, Cx cortex, Hip hippocampus, ST striatum, OT olfactory tubercle, Th thalamus. Koudih et al. (2012)

Recently, Fu et al. reported several  $^{18}\text{F}$ -labeled methoxypyrimidines as PET tracers for GluN2B. The 2-((1-(4-[ $^{18}\text{F}$ ]fluoro-3-methylphenyl)-1H-1,2,3-triazol-4-yl)methoxy)-5-methoxypyrimidine ([ $^{18}\text{F}$ ]N2B-0518) strongly inhibited the current responses of the GluN1/GluN2B receptors with an  $\text{IC}_{50}$  value of 28 nM. In vitro autoradiography demonstrated that the localization of [ $^{18}\text{F}$ ]N2B-0518 was consistent with the expression of the GluN2B subunit. In vivo biodistribution study showed that [ $^{18}\text{F}$ ]N2B-0518 showed moderate brain uptake (3.60% ID/g at 2 min); however, no detailed brain distribution studies have been reported (Fu et al. 2019b).

### 19.1.5 Conclusion and Perspectives

Despite considerable efforts, no radioligands have been confirmed as potential in vivo imaging agents for NMDARs by PET or SPECT imaging. Because NMDARs form complex systems with ligand-gated ion channels including two

agonist-binding sites, a transmembrane ion permeation pore, and various allosteric modulator binding sites, it may be difficult to apply the strategy of developing radioligands for other neurotransmitter receptors that can be successfully visualized by PET or SPECT imaging.

Initially developed radiolabeled dissociative anesthetic derivatives, which are non-competitive NMDAR open channel blockers, were shown to exhibit high non-specific in vivo accumulation probably due to high lipophilicity and/or lack of selectivity for NMDARs. Among these dissociative anesthetics, only MK-801 has been confirmed as an NMDAR specific ligand (Wong et al. 1986). Although several radiolabeled MK-801 derivatives showed in vitro brain localization consistent with NMDAR expression, these ligands exhibited high non-specific binding under in vivo conditions. It should be noted that [ $^3\text{H}$ ]MK-801 exhibited high specific binding (78%) in vivo if the brain tissue was homogenized, filtered, and washed. In addition, the specific binding was abolished by omission of the washing procedures (Price et al. 1988). These results indicated that although [ $^3\text{H}$ ]MK-801 has a certain in vivo specific binding component to NMDARs, the high adsorption property of this ligand on brain tissue may hamper in vivo imaging of NMDARs, probably owing to high lipophilicity of [ $^3\text{H}$ ]MK-801. Radiolabeled MK-801 derivatives with optimized lipophilicity and binding affinity for the PCP-binding site may be potential imaging agents for NMDARs. Radiolabeled diarylguanidine derivatives were developed as more hydrophilic imaging agents for the PCP-binding site of NMDARs. In vivo studies of ischemic rats demonstrated that pathological activation of NMDARs led to an increase in [ $^{125}\text{I}$ ]CNS 1261 binding (Owens et al. 2000). However, the possibility that this change was caused by change in cerebral blood flow and/or input function of these tracers could not be excluded. In vivo displacement effect of open channel blockers for [ $^{123/125}\text{I}$ ]CNS 1261 binding in the brain tissues has not yet been confirmed. In addition, rapid in vivo metabolism of [ $^{125}\text{I}$ ]CNS 1261 (half-life, 2.2 min in rat plasma) should be taken into consideration (Owens et al. 2000). Nevertheless, altered in vivo pharmacokinetics of [ $^{123}\text{I}$ ]CNS 1261 in patients with schizophrenia compared with healthy controls was noteworthy (Pilowsky et al. 2006). On the other hand, there are several prospective reports regarding the development of PET tracers for the ion channel site.  $^{18}\text{F}$ -GE-179 showed high affinity and selectivity for the ion channel site, although in vivo pharmacokinetic studies have not yet been optimized (Robins et al. 2010; McGinnity et al. 2014; Schoenberger et al. 2018). One limitation of this chemical backbone is the rapid metabolism which could not be ignored for specific PET imaging. Attempts to improve the metabolic stability, such as reports on the fluorovinylloxy analogs (Klein et al. 2017), could facilitate the development of promising PET tracers for this site. Because it is suggested that only 30% NMDARs are under open states (Jahr 1992), imaging PCP-binding site by such activity-dependent-binding radioligands would be difficult. Nonetheless, further studies on the development of new imaging probes for the PCP-binding site should be undertaken. Radioligands for this site could detect activated NMDARs in the living brain including various brain disorders. Indeed, PET signals of  $^{18}\text{F}$ -GE-179 increased in the disease states, such as focal epilepsies (McGinnity et al. 2015) and brain ischemia (Zhou et al. 2018).

Another possibility is that the poor *in vivo* results of radioligands for the PCP-binding site could be caused by their restricted *in vivo* access to the inside of the ion channel. Compared with the PCP-binding site, glycine- and ifenprodil-binding sites were reported to be located outside of the receptor. Accordingly, these two sites have been considered as attractive targets for visualization of NMDARs by PET or SPECT imaging.

Several radioligands for the glycine-binding site showed promising *in vitro* results. In contrast, *in vivo* results revealed quite low brain uptake and unusual cerebellar localization of these tracers. The difference between *in vitro* and *in vivo* studies could be explained by the influence by endogenous glycine site agonists. The *in vivo* binding of [<sup>11</sup>C]L-703,717 was strongly affected by endogenous d-serine concentration (Haradahira et al. 2003). On the other hand, in most of the *in vitro* experiments, brain slices or homogenates were well washed, which resulted in the removal of endogenous glycine and d-serine. Considering these results, we hypothesized that the agonist-insensitive radioligands with high affinity for glycine site radioligands could be useful for *in vivo* imaging of NMDARs. Fuchigami and Magata et al. developed high-affinity “agonist-insensitive” glycine site ligand [<sup>11</sup>C]**23**, but BBB passage was too low to obtain reliable *in vivo* results (Fuchigami et al. 2008). Further investigations should be carried out for the development of new imaging probes with high BBB penetration to allow for the binding mechanism of glycine antagonists to be identified.

Radioligands for the ifenprodil-binding site showed consistent localization with the expression of GluN2B and high specificity to GluN2B-rich region only under *in vitro* condition but not *in vivo*. It is suggested that ifenprodil could exhibit higher affinity for the agonist-bound activated or desensitized state of NMDARs relative to the agonist-unbound state (Kew et al. 1996). It is demonstrated that neuroprotective effects of GluN2B antagonists were much more potent at pH = 6.5 than at pH = 7.5 (Mott et al. 1998). In the ischemic tissue with NMDAR activated state, pH value was reported to be around 6.5 (Silver and Erecińska 1992). Therefore, it is possible that the radioligands for the ifenprodil-binding site could recognize only agonist-bound and/or activated NMDAR like PCP-binding site ligands. Nonetheless, new radioligands with optimized lipophilicity, metabolic stability, and binding affinity could show different pharmacokinetics from reported radioligands. Continuous efforts should be undertaken in order to develop promising *in vivo* imaging agents for the ifenprodil-binding site. Although benzimidazole derivative [<sup>125</sup>I]**33** showed inconsistent *in vivo* localization with GluN2B expression, this tracer could exhibit partial specific binding for GluN2B (Fuchigami et al. 2010). Further SAR studies of benzimidazoles could provide promising radioligands for the ifenprodil-binding site of NMDARs. Radiolabeled benzylamidines showed excellent high specific binding for the GluN2B under *in vitro* conditions (Arstad et al. 2006). Optimized benzylamidine derivatives with high metabolic stability and moderate lipophilicity could be attractive imaging probe candidates for the ifenprodil-binding site of NMDARs. Recently, new series of antagonists for the ifenprodil-binding site, such as MK-0657 and HON0001, were reported (Suetake-Koga et al. 2006; Addy et al. 2009). Indeed, [<sup>18</sup>F]-labeled MK-0657 derivatives showed promising *in vitro* results



(Koudih et al. 2012). More recently, there are several innovative reports on the development of PET tracers for imaging of GluN2B. Two enantiopure 3-benzazepine derivatives, (*R*)- $^{11}\text{C}$ -Me-NB1 and (*R*)- $^{18}\text{F}$ -OF-Me-NB1, were demonstrated to be promising radioligands for the specific imaging of GluN2B receptors as confirmed both by in vitro and in vivo studies (Haider et al. 2019). Further investigations are needed to confirm the feasibility of these compounds as clinically useful PET tracers.

In general, higher  $B_{\text{max}}/K_d$  of radioligands could give higher target-to-non-target (T/N) binding ratios. Eckelman et al. suggested that PET tracers should have  $B_{\text{max}}/K_d$  ratio greater than 10 to provide a useful specific in vivo signal (Eckelman et al. 1979, 2006).  $B_{\text{max}}$  values of the PCP-binding site in the dentate gyrus of the rat brain and CA1 stratum radiatum of the human brain using [ $^3\text{H}$ ]MK-801 were demonstrated to be 173 fmol/mg and 141 fmol/mg protein, respectively (Bowery et al. 1988). Accordingly,  $K_d$  values below 1.73 and 1.41 nM may be required for PCP-binding site ligands in the rat and human brain, respectively, to obtain a  $B_{\text{max}}/K_d$  ratio over 10. The  $B_{\text{max}}$  value of the glycine-binding site in rat brain membranes using [ $^3\text{H}$ ]MDL 105,519 was calculated to be 12.1 pmol/mg protein (Baron et al. 1996). Therefore, a  $K_d$  below 121 nM may be required for glycine-binding site ligands in the rat brain to obtain a  $B_{\text{max}}/K_d$  ratio over 10. The  $B_{\text{max}}$  value of the ifenprodil-binding site in rat brain membranes using [ $^3\text{H}$ ]Ro 25-6981 was estimated to be 1.6 pmol/mg protein (Mutel et al. 1998). Thus, a  $K_d$  below 16 nM may be required for glycine-binding site ligands in the rat brain to obtain a  $B_{\text{max}}/K_d$  ratio over 10. Although several radioligands for NMDARs meet the above conditions, numerous factors decrease the reliability of in vivo images such as the low plasma-free fraction of radioligands and Pgp efflux. It has been suggested that radiotracers with low molecular weight (MW) (<500 Da) and moderate lipophilicity ( $\log D_{7.4}$  values; 2.0–3.5) could have optimal BBB permeability (Waterhouse 2003; Pike 2009). Recently, a central nervous system PET multiparameter optimization (CNS PET MPO) algorithm has been developed for the prediction of the required physicochemical properties of the clinically useful CNS PET agents (Wager et al. 2010; Zhang et al. 2013). The researchers have used a set of six parameters, including ClogP (calculated logP), ClogD (calculated distribution coefficient at pH = 7.4), MW, TPSA (topological polar surface area), HBD (number of hydrogen bond donors), and pKa (ionization constant of the most basic center) as important factors for CNS PET agents. In their algorithm, desirable ranges were set as ClogP  $\leq 2.8$ , ClogD  $\leq 1.7$ , MW  $\leq 305.3$ ,  $44.8 < \text{TPSA} \leq 63.3$ , HBD  $\leq 1$ , and pKa  $\leq 7.2$ . The scores of each parameter can be calculated ranging from 0.0 to 1.0. The most successful CNS PET tracers in the previous database had CNS PET MPO scores of  $>3$  (Zhang et al. 2013). Therefore, it would be desirable that the newly developed radioligands fulfill such conditions. Furthermore, exhaustive binding assays of these ligands for various receptors and transporters expressed in the brain should be carried out to screen NMDAR-specific radioligands.

Another interpretation is that the discrepancy between the in vitro and in vivo brain localization of radioligands for NMDARs could be caused by dynamic plasticity of NMDARs. It has been suggested that considerable structural change of

NMDARs occurs following binding of both endogenous agonists and exogenous ligands (Traynelis et al. 2010). It is reported that the number of NMDARs on the cell surface is frequently changed by tyrosine phosphorylation and/or cysteine palmitoylation of C-terminus of NMDARs with an alteration in subunit composition (Vallano et al. 1996; Roche et al. 2001; Hayashi et al. 2009). Further investigations such as in vivo imaging studies of genetically modified or drug-treated animals with various conditions of NMDARs would be needed to clarify the discrepancy between the in vitro and in vivo experiments. In addition, novel screening systems other than traditional in vitro assays should be used. Using living brain slices as a screening method (Matsumura et al. 1995) may lead to the development of promising radioligands for NMDARs.

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# Progress in PET Imaging of the Norepinephrine Transporter System

# 20

Yu-Shin Ding

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## Abstract

The norepinephrine transporter (NET) has long been recognized to play a role in various neurological and psychiatric disorders, e.g., ADHD, substance abuse, depression, alcoholism, obesity, addiction, and Alzheimer's and Parkinson's diseases. However, many of the important findings resulting from studies in vitro

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using postmortem tissues have never been verified via in vivo methods due to the lack of suitable radioligands, preventing the brain imaging of NET in living systems. We have identified the superiority of (*S,S*)-[<sup>11</sup>C]MRB and the suitability of the MRB analogs as potential NET ligands for PET translational studies from preclinical investigation in animals to clinical research in humans. In this review article, progress in these translational research studies will be discussed, including the role of NET in ADHD, substance abuse, depression, post-traumatic stress disorder (PTSD), alcohol dependence, obesity, aging, Parkinson's disease, and Alzheimer's disease.

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## 20.1 Introduction

In a previous review article (Ding et al. 2006), we provided background and clinical relevance of the norepinephrine transporter (NET) to support the need to develop novel radiotracers to study the brain NET system in order to better understand its role in brain function and diseases. We also discussed the design and biological evaluation of several radioligands for imaging the brain NET system with PET. Based on these characterization studies (Ding et al. 2003, 2005; Lin et al. 2005), including C-11 labeled desipramine (DMI), 2-hydroxy-desipramine (HDMI), talopram, talsupram, nisoxetine (Nis), oxaprotiline (Oxap), and lortalamine, and C-11 and F-18 derivatives of reboxetine, methylreboxetine (MRB), and their individual (*R,R*) and (*S,S*) enantiomers, in conjunction with studies using radiolabeled 4-iodo-tomoxetine and 2-iodo-nisoxetine, we have identified the superiority of (*S,S*)-[<sup>11</sup>C]MRB and the suitability of the MRB analogs as potential NET ligands for PET translational studies from preclinical investigation in animals to clinical research in humans. In this review article, progress in this translational research will be discussed.

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## 20.2 Challenges in NET Imaging of the Brain

NET possesses several intrinsic disadvantages as a molecular imaging target as compared to the other monoamine transporters (dopamine transporter (DAT) and serotonin transporter). These disadvantages have presented drawbacks, described below, and hindered for decades of progress in ligand development and imaging of NET in the living human brain.

### 20.2.1 NET Has a Relatively Low Density in the Brain

B<sub>max</sub> for NET is ~5–150 fmol/mg tissue (Smith et al. 2006; Tejani-Butt 1992); however, DAT has a more than 50-fold higher density, and SERT is over five-tenfold higher (Kaufman et al. 1991; Zeng et al. 2006).

### **20.2.2 NET Has a Widespread Distribution in the Brain**

Studies in rodents, cats, non-human primates, and humans have indicated that noradrenergic terminals in the central nervous system (CNS) are widespread. Highest levels occur in the locus coeruleus complex (LC) and raphe nuclei, with moderate density in the hypothalamus, midline thalamic nuclei, bed nucleus of the stria terminalis, and central nucleus of the amygdala and brainstem nuclei, with lowest levels in basolateral amygdala and cortical and striatal regions (Ding et al. 2003, 2010; Tejani-Butt 1992; Charnay et al. 1995; Ding and Fowler 2005; Gallezot et al. 2007, 2010; Hannestad et al. 2010a).

### **20.2.3 NET Has a Lower Contrast Between NET-Poor and NET-Rich Regions**

Two detrimental effects on NET imaging as the result of the widespread distribution of NET in the brain are (1) lower contrast and (2) lack of an obvious reference region. In contrast, a significant difference in the density between the transporter-rich and transporter-poor regions for both DAT and SERT systems offers much better contrast for PET imaging of DAT/SERT and simplifies data analysis methods by using a reference region approach.

### **20.2.4 The Locus Coeruleus (LC), the Highest NET Density Region, Has a Small Structure That Makes It Difficult to Be Delineated and Quantitated**

We and other research groups have been making progress toward ligand development for PET imaging of NET. After over two decades of effort, (*S,S*)-[<sup>11</sup>C]MRB and its analogs are considered by far the most promising NET ligands for PET imaging (Ding et al. 2006). This ligand has been used to study the role of NET in humans. How best to carry out these translational research studies and how best to normalize data from the NET-PET studies to obtain a measure related to NET availability have been the subject of our recent work. Specific examples of tracer applications in the neurosciences (mainly in the brain) are given in the following sections.

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## **20.3 Translational PET Imaging Studies of NET**

The NET has been suggested to play a role in numerous disorders such as substance abuse and withdrawal, ADHD, depression, schizophrenia, Alzheimer's disease, Parkinson's Disease, as well as aging; however, in vivo brain imaging studies of NET have not been possible due to the lack of suitable radioligands (Ding et al. 2006). We have evaluated the suitability of (*S,S*)-[<sup>11</sup>C]MRB in preclinical models and further evaluated it in humans. The results show that (*S,S*)-[<sup>11</sup>C]MRB possesses

properties that are highly favorable for the imaging of brain NET as a PET tracer, including (a) a regional distribution consistent with the known distribution of NET in the brain; (b) high test/retest reproducibility in nonhuman primates and humans; (c) excellent specificity and selectivity for NET over other monoamine transporters, namely, DAT and SERT; and (d) the short half-life of its  $^{11}\text{C}$  label ( $t_{1/2} = 20$  min), enabling multiple assessments (e.g., baseline and drug intervention) in the same subject on the same day. So far, we have carried out over 300 scans using (*S,S*)-[ $^{11}\text{C}$ ]MRB, including preclinical and clinical studies, to investigate cocaine abuse, alcoholism, depression, post-traumatic stress disorder (PTSD), obesity, brown adipose tissue (brown fat), and diabetes-related studies (studies related to drug occupancy of several novel pharmaceuticals targeting the NET system during their Phase I clinical trials are not included). These studies have facilitated our better understanding of the role of NET in brain function and diseases. Furthermore, the use of PET with a suitable NET ligand to study drug occupancy and drug mechanisms will assist in the development of the next generation of effective drugs for the treatment of various NET-related disorders. Here are presented a few examples to illustrate the progress in PET imaging of the NET system using (*S,S*)-[ $^{11}\text{C}$ ]MRB (abbreviated henceforth as [ $^{11}\text{C}$ ]MRB).

### 20.3.1 NET Imaging in Substance Abuse

Cocaine inhibits neurotransmitter uptake of human monoamine transporters, including the NET, DAT, and SERT, with comparable potency. Alterations in NET concentration in the CNS following cocaine exposure have been identified in rats, nonhuman primate models, and postmortem human studies in cocaine users (Beveridge et al. 2005; Macey et al. 2003; Mash et al. 2005). Although these observations point to potentially important dysregulations in NET and NET-mediated neuroenergetics in response to chronic cocaine use (Weinshenker and Schroeder 2007), *in vivo* studies of NET regulation in clinical populations have yet to be explored. Thus, our aim was to use [ $^{11}\text{C}$ ]MRB to determine the alterations in brain NET concentration in chronic cocaine users (Ding et al. 2010).

#### 20.3.1.1 NET Abnormalities in Cocaine Dependence

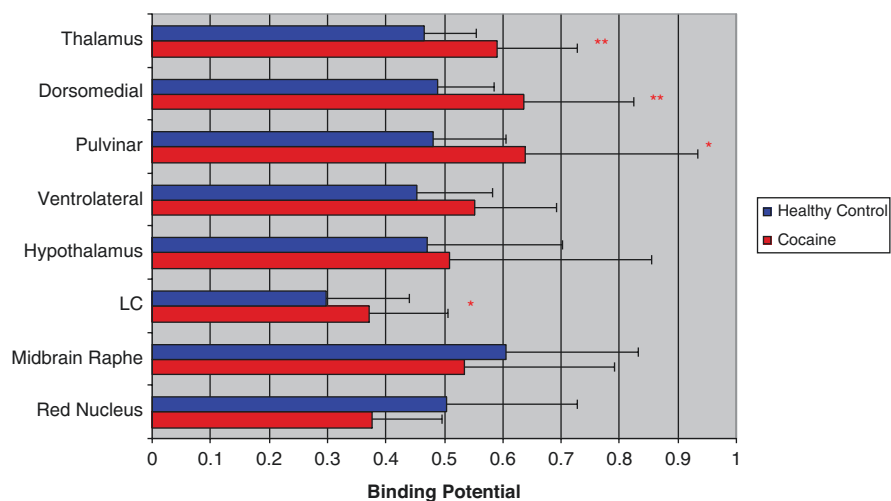
In a pilot study, 12 healthy control subjects (HC) (6 M and 6 F; range 25–54) and 10 cocaine-dependent individuals (COC) (7 M, 3F; range 39–49) were recruited (Ding et al. 2010). COC subjects were studied either as inpatients or outpatients, with cocaine abstinence at the time of PET scanning established by 24-h inpatient supervision and/or negative urine toxicology testing, respectively. All subjects participated in magnetic resonance imaging (structural MRI, 3 T) and PET scanning (high-resolution research tomograph, HRRT, 207 slices, resolution  $\sim 2.5$  mm FWHM in 3D acquisition mode) on two different days. Binding potential ( $\text{BP}_{\text{ND}}$ ) parametric images were computed using the simplified reference tissue model (SRTM2) with the occipital cortex as reference region.  $\text{BP}_{\text{ND}}$  values were compared between the two groups.

High-resolution imaging of brain NET using [ $^{11}\text{C}$ ]MRB and HRRT clearly delineated the fine-scale regional distribution of [ $^{11}\text{C}$ ]MRB in small brain regions known to have high NET concentrations, including the hypothalamus, red nucleus, midbrain raphe, pontine raphe, thalamus and its subnuclei (dorsomedial thalamic nucleus, ventrolateral thalamic nucleus, pulvinar), and the area of locus coeruleus (LC). The  $\text{BP}_{\text{ND}}$  was significantly increased in the thalamus (27%;  $P < 0.02$ ) and dorsomedial thalamic nuclei (30%;  $P < 0.03$ ) in COC as compared to HC. We also found that there was an age effect on NET concentrations in HC. Thus, group comparison in a covariance analysis to account for the differences in age between groups was also performed. Upon age normalization, the upregulation of NET in COC also reached significance in LC (63%,  $P < 0.01$ ) and pulvinar (55%,  $P < 0.02$ ) regions (Fig. 20.1).

### 20.3.1.2 Duration of Abstinence Effect on NET Binding

In a subsequent study, we looked into the duration of the effect of abstinence on NET binding. Regional brain NET binding potential ( $\text{BP}_{\text{ND}}$ , MRTM2) was measured with [ $^{11}\text{C}$ ]-MRB PET scanning in healthy controls ( $n = 12$ ,  $38 \pm 11$  years) and patients with chronic cocaine dependence (total  $n = 15$ ,  $43 \pm 6$  years, with  $22 \pm 7$  years of cocaine use) after 1 week ( $n = 4$ ), 2 week ( $n = 6$ ), 3 to 5 weeks ( $n = 5$ ) of abstinence.

Our results served to reproduce our previous finding that NET is upregulated in chronic cocaine users after recent abstinence, as compared with controls. In addition, we found that subjects with cocaine dependence had higher NET  $\text{BP}_{\text{ND}}$  values in the ventral lateral thalamus (70%;  $p < 0.05$ ) after 1 week of abstinence; that level



\*\* Reached statistically significant

\* Reached statistically significant upon age-normalization

**Fig. 20.1** Effect of cocaine dependence on [ $^{11}\text{C}$ ]MRB binding in humans.  $\text{BP}_{\text{ND}}$  values (binding potential) were compared between healthy control and cocaine-dependent individuals



remained elevated (85%;  $p < 0.01$ ) during the second week of abstinence. By the third week, NET levels slowly declined (11%;  $p = 0.60$ ) (Li et al. 2006). Similar significant patterns were observed in the measurement of the whole thalamus and the midbrain raphe nuclei.

The observations of an increased binding of [ $^{11}\text{C}$ ]MRB in our in vivo PET imaging studies of chronic COC subjects is consistent with previously published in vitro data in rodents, rhesus monkeys, and postmortem human brains (Beveridge et al. 2005; Macey et al. 2003; Mash et al. 2005). Our results are also in agreement with an in vitro study in which an upregulation of NET mRNA in the LC was observed in rats following chronic binge administration of cocaine (Burchett and Bannon 1997). Interestingly, in this rat study, an alteration was observed in an opposite direction for the other two monoamine transporters, i.e., a decrease in SERT mRNA (in raphe) and DAT mRNA (in midbrain) following a binge regimen of chronic COC.

### 20.3.2 NET Imaging in ADHD

Attention deficit hyperactivity disorder (ADHD), a complex developmental behavioral and cognitive disorder that affects approximately 3–7% of school-aged boys and 2–4% of adults, is associated with impaired academic and occupational achievements, problem behaviors, increased risk for accidents, and substance abuse. (Dopheide and Pliszka 2009; Barkley 1997) The core features of ADHD—inattention, hyperactivity, and impulsivity—can be attributed to dysfunction in neural systems that regulate attention, executive function, motor control, and reward (Bush 2010; Makris et al. 2009); however, effective treatments remain elusive, and the neurochemical mechanisms of this disorder are poorly understood (Bush 2010; Makris et al. 2009). The three therapeutic classes of pharmacologic treatments for ADHD—(a) psychostimulants (e.g., methylphenidate hydrochloride; MPH); (b) non-stimulant catecholamine reuptake inhibitors (e.g., atomoxetine, ATX); and (c)  $\alpha 2$  adrenergic receptor agonists (e.g., guanfacine)—all modulate DA and NE neurotransmission (Makris et al. 2009; Madras et al. 2005). Small changes in extracellular NE or DA concentration affect networks of pyramidal cells in the prefrontal cortex (PFC) which regulates and sustains attention. The beneficial effects of DA occur at D1 receptors, while those of NE are believed to occur at  $\alpha 2$  receptors (Arnsten 2009). Here we present several of our translational research studies of drug occupancy designed directly toward developing a better understanding of the mechanisms that contribute to the therapeutic effects of pharmacologic treatments for ADHD.

#### 20.3.2.1 ATX Occupancy Study on NET in Humans and Nonhuman Primates

Atomoxetine (ATX, Strattera), a potent NET reuptake inhibitor ( $K_i \sim 5$  nM), is the first non-stimulant drug that was approved for the treatment of ADHD (Arnsten 2006; Bymaster et al. 2002; Chamberlain et al. 2007a; Del Campo et al. 2011; Michelson et al. 2001; Spencer et al. 2002; Swanson and Volkow 2009; Volkow

et al. 2001). When [ $^{11}\text{C}$ ]MRB was initially tested in a drug occupancy study in humans using atomoxetine (ATX), a reduction in [ $^{11}\text{C}$ ]MRB binding was observed, but there was no obvious dose-dependent effect (Logan et al. 2007). One hypothesis was that the doses of ATX examined in the human PET study (oral, 25, 50 or 100 mg), though clinically relevant, were too high. That is, the three tested doses all reached saturation, the plateau portion of the corresponding dose-occupancy curve, and as a result there was no obvious dose-dependent effect. Another occupancy study of ATX using [ $^{18}\text{F}$ ]F-MeNER-D2 in rhesus monkeys showed a reasonable dose-dependent occupancy, though only in one of the two monkeys studied and the  $\text{IC}_{50}$  (or  $\text{ED}_{50}$ ) was not determined (Seneca et al. 2006).

Thus, the suitability of [ $^{11}\text{C}$ ]MRB for occupancy studies was further evaluated by carrying out investigations in a nonhuman primate model (rhesus macaque) using lower doses of ATX than those used in the previous PET study with [ $^{11}\text{C}$ ]MRB (Logan et al. 2007) and with a continuous ATX infusion paradigm to maintain a constant concentration of ATX in the plasma during the PET acquisition time (120 min). Ten PET studies were performed in three anesthetized rhesus monkeys following an infusion of ATX or placebo.

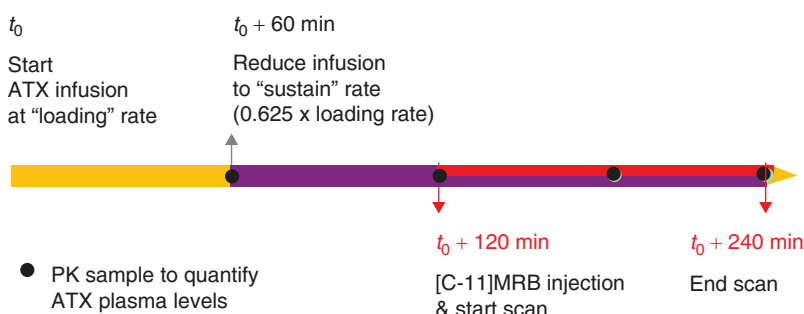
For each PET study, two infusion rates were used: ATX infusion at a “loading” rate was first given for 1 h starting at 2 h before the injection of [ $^{11}\text{C}$ ]MRB; then ATX was given at a “maintenance” rate (37.5% lower than the “loading” rate) at 1 h before the injection of [ $^{11}\text{C}$ ]MRB. This infusion rate was maintained until the end of the 2-h PET scan (i.e., 3 h at “maintenance” rate) (Fig. 20.2). Three doses of “maintenance” rate were used: 0.01 mg/kg/h (low dose), 0/03 mg/kg/h (medium dose), and 0.09 mg/kg/h (high dose) mg/kg/h (i.e., each monkey had four PET scans, including the baseline scan with placebo). The design was to have each monkey have four PET scans on 2 study days. These 2 study days were scheduled at least

### Study Design:

**Each monkey was scanned four times, twice a day**

**Day 1: baseline and medium dose scans**

**Day 2: low and high dose scans**

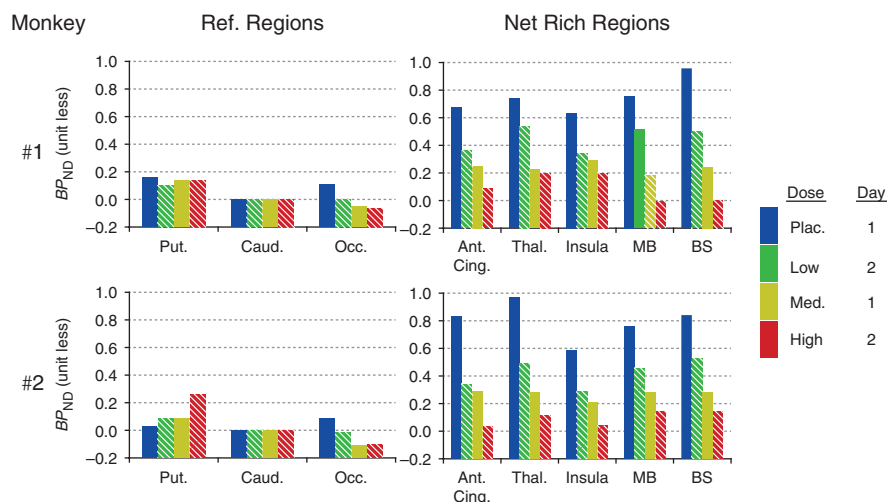


**Fig. 20.2** Study design for the two-day, four-dose studies to measure ATX occupancy of NET in non-human primates using [ $^{11}\text{C}$ ]MRB

3 weeks apart to allow recovery from the effect of the anesthesia and blood drawing. On each study day, two PET scans were acquired, beginning 4 h apart. The first study day consisted of the baseline and medium-dose scans, and the second study day consisted of the low- and high-dose scans.

A dose-dependent reduction of [ $^{11}\text{C}$ ]MRB volume of distribution was observed after correction for [ $^{11}\text{C}$ ]MRB plasma free fraction. Although the slow clearance of [ $^{11}\text{C}$ ]MRB in NET-rich regions such as the thalamus posed a concern for kinetic modeling, the combination of the suitably designed infusion study paradigm and refined kinetic modeling methods an unequivocal dose-dependent occupancy of atomoxetine was obtained (Fig. 20.3) clearly demonstrating the suitability of [ $^{11}\text{C}$ ]MRB for drug occupancy studies of the NET (Gallezot et al. 2010).

Based on this study, ATX  $\text{IC}_{50}$  was estimated to be  $32 \pm 13$  ng/mL plasma, corresponding to an effective dose ( $\text{ED}_{50}$ ) of 0.13 mg/kg, which is much lower than the therapeutic dose of ATX in ADHD (1.0–1.5 mg/kg). In the previous ATX occupancy study in humans, three clinically relevant doses were used (oral, 25, 50, and 100 mg), and the blood levels of ATX (in ng/mL plasma) averaged over subjects were 116 (25 mg ATX), 206 (50 mg ATX), and 421 (100 mg ATX), which was significantly higher than the estimated  $\text{IC}_{50}$  for ATX in plasma ( $\sim 30$  ng/mL). This suggests that the lack of an obvious dose-dependent effect in the previous study was indeed because the three clinically relevant doses all reached saturation with over 78–93% occupancy, i.e., the plateau portion of the corresponding dose-occupancy curve (Logan et al. 2007). Taken together, our results clearly indicate the importance of dose selection for PET occupancy studies, especially when observation of dose-dependency of the study drug is one of the goals for the outcome measurement.



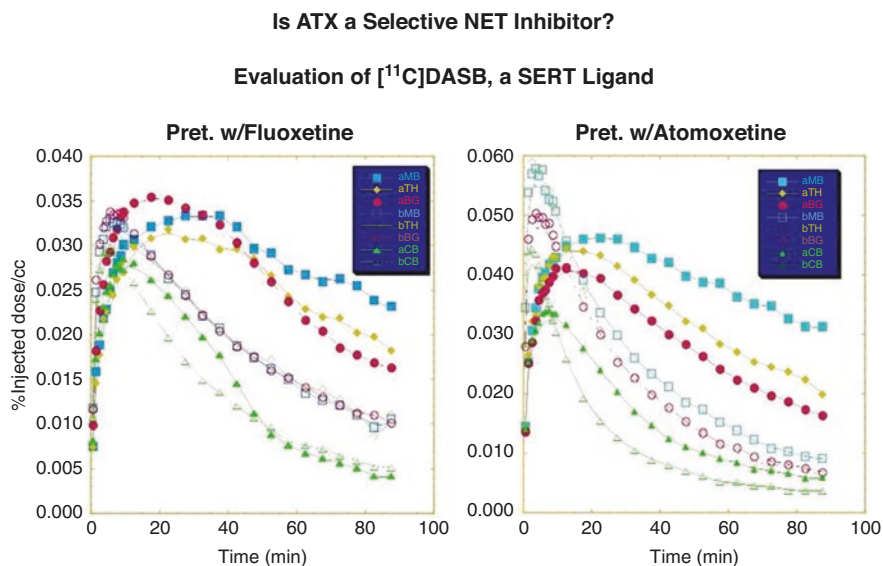
**Fig. 20.3** Dose-dependent occupancy of ATX in NET-rich regions in two monkeys using [ $^{11}\text{C}$ ]MRB after the placebo (Plac.), low, medium, and high ATX doses.  $\text{BP}_{\text{ND}}$  values were measured for NET-poor regions [putamen (put.) and occipital cortex (Occ.)] and NET-rich regions [anterior cingulate (Ant. Cing.), thalamus (Thal.), Insular, mid brain (MB) and brain stem (BS)], using caudate as a reference region

### 20.3.2.2 ATX Occupancy Study on SERT: Implications for Treatment of Depression and ADHD

The development of ATX and its approval for the treatment of ADHD by the Food and Drug Administration (Faraone et al. 2005) was based on the rationale that it selectively blocks the NET (Barton 2005) and increases the availability of intrasynaptic NE, resulting in the improvement of high-level cognitive functions that are often impaired in ADHD, such as working memory and inhibitory response control (Robbins and Arnsten 2009). In principle, ATX may offer clinical advantages by virtue of its limited effects on subcortical dopamine, which is thought to be responsible for the abuse potential of psychostimulant treatment (Volkow 2006). Furthermore, this single NE pathway of drug mechanism may facilitate further drug research and development.

The noradrenergic action also exerts important clinical effects in different antidepressants such as desipramine and nortriptyline (tricyclics with prevalent noradrenergic effects). Thus, ATX has also been used to improve responses to treatment of depression (Dell'Osso et al. 2010; O'Sullivan et al. 2009).

Serendipitously, we discovered that ATX displayed a blocking effect similar to fluoxetine (a selective SERT inhibitor) (Ding and Fowler 2005) on binding of [ $^{11}\text{C}$ ]DASB, a selective SERT ligand (Houle et al. 2000) (Fig. 20.4). These results suggested that ATX not only binds to NET but also binds to SERT with high affinity. To date, whether the therapeutic effects of ATX in the treatment of either ADHD or depression are due to inhibitory effects on one (NET) or two transporters (NET and SERT) is not known. Therefore, we took on this investigation by carrying out comparative occupancy studies with [ $^{11}\text{C}$ ]MRB (a NET ligand, as described above



**Fig. 20.4** ATX displayed a blocking effect similar to fluoxetine (a selective SERT inhibitor) on binding of [ $^{11}\text{C}$ ]DASB (a selective SERT ligand), questioning the *in vivo* selectivity of ATX

(Gallezot, 2007 #11;Gallezot, 2010 #12) and [ $^{11}\text{C}$ ]AFM (a SERT ligand) (Huang et al. 2002) to evaluate the in vivo  $\text{IC}_{50}$  of ATX for both NET and SERT in nonhuman primates using PET.

The continuous infusion protocol with [ $^{11}\text{C}$ ]AFM was similar to our previous study with [ $^{11}\text{C}$ ]MRB (see above) (Gallezot et al. 2007, 2010). Two monkeys were scanned at different maintenance rates: 0.045 mg/kg/h (low dose), 0.135 mg/kg/h (medium dose), and 0.405 mg/kg/h (high dose). One monkey was scanned with the higher maintenance rates: 0.135 mg/kg/h (medium dose), 0.405 mg/kg/h (high dose), and 1.054 mg/kg/h (very high dose) (details of this study will be published elsewhere).

### **20.3.2.3 Summary of the Comparative Occupancy Studies of ATX for NET (Using [ $^{11}\text{C}$ ]MRB) and SERT (Using [ $^{11}\text{C}$ ]AFM)**

ATX displayed dose-dependent occupancy of both NET and SERT, with a higher occupancy of NET:  $\text{IC}_{50}$  of  $31 \pm 10$  and  $99 \pm 21$  ng/mL plasma for NET and SERT, respectively. At a clinically relevant dose (1.8 mg/kg,  $\sim 600$  ng/mL plasma), ATX would occupy  $>95\%$  of NET and occupy  $>85\%$  of SERT. It is worth noting that this value of 3.2 for the selectivity ratio of ATX toward NET vs. SERT measured via in vivo PET imaging in non-human primates (NHP) is similar to a value of 4.5 for the selectivity ratio derived from the previously reported in vitro  $K_d$  values using human NET and SERT tissue homogenates (8.9 and 2 nM for SERT and NET, respectively) (Tatsumi et al. 1997). Based on these data, we propose that ATX at clinically relevant doses occupies high proportions of both NET and SERT. Thus, the interpretation of the therapeutic mode of action of ATX for treatment of depression and ADHD may be more complex than selective blockade of the NET.

### **20.3.2.4 The Mechanisms of Action of ATX in ADHD and Depression**

It has been shown that response inhibition (the ability to exert high-level inhibitory control over motor responses so as to suppress unwanted actions) is impaired in subjects with ADHD and is one of the most robust cognitive findings in ADHD research (Lijffijt et al. 2005). Response inhibition (a.k.a. impulsivity) can be measured by the stop-signal task (SST) and quantified by the stop-signal reaction time (SSRT), an estimate of the time taken to inhibit the prepotent motor response (Aron et al. 2003a). The research group at Cambridge, UK, found that single doses of atomoxetine (60 mg) were able to improve response inhibition in humans and in rats (Chamberlain et al. 2006, 2007a; Bari et al. 2009; Eagle et al. 2008). By contrast, serotonin manipulations with citalopram, a SERT blocker (Del Campo et al. 2011; Chamberlain et al. 2006), or with buspirone, a serotonin 5-HT $_{1A}$  receptor agonist (Chamberlain et al. 2006, 2007b), appear to have no behavioral effects on the SSRT. These results suggest that atomoxetine enhances stopping selectively via actions on NE uptake and that NET drugs may be more suited to ameliorating impaired response inhibition as an ADHD therapeutic target.

Taken together, the inhibitory effect of ATX on NET plays a crucial role in its therapeutic effect on ADHD, while the inhibitory effect of ATX on SERT may play a more important role in its therapeutic effect on depression. Further studies to

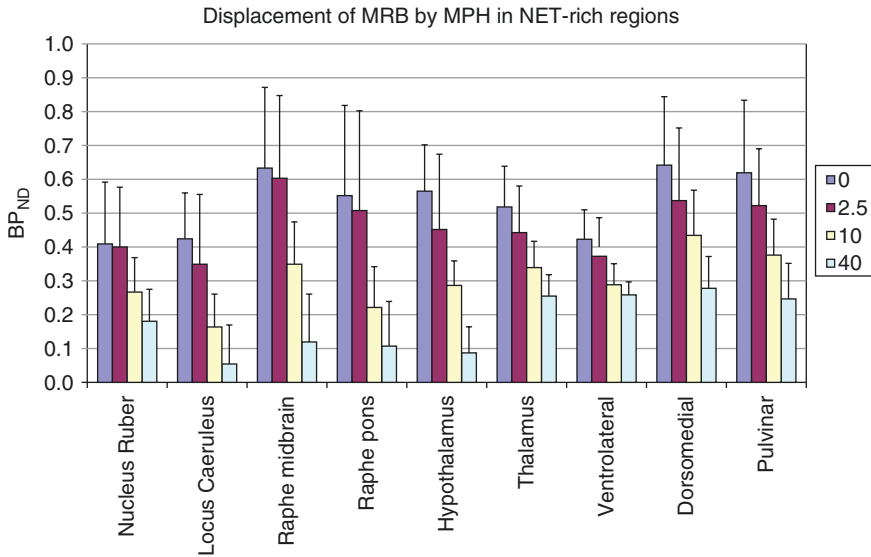
elucidate the actual relationship between the drug mechanisms and drug therapeutic effects are needed.

### 20.3.2.5 MPH Occupancy Study on NET in Humans

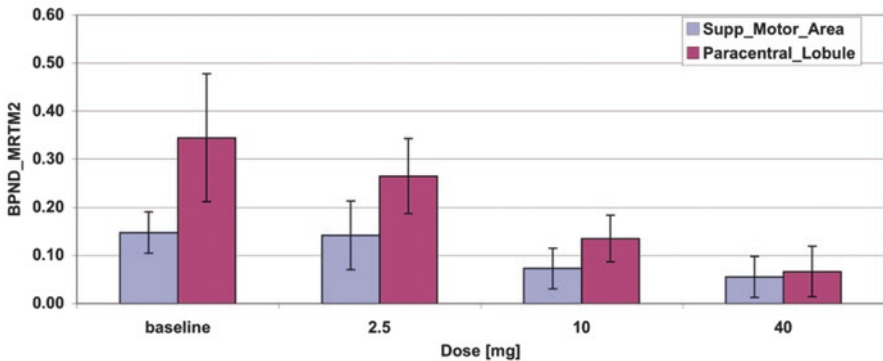
The effects of MPH on the DA system have been studied extensively. Our previous PET studies have demonstrated that MPH's pharmacological effects reside in *d-threo*-MPH, the active enantiomer (Ding et al. 1997), and that at clinical doses, MP significantly occupies DAT in vivo in humans (Swanson and Volkow 2009; Volkow et al. 1998), suggesting that DA reuptake inhibition may be an important therapeutic mechanism in ADHD. However, MPH has shown higher in vitro affinity for NET than DAT ( $K_i = 38$  nM vs. 193 nM) (Eshleman et al. 1999). The potential role of NE in ADHD was supported by animal and human data (Biederman et al. 2008). Low doses of MPH increase levels of NE in the PFC in animals (Berridge et al. 2006), and low doses of MPH improve performance of PFC tasks in subjects with ADHD (Aron et al. 2003b). The absence of a suitable NET tracer has delayed similar in vivo PET studies of NET until now.

We used [ $^{11}\text{C}$ ]MRB to determine the effective dose 50 ( $\text{ED}_{50}$ ) of MPH for NET (Hannestad et al. 2010b). We first conducted a preliminary duration study to determine how much time it takes from oral MPH administration to maximum occupancy of the brain NET. As peak displacement of [ $^{11}\text{C}$ ]MRB by MPH (oral 40 mg, i.e., 0.35–0.55 mg/kg, the average clinical maintenance dose of MPH) was achieved by 75 min, we chose 75 min as the time between oral MPH dosing and [ $^{11}\text{C}$ ]MRB injection for the subsequent occupancy study. Based on our previous experience with ATX occupancy studies in humans, dose selection is critical if dose-dependent occupancy is one of the goals to be achieved. Thus, in this within-subject design study, healthy subjects ( $n = 11$ ) received oral, single-blind placebo and 2.5 mg, 10 mg, and 40 mg of MPH 75 min prior to [ $^{11}\text{C}$ ]MRB injection. Dynamic PET imaging was performed for 2 h with HRRT. Three regions with known lowest density of NET (occipital cortex, caudate and putamen) were investigated (Logan 2005) further as reference regions by comparing the distribution volume ( $V_T$ ) in these three regions between the placebo and 40 mg MPH conditions (Hannestad et al. 2010b). We found no significant change of  $V_T$  (or  $V_T/f_p$ , where  $f_p$  = plasma free fraction) in these three regions between the placebo and 40 mg MPH conditions, and the putamen and occipital had similar  $V_T$  values. Thus, the multilinear reference tissue model with occipital cortex as the reference region was used to estimate binding potential ( $\text{BP}_{\text{ND}}$ ) in the thalamus and other NET-rich regions.

We found that  $\text{BP}_{\text{ND}}$  was reduced by MPH in a dose-dependent manner in thalamus and other NET-rich regions (Fig. 20.5). The global  $\text{ED}_{50}$  was estimated to be 0.14 mg/kg; therefore the average clinical maintenance dose of MPH (0.35–0.55 mg/kg) produces 70–80% occupancy of NET. This is the first in vivo study in humans showing that clinically relevant doses of MPH occupy significant levels of NET, despite the fact that MPH has been used for the treatment of ADHD for decades. The  $\text{ED}_{50}$  is lower than that for DAT (0.25 mg/kg), suggesting the potential relevance of NET inhibition in the therapeutic effects of MPH in ADHD.



**Fig. 20.5** BP<sub>ND</sub> values were reduced by methylphenidate in a dose-dependent manner in NET-rich regions after four doses of oral methylphenidate (0, 2.5, 10, 40 mg)



**Fig. 20.6** Dose-dependent displacement of [<sup>11</sup>C]MRB by methylphenidate in two cortical regions (paracentral lobule and the supplementary motor area (SMA)) were also detected after four doses of oral methylphenidate (0, 2.5, 10, 40 mg)

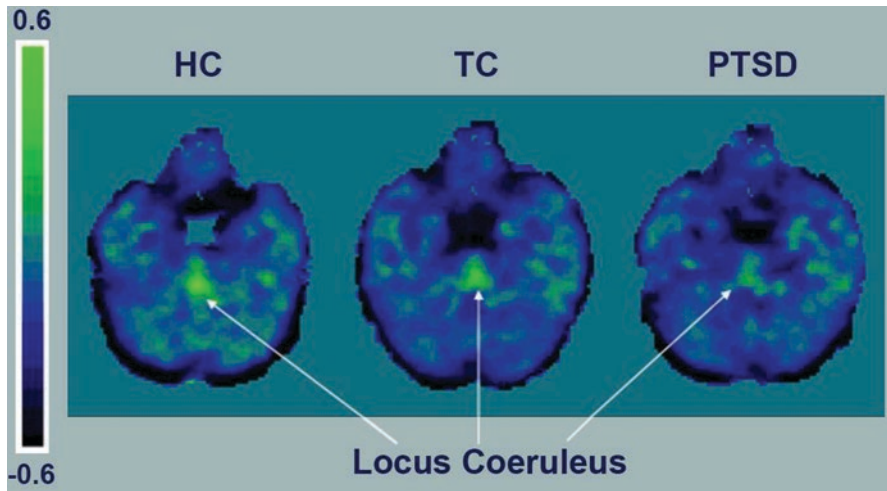
Previous studies also suggested that effects on delayed responding and working memory are mediated by noradrenergic afferents from LC (the highest NET density region) to PFC (Biederman et al. 2008; Solanto 1998). However, limited studies were able to link brain processes with attention and ADHD. It has also been shown that the temporal, parietal, and prefrontal (PFC) association cortices mediate different aspects of attention and that treatments for ADHD enhance catecholamine actions in PFC and improve PFC functions (Arnsten 2009). Although the density of NET in PFC is very low, we were able to detect dose-dependent displacement of [<sup>11</sup>C]MRB by MP in two cortical regions in which NET densities were high enough: paracentral lobule and the supplementary motor area (SMA) (Fig. 20.6), which are

areas involved in the response to MPH and inhibition control during cognitive tasks, respectively (Li et al. 2006; Hannestad et al. 2010b; Volkow et al. 2008). Although we cannot assess the effect of MPH in the PFC directly, our assumption is that the global  $ED_{50}$  calculated using NET-rich regions, including the neocortical paracentral lobule, is similar to the  $ED_{50}$  in regions with lower NET density, such as the PFC. It is also important to fill the gap by combining PET (offers specificity) and other modality such as fMRI (offers spatial and temporal resolution) to explore the underlying molecular processes.

### 20.3.3 NET Imaging in PTSD

Termination of the action of NE on adrenergic receptors is accomplished primarily by NE reuptake into the presynaptic terminal via the NE transporters (NET). Animal studies and postmortem studies in humans have implicated reductions in NET expression in depression, which is often comorbid with PTSD. While acute stress does not alter NET density, there is evidence from rodent studies that repeated stress decreases NET density in the locus coeruleus (LC) and limbic brain regions (Zafar et al. 1997). These important findings, however, have not yet been confirmed in humans using *in vivo* methods because brain imaging of NET in living humans has been hampered for over 20 years by the lack of suitable radioligands. Given that the NET is a suitable target for antidepressants and the established role of noradrenergic mechanisms in PTSD, additional research on the distribution and function of NET in the brain of patients with PTSD is important to better understand the hypothesized dysregulations in NE function in PTSD. Thus, the hypothesis that lower NET density may be related to the development of mood and anxiety disorders (Arnsten and Li 2005; Liprando et al. 2004; Miner et al. 2006; Moron et al. 2002) was investigated using the most promising NET ligand, [ $^{11}C$ ]MRB. In order to evaluate whether trauma exposure and PTSD would be associated with lower NET density in the locus coeruleus (LC), we conducted PET studies using [ $^{11}C$ ]MRB in three groups of individuals: 18 healthy adults (healthy controls: HC); 16 adults exposed to trauma who did not develop PTSD (trauma control: TC), and 22 adults exposed to trauma who developed PTSD. As a secondary aim, we evaluated the relation between NET availability in LC and a contemporary phenotypic model of PTSD symptomatology, namely, a five-factor model comprised of reexperiencing, avoidance, numbing, dysphoric arousal, and anxious arousal (Armour et al. 2012; Elhai et al. 2011; Pietrzak et al. 2012). Our study results revealed that both the PTSD group and the TC group had significantly lower NET density than the HC group (41% lower with Cohen's  $d = 1.04$  for the PTSD group, and 31% lower with Cohen's  $d = 0.77$  for the TC group, respectively), while NET density was not significantly different between the TC and PTSD groups (15% lower in PTSD versus TC group; Cohen's  $d = 0.27$ ). In the PTSD group, NET density in the LC was independently positively associated with severity of anxious arousal (i.e., hypervigilance symptoms,  $r = 0.52$ ), but not





**Fig. 20.7** Three representative participants were chosen with binding potential (BP) values similar to the mean of each study group. PET images are in Montreal Neurological Institute space. HC indicates healthy adults; PTSD, adults exposed to trauma who developed posttraumatic stress disorder; and TC, adults exposed to trauma who did not develop PTSD

with any of the other symptom clusters. Results of this study suggest that TC and PTSD are associated with significant reductions in NET availability in LC and that NET density in this brain region is associated with increased severity of anxious arousal symptoms among individuals with PTSD (Pietrzak et al. 2013) (Fig. 20.7).

#### 20.3.4 NET Imaging in Alcohol Dependence

We investigated the role of NET in alcohol dependence in human subjects, *in vivo*, using [ $^{11}\text{C}$ ]MRB and PET. Unmedicated, recently abstinent (4–6 weeks abstinent to compensate for acute alcohol withdrawal effects) nonobese men and women meeting DSM-IV (Pincus et al. 1996) [72] criteria for current alcohol dependence ( $n = 15$ ) and age- and sex-matched healthy control subjects ( $n = 15$ ) were studied and compared. Our hypothesis was that in areas of highest NET density, we would detect lower ligand binding commensurate with NET downregulation due to chronic alcohol exposure. We identified two brain regions in alcoholics, thalamus and locus coeruleus, that exhibited reduced NET binding potential ( $\text{BP}_{\text{ND}}$ ). We have postulated that these regions are involved in alcohol craving, acquisition, and ingestion and that they may serve as biomarkers of altered NET expression in alcohol dependence disorders and as potential therapeutic targets for drug development (unpublished results; details of this study will be published elsewhere).

### 20.3.5 NET Imaging in Brown Adipose Tissue (Brown Fat)

We have conducted studies in humans to investigate the role of NET in obesity. The preliminary data suggest that there is lower NET binding in the thalamus (including pulvinar) in obese subjects (body mass index (BMI) of  $34.7 \pm 2.6$ ) as compared to matched lean subjects (BMI of  $23.1 \pm 1.4$ ) (unpublished results, details of this study will be published elsewhere).

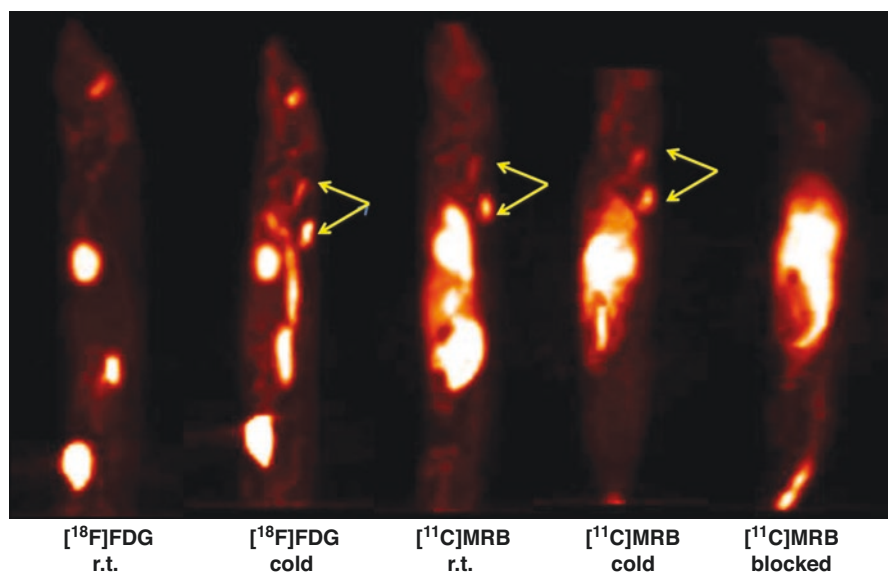
Obesity is characterized by impaired energy expenditure associated with adaptive thermogenesis (Camastra et al. 1999; Golay et al. 1986; Jequier and Schutz 1985; Yeckel et al. 2009). Brown adipose tissue (BAT, brown fat), one of the primary tissues responsible for adaptive non-shivering thermogenesis in mammals (Wijers et al. 2009), was previously thought to disappear in humans as part of development into adulthood. The recent confirmation that BAT still exists in adult humans provides a renewed interest in this tissue; however, the relative importance of BAT activity to adult human whole body energy expenditure is still unknown. Lessons learned from steps taken to reduce undesirable FDG-labeling during many false-positive clinical scans (Christensen et al. 2006; Hany et al. 2002; Nedergaard et al. 2007) demonstrate that repeat scans after beta-adrenergic blockade eliminate the interfering BAT signal (Agrawal et al. 2009). These data suggest the presence of strong sympathetic nervous system (SNS) activation of BAT in humans and the potential involvement of noradrenergic system in the regulation of BAT activity. Thus, in addition to a NET obesity study in the brain, we also carried out a pilot study to look into the NET involvement in both central and peripheral SNS function and to determine whether BAT can be visualized with NET-PET imaging and whether NET binding in BAT can be quantified under both basal and activated conditions in humans.

Based on [ $^{18}\text{F}$ ]FDG-PET imaging data, [ $^{18}\text{F}$ ]FDG-imaging requires subjects under cold stimulation during the PET imaging, and BAT does not appear to be visualized with [ $^{18}\text{F}$ ]FDG-PET at room temperature (Saito et al. 2009; van Marken Lichtenbelt et al. 2009). The BAT metabolic activity is related to age, gender, BMI, cold exposure, and medications (Cypess et al. 2009). Thus, the [ $^{18}\text{F}$ ]FDG-PET approach may not be suitable for imaging BAT at the basal state, i.e., room temperature. Since BAT activation appears to be dynamic, expressing large variability due to thermal sensitivity, individual differences in BAT activation due to variations in thermal response are likely to make comparisons between obese and normal-weight subjects difficult. Moreover, [ $^{18}\text{F}$ ]FDG is not specific to BAT, nor is glucose the primary substrate for BAT heat production. The [ $^{18}\text{F}$ ]FDG-PET strategy will not address regulatory mechanisms that are related to the specific SNS modulation features of BAT, which are likely to be key features to understanding potential dysfunction. Therefore, it is important to develop new and more specific methods to address both the plasticity of BAT activation and basal characteristics that do not require stimulation or that could alternatively be used in combination with strategies that highlight BAT stimulated activity.

### 20.3.5.1 Novel Strategy to Imaging BAT in the Basal State Using NET- $^{11}\text{C}$ MRB: Preclinical Evaluation

We first conducted both ex vivo and in vivo evaluation in rats using  $^{11}\text{C}$ MRB to determine BAT imaging in the basal state of BAT (at room temperature; r.t.), which is not achievable with  $^{18}\text{F}$ FDG (Lin et al. 2012). In the ex vivo study, awake male Sprague-Dawley rats were administered intravenous  $^{18}\text{F}$ FDG or  $^{11}\text{C}$ MRB after exposure to cold ( $4\text{ }^{\circ}\text{C}$  for 4 h,  $n = 9$ ) or remaining at r.t. ( $n = 9$ ) and were sacrificed at 20, 40, and 60 min postinjection. The results demonstrate that the uptake of  $^{11}\text{C}$ MRB in BAT was three times higher than  $^{18}\text{F}$ FDG at r.t. ( $P = 0.0088$ ), and the cold exposure increased  $^{18}\text{F}$ FDG uptake in BAT tenfold compared to the r.t. control ( $1.6 \pm 0.3$  vs.  $0.2 \pm 0.05\% \text{ID/g}$ ,  $P = 0.0009$ ), while no significant thermal effect was observed with  $^{11}\text{C}$ MRB BAT uptake ( $0.87 \pm 0.18$  vs.  $0.63 \pm 0.09\% \text{ID/g}$ ,  $P = 0.082$ ). In addition to the brain and heart, BAT exhibited specific  $^{11}\text{C}$ MRB uptake that was significantly reduced to near baseline levels ( $P = 0.0013$ ) by pre-treatment with unlabeled MRB or nisoxetine. These results were concordant with the in vivo PET imaging of anesthetized rats, which clearly demonstrated intense  $^{11}\text{C}$ MRB uptake in the interscapular BAT of both r.t. and cold-exposed rats; in contrast,  $^{18}\text{F}$ FDG in BAT was only detected in rats treated with cold (Fig. 20.8).

HPLC analysis revealed that 94–99% of total radioactivity in BAT represented unchanged  $^{11}\text{C}$ MRB, which further supports the NET-PET strategy for imaging BAT in humans at basal conditions. This NET-PET imaging strategy for BAT will



**Fig. 20.8** PET images (50–70 min post-injection) on isoflurane-anesthetized rats from left to right:  $^{18}\text{F}$ FDG room temperature control,  $^{18}\text{F}$ FDG cold-exposed rat,  $^{11}\text{C}$ MRB room temperature control,  $^{11}\text{C}$ MRB cold-exposed rat, and  $^{11}\text{C}$ MRB following blockade by unlabeled MRB. Brown adipose tissue (BAT) is denoted by arrows

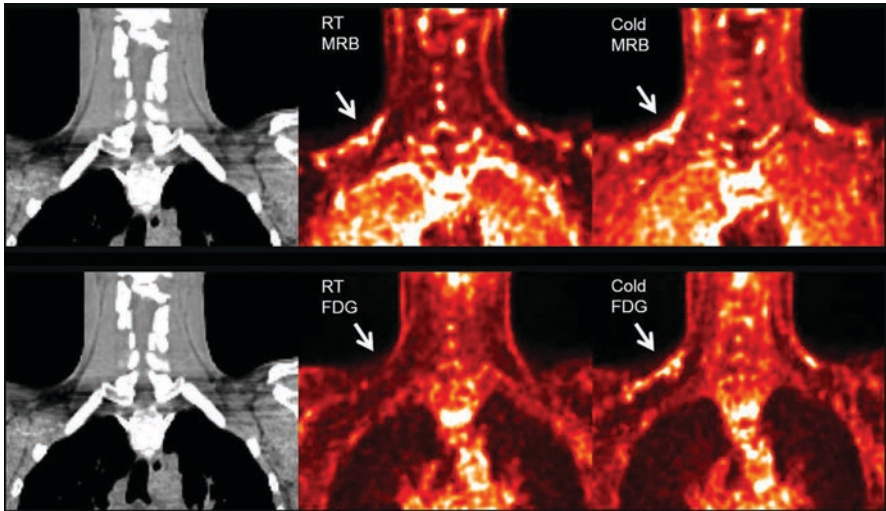
help to provide insight into whether BAT is a pivotal target tissue for therapeutic interventions designed to reduce human obesity and will also help to elucidate mechanisms related to its function/dysfunction. Our choice of the [ $^{11}\text{C}$ ]MRB ligand also has other specific advantages. Studies have shown that disruption of NET function directly impacts both autonomic function and mental health (Esler et al. 1991; Ganguly et al. 1986; Haenisch et al. 2009; Hahn et al. 2008, 2005, 2003, 2009) and that NET function is regulated both centrally and peripherally by insulin (Robertson et al. 2010). Unlike many other NET ligands, [ $^{11}\text{C}$ ]MRB is able to cross the blood-brain barrier, allowing for simultaneous central and peripheral imaging. Thus, using [ $^{11}\text{C}$ ]MRB may help to further elucidate mechanisms of BAT action and the role of NET in energy balance of obesity and insulin resistance by simultaneously correlating the CNS and peripheral SNS functions.

### 20.3.5.2 Novel Strategy to Imaging BAT in the Basal State Using NET-[ $^{11}\text{C}$ ]MRB in Humans

Ten healthy, Caucasian subjects (5 M, age  $24.6 \pm 2.6$ , BMI  $21.6 \pm 2.7$  kg/m<sup>2</sup>; 5 F, age  $25.4 \pm 2.1$ , BMI  $22.1 \pm 1.0$  kg/m<sup>2</sup>) underwent  $^{11}\text{C}$ -MRB PET-CT imaging (mCT PET/CT scanner, Siemens/CTI, USA) for cervical/supraclavicular BAT under r.t. and cold-stimulated conditions (RPCM Cool Vest; enthalpy 15 °C) compared to [ $^{18}\text{F}$ ]FDG PET-CT imaging (Hwang et al. 2015). Cold condition scans were performed while subjects wore a climate vest loaded with cold packs, whereas room temperature scans were conducted using the same climate vest loaded with room temperature packs, to avoid potential biases from changes in attenuation between the two conditions. For [ $^{18}\text{F}$ ]FDG studies, subjects wore the climate vest for half an hour before [ $^{18}\text{F}$ ]FDG injection, during the 60-min period between the injection and scan, and during the 25–30 min scan. For [ $^{11}\text{C}$ ]MRB PET-CT scans, the climate vest was placed on the subjects 30 min prior to intravenous [ $^{11}\text{C}$ ]MRB administration and remained on the subject for the duration of the 120–125 min dynamic scan. PET-CT scanning began with the administration of [ $^{11}\text{C}$ ]MRB. All scans were performed under fasting conditions.

For analysis of tracer uptake, we chose an [ $^{18}\text{F}$ ]FDG SUV  $>1.5$  and Hounsfield units ( $-200$  to  $-50$ ) as has been used by others (Blondin et al. 2014; Lee et al. 2014) to define BAT tissue. Uptake of [ $^{11}\text{C}$ ]MRB was quantified as the distribution volume ratio (DVR) using the occipital cortex as the reference region since the tracer binding has been well characterized in our previous brain studies in humans (Ding et al. 2010; Hannestad et al. 2010a; Pietrzak et al. 2013; Li et al. 2014), and uptake of metabolites of [ $^{11}\text{C}$ ]MRB was minimal in rodent BAT. Muscle was also chosen as a peripheral reference, and a ratio of BAT-DVR to muscle-DVR (BAT/muscle) was compared. Total body fat and lean body mass were assessed via bioelectrical impedance analysis.

Consistent with previous studies, we found [ $^{18}\text{F}$ ]FDG uptake in BAT was difficult to identify at r.t. but easily detected with cold stimulation ( $p = 0.01$ ). In contrast, BAT [ $^{11}\text{C}$ ]MRB uptake (also normalized for muscle) was equally evident under both r.t. and cold conditions (BAT DVR, r.t.  $1.0 \pm 0.3$  vs. cold  $1.1 \pm 0.3$ ,  $p = 0.31$ ; BAT/muscle DVR, r.t.  $2.3 \pm 0.7$  vs. cold  $2.5 \pm 0.5$ ,  $p = 0.61$ ) (Fig. 20.9). Importantly, BAT

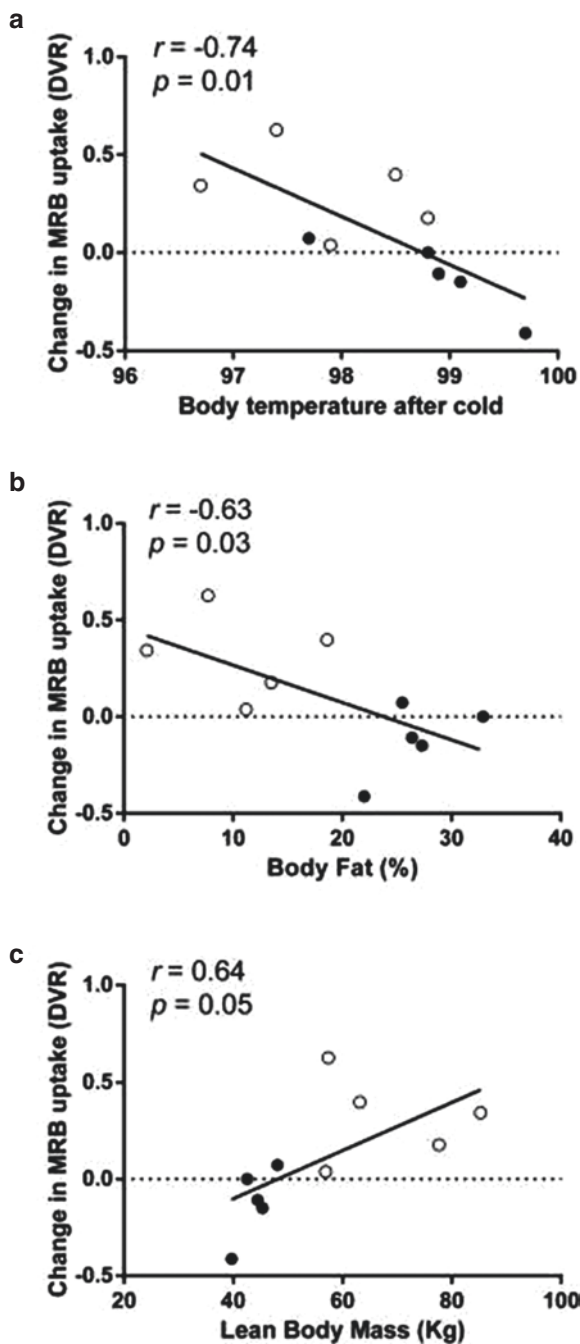


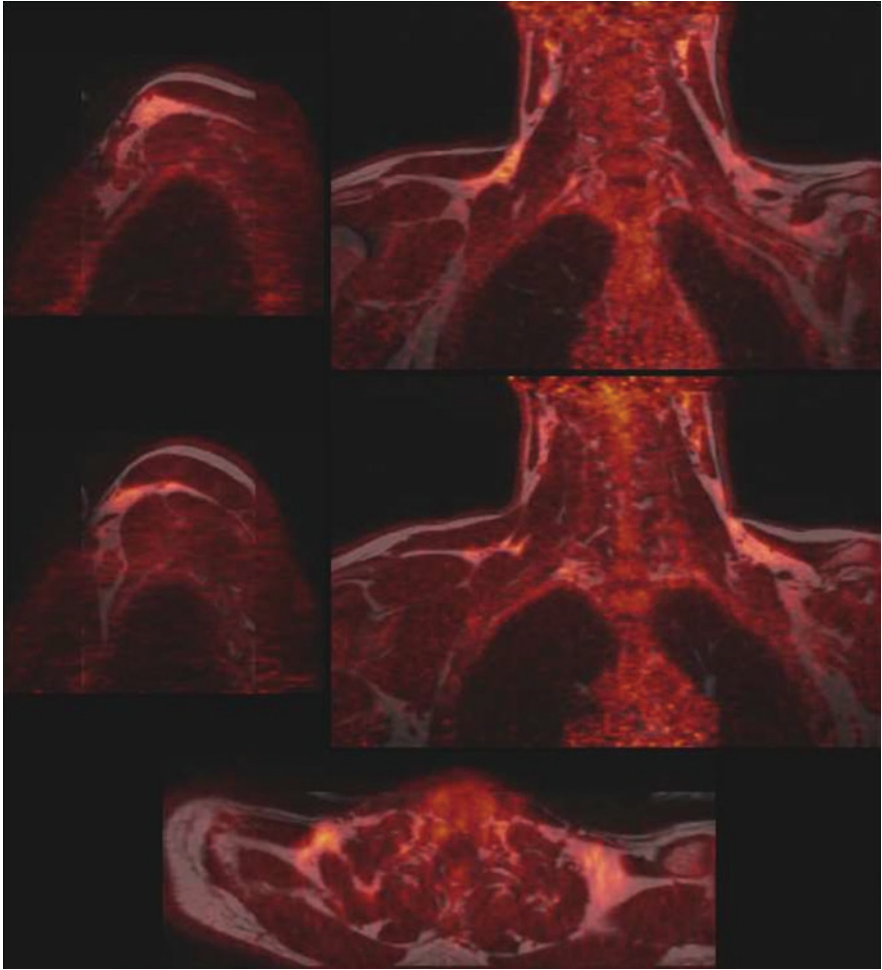
**Fig. 20.9** PET and CT images in one representative male subject showed that [ $^{18}\text{F}$ ]FDG uptake in BAT was observed under mild cold conditions but not at room temperature, while [ $^{11}\text{C}$ ]MRB uptake in BAT was observed under both room temperature and cold conditions. [ $^{18}\text{F}$ ]FDG and [ $^{11}\text{C}$ ]MRB are scaled from SUV 0 (black) to SUV 2 (white). [ $^{11}\text{C}$ ]MRB images are computed from the average of frames acquired between 40 to 60 min post-injection

DVR and BAT/muscle DVR of [ $^{11}\text{C}$ ]MRB at r.t. correlated positively with core body temperature ( $r = 0.76$ ,  $p = 0.05$  and  $r = 0.92$ ,  $p = 0.004$ , respectively), a relationship not observed with [ $^{18}\text{F}$ ]FDG ( $p = 0.63$ ). There were no relationships between [ $^{11}\text{C}$ ]MRB uptake and heart rate or BMI. However, the magnitude of change in [ $^{11}\text{C}$ ]MRB BAT uptake in response to cold correlated negatively with percent body fat ( $r = -0.74$ ,  $p = 0.04$ ) and positively with lean mass ( $r = 0.71$ ,  $p = 0.05$ ), which showed a gender difference (Fig. 20.10). This gender difference most likely reflected the difference in body composition and body temperature between men and women. There were no relationships between the change in [ $^{18}\text{F}$ ]FDG uptake in response to cold and body composition, BMI, body temperature, heart rate, or blood pressure.

We recently assessed BAT using a state-of-the-art PET/MR combined scanner with simultaneous acquisition (Biograph mMR, Siemens). Representative images of the overlaid PET data on MR fat images acquired with a 3D two-point Dixon sequence showed excellent co-localization of BAT area from the two imaging modalities (Fig. 20.11). The simultaneously acquired PET and MR-fat images significantly facilitated the localization and quantification of BAT. Further, using a 3D multiple gradient echo sequence, MR quantification of BAT allows mapping of percentage water for quantifying fat, water, and  $R2^*$  ( $= 1/T2^*$ ) in BAT. The fat fraction (FF) can be calculated from  $\text{PDFF} = F/(F + W)$ , where PDFF is the proton density FF, F is the fat signal, and W is the water signal. The FF is known to be lower in BAT than in WAT (Hu et al. 2013), while  $R2^*$  is expected to be higher. Using this strategy, the changes in fat fraction and fat volume in BAT, measured with MR, can be

**Fig. 20.10** Relationship between change in MRB uptake in response to cold and (a) body temperature after cold; (b) percentage body fat; (c) lean body mass (open circle = men; closed circle = women)





**Fig. 20.11** PET-BAT image overlaid on MR-fat images acquired with a 3D two-point Dixon sequence. Images have been reformatted into three planes: sagittal through each shoulder (left), coronal (right), and axial (bottom)

correlated with the BAT activation measured with PET after various interventions. Combination of the high specificity of PET and high spatial resolution of MR will provide unique information on BAT quantification.

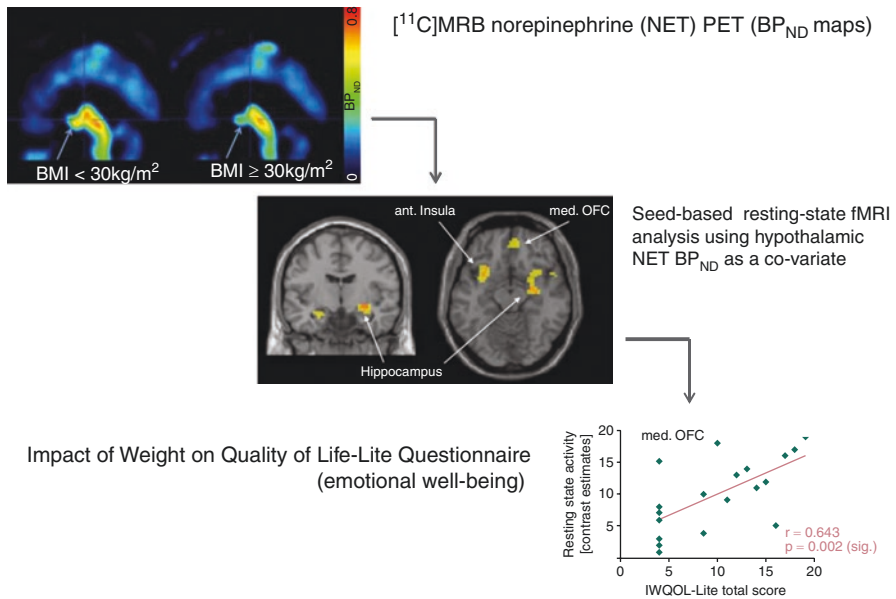
### 20.3.6 NET Imaging in Obesity

In collaboration with University Medical Centre in Leipzig, the central NET system was investigated using [ $^{11}\text{C}$ ]MRB, including NET imaging in obesity.

### 20.3.6.1 Linking NET to Emotional Distress and Obesity

In a group of lean to morbidly obese individuals ( $n = 20$ ), Melasch et al. (Melasch et al. 2016) showed that an increased body mass index (BMI) is related to a lowered NET availability within the hypothalamus, known as the brain’s homeostatic control site. The hypothalamus displayed a strengthened connectivity in relation to the individual hypothalamic NET availability to the anterior insula/frontal operculum, as well as the medial orbitofrontal cortex, assumed to host the primary and secondary gustatory cortex, respectively ( $n = 19$ ). The resting-state activity in these two regions was correlated positively to the BMI and the scores of IWQOL-Lite (the impact of weight on quality of life-Lite questionnaire) (Fig. 20.12), but not to the Beck depression inventory (Pattinson et al. 2009), suggesting that the higher the resting-state activity in these regions, and hence the higher the BMI, the stronger the negative impact of the body weight on the individual’s emotional well-being was. This pilot study suggests that the loss in emotional well-being with weight is embedded within the central norepinephrine network.

#### NET-Dependent Changes and Neurobehaviour in Obesity

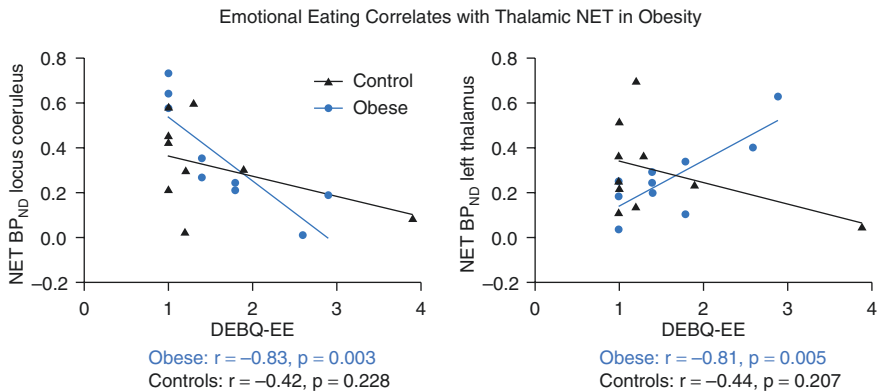


**Fig. 20.12** NET-Dependent Changes and Neurobehaviour in Obesity: Seed-based resting-state fMRI analysis using hypothalamic NET  $BP_{ND}$  as a co-variate showing stronger connectivity between the hypothalamus and limbic brain areas that determines emotional well-being



### 20.3.6.2 Central In Vivo NET Availability Is Altered in Emotional Eating of Individuals with Obesity

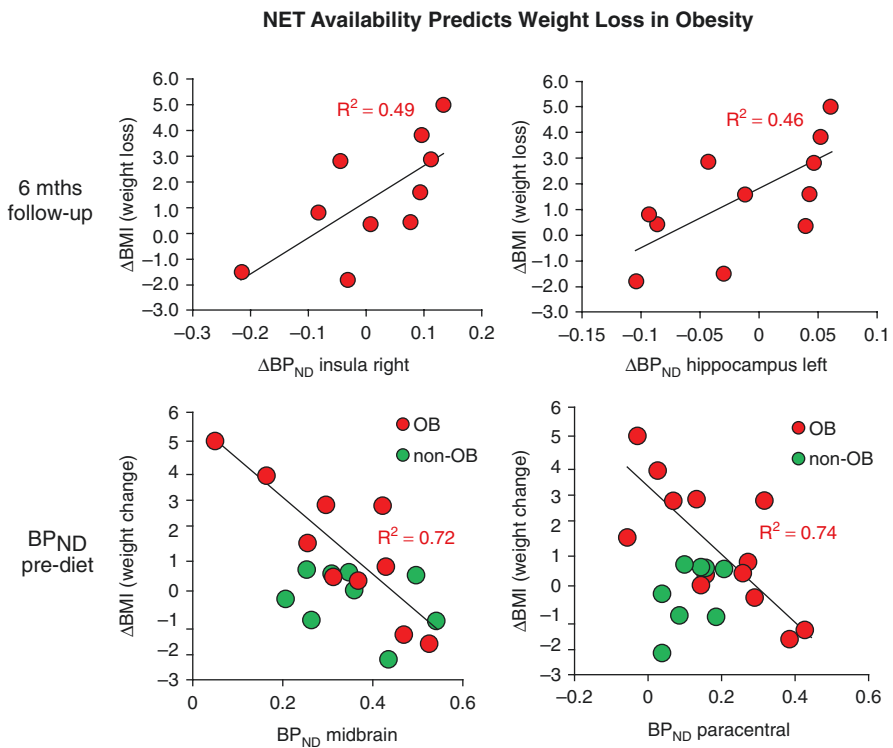
Our previous studies in cocaine abuse using [ $^{11}\text{C}$ ]MRB indicated significant upregulation of NET in the thalamus of cocaine users, as compared to healthy controls (Ding et al. 2010). However, when comparing PET imaging using [ $^{11}\text{C}$ ]MRB for healthy controls and morbidly obese subjects, a decrease in NET availability was found in the thalamus in obese subjects (Li et al. 2014). These findings present a notable difference in neurotransmission between individuals with substance use disorder and those with obesity. To investigate an association between NET availability in obese individuals and emotional eating (EE), Bresch et al. (Bresch et al. 2017) sought to measure EE through the EE subscale of the Dutch Eating Behavior Questionnaire for obese subjects and healthy controls before performing PET scans with [ $^{11}\text{C}$ ]MRB. This study did not find significant differences in EE scores and regional NET levels between healthy controls (BMI  $23.9 \pm 2.5 \text{ kg/m}^2$ ) and obese participants (BMI  $42.4 \pm 3.7 \text{ kg/m}^2$ ). However, for obese individuals only, correlative data analyses pointed to a sinoidal distribution pattern as a higher degree of EE related to lower NET availability in the locus coeruleus and to higher NET availability in the left thalamus (Fig. 20.13). Collectively, these studies suggest that NE transmission is regionally impaired in substance use disorder, obese, and EE populations and that regional fluctuations in NET availability determine the deviations in motivated behavior, such as emotional eating, which are seen in substance use disorder and obesity (Lindgren et al. 2018).



**Fig. 20.13** Emotional Eating Correlates with Thalamic NET in Obesity: for obese individuals higher EE scores correlated with lower NET availability in the locus coeruleus and higher NET availability in the left thalamus

### 20.3.6.3 NET Availability and Success of Weight Loss in Obese Individuals

A recent study was carried out by Vettermann et al. to compare central NET availability in ten obese, otherwise healthy individuals with a body mass index (BMI) of  $42.4 \pm 3.7 \text{ kg/m}^2$  (age  $34 \pm 9$  years, four women) and ten matched nonobese, healthy controls (BMI  $23.9 \pm 2.5 \text{ kg/m}^2$ , age  $33 \pm 10$  years, four women) using [ $^{11}\text{C}$ ]MRB before and 6 months after dietary intervention. The results indicated that an increase in binding potential ( $\text{BP}_{\text{ND}}$ ) in the insula and the hippocampus of obese individuals correlated well with changes in BMI ( $-3.3 \pm 5.3\%$ ;  $p = 0.03$ ) following completion of the 6-mo dietary intervention. Furthermore, NET availability before intervention could help to predict the amount and success of weight loss in obese individuals. Lower  $\text{BP}_{\text{ND}}$  in these regions (but also in the midbrain and the paracentral cortex) at baseline was associated with higher achieved weight loss (e.g., hippocampal area  $R^2 = 0.80$ ;  $p < 0.0001$ ) (Fig. 20.14).



Vettermann F, et al. Eur J Nucl Med Mo Imaging. 2018 Jul;45(9):1618-1625.

**Fig. 20.14** NET Availability Predicts Weight Loss in Obesity: NET availability before intervention could help predict the amount and success of weight loss in obese individuals

These first longitudinal interventional data on NET availability in highly obese individuals indicate that the central NA system is modifiable and that weight loss may be predicted by measuring regional NAT availability prior to intervention. If confirmed, such measurement of NAT availability with PET may serve as an individual biomarker to predict the likelihood of achieving weight loss after dietary intervention and offer the possibility of identifying patients who will or will not benefit from dietary intervention. Whether measurement of NET availability with PET is also able to predict the success of other anti-obesity treatments such as bariatric surgery and might therefore be used in the future in formulating personalized interventional strategies in individuals with obesity needs to be evaluated in prospective comparative studies.

### 20.3.7 NET in Parkinson's Disease (PD)

Although postmortem studies have detected a substantial noradrenergic deficit in Parkinson's disease patients (Kish et al. 1984), in vivo assessment of the noradrenergic system has been hampered by the lack of suitable imaging tools. Through our extensive development and validation studies in both nonhuman primates and humans, [ $^{11}\text{C}$ ]MRB has now been adopted by many other researchers for NET imaging, including in PD research (Nahimi et al. 2018; Sommerauer et al. 2018a, b).

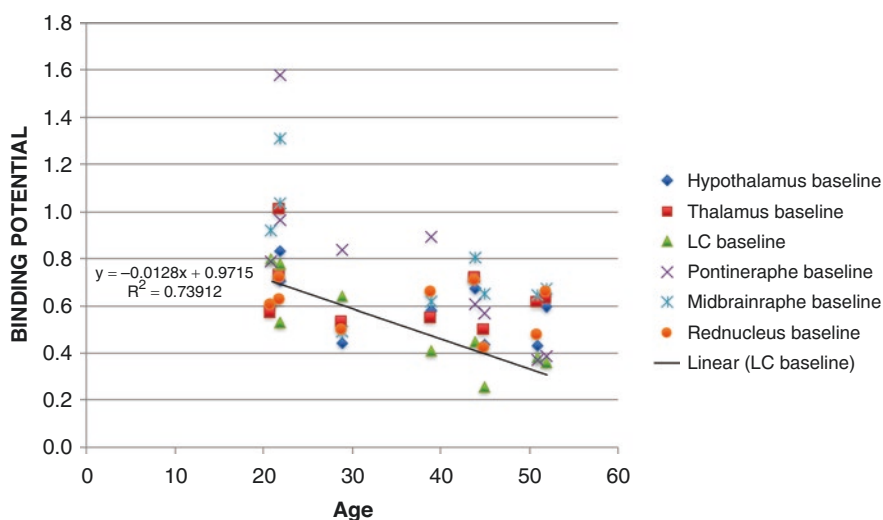
Rapid eye movement sleep behavior disorder (RBD) accompanies Parkinson's disease and its presence predicts an unfavorable disease course with a higher propensity to cognitive impairment and orthostatic hypotension. Noradrenergic innervation of the brain arises almost exclusively from the neuromelanin-containing cells of the locus coeruleus. Using the neuromelanin-sensitive MR and NET availability biomarker ([ $^{11}\text{C}$ ]MRB), the association of RBD, cognition, and autonomic dysfunction in PD with loss of noradrenergic function was examined (Sommerauer et al. 2018a). The authors found that Parkinson's disease patients with RBD showed decreased locus coeruleus neuromelanin signal on MRI ( $P < 0.001$ ) and widespread reduced binding of [ $^{11}\text{C}$ ]MRB ( $P < 0.001$ ), which correlated with amount of REM sleep without atonia. Parkinson's disease with RBD was also associated with a higher incidence of cognitive impairment, slowed EEG activity, and orthostatic hypotension.

Another recent study (Nahimi et al. 2018) using [ $^{11}\text{C}$ ]MRB showed reduced NET binding in patients with PD, as compared to controls, with significant declines in red nucleus and thalamus. They found no associations between [ $^{11}\text{C}$ ]MRB  $\text{BP}_{\text{ND}}$  estimates and disease severity in terms of disease duration, disease stage, or severity of motor symptoms. However, tremor symptoms in PD patients were associated with higher [ $^{11}\text{C}$ ]MRB  $\text{BP}_{\text{ND}}$  in the thalamus compared to patients without tremor. It is known that PD patients with tremor-dominant disease represent a subgroup of a more benign course of disease with slower progression of motor and non-motor symptoms and lower frequency of cognitive impairment. So this better-preserved NET availability (noradrenergic neurons) when compared to the akinetic-rigid subtype could be potentially an important biomarker, though a larger sample size is needed to reproduce this interesting result.

### 20.3.8 NET in Aging and Alzheimer's Disease (AD)

There is little published data on how in vivo NET availability varies with age, but our preliminary data have documented declines in NET availability in healthy adults in the pulvinar, hypothalamus, and LC ( $P = 0.03, 0.009, 0.04$ , respectively; age range = 25–54) (Fig. 20.15). Similar results were reproduced in a collaborative project with a Vancouver group (Yatham et al. 2018) and in a recent study (Vanicek et al. 2014). According to our previous study, the % BP<sub>ND</sub> change for DAT was approx. 0.7% per year (Volkow et al. 1996); however, the %BP<sub>ND</sub> decline per year for NET was about threefold faster based on our pilot study (Ding et al. 2010). These intriguing results warrant further investigation with a larger sample size.

Alzheimer's disease (AD) is a major health challenge in our aging society. An improved understanding of its underlying pathological mechanisms is urgently needed to enable the development of effective treatments. Postmortem findings indicate that tau pathology emerges decades before amyloid pathology, appearing first in the brainstem—in particular in the locus coeruleus (LC, the major source of the brain's norepinephrine [NE]) (Baker et al. 1989; Sharma et al. 2010; Aston et al. 1986)—before slowly spreading to the entorhinal cortex and other LC targets, including hippocampus (Braak et al. 2011). The LC is particularly vulnerable to toxins and infection and is often the first place AD pathology appears. Postmortem studies suggest that loss of LC neurons predicts onset and severity of AD symptoms better than A $\beta$ /neurofibrillary tangle pathology or cell loss in any other brain region implicated in AD (Bondareff et al. 1981, 1987; Burke et al. 1988; Busch et al. 1997; Chan-Palay and Asan 1989; Gannon et al. 2015; German et al. 1992; Hardy et al. 1985; Iversen et al. 1983; Mann et al. 1982; Marcyniuk et al. 1986; Strong et al.



**Fig. 20.15** Age effect on NET availability (binding potential) in HC

1991; Wilson et al. 2013). However, *it is not yet possible to measure the early progression of tau pathology in the LC of living humans* due to the off-target binding of the current tau radioligands to neuromelanin, which is found at high levels in the LC (Hansen et al. 2016) (Lemoine et al. 2018; Saint-Aubert et al. 2017).

Our studies in humans using NET- $[^{11}\text{C}]$ MRB imaging demonstrated a special vulnerability of LC to aging and stress (PTSD), suggesting that NET dysfunction plays an important role in aging. We and others have recently implemented a specific neuromelanin-MR sequence (Clewett et al. 2016; Keren et al. 2009, 2015; Sasaki et al. 2006) that makes it possible to reliably delineate and quantitate the NET-MRB binding (NET availability) and MR (neuromelanin) signal intensities in LC (indexing functional and structural integrity) by using state-of-the-art PET-MR combined imaging with simultaneous acquisition. We hypothesize that NET-MRB imaging may provide a new noninvasive approach for preclinical diagnosis, staging, and assessment of therapeutic interventions in AD.

### 20.3.9 Kinetic Modeling for NET Imaging Studies in the Brain

It is important to point out that arterial input function measurement was conducted for almost all of our PET-NET brain imaging studies using  $[^{11}\text{C}]$ MRB (Ding et al. 2010; Gallezot et al. 2007, 2010; Logan et al. 2007, 2005; Hannestad et al. 2010b; Li et al. 2005). This made it possible to quantify the volume of distribution ( $V_T$ ) of  $[^{11}\text{C}]$ MRB using multilinear analysis MA1 (Ichise et al. 2002) and a direct comparison of the quantification of  $[^{11}\text{C}]$ MRB binding potential ( $\text{BP}_{\text{ND}}$ ) without arterial blood sampling using either MRTM2 or SRTM2 (Ichise et al. 2002; Innis et al. 2007). With these validation techniques (Gallezot et al. 2010; Hannestad et al. 2010b), we evaluated several candidate reference regions so that placement of arterial lines may not be needed for future NET imaging studies using  $[^{11}\text{C}]$ MRB.

As described above, one of the intrinsic challenges for imaging NET is its wide distribution throughout the brain, with relatively low contrast between transporter-poor and transporter-rich regions. It has been shown that the lowest NET densities measured by autoradiography in rhesus monkeys were in the putamen and caudate (Smith et al. 2006), while the lowest concentration of NET in humans was found in the occipital cortex, based on one in vitro autoradiography study in humans using  $[^{18}\text{F}]$ F-MeNER-D2 (Schou et al. 2005). Thus, evaluation of caudate, putamen, and occipital cortex as potential reference regions was conducted in both rhesus monkeys and humans.

Based on our  $[^{11}\text{C}]$ MRB studies in nonhuman primates (NHP), we found that the use of caudate as the reference region in NHP leads to the least bias. This bias is not significant as compared to the “gold standard”  $\text{IC}_{50}$  estimate obtained using only  $V_T$  or  $V_T/f_p$  values (Gallezot et al. 2010). Moreover, similar results were obtained while using the binding potentials  $\text{BP}_F$  and  $\text{BP}_{\text{ND}}$  (Innis et al. 2007). The latter could be estimated without arterial blood sampling by using MRTM2 and provided results equivalent to those obtained with the arterial data, i.e., without introducing further bias. Based on our  $[^{11}\text{C}]$ MRB studies in both NHP and humans, it seems that the

fraction of displaceable binding in the occipital cortex for [ $^{11}\text{C}$ ]MRB is higher in rhesus monkeys than in humans.

Interestingly, the issue of selecting a reference region seems to be less troublesome in humans since all three NET-poor regions (caudate, putamen and occipital) were considered as reasonable reference regions based on the results of our occupancy study of methylphenidate in humans (Hannestad et al. 2010b) (see above discussion). The occipital cortex has lower levels of NET than the striatum (Schou et al. 2005) and is a relatively bigger region with a simpler shape than the caudate and putamen, making it more suitable as the reference region for human studies, especially when ROIs are delineated with an automated template. Thus, using MRTM2 with a reference region to quantify [ $^{11}\text{C}$ ]MRB binding without arterial sampling for occupancy studies is feasible in nonhuman primates and adequate in humans.

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## 20.4 Summary and Outlook

The main purpose of this review article is to outline the progress that has been made on PET imaging of the NET system during the past decade, starting from ligand development, ligand characterization, toxicology evaluation, and ligand validation and approval for human studies (Ding et al. 2006), leading to translational research in humans. A few specific applications using [ $^{11}\text{C}$ ]MRB in the scope of evaluation of disease abnormality, determination of drug occupancy, understanding biological processes, and interpretation of drug mechanisms have been explored. Our pilot investigations clearly indicated how little we know about the role of NET in various CNS disorders. As previously mentioned, many of the important findings resulted from studies *in vitro* using postmortem tissues and have suggested the implication of NET in various neurological and psychiatric disorders, but they have never been verified via *in vivo* methods due to the lack of suitable radioligands.

We have just begun to learn the role of NET in the pathophysiology and the treatment of several disorders, including cocaine abuse, ADHD, depression, PTSD, alcohol dependence, obesity, and also in normal aging. For example, cocaine is the most abused drug with the highest reinforcing ability. Its effect on the dopamine system has been well characterized; however, its effect on NET in humans has never been investigated. Our studies on cocaine abusers and the effect of duration of abstinence on NET binding provide the first evidence of NET abnormalities in cocaine dependence.

Our occupancy studies of MPH demonstrate that clinically relevant doses block 70% to 80% of NET, whereas only 60% to 70% of DAT is blocked at similar doses. Despite of the fact that MPH has been the most prescribed drug for the treatment of ADHD over the last 50 years, our study provides the first evidence questioning the traditional thinking that the therapeutic effect of MPH is mainly due to its inhibition of DAT.

Our PET imaging studies on ATX unveiled its high *in vivo* affinity toward both NET and SERT, questioning the traditional thinking that ATX is a selective NET inhibitor. These results suggest a cautious interpretation of the therapeutic mode of

action of ATX for treatment of depression and ADHD, as the therapeutic effect of ATX may be more complex than selective blockade of the NET.

It is important to point out that most of our investigations to evaluate disease abnormality are pilot studies with relatively small sample sizes. Further studies employing larger samples are needed to confirm the results. Furthermore, other important aspects (e.g., gender) that have been linked to various disorders may be revealed.

Though the major focus of this chapter is on our translational research using [ $^{11}\text{C}$ ]MRB, its F-18 analogues, such as [ $^{18}\text{F}$ ]fluoromethyl ((*S,S*)-[ $^{18}\text{F}$ ]FMeNER) (Schou et al. 2004) and [ $^{18}\text{F}$ ]fluoroethyl analogs, ((*S,S*)-[ $^{18}\text{F}$ ]FRB) (Ding et al. 2006, 2005; Lin et al. 2005), also showed promising results. Unfortunately, their relatively lower binding affinity and defluorination problems precluded their further translational research in humans (Ding et al. 2006; Takano et al. 2008a, b). NET ligand development work is ongoing. Future studies using PET ligands with greater affinity and specificity toward NET may help to further examine the role of noradrenergic function in various CNS disorders and facilitate our understanding of the role of NET in living humans, without having to rely solely on results from postmortem studies.

New strategies are being developed for synergistic PET/MR imaging using the simultaneous data acquisition capabilities of a combined PET/MR scanner (e.g., Siemens mMR) in order to take advantage of the spatial resolution of MRI with the biochemical and functional information from PET. This state-of-the-art PET/MR combined modality would facilitate co-registration; for example, the locus coeruleus (LC), which plays an important role in various CNS disorders and is the region of highest NET density, has a very small structure that poses a challenge to NET imaging due to difficulties encountered during PET and MRI co-registration when separate PET and MRI scans are collected—a problem that can be addressed by simultaneous MR and PET data acquisition. Furthermore, it would permit simultaneous imaging of physiologic and pathophysiologic processes and avoid the potential confounding effects of day-to-day and status variations. It would also provide an unprecedented opportunity to correlate PET measures with MR-detected patterns of neural synchrony in both cortical and subcortical regions, facilitating the search process for optimal biomarkers for various CNS disorders. For example, we have demonstrated the dose-dependent occupancy of MPH in LC, subcortical and cortical regions by using NET-PET imaging (see above Sect. 3.2.5). The use of PET/MR with simultaneous acquisition after the treatment with MPH may facilitate our understanding of the functional connectivity among the brain regions by linking the effect of MPH from LC to its projection into cortical regions, including PFC.

Further, NET imaging in PD and AD may provide a new noninvasive approach for preclinical diagnosis, staging, and assessment of therapeutic interventions in these neurodegenerative diseases.

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# Positron Emission Tomography (PET) Imaging of Opioid Receptors

# 21

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## Abstract

The opioid system consists of opioid receptors (which mediate the actions of opium), their endogenous ligands (the enkephalins, endorphins, endomorphins, dynorphin, and nociceptin), and the proteins involved in opioid production, transport, and degradation. PET tracers for the various opioid receptor subtypes are available, and changes in regional opioidergic activity have been assessed during both sensory and affective processing in healthy individuals and in various disease conditions such as chronic pain, neurodegeneration, epilepsy, eating disorders, behavioral addiction, and substance abuse. It is not always clear whether observed changes of tracer binding reflect altered release of endogenous opioids or altered opioid receptor expression. This issue may be resolved by studies in experimental animals that combine *in vivo* PET imaging with *ex vivo* immunohistochemistry. Some radioligands for opioid receptors have suboptimal kinetics (i.e., slow dissociation from their target protein) or can induce undesired side effects even at low administered doses (sedation, respiratory arrest). Yet, PET offers the unique opportunity of quantifying opioid receptor-mediated signaling in the living human brain. PET imaging has provided evidence for a link between opioid neurotransmission and peripheral immune activation.

## 21.1 Introduction

The term “opioid” is derived from opium: dried latex from the opium poppy (*Papaver somniferum*) which has been used since ancient times as an analgesic and recreational drug (Kalant 1997). An opioid is either a synthetic narcotic that has similar activities as opium or a naturally occurring peptide that has such activities. Several classes of endogenous opioids have been discovered, viz., the enkephalins



(Hughes et al. 1975), endorphins (Cox et al. 1976; Simantov and Snyder 1976), endomorphins (Zadina et al. 1997), dynorphins (Goldstein et al. 1979), and nociceptin (Meunier et al. 1995; Reinscheid et al. 1995). These messenger substances bind to G-protein-coupled opioid receptors in the brain, the spinal cord, and the digestive tract. Agonist binding to opioid receptors results in a decrease of the activity of adenylyl cyclase (Snyder and Pasternak 2003; Trescot et al. 2008; Waldhoer et al. 2004).

At least four broad subclasses of opioid receptors have been identified by the use of selective ligands (Dhawan et al. 1996). Assessment of antagonist potencies against the actions of various opioid agonists in different tissue preparations resulted in the identification of the mu, delta, and kappa subclasses (Lord et al. 1977; Martin et al. 1976). Later, mRNAs for these three subtypes were cloned and characterized (Chen et al. 1993; Evans et al. 1992; Kieffer et al. 1992; Yasuda et al. 1993). Mu opioid receptors ( $\mu$ OR) are named after morphine, the alkaloid which is the active compound in opium (Martin et al. 1976). Kappa opioid receptors ( $\kappa$ OR) derive their name from the drug ketocyclazocine (Martin et al. 1976). Delta opioid receptors ( $\delta$ OR) were initially characterized in the murine vas deferens (Lord et al. 1977).  $\mu$ OR are  $\beta$ -endorphin preferring,  $\kappa$ OR dynorphin preferring, and  $\delta$ OR enkephalin preferring (Minami and Satoh 1995). Nonclassical, nociceptin- or opiate-like receptors 1 (ORL1) bind nociceptin (Meunier et al. 1995; Reinscheid et al. 1995).

$\beta$ -Endorphin binds with nM affinity to  $\mu$ OR and  $\delta$ OR ( $K_i$  about 1 nM) but with low affinity to  $\kappa$ OR ( $K_i = 52$  nM) (Raynor et al. 1994). In contrast, dynorphin A has sub-nM affinity to  $\kappa$ OR ( $K_i = 0.5$  nM), low affinity to  $\mu$ OR ( $K_i = 32$  nM), and negligible affinity to  $\delta$ OR ( $K_i > 1$   $\mu$ M) (Raynor et al. 1994). Enkephalins have sub-nM or nM affinity to  $\delta$ OR and  $\mu$ OR ( $K_i$  values 0.6–4 nM) but negligible affinity to  $\kappa$ OR ( $K_i > 1$   $\mu$ M) (Raynor et al. 1994). Endomorphins are very potent and selective  $\mu$ OR agonists ( $K_i = 0.3$ – $0.7$  nM) (Hackler et al. 1997). Stimulation of  $\mu$ OR or  $\delta$ OR generally results in analgesia and reward, whereas  $\kappa$ OR stimulation may have negative consequences, such as dysphoria and hallucinations.

Opioids and their receptors play important roles in a large spectrum of physiological processes, including learning and memory, reward, mood, appetite, circadian rhythms, sexual activity, pregnancy, locomotion, cardiovascular, gastrointestinal, renal and hepatic function, respiration, regulation of body temperature, and immunological response (Vaccarino and Kastin 2001). Well known is their involvement in regulating the sensation of pain; opiates are clinically employed to produce analgesia.

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## 21.2 Opioid Receptor Ligands for PET: A Historical Overview

Several radioligands for PET imaging of opioid receptors have been prepared. These include subtype-selective and non-subtype-selective agonists and antagonists (see Table 21.1). Many comprehensive reviews on opioid receptor imaging were published between 2000 and 2008 (Frost 2001; Hammers and Lingford-Hughes

**Table 21.1** PET probes for the opioid system

Name	Action	Abbreviation	Initial publication
[ <sup>18</sup> F]Cyclofoxy	Antagonist, $\mu$ OR, and $\kappa$ OR	–	Pert et al. (1984)
[ <sup>11</sup> C]Diprenorphine	Antagonist, non-subtype selective	DPN	Luthra et al. (1985)
[ <sup>18</sup> F]Diprenorphine	Antagonist, non-subtype selective	FDPN	Wester et al. (2000)
[ <sup>11</sup> C]Buprenorphine	Mixed agonist/antagonist, non-subtype selective	BPN	Luthra et al. (1987)
[ <sup>18</sup> F]-FE-PEO	Full agonist, non-subtype selective	–	Riss et al. (2013)
[ <sup>11</sup> C]Carfentanil	Agonist, $\mu$ OR	CFN	Dannals et al. (1985)
[ <sup>11</sup> C]AH7921	Agonist, $\mu$ OR	–	Rafique et al. (2017)
[ <sup>11</sup> C]Methylnaltrindole	Antagonist, $\delta$ OR	MeNTI	Lever et al. (1992)
[ <sup>64</sup> Cu]TIPP conjugate	Antagonist, $\delta$ OR	–	Pirisedigh et al. (2017)
[ <sup>11</sup> C]GR103545	Agonist, $\kappa$ OR	–	Talbot et al. (2005)
[ <sup>11</sup> C]EKAP	Agonist, $\kappa$ OR	–	Li et al. (2019a)
[ <sup>11</sup> C]FEKAP	Agonist, $\kappa$ OR	–	Li et al. (2019b)
[ <sup>11</sup> C]MeJDTic	Antagonist, $\kappa$ OR	–	Poisnel et al. (2008)
[ <sup>11</sup> C]JLY2795050	Antagonist, $\kappa$ OR	–	Zheng et al. (2013)
[ <sup>11</sup> C]JLY2459989	Antagonist, $\kappa$ OR	–	Zheng et al. (2014)
[ <sup>18</sup> F]JLY2459989	Antagonist, $\kappa$ OR	–	Cai et al. (2017), Li et al. (2018)
[ <sup>11</sup> C]NOP-1A	Antagonist, ORL1	–	Pike et al. (2011)
[ <sup>18</sup> F]MK-0911	Antagonist, ORL1	–	Hostetler et al. (2013)

2006; Henriksen and Willoch 2008; Koeppe and Duncan 2000; Lever 2007; Ravert et al. 2004; Sprenger et al. 2005).

The first attempt at *in vivo* labeling of opioid receptors was made in 1975. Investigators at Johns Hopkins University administered a labeled opioid receptor antagonist, [<sup>3</sup>H]naloxone, to living rats. In the excised brain tissue of these animals, they observed a regional distribution of radioactivity which corresponded to the known distribution of opioid receptors in the rodent brain (striatum > hind-brain > cerebellum). Because of strong nonspecific binding of [<sup>3</sup>H]naloxone, acceptable target-to-nontarget ratios of radioactivity could only be reached after *ex vivo* washing of brain slices to remove nonspecifically bound tracer. Yet, this seminal study paved the way for all later imaging of neuroreceptors in the living brain (Pert and Snyder 1975).

The first successful PET probe for opioid receptors was prepared in 1984 (Pert et al. 1984; Channing et al. 1985). After intravenous administration of [<sup>18</sup>F]3-acetylcyclofoxy (3-acetyl-6-deoxy-6- $\beta$ -[<sup>18</sup>F]-fluoronaltraxone), a regional distribution of radioactivity was observed which corresponded to the regional distribution of opioid receptors in the rat and baboon brain. Bound tracer in the baboon brain was completely displaced by the active (–)enantiomer of naloxone, whereas an

identical dose of the inactive (+)enantiomer (0.13 mg/kg) had no effect (Pert et al. 1984).

Since [ $^3\text{H}$ ]naloxone showed a low ratio of specific-to-nonspecific binding, an antagonist with a higher affinity ([ $^3\text{H}$ ]diprenorphine, [ $^3\text{H}$ ]DPN) was prepared and successfully applied in autoradiography (review in (Frost 2001)). However, labeling DPN with  $^{11}\text{C}$  proved difficult. Because of these difficulties, radiochemists focused their attention on other opioid ligands, and a method was developed to label carfentanil (CFN) with  $^{11}\text{C}$ , using a carboxylic acid analog of carfentanil as a precursor (Dannals et al. 1985). The resulting tracer, [ $^{11}\text{C}$ ]CFN, was used to acquire the very first images of opioid receptors in the human brain (Frost et al. 1985). At baseline, a regional distribution of radioactivity was observed which corresponded to the known regional densities of  $\mu\text{OR}$ . After treatment of the volunteer with naloxone (1 mg/kg), cerebral uptake of the tracer was strongly suppressed, indicating that [ $^{11}\text{C}$ ]CFN binds to  $\mu\text{OR}$  in the living brain in vivo (Frost et al. 1985). A regional distribution of radioactivity consistent with  $\mu\text{OR}$  binding and blockade of tracer uptake by naloxone were also observed in the mouse brain. Thus, [ $^{11}\text{C}$ ]CFN is suitable for studies of  $\mu\text{OR}$  in rodents (Saji et al. 1992).

In the same year that the first images of opioid receptors in the human brain were acquired, radiochemists developed a method for preparation of [ $^{11}\text{C}$ ]DPN. The compound was labeled at the N-carbon atom of its cyclopropylmethyl group (Luthra et al. 1985). A study with [ $^{11}\text{C}$ ]DPN in four healthy volunteers indicated that the tracer is rapidly washed out from brain regions with a low density of opioid receptors, whereas activity is retained in receptor-rich target areas. Eighty to ninety percent of tracer binding in target regions is naloxone reversible, suggesting that [ $^{11}\text{C}$ ]DPN is suitable for studies of opioid receptors in man (Jones et al. 1988).

A different method to prepare [ $^{11}\text{C}$ ]DPN was later published. That method involves [ $^{11}\text{C}$ ]-*O*-methylation of 3-*O*-*t*-butyldimethylsilyl-(6-*O*-desmethyl) diprenorphine with [ $^{11}\text{C}$ ]methyl iodide, followed by acidic deprotection of the reaction product (Lever et al. 1987). The advantage of this approach compared to the previously published method is its greater radiochemical yield.

The non-subtype-selective opioid ligand buprenorphine (BPN) has also been labeled with  $^{11}\text{C}$ . The first published method involved the reaction of an *N*-(decyclopropylmethyl) precursor of buprenorphine with [ $^{11}\text{C}$ ]cyclopropanecarbonyl chloride followed by reduction with lithium aluminum hydride (Luthra et al. 1987). Later, a more facile synthesis of [ $^{11}\text{C}$ ]BPN was reported, in which the drug was labeled with  $^{11}\text{C}$  at the 6-methoxy position, using a methylation reaction with [ $^{11}\text{C}$ ]iodomethane (Lever et al. 1990). In a preclinical study in a baboon, the cerebral kinetics of [ $^{11}\text{C}$ ]BPN and [ $^{11}\text{C}$ ]DPN were directly compared. (–)Naloxone-sensitive binding of both ligands was observed in the striatum but not in the cerebellum. Uptake values and time courses in the target region were similar. The investigators concluded that [ $^{11}\text{C}$ ]DPN shows better kinetics than [ $^{11}\text{C}$ ]BPN since [ $^{11}\text{C}$ ]DPN is cleared more rapidly from brain areas devoid of opioid receptors (Shiue et al. 1991). [ $^{11}\text{C}$ ]BPN shows a regional distribution in the baboon brain consistent with regional opioid receptor density, whereas bound radioactivity in target areas can be displaced

by naloxone. Moreover, [ $^{11}\text{C}$ ]BPN binding in the striatum shows a good ( $\sim 10\%$ ) short-term test-retest variability (Galynker et al. 1996).

A non-subtype-selective agonist radioligand for opioid receptor imaging has become available. This compound, called [ $^{18}\text{F}$ ]FE-PEO, binds with  $K_i$  values of 0.4–1.6 nM to all opioid receptor subtypes. The distribution of radioactivity in the rat brain corresponds to regional opioid receptor levels, but nonspecific binding is rather high ( $\geq 33\%$  even in target areas). [ $^{18}\text{F}$ ]FE-PEO is relatively slowly metabolized (60% parent in rat plasma after 3 h), and radioactive metabolites do not enter the brain (Riss et al. 2013).

A benzamide radioligand for  $\mu$ -opioid receptors with agonist activity ([ $^{11}\text{C}$ ]AH7921) was recently labeled with  $^{11}\text{C}$  and tested in rats as a potential alternative to [ $^{11}\text{C}$ ]CFN (Rafique et al. 2017). Although this agent showed several drawbacks, such as a low binding potential and entry of a radioactive metabolite in the brain, the authors claim that it may be used for quantitative imaging of  $\mu\text{OR}$  availability and competition with endogenous opioids.

The first successful radioligand for  $\delta\text{OR}$  was the antagonist  $\text{N}1'-(^{11}\text{C})\text{methyl}$  naltrindole ([ $^{11}\text{C}$ ]MeNTI). [ $^{11}\text{C}$ ]MeNTI was synthesized by alkylation of 3-*O*-benzylnaltrindole with [ $^{11}\text{C}$ ]methyl iodide followed by removal of the protecting benzyl moiety. [ $^{11}\text{C}$ ]MeNTI binding in the brain of living mice was blocked (up to 75%) by naltrindole ( $\delta\text{OR}$  antagonist), but not by cyprodine ( $\mu\text{OR}$  antagonist) nor by U50,488 ( $\kappa\text{OR}$  agonist). Regional tracer uptake showed a linear, close correlation to  $\delta\text{OR}$  densities known from autoradiography. These data indicated that [ $^{11}\text{C}$ ]MeNTI binds selectively to  $\delta\text{OR}$  in the mouse brain in vivo (Lever et al. 1992).

An attempt to label the delta-1 subtype of  $\delta\text{OR}$  in the mouse brain with the selective ligand [ $^3\text{H}$ ]benzylidenenaltrexone (BNTX) was only partially successful. Although some specific binding of the compound was observed in vivo and tracer uptake could be blocked with the appropriate compounds, ratios of specific-to-nonspecific binding were very poor (Lever et al. 1996). Fluorinated analogs of naltrindole ( $\text{N}1'$ -fluoroethylnaltrindole or BU97001 and  $\text{N}1'$ -fluoroethyl-(14-formylamino)-naltrindole or BU97018) have been proposed for PET imaging. BU97001 has been tritiated and used successfully for autoradiography of the rat brain; it shows sub-nM affinity to  $\delta\text{OR}$  ( $K_d$  0.42 nM) (Tyacke et al. 2002). However, PET data for [ $^{18}\text{F}$ ]BU97001 have not yet been reported.

A  $^{11}\text{C}$ -labeled *N*-substituted quinolinimide derivative with much higher subtype selectivity than [ $^{11}\text{C}$ ]MeNTI has been tested as a potential ligand for in vivo imaging of  $\delta\text{OR}$ . The compound was rapidly degraded in both plasma and brain tissue after intravenous injection in rats, and specific binding could not be detected in vivo (Bourdier et al. 2007). A radiofluorinated  $\delta\text{OR}$  antagonist based on the Dmt-Tic pharmacophore has also been prepared. Although the compound showed significant binding to  $\delta\text{OR}$  in brain slices, it failed to cross the blood–brain barrier in living rats (Ryu et al. 2008). Three benzamide  $\delta\text{OR}$  agonists have been labeled with  $^{11}\text{C}$  and subjected to preclinical testing. All compounds showed a very low brain uptake and no regional specific binding in mice. One of the compounds, called [ $^{11}\text{C}$ ]SNC80, was also tested in a monkey. Results in this primate were similar to those seen

previously in mice, i.e., low blood–brain barrier permeability and a uniform distribution of radioactivity throughout the brain, indicating nonspecific binding (Pichika et al. 2010).

Recently, a conjugate of the  $\delta$ OR antagonist TIPP was labeled with  $^{64}\text{Cu}$  (Pirisedigh et al. 2017). This tracer was only tested by ex vivo autoradiography in rat brain sections. It showed a regional distribution corresponding to the known expression levels of the  $\delta$ OR.

Initial attempts at developing subtype-selective radioligands for  $\kappa$ OR were not successful.  $^{18}\text{F}$ -labeled analogs of U-50488 were prepared by fluoroalkyl substitution, but the labeling procedure resulted in a very strong ( $\geq 100$ -fold) loss of affinity with reference to the parent compound (Chesis and Welch 1990). A successful agonist radioligand for  $\kappa$ OR,  $[^{11}\text{C}]\text{GR}89696$ , was finally prepared in 1999 (Ravert et al. 1999). The R enantiomer, later called  $[^{11}\text{C}]\text{GR}103545$ , shows an excellent hypothalamus/cerebellum ratio of 11.4 in the mouse brain at 90 min after injection, in contrast to the S enantiomer which is inactive (target/nontarget ratio = 1) (Ravert et al. 2002). Using subtype-selective opioid receptor ligands in blocking studies, the binding of  $[^{11}\text{C}]\text{GR}103545$  was shown to be  $\kappa$ OR selective and saturable. Preclinical evaluation in baboons also indicated that  $[^{11}\text{C}]\text{GR}103545$  is a promising tracer for imaging of  $\kappa$ OR. A regional distribution of radioactivity is observed that is consistent with the regional density of  $\kappa$ OR. Naloxone pretreatment reduces tracer uptake in all brain regions to the level observed in the cerebellum. In vivo binding of the tracer is stereoselective, and significant washout occurs within the time frame of a PET scan (Talbot et al. 2005). A one-pot radiosynthesis for preparation of  $[^{11}\text{C}]\text{GR}103545$  with high specific radioactivity has recently been described (Nabulsi et al. 2011). Two other agonist radioligands for  $\kappa$ OR, FEKAP and EKAP, have recently been labeled with  $^{11}\text{C}$  and evaluated in non-human primates (Li et al. 2019a; Li et al. 2019b). Both ligands show sub-nM affinity to  $\kappa$ OR, reasonable to excellent selectivity for  $\kappa$ OR over  $\mu$ OR and  $\delta$ OR, a fairly rapid metabolism in monkey plasma, fast and reversible kinetics in the brain and regional binding potential values consistent with binding to  $\kappa$ OR. Blocking and receptor occupancy studies were also performed, using the opioid drugs naloxone, LY2795050 and LY2456302. Thus, these ligands seem suitable for in vivo studies of cerebral  $\kappa$ OR.

Antagonist radioligands for  $\kappa$ OR have also been proposed.  $[^{11}\text{C}]\text{MeJDTic}$  was prepared by methylation of the precursor JDTic with  $[^{11}\text{C}]\text{methyl triflate}$ . In a pre-clinical study in mice, high uptake of the tracer was noted in target organs (lung, hypothalamus). Blocking and displacement studies with nonradioactive kappa, mu, and delta ligands suggested that  $[^{11}\text{C}]\text{MeJDTic}$  binds specifically to  $\kappa$ OR in the brain of living mice in vivo (Poisonel et al. 2008). Yet, no follow-up studies with  $[^{11}\text{C}]\text{MeJDTic}$  have been reported. A fluoropropyl derivative of JDTic was later labeled with  $^{18}\text{F}$  and evaluated in intact mice. This tracer failed because radioactive metabolites were formed that entered the brain and significant specific binding in target regions could not be detected (Schmitt et al. 2017). Another antagonist,  $[^{11}\text{C}]\text{LY}2795050$ , was evaluated in Sprague–Dawley rats, wild-type and  $\kappa$ OR knockout mice, and rhesus monkeys. Specific binding of the probe to cerebral  $\kappa$ OR

was observed; this binding was completely blocked by pretreatment of monkeys with naloxone (1 mg/kg) and dose-dependently inhibited by the selective  $\kappa$ OR antagonist LY2456302. Tracer metabolism in primates is moderate (40% parent remaining in monkey plasma at 30 min), and tracer pharmacokinetics appears favorable; thus, [ $^{11}\text{C}$ ]LY2795050 is the first successful antagonist radioligand for  $\kappa$ OR (Zheng et al. 2013).

Still another  $\kappa$ OR antagonist, LY2459989, was labeled in recent years, both with  $^{11}\text{C}$  (Zheng et al. 2014) and  $^{18}\text{F}$  (Li et al. 2018). The two tracers show sub-nM affinity to  $\kappa$ OR and high selectivity for  $\kappa$ OR over  $\mu$ OR and  $\delta$ OR. They have been evaluated in rats and rhesus monkeys. A rapid uptake in the brain is observed and regional binding potential values are consistent with the known distribution of  $\kappa$ OR in primates. Pretreatment of animals with naloxone results in a homogeneous background radioactivity throughout the brain, and tracer uptake in target regions is dose-dependently reduced by the selective  $\kappa$ OR antagonists LY2456302 and unlabeled LY2459989. Thus, [ $^{11}\text{C}$ ]- and [ $^{18}\text{F}$ ]LY2459989 should also be considered as useful  $\kappa$ OR ligands.

The last subclass of the opioid system for which imaging probes have been developed is the nociceptin/orphanin or opiate-like receptor (ORL1). [ $^{11}\text{C}$ ]Methyl-Ro64-6198 failed as a radioligand for this subtype since its *in vivo* binding in the mouse brain was largely nonspecific (Ogawa et al. 2001). A second probe, [ $^{11}\text{C}$ ]CPEB, showed some saturable *in vivo* binding but lack of specificity to ORL1 (Ogawa et al. 2003).

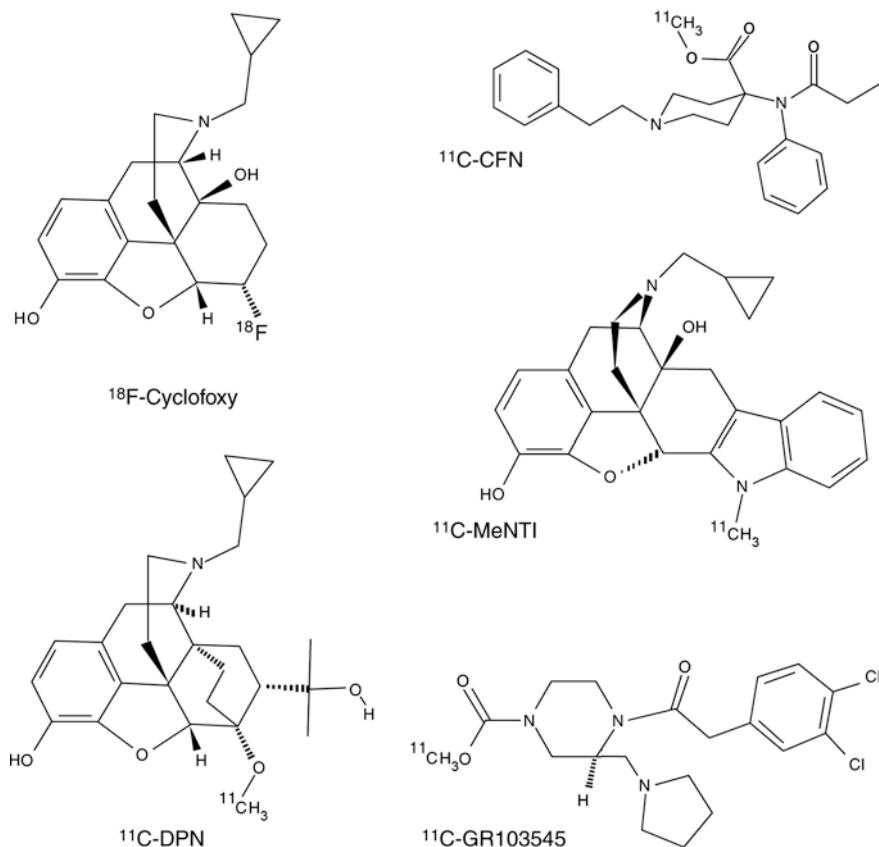
The first successful ORL1 ligand was [ $^{11}\text{C}$ ]NOP-1A, a specific antagonist with sub-nM affinity (0.15 nM). This compound was selected from a library of ORL1 antagonists developed by Eli Lilly & Co. (Pedregal et al. 2012). After intravenous administration of [ $^{11}\text{C}$ ]NOP-1A to rhesus monkeys, a regional cerebral distribution of radioactivity is observed consistent with binding of the probe to its target. Brain uptake is reduced by 50–70% after pretreatment of animals with the ORL1 antagonist SB-612111 (Pike et al. 2011; Kimura et al. 2011). The regional distribution of [ $^{11}\text{C}$ ]NOP-1A in the human brain resembles that in the monkey brain. Regional distribution volume of the tracer can be reliably estimated using a two-tissue compartment model with arterial sampling or Ichise's noncompartmental bilinear analysis (Lohith et al. 2012).

Another ORL1 antagonist for PET imaging has been developed by Merck. Their compound [ $^{18}\text{F}$ ]MK-0911 shows a regional distribution in the monkey and human brain consistent with the regional expression of ORL1. Binding in the monkey brain is inhibited after pretreatment of animals with several structurally diverse ORL1 antagonists (MK-0584, MK-0337, MK-5757). Dose-dependent occupancy of ORL1 in the human brain by nonradioactive MK-5757 could be assessed with [ $^{18}\text{F}$ ]MK-0911 and PET (Hostetler et al. 2013).

## 21.3 Characteristics of Widely Used Radioligands

### 21.3.1 Cyclofoxy

[ $^3\text{H}$ ]Cyclofoxy (Fig. 21.1), the parent compound of [ $^{18}\text{F}$ ]cyclofoxy, labels both  $\mu\text{OR}$  and  $\kappa\text{OR}$  in autoradiography of the rat and guinea pig brain (McLean et al. 1987; Rothman and McLean 1988). The regional pattern of radioactivity in the rat brain after in vivo injection of [ $^3\text{H}$ ]cyclofoxy corresponds to the regional distribution of these opioid receptor subtypes (Rothman and McLean 1988; Ostrowski et al. 1987), and brain radioactivity is reduced to background levels after pretreatment of animals with unlabeled naloxone (Ostrowski et al. 1987).  $K_i$  values of (–)cyclofoxy for  $\mu\text{OR}$ ,  $\kappa\text{OR}$ , and  $\delta\text{OR}$  in rat brain membranes are 2.6, 9.3, and 89 nM, respectively, whereas the (+)enantiomer is essentially inactive ( $K_i$  values  $> 10 \mu\text{M}$ ) (Rothman et al. 1988). Consistent with these in vitro data, uptake of the radiofluorinated (–) enantiomer in the rat brain shows striking regional differences consistent with tracer



**Fig. 21.1** Chemical structures of widely used opioid radioligands

binding to opioid receptors, whereas the tritiated (+)enantiomer is rather homogeneously distributed and shows low brain uptake. Pretreatment of animals with unlabeled (-)cyclofoxy reduces the uptake of [ $^{18}\text{F}$ ](-)cyclofoxy in all regions to the level of the tritiated (+) enantiomer (Kawai et al. 1990). Thus, [ $^{18}\text{F}$ ]cyclofoxy visualizes  $\mu\text{OR}$  and  $\kappa\text{OR}$  in the living brain.

Kinetic analysis of [ $^3\text{H}$ ]cyclofoxy uptake in the rat brain suggested an in vivo  $K_d$  of 2.1–5.2 nM in different brain areas and a  $B_{\text{max}}$  ranging from 1 pmol/g in the cerebellum to 78 pmol/g in the thalamus. Binding was reversible on a PET time scale (Sawada et al. 1991). In a follow-up study, a combination of bolus injection and constant infusion of [ $^{18}\text{F}$ ]cyclofoxy was applied to estimate the receptor-ligand parameters in a condition of “true” tissue-blood equilibrium, which was reached after 60 min.  $K_d$  values of 1.4–2.9 nM were measured and  $B_{\text{max}}$  in target areas ranged from 15 to 74 pmol/g tissue. Rodent cerebellum could not be used as a reference region since the level of nonspecific binding in the cerebellum was different from that in other parts of the brain (Kawai et al. 1991). Bolus plus continuous infusion of [ $^{18}\text{F}$ ]cyclofoxy was also advantageous in baboons, as it resulted in shorter scan times, a simplified protocol for blood sampling, and more accurate receptor measurements (Carson et al. 1993).

### 21.3.2 CFN (Fig. 21.1)

In vitro binding assays in human thalamic membranes indicated a  $K_d$  of 0.08 nM for [ $^3\text{H}$ ]CFN (Titeler et al. 1989). In the rat brain,  $K_i$  values of 0.051, 4.7, and 13 nM were measured at  $\mu\text{OR}$ ,  $\delta\text{OR}$ , and  $\kappa\text{OR}$ , respectively (Frost et al. 1985). For the guinea pig brain, the corresponding values were 0.024, 3.28, and 43.1 nM (Cometta-Morini et al. 1992). In vitro competition studies using opioid ligands with a wide range of subtype selectivities confirmed that [ $^3\text{H}$ ]CFN selectively labels  $\mu\text{OR}$ , in both the rat and human brain (Titeler et al. 1989). Thus, CFN is a potent and selective  $\mu\text{OR}$  agonist. Later in vitro binding studies on rat brain sections demonstrated that [ $^{11}\text{C}$ ]CFN has higher affinity for the  $\mu_1$  subtype than for  $\mu_2$ . Small animal imaging studies in which the  $\mu\text{OR}$  ligand cyprodime and the  $\mu_1$ -subtype-selective ligand naloxonazine were used have indicated that the in vivo binding of [ $^{11}\text{C}$ ]CFN involves only the  $\mu_1$  subtype, whereas binding to  $\mu_2$  receptors cannot be detected (Eriksson and Antoni 2015). Thus, [ $^{11}\text{C}$ ]CFN images of the human brain may primarily show the distribution of the  $\mu_1$  subtype.

A study which applied compartment modeling for the analysis of [ $^{11}\text{C}$ ]CFN binding in the human brain showed that the occipital cortex can be considered as a region with low or negligible receptor density. Target region-to-occipital cortex ratios of radioactivity at late time points can be used instead of ratios of  $k_3/k_4$  when tracer kinetic modeling is not feasible, e.g., in clinical research protocols (Frost et al. 1989). However, Logan graphical analysis or a simplified reference tissue model approach can provide more accurate estimates of [ $^{11}\text{C}$ ]CFN binding to  $\mu\text{OR}$  in the human brain than the ratio method (Endres et al. 2003). A test-retest study which assessed [ $^{11}\text{C}$ ]CFN binding in the brain of healthy volunteers twice, on a



single day, indicated that both the regional two-tissue compartment model distribution volume ( $V_T$ ) of the tracer and binding potential ( $BP_{ND}$ ) are reproducible (variability  $<6$  and  $<10\%$ , respectively) (Hirvonen et al. 2009).

Administration of [ $^{11}C$ ]CFN can induce somnolence and sedation and, at higher dose, respiratory arrest. These pharmacological effects are related to the fact that CFN is an opioid agonist. The upper limit for the administered mass of CFN is about  $0.1 \mu\text{g}/\text{kg}$  (Frost 2001). [ $^{11}C$ ]CFN binding in the human brain is affected by polymorphisms of the  $\mu\text{OR}$  gene. Carriers of the G allele (Asn40Asp variant, which may contribute to the development of alcohol dependence) have lower global binding potential than subjects homozygous for the A allele (Weerts et al. 2013).

Radiofluorinated analogs of carfentanil with subnanomolar affinities to  $\mu\text{OR}$  have been prepared (Henriksen et al. 2005a). *N*-[4-(methoxymethyl)-1-[2-(2-thienyl)ethyl]-4-piperidinyl]-*N*-phenyl-2-( $\pm$ )-[ $^{18}F$ ]fluoropropanamide ([ $^{18}F$ ]sufentanil) showed a high brain uptake in living mice and a distribution of radioactivity corresponding to regional  $\mu\text{OR}$  expression. The compound was rapidly metabolized in plasma (only 21% parent remaining at 40 min), but the metabolites appeared to not enter the brain in significant amounts (Henriksen et al. 2005b).

An updated (modernized) synthetic procedure for the preparation of [ $^{11}C$ ]CFN (by reaction of a carboxylate precursor with [ $^{11}C$ ]methyltriflate) has recently been described (Blecha et al. 2017). This procedure has the additional advantage that organic solvents could be replaced by ethanol.

### 21.3.3 DPN (Fig. 21.1)

Another widely applied radioligand for opioid receptor mapping is [ $^{11}C$ ]DPN. [ $^3H$ ]DPN binds with subnanomolar ( $K_d$  0.22–0.23 nM) affinity to opioid receptors in the rat brain (Chang et al. 1981) and in the human frontal cortex (Pfeiffer et al. 1982).  $K_i$  values of DPN for  $\mu\text{OR}$ ,  $\delta\text{OR}$ , and  $\kappa\text{OR}$  in the rat brain are 0.20, 0.18, and 0.47 nM, respectively (Chang et al. 1981). Thus DPN is a non-subtype-selective opioid antagonist. Automated procedures for radiosynthesis of [ $^{18}F$ ]FDPN and [ $^{11}C$ ]DPN have been described (Schultz et al. 2013; Fairclough et al. 2014).

Using RB101, an inhibitor of enkephalin-degrading enzymes, the *in vivo* binding of [ $^3H$ ]DPN to opioid receptors in the mouse brain was shown to be sensitive to competition by endogenous enkephalins. Although RB101 is devoid of affinity to opioid receptors, the compound dose-dependently inhibits [ $^3H$ ]DPN binding in the unstressed mouse brain up to a maximum of 30%. When mice were stressed by a warm-swim test, a greater inhibition of [ $^3H$ ]DPN binding by RB101 was observed (maximally 45%). These effects of RB101 are probably related to increased levels of endogenous enkephalins (Ruiz-Gayo et al. 1992).

A validation study using [ $^3H$ ]DPN in rats showed that compartment modeling of PET data with the cerebellum as reference region can provide accurate measurements of regional opioid receptor  $B_{max}$  and  $K_d$ , although  $K_d$  values estimated *in vivo* are an order of magnitude higher than those determined *in vitro*, probably due to diffusion gradients in tissue or protein binding of the radioligand (Cunningham

et al. 1991). However, the cerebellum cannot be considered as a region devoid of opioid receptors in human studies since  $\mu$ OR and  $\kappa$ OR are expressed at low levels in this area of the human brain (Schadrack et al. 1999).

The kinetics of [ $^{11}$ C]DPN and [ $^{11}$ C]CFN in the brain of human volunteers have been directly compared (Frost et al. 1990; Villemagne et al. 1994). Binding of both ligands in the thalamus (an area containing mainly  $\mu$ OR) was not significantly different, but [ $^{11}$ C]DPN showed significantly greater binding than [ $^{11}$ C]CFN in the striatum, cingulate, and frontal cortex. Dissociation of [ $^{11}$ C]CFN from its binding sites was observed, but dissociation of [ $^{11}$ C]DPN was hardly noticeable. [ $^{11}$ C]CFN binding reached a plateau, whereas [ $^{11}$ C]DPN uptake showed a steady increase (Frost et al. 1990). Total radioactivity in the human brain (SUV) and the ratio of total/nonspecific binding were twice as great after administration of [ $^{11}$ C]DPN than [ $^{11}$ C]CFN. The dose of naloxone required to fully block [ $^{11}$ C]DPN binding was ten times greater than that required for [ $^{11}$ C]CFN (Villemagne et al. 1994). Thus, [ $^{11}$ C]DPN appeared to label  $\delta$ OR and KOR in addition to the  $\mu$ OR which are selectively labeled by [ $^{11}$ C]CFN. In a later animal study, the sensitivity of [ $^{11}$ C]CFN and [ $^3$ H]DPN binding to endogenous opioid release was directly compared, opioid release being stimulated by administering amphetamine and methadone to rats. The binding of [ $^{11}$ C]CFN but not [ $^3$ H]DPN was significantly reduced after such challenges. In vitro binding and subcellular fractionation studies indicated that this may be due to the fact that [ $^{11}$ C]CFN binds preferentially to receptors on the cell surface, whereas [ $^3$ H]DPN binds to both surface and internalized receptors. Thus, [ $^{11}$ C]CFN may be a better ligand than [ $^{11}$ C]DPN for in vivo imaging of endogenous opioid peptide release (Quelch et al. 2014).

Both the regional density ( $B_{\max}$ ) and radioligand affinity ( $K_d$ ) of opioid receptors can be assessed by applying a two-injection protocol, in which [ $^{11}$ C]DPN is administered at high (20–160 TBq/mmol) and low (1–3 TBq/mmol) specific radioactivities (Sadzot et al. 1991) or by a pulse-chase protocol (Jones et al. 1994a). Parametric images of [ $^{11}$ C]DPN binding can be prepared, using both conventional spectral analysis and rank shaping (Hammers et al. 2007a).

Radiofluorinated analogs of DPN have been prepared. *N*-(3-[ $^{18}$ F]Fluoropropyl)-*N*-nordiprenorphine (FPND) showed rather limited brain uptake (Bai et al. 1990) but a striatum/cerebellum ratio of  $3.3 \pm 0.7$  at 30 min postinjection which was reduced to unity after pretreatment of animals with naloxone (1 mg/kg). The tracer was extensively metabolized (<25% parent remaining in blood at 30 min), but not significantly defluorinated (Chesis et al. 1990). 6-*O*-(3-[ $^{18}$ F]Fluoroethyl)-6-*O*-desmethyldiprenorphine ([ $^{18}$ F]DPN) showed a regional binding pattern in the human brain similar to that of [ $^{11}$ C]DPN (Wester et al. 2000). FDPN binds to  $\mu$ OR,  $\delta$ OR, and  $\kappa$ OR in rat brain slices like the parent tracer, DPN (Wester et al. 2000). A scan duration of at least 90 min is required for reliable estimates of tracer distribution volume (Boecker et al. 2005). If metabolite-corrected plasma data are used as input function, individualized metabolite correction should be performed since women metabolize FDPN significantly faster than men (Henriksen et al. 2006).

### 21.3.4 MeNTI

$K_i$  values of MeNTI (Fig. 21.1) at  $\delta$ OR,  $\mu$ OR, and  $\kappa$ OR in guinea pig brain membranes are 0.02, 14, and 65 nM, respectively (Portoghese et al. 1990). Thus, MeNTI is a very potent and subtype-selective  $\delta$ OR antagonist.

In the human brain, [ $^{11}\text{C}$ ]MeNTI shows regional kinetics concordant with selective binding of the tracer to  $\delta$ OR. Prolonged retention of radioactivity is observed in receptor-rich areas, whereas the ligand is rapidly washed out from nontarget areas of the brain. The regional distribution of radioactivity is closely correlated with local  $\delta$ OR densities but not with regional densities of  $\mu$  or  $\kappa$  binding sites. Tracer uptake is blocked after pretreatment of volunteers with naltrexone (up to 73%) (Madar et al. 1996).

[ $^{11}\text{C}$ ]MeNTI shows irreversible binding characteristics in the human brain during a 90-min scanning period. A constrained three-compartment kinetic model is superior to other data-analytical techniques for quantification of the in vivo binding of the tracer to  $\delta$ OR (Smith et al. 1999).

### 21.3.5 GR103545

The subtype selectivity of [ $^{11}\text{C}$ ]GR103545 (Fig. 21.1) has been determined in HEK-239 cells expressing human  $\kappa$ OR, CHO-K1 cells expressing human  $\mu$ OR, and CHO cells expressing human  $\delta$ OR.  $K_i$  values at the three OR subtypes were 0.02, 16.2, and 536 nM, respectively (Schoultz et al. 2010). Thus, [ $^{11}\text{C}$ ]GR103545 is an extremely potent and subtype-selective  $\kappa$ OR agonist.

In a preclinical study in awake rhesus macaques, [ $^{11}\text{C}$ ]GR103545 showed a regional volume of distribution in the brain which corresponded to regional  $\kappa$ OR densities (Schoultz et al. 2010). In a more recent study, the use of a bolus-plus-continuous-infusion protocol was shown to permit estimation of both  $B_{\max}$  and  $K_d$  of  $\kappa$ OR in the monkey brain in vivo (Tomasi et al. 2013). The cerebellum was used as the reference region in kinetic analysis.

A low injected mass of [ $^{11}\text{C}$ ]GR103545 is essential since this  $\kappa$ OR agonist can induce respiratory depression, reductions of heart rate, and sedation even at very low administered doses of 0.3  $\mu\text{g}/\text{kg}$ . Based on the  $K_d$  value estimated for the radioligand in nonhuman primates, a dose of 1.4  $\mu\text{g}$  (3.38 nmol) was estimated as an acceptable dose limit in human studies for subjects with a body mass of 70 kg (Tomasi et al. 2013). In a preclinical study in rats, [ $^{11}\text{C}$ ]GR103545 was found to be suitable for estimation of the dose-dependent occupancy of the cerebral  $\kappa$ OR population by salvinorin A (Placzek et al. 2015).

A first-in-man study with [ $^{11}\text{C}$ ]GR103545 has been performed. All subjects were scanned at baseline. Seven subjects were also scanned 75 min after oral administration of naltrexone and six subjects at two time points (1.5 h and 8 h) after intake of the selective  $\kappa$ OR antagonist PF-04455242. Drug treatment resulted in a decrease of tracer distribution volume ( $V_T$ ) in all brain regions. A two-tissue compartment model with multilinear analysis method was identified as the method of choice for

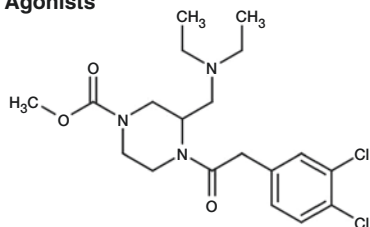
estimation of  $V_T$ . The in vivo affinity of the radiotracer for its target was estimated as 0.069 nM. [ $^{11}\text{C}$ ]GR103545 seems suitable for quantification of regional  $\kappa\text{OR}$  density and  $\kappa\text{OR}$  occupancy in the human brain, although the radioligand has slow kinetics and variability in the estimated model parameters is rather high (Naganawa et al. 2014a).

### 21.3.6 LY2795050

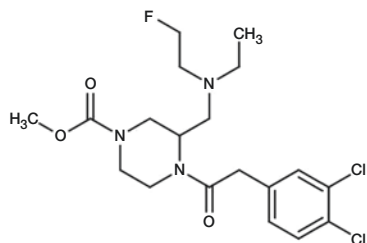
LY2795050 (Fig. 21.2) is an opioid antagonist with  $K_i$  values of 0.72, 25.8 and 153 nM at cloned human  $\kappa$ ,  $\mu$  and  $\delta$  receptors (Zheng et al. 2013). In vivo competition studies in the monkey brain confirmed that [ $^{11}\text{C}$ ]LY2795050 binds specifically and selectively to  $\kappa\text{OR}$ . Although the in vivo selectivity of the ligand for  $\kappa\text{OR}$  over  $\mu\text{OR}$  is smaller than that observed in vitro (7.6 rather than 36), the PET signal of a region expressing similar densities of  $\kappa\text{OR}$  and  $\mu\text{OR}$  will predominantly (for 88%) reflect the presence of the  $\kappa\text{OR}$  (Kim et al. 2013).

Blocking and kinetic modeling studies of [ $^{11}\text{C}$ ]LY2795050 binding in the human brain indicated regional distribution volume ( $V_T$ ) values corresponding to known regional densities of the  $\kappa\text{OR}$  (Naganawa et al. 2014b). Since a reference region without specific binding is lacking in human brain, the authors proposed the use of a fixed value for  $V_T$  in cerebellum in model fits. This approach assumes that binding of the ligand in cerebellum is unaltered under the study conditions. A later

#### Agonists

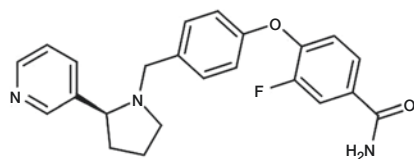


#### EKAP

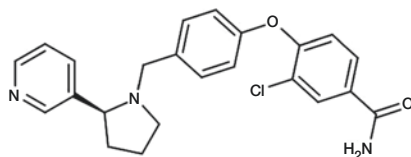


#### FEKAP

#### Antagonists



#### LY2459989



#### LY2795050

**Fig. 21.2** Structures of some  $\kappa\text{OR}$  agonists and antagonists that have been radiolabeled for PET imaging

publication examined the reproducibility of [ $^{11}\text{C}$ ]LY2795050 scans in human brain, using a two-tissue compartment model fit with fixing of cerebellar  $V_T$ . The test-retest reproducibility of  $V_T$  in all brain regions was better than 10% with exception of the amygdala where it was 12%. In contrast to  $V_T$ , binding potential values showed a good reproducibility in regions with moderate or high receptor densities ( $\text{BP}_{\text{ND}} > 0.4$ ) but a poor reproducibility in areas with small receptor populations (Naganawa et al. 2015).

An automated GMP procedure for preparation of [ $^{11}\text{C}$ ]LY2795050 has been described (Yang et al. 2018).

### 21.3.7 LY2459989

LY2459989 (Fig. 21.2) is an opioid antagonist with  $K_i$  values of 0.18, 7.68, and 91.3 nM at cloned human  $\kappa$ ,  $\mu$ , and  $\delta$  receptors (Zheng et al. 2014). Ex vivo biodistribution studies in rodents and imaging studies in monkeys have confirmed that [ $^{11}\text{C}$ ]LY2459989 binds specifically to  $\kappa\text{OR}$ . Pretreatment of monkeys with naloxone (1 mg/kg) results in a uniform distribution of radioactivity in the brain and tracer uptake is dose-dependently reduced after pretreatment of animals with the subtype-selective  $\kappa\text{OR}$  antagonists LY2456302 and unlabeled LY2459989. The observed binding potential values of the tracer are high, and regional binding potential is an accurate reflection of known expression levels of the  $\kappa\text{OR}$  (Zheng et al. 2014).

In a recent paper, sensitivities of the binding of [ $^{11}\text{C}$ ]GR103545 and [ $^{11}\text{C}$ ]LY2459989 in rat brain to competition by unlabeled  $\kappa\text{OR}$  antagonists and agonists were directly compared. The binding potential values of both tracers were similarly affected when animals were pretreated with kappa antagonists (naloxone, naltrexone, LY2795050, JDTic, nor-BNI) and also after treatment of rats with the kappa agonists butorphan and GR89696. However, pretreatment of animals with the  $\kappa\text{OR}$  agonists salvinorin A and U-50488 resulted in a strong decrease of [ $^{11}\text{C}$ ]GR103545 binding, but did not affect the binding potential of [ $^{11}\text{C}$ ]LY2459989 (Placzek et al. 2019). This surprising discrepancy suggests that agonists may interact in different ways with  $\kappa\text{OR}$  and the target occupancy of some  $\kappa\text{OR}$  agonists cannot be detected with [ $^{11}\text{C}$ ]LY2459989.

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## 21.4 PET Studies in Healthy Volunteers

### 21.4.1 Influence of Gender, Hormonal Status, and Age

[ $^{11}\text{C}$ ]CFN-PET has been applied for the study of changes of opioid receptor binding in the brain of women during the menstrual cycle. No significant differences in tracer binding were noted between the follicular and luteal phases of the cycle, but follicular levels of estradiol in the circulation were negatively correlated with [ $^{11}\text{C}$ ]CFN binding potential in the amygdala and hypothalamus. These results, combined with measurements of luteinizing hormone pulse, suggested that  $\mu\text{OR}$  in the

amygdala modulate gonadal steroid hormone pulsatility, whereas circulating estradiol regulates  $\mu$ OR function in the brain (Smith et al. 1998). Neuroimaging has shown that black cohosh, a herbal extract used to treat menopausal symptoms, affects the endogenous opioid receptor system. Significant increases in [ $^{11}$ C]CFN binding potential (10–61%) were noted in the cortical areas, thalamus, and nucleus accumbens of the brain of postmenopausal women (Reame et al. 2008). In the thalamus and nucleus accumbens, similar effects were observed after estrogen replacement therapy (Smith et al. 2006).

A PET study with [ $^{11}$ C]CFN, involving 50 subjects, indicated that the cerebral binding of [ $^{11}$ C]CFN is affected both by gender and by age. Binding potential of the radioligand increased with age in the neocortex and putamen. Women showed higher [ $^{11}$ C]CFN binding than men of the same age in several cortical and subcortical areas; however, in postmenopausal women,  $\mu$ OR binding in the thalamus and amygdala declined to levels lower than in age-matched men (Zubieta et al. 1999). A PET study with [ $^{18}$ F]cyclofoxy confirmed that opioid receptor binding in the thalamus of females is lower than in males, both in healthy aged subjects and in patients with Alzheimer disease (Cohen et al. 2000). MRI-based partial volume correction is required for the accurate assessment of receptor densities in small structures of the human brain. Applying such correction to [ $^{11}$ C]CFN data for the left temporal cortex, binding of the radioligand appeared to increase with age (0.9% per year), consistent with  $B_{\max}$  values from postmortem autoradiography. Without partial volume correction, the effect of aging could not be observed (Bencherif et al. 2004a).

Correlations between regional serotonin transporter and  $\mu$ OR binding were examined in a PET study involving 21 healthy volunteers. The subjects were first injected with the 5-HT transporter tracer [ $^{11}$ C]MADAM. When radioactivity from the first injection had decayed, the  $\mu$ OR tracer [ $^{11}$ C]carfentanil was injected and a second PET scan was made, with the subject's brain in exactly the same position in the camera. Voxel-based correlation analysis was used to identify brain regions with positive correlations between serotonin transporter and  $\mu$ OR binding. Significant correlations were observed in anteromedial thalamus, dorsal anterior cingulate cortex, dorsolateral prefrontal cortex, amygdala, left parietal cortex, and medial temporal cortex. This finding may indicate that in these regions, endogenous opioids are modulating serotonin release from nerve endings (Tuominen et al. 2014).

A PET study with the  $\kappa$ OR ligand [ $^{11}$ C]LY2795050 in 27 healthy volunteers reported opposite sex differences in  $\kappa$ OR availability as were reported for total opioid receptor binding with [ $^{11}$ C]CFN. Men had significantly higher  $V_T$  values than women in several brain regions (Vijay et al. 2016).

### 21.4.2 Feeding

An interesting PET study in ten healthy male volunteers examined whether feeding releases endogenous opioids in the brain and if the magnitude of this response is related to the pleasure of eating. Each volunteer was scanned three times with [ $^{11}$ C]CFN: (i) after an overnight fast, (ii) after an overnight fast followed by a

pleasant meal (preferred type of pizza), and (iii) after an overnight fast plus the consumption of an unpleasant meal (nutrient drink). Feeding was associated with a significant release of endogenous opioids (a decrease of radioligand binding potential throughout the brain), but this release was more pronounced after the unpleasant than after the pleasant meal. Thus, opioid release appears to be not directly related to the subjective pleasure but rather to metabolic and homeostatic responses of the body to food consumption (Tuulari et al. 2017).

### 21.4.3 Personality Traits

Several PET studies have explored relationships between opioid receptor availability in the human brain and personality traits. Using the tracer [ $^{18}\text{F}$ ]DPN, German investigators scanned 23 healthy male volunteers under resting conditions. Subsequently, the subjects were psychologically tested for novelty seeking, harm avoidance, reward dependence, and persistence. A significant correlation was detected between the score for reward dependence (a personality trait predisposing for addictive behavior) and tracer binding potential in the bilateral ventral striatum with nucleus accumbens (Schreckenberger et al. 2008).

A later study from the USA involved scanning of 19 young healthy male volunteers with the  $\mu\text{OR}$  ligand [ $^{11}\text{C}$ ]CFN and PET. Subjects were scanned both at baseline and during receipt of a painful stress challenge (infusion of hypertonic saline solution in the left masseter muscle). High scores for impulsiveness and low deliberation scores were associated with significantly higher baseline  $\mu\text{OR}$  availability and greater stress-induced endogenous opioid system activation, measured as a reduction of [ $^{11}\text{C}$ ]CFN binding potential. These differences were noted in several brain areas (prefrontal and orbitofrontal cortices, anterior cingulate, thalamus, nucleus accumbens, and basolateral amygdala). Thus,  $\mu\text{OR}$  binding potential appears to predict an individual's vulnerability to risky behaviors, such as the development of substance abuse (Love et al. 2009).

A study from Finland measured brain  $\mu\text{OR}$  availability with [ $^{11}\text{C}$ ]CFN under resting conditions in 22 healthy volunteers: 10 individuals scoring in the upper and 12 individuals scoring in the lowest quartile for harm avoidance. A high score for harm avoidance (particularly in the subscales shyness with strangers, fatigability, and asthenia) was associated with high  $\mu\text{OR}$  availability (or low endogenous  $\mu$ -opioid drive) in the anterior cingulate cortex, ventromedial and dorsolateral prefrontal cortices, and anterior insular cortex. Thus, high  $\mu\text{OR}$  availability may be related to vulnerability for affective disorders (Tuominen et al. 2012). In a later study from the same group, 49 healthy volunteers were scanned with [ $^{11}\text{C}$ ]CFN, and the avoidance and anxiety dimensions of their attachment were measured with the Experiences in Close Relationships-Revised scale. In this study, the avoidance dimension was found to be negatively (not positively) correlated with  $\mu\text{OR}$  availability (i.e., [ $^{11}\text{C}$ ]CFN binding potential) in the thalamus, anterior cingulate cortex, frontal cortex, amygdala, and insula, whereas the anxiety dimension was not correlated with  $\mu\text{OR}$  availability in any brain region. The authors suggested that

inter-individual differences in  $\mu$ OR availability may be related to differences in avoidant attachment style (Nummenmaa et al. 2015). In a study from the following year, measured [ $^{11}$ C]CFN binding potentials were compared to individual differences in behavioral dispositions when subjects encountered signals of reward and harm. Sensitivity of the behavioral activation system but not the behavioral inhibition system was found to be positively associated with [ $^{11}$ C]CFN binding potential in frontal cortex, amygdala, ventral striatum, brainstem, cingulate cortex, and insula. Particularly the score for “fun seeking” was strongly associated with  $\mu$ OR availability (Karjalainen et al. 2016).

In an American PET study with [ $^{11}$ C]DPN in 12 healthy subjects, a high score of neuroticism was found to be associated with high opioid receptor binding in the right anterior insula. On the other hand, a high score of extraversion was associated with low tracer binding in the left posterior insula (Rodman et al. 2017).

#### 21.4.4 Affective Responses

The opioid system appears to play an important role in the regulation of affective responses. An initial PET study examined [ $^{11}$ C]CFN binding and blood flow (uptake of  $^{15}$ O-water) in the brain of 12 healthy male volunteers, which viewed either aversive images (facial mutilation, wounds, dead bodies) or neutral images (common objects, benign scenes, normal faces). Higher baseline  $\mu$ OR binding potential in the left temporal pole of the subjects was associated with lower blood flow in this region during presentation of the aversive emotional stimuli. This finding was interpreted as evidence for an inhibitory or anxiolytic role of the opioid system in limbic regions of the brain (Liberzon et al. 2002).

A second study examined healthy female volunteers, using [ $^{11}$ C]CFN and PET. Sustained neutral and sadness states were elicited by the recall of an autobiographical event associated with these emotions. Sustained sadness caused an increase of  $\mu$ OR availability in the rostral anterior cingulate, ventral pallidum, amygdala, and inferior temporal cortex which was interpreted as reflecting a reduction of the release of endogenous opioids due to deactivation of opioid neurotransmission (Zubieta et al. 2003).

Whereas sustained sadness in healthy volunteers is associated with a reduction in the release of endogenous opioids, positive emotion appears to be associated with an increase of that release. An interesting study from London scanned 25 healthy male volunteers with [ $^{11}$ C]DPN and PET, both during positive mood induction (sexually oriented video clip, listening to favorite music, reading positive statements, unexpected gift of 30 GBP) and during neutral mood (nature movie, listening to unknown classical music, reading neutral statements, no gift). Significant reductions of [ $^{11}$ C]DPN binding were noted in the hippocampus during positive mood induction, and the magnitude of positive mood change was negatively correlated with DPN binding in the amygdala (Koepp et al. 2009). An interesting PET study with [ $^{11}$ C]CFN evaluated the response of the  $\mu$ -opioid system to social feedback. Eighteen subjects selected at least 40 profiles of preferred sex-individuals from a



database of 500. During a subsequent scanning session with [ $^{11}\text{C}$ ]CFN, images of the faces of these individuals and their responses to the question: Would you like me? (Definitely no, or very likely yes) were presented to them on a computer screen (35 min of acceptance followed by 35 min of rejection, or vice versa, in counterbalanced order). During a separate scanning session with [ $^{11}\text{C}$ ]CFN, the same subjects were scanned at baseline. The human faces on the screen were then replaced by gray blocks and in stead of feedback, the screen said: “non applicable.” Social rejection was found to be associated with significant opioid release in the ventral striatum, amygdala, midline thalamus, and periaqueductal gray, a response that may be aimed at reducing the experience of social pain. This interpretation was supported by the observation that a higher score for the character trait resiliency was associated with greater opioid release in the amygdala, periaqueductal gray and subgenual anterior cingulate cortex (ACC), whereas reduced negative feelings during rejection were associated with greater opioid release in the pregenual ACC. In contrast to social rejection, acceptance resulted in activation of opioid release in the amygdala and anterior insula but deactivation in the midline thalamus and subgenual ACC. Opioid release in the left ventral striatum during acceptance was associated with the desire for social interaction. Thus, neurotransmission via  $\mu\text{OR}$  may be involved in the preservation and promotion of emotional well-being of human subjects in their social environment (Hsu et al. 2013).

A PET study with [ $^{11}\text{C}$ ]CFN examined the effect of social touch on the activity of the endogenous opioid system. Eighteen male volunteers were scanned twice with this tracer. During the baseline scan, the subjects were left alone, whereas during the social touch scan, their partners were present and were caressing them in a non-sexual way. The scans were separated by an interval of 2 h, and the order of the scans was counterbalanced, i.e., in half of the subjects, the baseline scan was made first, whereas in the other half, it was made last. The subjects reported pleasant sensations during the social touch scan, and the binding potential of the tracer was increased in various regions of the cortex (frontal, cingulate, insular), the thalamus, and the striatum, indicating deactivation of the endogenous opioid system (Nummenmaa et al. 2016).

Another study from the same group focused on the impact of social laughter on the opioid system. Twelve male volunteers were scanned twice with [ $^{11}\text{C}$ ]CFN. Before the test scan, their closest friends were present in the room and they were watching laughter-inducing comedy clips together, but before the baseline scan the subjects spent 30 min alone in the testing room. After the laughing session, the binding potential of the tracer was decreased in the thalamus, caudate nucleus, and anterior insula. In cingulate and orbitofrontal cortex, the magnitude of this decrease was correlated to the rate of social laughter. These data suggest that endogenous opioids are released after social laughter (Manninen et al. 2017).

In a study with the PET tracers [ $^{11}\text{C}$ ]CFN and [ $^{11}\text{C}$ ]raclopride, regional  $\mu\text{OR}$  and  $\text{D}_2\text{R}$  availabilities were quantified in the brains of 35 healthy adult females. During subsequent fMRI sessions, the subjects watched movie scenes with varying emotional content, ranging from very pleasant to vary scary, and regional increases of cerebral blood flow were quantified. The availability of  $\mu\text{OR}$  in the brain was found

to be negatively correlated to fMRI-quantified blood flow responses to arousing scenes in amygdala, hippocampus, thalamus, and hypothalamus. Measured binding potentials of D<sub>2</sub>R were not correlated with arousal or pleasantness in any brain region. Thus, emotional arousal appears to be regulated by the  $\mu$ OR system (Karjalainen et al. 2019).

A PET study with the  $\kappa$ OR ligand [<sup>11</sup>C]EKAP in a group of 18 healthy middle-aged volunteers examined the relationship between social status (measured with the Barratt Simplified Measure of Social Status test) and tracer volume of distribution in various areas of the brain. Social status was found to be inversely correlated to regional  $\kappa$ OR levels in amygdala, anterior cingulate cortex, caudate, frontal cortex, hippocampus, pallidum, putamen, and ventral striatum, i.e., in regions related to reward and aversion, whereas in other brain regions, no correlation was observed (Matuskey et al. 2019). This finding was interpreted as evidence for the hypothesis that  $\kappa$ OR may mediate the negative effects of social behaviors.

### 21.4.5 Physical Exercise

Long-distance runners have often reported to enter a state of euphoria while running, commonly referred to as “runner’s high.” In order to examine possible mechanisms underlying this phenomenon, ten athletes were scanned with [<sup>18</sup>F]DPN and PET, both at rest and after 2 h of endurance running (in random order). By SPM2 analysis, reductions of tracer binding were noted in several brain areas after exercise. Reported levels of euphoria were inversely correlated with [<sup>18</sup>F]DPN binding in the prefrontal/orbitofrontal cortices, anterior cingulate cortex, bilateral insula, parainsular cortex, and temporoparietal regions. These findings support the idea that running is associated with increased release of endorphins, resulting in positive mood changes (Boecker et al. 2008).

A more recent PET study with [<sup>11</sup>C]CFN in seven healthy male subjects performing cycling exercise has indicated that the pattern and emotional impact of  $\mu$ OR activation in the human brain is dependent on the intensity of the workload. Subjects performed either heavy or severe exercise. After both forms of exercise, reduced tracer binding was noted in several brain regions (particularly insular cortex and cerebellum), indicating activation of the opioid system. In heavy exercise, such decreases were correlated with increased positive sensations. However, in severe exercise, they were correlated with negative sensations. In severe but not in heavy exercise, an increase of tracer binding was noted in the pituitary gland, indicating deactivation of the  $\mu$ OR system in that organ. This last finding may be related to the development of fatigue and exhaustion in exercising subjects (Hiura et al. 2017). Similar findings were reported in a later PET study with [<sup>11</sup>C]CFN. In that study, 22 healthy subjects were scanned on 3 different occasions: after rest, a 60 min session of moderate exercise and a 60 min session of high-intensity exercise, using a bicycle ergometer. Moderate exercise did not result in significant changes of  $\mu$ OR binding, although increased euphoria was associated with decreased  $\mu$ OR availability. High-intensity exercise resulted in a significant decrease of  $\mu$ OR binding in thalamus,

insula, orbitofrontal cortex, hippocampus, and anterior cingulate, i.e., in brain regions involved in pain, reward, and emotional processing. These regional decreases were associated with increased negative emotions, such as irritation, exhaustion, and dissatisfaction (Saanijoki et al. 2018a).

Since the endogenous opioid system is involved in the rewarding aspects of both food and physical exercise, a combined PET-fMRI study in 24 healthy lean male volunteers examined the interaction between endogenous opioid release following exercise and anticipatory food reward. Two [ $^{11}\text{C}$ ]CFN-PET scans were made of each subject on separate days: after 1 h of rest and after 1 h of moderate-intensity aerobic cycling exercise. Immediately after each PET scan, a fMRI scan was made during which images of palatable food and of nonpalatable items (cars) were presented to the subjects. The order of the rest and exercise scans was counterbalanced among the participants, and all participants fasted 3 h before each PET scan. Changes of  $\mu\text{OR}$  availability after exercise showed a large variability among subjects. A larger exercise-induced decrease of  $\mu\text{OR}$  binding was associated with larger increases in anticipatory food reward responses (food-induced increases of cerebral blood flow) following exercise. Thus, differences in  $\mu\text{OR}$  activation following exercise may contribute to inter-individual variation in food craving and food consumption (Saanijoki et al. 2018b).

### 21.4.6 Pain

Opioid receptor expression ([ $^{11}\text{C}$ ]DPN or [ $^{18}\text{F}$ ]DPN binding) is high in projections of the medial and part of the lateral pain system of the human brain (Baumgartner et al. 2006; Jones et al. 1991). The lateral network is believed to be associated with the sensory aspects and the medial network with the affective aspects of pain perception. Reductions in receptor availability have been observed in healthy volunteers suffering pain. These data can be interpreted as decreases of regional receptor availability because of release of endogenous opioids, although internalization or downregulation of opioid receptors can also occur. Released opioids may mediate antinociception.

Sustained masseter muscle pain in 20 healthy volunteers was associated with significant declines of the binding of [ $^{11}\text{C}$ ]CFN in several cortical and subcortical brain regions (Zubieta et al. 2001). When capsaicin was applied to the dorsal side of the left hand of eight healthy volunteers, a pain-related decrease in [ $^{11}\text{C}$ ]CFN binding was measured in the contralateral thalamus (Bencherif et al. 2002). In eight healthy volunteers who received painful heat stimulation through the right forearm, a significant reduction of [ $^{18}\text{F}$ ]DPN binding was noted in the limbic and paralimbic areas of the brain, including the rostral anterior cingulate cortex and the insula (Sprenger et al. 2006a).

The duration of pain-induced changes in  $\mu\text{OR}$  binding potential has been examined in a PET study with [ $^{11}\text{C}$ ]CFN. Three subsequent 90-min scans were made in 14 healthy male volunteers; the first scan consisted of two 45-min baseline intervals, the second scan involved a 45-min period of sustained muscle pain followed by a

45-min baseline interval, and the third scan consisted again of two 45-min baseline intervals. Robust decreases of [ $^{11}\text{C}$ ]CFN binding were noted in several cortical and subcortical regions during the pain challenge. However, these changes did not persist in a subsequent scan. Pain-induced alterations of opioid release and/or  $\mu\text{OR}$  internalization are therefore short lasting (Scott et al. 2007a).

In a study involving 14 male and 14 female healthy individuals subjected to sustained masseter muscle pain (infusion of hypertonic saline), men demonstrated larger declines of [ $^{11}\text{C}$ ]CFN binding in the anterior thalamus, ventral basal ganglia, and amygdala. Conversely, women showed greater declines of the PET signal in the nucleus accumbens. Thus, opioid receptor-mediated antinociceptive responses show gender differences (Zubieta et al. 2002).

Opioid release is also involved in placebo analgesia. In a PET study with [ $^{11}\text{C}$ ]CFN, healthy subjects were scanned under four different conditions: painful skin heat with placebo cream, painful skin heat under control conditions, nonpainful warmth with placebo cream, and nonpainful warmth under control conditions. In all cases, the volunteers were told that the placebo cream was a powerful analgesic. Placebo treatment resulted in significant decreases of [ $^{11}\text{C}$ ]CFN binding in several brain regions known to be involved in pain and affect; thus, endogenous opioid release appears to be part of the mechanism by which expectancies regulate the perception of pain (Wager et al. 2007). A later study from the same group examined placebo and nocebo effects, i.e., the subjects expected the placebo cream to either ameliorate or exacerbate their pain. Placebo and nocebo effects were found to be associated with opposite opioid and dopaminergic responses. A placebo activated opioid neurotransmission in the anterior cingulate, orbitofrontal and insular cortices, nucleus accumbens, amygdala, and periaqueductal gray, whereas dopaminergic neurotransmission was activated in the ventral basal ganglia including the nucleus accumbens. Nocebo responses consisted of a deactivation of opioid and dopamine release. The magnitude of opioid and dopamine activation was correlated with the anticipated and subjectively perceived placebo effect (Scott et al. 2008). However, a subsequent [ $^{11}\text{C}$ ]CFN-PET study in a large number of healthy volunteers ( $n = 48$ ) indicated that a priori expectations (anticipations) are not correlated with placebo-induced decreases in pain ratings during placebo administration. Subjects with higher expectations showed greater  $\mu\text{OR}$  activation in the dorsolateral prefrontal cortex after placebo administration, but this did not correspond to subjective analgesic effects. The largest placebo responses were noted in individuals with low expectations and high subjective effectiveness. Activation of  $\mu\text{OR}$  in anterior cingulate cortex was correlated with the effectiveness of the placebo. In subjects with high expectations but low reported effectiveness of the placebo, placebo administration resulted in a nocebo, hyperalgesic response (Pecina et al. 2014).

A PET study in an animal model of placebo analgesia has confirmed that the intrinsic  $\mu\text{OR}$  system in the medial prefrontal cortex is involved in the placebo effect (Zeng et al. 2018).

When heat pain was combined with either acupuncture or placebo needle stimulation in healthy volunteers, acupuncture was shown to result in altered binding of [ $^{11}\text{C}$ ]DPN (mainly decreases) in the orbitofrontal cortex, medial prefrontal cortex, insula, thalamus, and anterior cingulate cortex (Dougherty et al. 2008).

Opioid receptor availability at baseline conditions appears to be a predictor of human sensitivity to both painful (cold pressor test) and mechanical stimuli (touch, vibration). Greater sensitivity was associated with a lower BP of [ $^{18}\text{F}$ ]DPN in various cortical areas (Mueller et al. 2010). In a later study on 12 healthy male volunteers, a significant correlation was observed between baseline binding potential of [ $^{11}\text{C}$ ]CFN ( $\mu\text{OR}$  binding) in the striatum and the cold pressor pain threshold (Hagelberg et al. 2012).

A Japanese study examined opioid mechanisms underlying heterotopic nociceptive counter-stimulation (HCNS). HCNS aims to activate endogenous pain inhibition processes, which can be beneficial in pain treatment. The right sural nerve of eight healthy volunteers was transcutaneously stimulated, resulting in leg pain. Immersion of the left hand in cold water was applied as HCNS. A late component of somatosensory evoked potentials, reflecting activity of the anterior cingulate cortex, was reduced by HCNS, and this reduction was associated with a higher binding potential of [ $^{11}\text{C}$ ]CFN in the right amygdala at baseline. Activation of  $\mu\text{OR}$  in the amygdala may thus be involved in the anti-nociceptive effects of HCNS (Piché et al. 2014).

Since peripheral inflammatory factors can influence central nervous system functioning, and, on the other hand, central mechanisms such as neural activity can modulate peripheral immune function, an American study examined relationships between IL-1 family cytokine levels in plasma and  $\mu\text{OR}$  availability in human brain. Thirty-four healthy volunteers (22 female, 12 male) were scanned with [ $^{11}\text{C}$ ]CFN in the presence and absence of a standardized pain challenge (intramuscular infusion of hypertonic saline by a computer-controlled pump). Cytokine levels in plasma were measured during both conditions. Higher baseline levels of IL-1 $\beta$  were shown to be associated with lower binding potentials of [ $^{11}\text{C}$ ]CFN in the amygdala and greater sensitivity to pain. Subjects with a greater pain-induced increase in IL-1 $\beta$  experienced less endogenous opioid analgesia. Activation of  $\mu\text{OR}$ s in ventral caudate and nucleus accumbens during the pain challenge was significantly associated with levels of IL-1 $\alpha$ , an anti-nociceptive cytokine. Thus, cytokine levels in human plasma and the endogenous opioid neurotransmitter system seem to interact (Prossin et al. 2015).

A PET study from Finland has indicated that vicarious pain (i.e., seeing others in pain) activates specific regions in the human brain and that  $\mu\text{OR}$  but not  $\text{D}_2\text{R}$  are involved in this response. Thirty-five healthy subjects were scanned, using the  $\mu\text{OR}$  tracer [ $^{11}\text{C}$ ]CFN and the  $\text{D}_2\text{R}$  tracer [ $^{11}\text{C}$ ]raclopride. Functional MRI scans of their brains were also made, during the watching of videoclips that depicted humans in painful or painless situations. The fMRI scans demonstrated increases of blood flow in various brain regions when the subjects watched painful scenes. In anterior and posterior insulae, thalamus, secondary and primary somatosensory cortices, primary motor cortex and superior temporal sulci, the magnitude of these increases was negatively correlated to measured [ $^{11}\text{C}$ ]CFN binding potentials, whereas the response in orbitofrontal cortex showed a positive correlation to  $\mu\text{OR}$  availability. In contrast to  $\mu\text{OR}$ , dopamine  $\text{D}_2\text{R}$  availability was not correlated to the blood flow response in any brain region (Karjalainen et al. 2017). Thus, inter-individual differences in  $\mu\text{OR}$  availability may explain why some humans react more strongly than others to the seeing of pain.

### 21.4.7 Vestibular Processing

An interesting study examined involvement of the opioid system in vestibular neurotransmission in humans. Ten right-handed healthy volunteers were scanned with the ligand [ $^{18}\text{F}$ ]DPN, both under baseline conditions and during caloric stimulation of the right ear. Each 90-min scan session consisted of three subsequent phases: a prestimulation period (30 min), a stimulation or sham period (30 min), and a follow-up period (30 min). During the stimulation period, warm (44 °C) and cold (30 °C) air were alternated at 5-min intervals in order to avoid habituation; during the sham period, the tubing was fixed to the subjects head in such a way that the air flow passed beside the ear. A decrease in tracer binding was noted in the right posterior insular cortex and the postcentral region during stimulation, indicating release of endogenous opioids and involvement of the opioid system in vestibular processing (Baier et al. 2010).

### 21.4.8 Myocardial Opioid Receptors

Both [ $^{11}\text{C}$ ]N-methyl-naltrindole ([ $^{11}\text{C}$ ]MeNTI) and [ $^{11}\text{C}$ ]CFN show specific binding in the human heart. However, the reduction of myocardial distribution volume and binding potential of these opioid ligands after pretreatment of volunteers with naloxone is only minor (14–21 and 19–25%, respectively), suggesting low levels of  $\delta\text{OR}$  and  $\mu\text{OR}$  expression and poor signal-to-noise ratios in PET (Villemagne et al. 2002).

### 21.4.9 Occupancy Studies

PET offers the unique opportunity of measuring the fraction of receptor populations occupied by nonradioactive drugs in the living brain and relating measured levels of occupancy to the magnitude of the therapeutic effect or to unwanted side effects. Receptor occupancy is estimated by assessing competition of a nonradioactive drug with the radioligand for the same binding sites. The basic idea was already published in 1980 (Homey et al. 1980).

Using the radioligand [ $^{11}\text{C}$ ]CFN, a half-life of  $\mu\text{OR}$  blockade of 72–108 h was measured in the brain of healthy volunteers after a single oral dosing of naltrexone, corresponding to the duration of pharmacological effects when the drug is used for treatment of heroin dependence (Lee et al. 1988).

A later study with [ $^{11}\text{C}$ ]CFN compared the duration of occupancy of  $\mu\text{OR}$  in the human brain by the antagonists nalmefene and naloxone. Clearance half-lives were 28.7 h after administration of 1 mg of nalmefene and 2.0 h for 2 mg of naloxone. The longer blockade by nalmefene could represent an advantage in the clinical use of this compound for reversal of opioid anesthesia or for treatment of drug addicts after an opioid overdose (Kim et al. 1997). Nalmefene has also been proposed for treatment of alcoholism. The decline of  $\mu\text{OR}$  occupancy in the human brain after single or repeated nalmefene dosing was found to be slower than the decline in

plasma concentration of the drug or its metabolites, indicating slow dissociation of receptor-bound nalmefene *in vivo* (Ingman et al. 2005).

Buprenorphine is used for the treatment of heroin dependence. Buprenorphine occupancy of  $\mu$ OR in the human brain has been examined, using [ $^{11}\text{C}$ ]CFN and PET. Four hours after administration of 2 mg buprenorphine, receptor occupancy ranged from 36 to 50%. When the buprenorphine dose was raised to 16 mg, levels of occupancy increased to 79–95%. Placebo-treated heroin abusers had greater  $\mu$ OR binding potential in the inferofrontal and anterior cingulate cortex compared to matched healthy controls (Zubieta et al. 2000). Increasing doses of buprenorphine were associated both with decreased  $\mu$ OR availability and with decreased withdrawal symptoms. The highest dose of buprenorphine (32 mg) caused virtually complete  $\mu$ OR occupancy (Greenwald et al. 2003). The duration of action of buprenorphine was examined in ten heroin-dependent subjects after termination of a 16 mg/day dose of the drug. Relative to a control group of heroin-dependent subjects maintained on placebo,  $\mu$ OR occupancy was 70, 46, 33, and 18% at 4, 28, 52, and 76 h after last use of the drug. Receptor occupancy was correlated with withdrawal symptoms; an occupancy of 50–60% appeared necessary for adequate symptom suppression (Greenwald et al. 2007).

Nasal spray formulations of naloxone may be useful in the treatment of victims of opioid overdose and may also be used for treating subjects with gambling disorder and alcohol addicts. Occupancy of the  $\mu$ OR population in the human brain after intranasal spray dosing of naloxone has been assessed in healthy volunteers with [ $^{11}\text{C}$ ]CFN and PET. Naloxone levels in the brain were found to peak within half an hour, a bit later than levels in plasma which peaked at 20 min. Receptor occupancies after intranasal administration of 2 and 4 mg naloxone were 67 and 85%, respectively, whereas the clearance half-life of the bound drug was about 100 min (Johansson et al. 2019).

The  $\kappa$ OR antagonist LY2456302 has been developed for treating neuropsychiatric disorders and substance abuse. Dose-dependent target occupancy by this drug has been measured in a PET study with [ $^{11}\text{C}$ ]LY2795050 in 30 healthy volunteers. An oral dose of 10 mg resulted in virtually complete  $\kappa$ OR occupancy and was well-tolerated (Naganawa et al. 2016).

Although the dose- and time-dependent opioid receptor occupancy by antagonists and partial agonists can be assessed with PET, measurements of receptor occupancy by full agonists may be problematic. No difference in [ $^{11}\text{C}$ ]DPN binding was found in the brain of opioid-dependent subjects stable on methadone (18–90 mg daily, PET scan performed at peak plasma levels of the agonist) as compared to healthy controls. In rats receiving increased doses of methadone, no dose-dependent reduction of the cerebral binding of [ $^{11}\text{C}$ ]DPN was observed. These negative findings were explained by assuming that methadone is already efficacious at very low levels of receptor occupancy (<10%, i.e., below the detection limit of PET) (Melichar et al. 2005). In a subsequent preclinical study, rats were pretreated with antinociceptive doses of oxycodone ( $\mu$ OR and  $\kappa$ OR agonist), morphine ( $\mu$ OR agonist), and buprenorphine ( $\delta$ OR and  $\kappa$ OR antagonist,  $\mu$ OR partial agonist). Full agonists did not reduce the brain uptake of [ $^{11}\text{C}$ ]DPN at all, whereas buprenorphine administration resulted in a strong reduction (up to 90%) of tracer binding (Hume et al. 2007).

Opioid receptor occupancy by the antagonist naloxone could also be measured with [ $^{11}\text{C}$ ]DPN and PET, a drug dose of 13  $\mu\text{g}/\text{kg}$  resulting in 50% occupancy (Melichar et al. 2003). The reason for the negative findings with exogenous full agonists is not clear, since endogenous enkephalins have been shown to compete with [ $^{11}\text{C}$ ]DPN for OR binding in the human brain (Ruiz-Gayo et al. 1992). Moreover, an early PET study reported 19–32% lower binding of [ $^{18}\text{F}$ ]cyclofoxy in various brain regions of long-term methadone-treated former heroin addicts as compared to controls (22 h after the last dose), and the reduction of tracer binding in caudate and putamen correlated with levels of methadone in plasma (Kling et al. 2000). Since the observed competition is both tracer and agonist dependent, these PET data may indicate the presence of multiple, different agonist binding sites on opioid receptor proteins or upregulation of subclasses of opioid receptors in heroin-dependent subjects.

Opioid receptor (particularly  $\mu\text{OR}$ ) agonists increase and antagonists decrease feeding and other rewarding behaviors in animal models. A [ $^{11}\text{C}$ ]CFN-PET study examined cerebral  $\mu\text{OR}$  occupancy in 26 healthy male volunteers after single oral doses of either GSK1521498 ( $\mu\text{OR}$  inverse agonist, candidate drug for the treatment of overeating) or naltrexone ( $\mu\text{OR}$  and  $\kappa\text{OR}$  antagonist). Activation of the amygdala by a palatable food stimulus (fruit drink) was examined in the same volunteers, using fMRI. Although a dose-dependent occupancy of  $\mu\text{OR}$  was observed for both compounds, GSK1521498 significantly attenuated activation of the amygdala by the food stimulus, in contrast to naltrexone. This difference may be related to the fact that  $\mu\text{OR}$  and  $\kappa\text{OR}$  signaling have different effects on feeding behavior and reward processing (Rabiner et al. 2011).

Naloxone administration triggers activity of the hypothalamic-pituitary-adrenal (HPA) axis by blocking opioid inhibitory tone at corticotropin-releasing factor (CRF) neurons in the paraventricular nucleus of the hypothalamus. These neurons regulate the secretion of adrenocorticotrophic hormone (ACTH) and cortisol. An interesting study measured  $\mu\text{OR}$  binding potential in the brain of 18 healthy subjects using [ $^{11}\text{C}$ ]CFN and PET. The following day, ACTH and cortisol responses to an incremental naloxone challenge were tested in the same subject group. Cortisol responses to naloxone were negatively correlated to [ $^{11}\text{C}$ ]CFN binding potential in several brain areas (ventral striatum, putamen, caudate, hypothalamus). Apparently, subjects with higher binding potential have lower endogenous occupancy of  $\mu\text{OR}$  by  $\beta$ -endorphin and thus place less inhibitory tone on the HPA axis; therefore these subjects show a smaller cortisol response to naloxone (Wand et al. 2011).

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## 21.5 PET Studies in Patients and Drug Addicts

### 21.5.1 Major Depressive, Borderline Personality, and Posttraumatic Stress Disorder

Using the sustained neutral and sadness state paradigm,  $\mu\text{OR}$  availability in the brain of healthy female volunteers and patients with major depressive disorder (MDD) has been compared. Healthy controls showed an increase of  $\mu\text{OR}$



availability in the anterior cingulate during the sustained sadness condition indicating deactivation of  $\mu$ OR-mediated neurotransmission, but in patients who did not respond to antidepressant treatment after PET imaging, sustained sadness was associated with a *decreased* binding of [ $^{11}\text{C}$ ]CFN in this region, i.e., activation of the opioid system. Patients with MDD showed significantly lower  $\mu$ OR binding potential in the posterior thalamus than healthy controls during the neutral state. These PET data indicated that endogenous opioid neurotransmission is altered in MDD (Kennedy et al. 2006). A later study from the same group showed that in MDD patients but not in healthy subjects, plasma levels of the inflammatory cytokine IL-18 are positively correlated with baseline  $\mu$ OR binding potential and with activation of  $\mu$ OR-mediated neurotransmission during a sadness challenge (Prossin et al. 2011). Thus, there may be a link between peripheral stress-activated proinflammatory mechanisms and central stress-activated neurotransmitter systems, such as the opioid system.

A study from the same group examined responses of the  $\mu$ OR system to sustained sadness in patients with borderline personality disorder. During the baseline (neutral) state, patients showed greater [ $^{11}\text{C}$ ]CFN binding than healthy controls in the bilateral orbitofrontal cortex, caudate, and nucleus accumbens besides the left amygdala but lower binding in the posterior thalamus. Induction of sadness caused greater declines of [ $^{11}\text{C}$ ]CFN binding potential in the pregenual anterior cingulate, left orbitofrontal cortex, left ventricular pallidum, left amygdala, and left inferior prefrontal cortex of the patients than of healthy controls and greater increases of binding potential in the left nucleus accumbens, the hypothalamus, and the right hippocampus/parahippocampus. Apparently, the opioid system of patients with borderline personality disorder differs from healthy, age- and sex-matched control subjects both at baseline and in its response to a negative emotional challenge (Prossin et al. 2010). Altered opioid neurotransmission may also be a mechanism underlying modified limbic system activity and mood disorders in polycystic ovary syndrome (PCOS). Insulin-resistant women with PCOS showed a greater blood flow response in limbic regions of the brain to unpleasant images than healthy controls, and the extent of this limbic activation was positively correlated to  $\mu$ OR availability in the right amygdala and left ventral anterior cingulate (Marsh et al. 2013).

An interesting study compared  $\mu$ OR binding in the brain of 14 healthy male volunteers, 15 males with combat exposure but without posttraumatic stress disorder (PTSD), and 16 male patients with PTSD, using [ $^{11}\text{C}$ ]CFN and PET. The two groups which had been exposed to war trauma showed lower  $\mu$ OR binding in the amygdala, nucleus accumbens, dorsal frontal cortex, and insular cortex than healthy volunteers but higher  $\mu$ OR binding in the orbitofrontal cortex. PTSD patients showed a significant reduction of  $\mu$ OR binding in the anterior cingulate cortex compared to both other groups. Combat-exposed subjects without PTSD had lower  $\mu$ OR binding in the amygdala but higher binding in the orbitofrontal cortex than either PTSD patients or healthy controls. Thus, the opioid system of the brain shows specific changes both after trauma and in PTSD (Liberzon et al. 2007).

Preclinical data have suggested that the dynorphin/ $\kappa$ OR system may be related to symptoms of fear and depression after exposure to trauma. An American study used

the  $\kappa$ OR ligand [ $^{11}\text{C}$ ]LY2795050 to scan the brains of 35 subjects with a wide range of exposure to trauma (ranging from healthy controls to subjects with PTSD, MDD, and generalized anxiety disorder). Measured distribution volumes of the tracer in amygdala, anterior cingulate cortex, and ventral striatum were significantly and negatively correlated with 24-h urinary cortisol levels and severity of loss (i.e., dysphoria, depressive symptoms, emotional numbing), but not with feelings of threat (i.e., fear) in the subject group (Pietrzak et al. 2014). In contrast to the findings of this initial study, a later pilot study with [ $^{11}\text{C}$ ]GR103545 did not observe any significant difference of regional tracer distribution volume (i.e.,  $\kappa$ OR availability) between 10 patients with MDD and 13 healthy volunteers (Miller et al. 2018).

If medication-free patients with MDD and healthy controls were scanned with [ $^{11}\text{C}$ ]CFN and subjected to a social rejection and acceptance protocol (as described above, Sect. 21.4.3 (Hsu et al. 2013)), the patients showed a reduced release of opioids in various brain regions in response to rejection and a slower emotional recovery than the healthy subjects. During acceptance, only the healthy controls reported an increase in self-esteem and an increased desire for social interaction, which was correlated with release of endogenous opioids in the nucleus accumbens. A reduced activity of the endogenous opioid system may thus be involved in the pathophysiology of MDD, resulting in a less effective recovery from negative and a decreased pleasure from positive social interactions (Hsu et al. 2015).

Another PET study from the same group examined changes of the regional binding potential of [ $^{11}\text{C}$ ]CFN and plasma levels of interleukin-18 in 28 human volunteers during the induction of sad or neutral mood, using a standard protocol. The subject group consisted of 15 healthy controls and 13 patients with unmedicated MDD. All subjects were female. IL-18 levels were increased during sadness, indicating peripheral immune activation and were reduced during neutral mood. IL-18 increases in the patients were significantly greater than in the control group. Moreover, these increases were found to be linearly proportional to sadness-induced  $\mu$ OR activation (i.e., reductions of tracer binding potential) in left ventral pallidum, bilateral anterior cingulate cortices, right hypothalamus, and bilateral amygdala. Apparently, changes in mood regulate human immune function, and this regulation may be related to central opioid neurotransmission (Prossin et al. 2016).

An interesting study with [ $^{11}\text{C}$ ]CFN examined the impact of placebo and antidepressant treatment in 35 medication-free patients with MDD. All subjects were treated with a placebo during two periods of 1 week. They were either informed that the placebo was a fast-acting antidepressant drug or that it was an inactive control. After each week of treatment, a [ $^{11}\text{C}$ ]CFN-PET scan was made. In half of the subjects, the “active” placebo was given first and was followed by the “inactive” placebo, whereas in the other half, the order of treatments was reversed. After both PET scans and placebo treatments, all subjects were treated for 10 weeks with a real antidepressant, in most cases a selective serotonin reuptake inhibitor.

Higher  $\mu$ OR binding in the nucleus accumbens at baseline was associated with better antidepressant response. A decrease of tracer binding potential in subgenual anterior cingulate cortex, nucleus accumbens, thalamus, and amygdala after 1 week of “active” placebo (compared to the amount of binding after 1 week of “inactive”

placebo) was associated with placebo-induced reductions in depressive symptoms and also with a better response to the real antidepressant at the end of the trial. Thus, a placebo is capable of activating the endogenous  $\mu$ -opioid system in patients with MDD and this system is also involved in the response of such patients to an antidepressant drug (Pecina et al. 2015).

### 21.5.2 Pain

In a group of four patients with rheumatoid arthritis, regional cerebral opioid receptor binding of [ $^{11}\text{C}$ ]DPN was quantified both during a period of pain and during a period when the subjects were out of pain. Significant decreases of [ $^{11}\text{C}$ ]DPN binding were seen in many brain areas when subjects suffered pain (particularly straight gyrus and frontal, cingulate, and temporal cortex) (Jones et al. 1994b). In a patient with central poststroke pain which developed after a small pontine hemorrhagic infarction, a reduction of the binding of [ $^{11}\text{C}$ ]DPN was also noted, and this reduction was more striking than the hypometabolism of [ $^{18}\text{F}$ ]FDG on the lateral cortical surface contralateral to the symptoms (Willoch et al. 1999).

Changes of [ $^{11}\text{C}$ ]DPN binding in the brain of six patients after surgical relief of trigeminal neuralgia pain were examined in a later study. The volume of distribution of the tracer in several cortical areas (prefrontal, insular, perigenual, midcingulate, inferior parietal), basal ganglia, and the thalamus was found to be significantly increased after surgery (Jones et al. 1999). These data and the findings in the previous studies were interpreted as evidence for the release of opioid peptides during various forms of pain resulting in a reduction of the fraction of the receptor population available for ligand binding. However, changes of opioid receptor numbers (upregulation after surgery or after the disappearance of pain) could not be ruled out.

In a later study involving four patients with central neuropathic pain (mainly poststroke) and age-matched controls, patients were found to have significantly less [ $^{11}\text{C}$ ]DPN binding in the dorsolateral, anterior cingulate, and insula cortices, the thalamus, and the inferior parietal cortex. These reductions were not a direct consequence of their cerebral lesions, since these were located in other brain areas. The authors interpreted the PET data as evidence for reduced damping of nociceptor activity in the patients and as a possible explanation for the fact that high doses of opiates are frequently required to achieve optimal analgesia in subjects with central neuropathic pain (Jones et al. 2004). Essentially similar findings were reported in a study from another institution, using the same tracer and involving 5 patients with central poststroke pain and 12 healthy volunteers (Willoch et al. 2004).

An interesting study compared changes of [ $^{11}\text{C}$ ]DPN binding in the brain of patients with peripheral neuropathic pain and central poststroke pain. Peripheral neuropathic pain patients showed bilateral and symmetric decreases of [ $^{11}\text{C}$ ]DPN binding, in contrast to central poststroke pain patients where an asymmetric decrease was noted, which was most prominent in the contralateral hemisphere. The symmetric and bilateral decline in the former patient group may reflect endogenous opioid release, whereas the lateralized decrease in the latter group suggests a loss of

opioid receptors (Maarrawi et al. 2007a). Thus, central and peripheral forms of neuropathic pain may have different effects on the opioid system of the brain.

In a group of ten patients with restless legs syndrome (RLS), significant negative correlations were observed between [ $^{11}\text{C}$ ]DPN binding in areas serving the medial pain system and RLS severity or pain scores. These findings suggested release of endogenous opioids within the medial pain system related to the severity of the syndrome (von Spiczak et al. 2005).

Changes of the opioid system in patients with cluster headache were also examined, using the tracer [ $^{11}\text{C}$ ]DPN and PET. When patients as a group were compared with healthy volunteers, a striking (>50%) decrease of tracer binding was noted in the pineal gland but not in any other brain area. Within the patient group, opioid receptor availability in the ipsilateral hypothalamus and anterior cingulate cortex was negatively correlated to the duration of the headache disorder. The patients did not experience acute pain during or immediately before the study; thus, the observed alterations of [ $^{11}\text{C}$ ]DPN binding appear to not represent an effect of acute pain processing but rather alterations in the opioid system related to the development of the disease. Opioidergic dysfunction in circuitries generating the biologic clock may be a mechanism underlying cluster headache (Sprenger et al. 2006b).

In patients with complex regional pain syndrome (CRPS), a chronic pain condition which can develop after limb trauma, binding of the non-subtype-selective OR ligand [ $^{18}\text{F}$ ]DPN was found to be reduced in the contralateral amygdala and parahippocampal gyri but increased in the contralateral prefrontal cortical areas. Ligand binding in the ipsilateral temporal cortex and midcingulate cortex was negatively correlated with pain scores, but ligand binding in the contralateral temporal cortex was positively correlated with anxiety and depression scales (Klega et al. 2010). These findings suggest that the endogenous opioid system is implied in both chronic pain and its psychiatric comorbidity.

A study with the tracer [ $^{11}\text{C}$ ]CFN compared  $\mu\text{OR}$  binding in the brain of four patients with trigeminal neuropathic pain and eight gender- and age-matched healthy controls. Patients showed reduced tracer binding in the left nucleus accumbens, and binding potential in this area was negatively correlated with their pain ratings. This finding was interpreted as evidence for downregulation of  $\mu\text{OR}$  after persistent activation of opioid neurotransmission in chronic pain (DosSantos et al. 2012a). A recent PET study in experimental animals has proven that reduced  $\mu\text{OR}$  availability in chronic neuropathic pain (measured with [ $^{18}\text{F}$ ]DPN and PET) reflects reduced  $\mu\text{OR}$  expression. In this study,  $\mu\text{OR}$  expression in various brain areas was quantified by ex vivo immunohistochemistry (Thompson et al. 2018).

Another PET study with [ $^{11}\text{C}$ ]CFN examined regional  $\mu\text{OR}$  binding potential values in the human brain during experimentally induced pain (application of 0.35 g of 10% capsaicin cream on a 6.25 cm<sup>2</sup> area of the left hand). The data from the PET study were compared to evaluations of the sleep quality of the subjects ( $n = 14$ ) during the previous month, using the Pittsburgh Sleep Quality Index. Poor sleep quality was significantly and positively correlated to greater  $\mu\text{OR}$  binding potential in the frontal lobes, and sleep duration was negatively associated with binding potential in these lobes, the temporal lobe and the anterior cingulate. Thus, individual variation

in the quality and duration of sleep appear to be associated with variation in cerebral  $\mu$ OR binding during tonic pain. Poor sleep may be related to impaired pain-suppressing mechanisms (less release of endogenous opioids in response to a painful stimulus) and an increased perception of pain (Campbell et al. 2013).

A more complicated study with [ $^{11}\text{C}$ ]CFN provided evidence for the hypothesis that both alterations in  $\mu$ OR availability and a reduced release of endogenous opioids contribute to the clinical symptoms of patients suffering from chronic back pain. Tracer binding potential was measured in 16 patients with chronic back pain and 16 age- and gender-matched healthy volunteers under three different conditions: at baseline, during pain expectation (intramuscular injection of isotonic saline), and during sustained moderate pain (intramuscular injection of hypertonic saline). Baseline  $\mu$ OR binding in the thalamus proved to be higher in the patients than in the control subjects, and the patients showed smaller reductions in  $\mu$ OR binding in several brain areas, both during the painful condition and the expectation of pain (Martikainen et al. 2013).

PET with the  $\mu$ OR tracer [ $^{11}\text{C}$ ]CFN has also been used to examine activity of the opioid system during migraine attacks. Seven patients with frequent migraine were scanned both during the ictal and interictal phase.  $\mu$ OR binding in the ipsilateral medial prefrontal cortex was found to be significantly reduced during the ictal phase, possibly due to increased release of endogenous opioids, in order to fight the ongoing pain. The magnitude of this reduction was negatively correlated to the combined extension and severity of the migraine attack (Dasilva et al. 2014).

Migraine, particularly in patients that have used opioids to suppress their pain, is frequently associated with allodynia, i.e., an increased sensitivity of the skin to stimuli that should not cause pain. Normal activities, such as washing the skin with warm water, or combing the hair, are painful for subjects suffering from allodynia. A report following on the previous PET study examined changes of [ $^{11}\text{C}$ ]CFN in patient's brains during migraine-associated allodynia. The skin was thermally challenged by a device that provided heat pulses every 10s during a period of 20 min. Patients were challenged and scanned both during the ictal and interictal phase of their migraine. A decrease of  $\mu$ OR binding potential was observed in the ventrolateral periaqueductal gray matter and the red nucleus area of the midbrain, when patients were thermally challenged and showed allodynia. The magnitude of this decrease was positively correlated to allodynic severity during a migraine attack (Nascimento et al. 2014).

Simultaneous PET (with the opioid receptor ligand [ $^{11}\text{C}$ ]DPN) and functional MRI (fMRI) has been used to examine the opioidergic pain system in the human brain. Eight healthy volunteers were scanned twice on the same day, during the application of a painful and a non-painful stimulus (pressure). The order of the scans was randomized. Pain-related activation of several brain regions was observed in fMRI, but in most of these regions, opioid receptor binding was not significantly altered. Co-localized fMRI and PET signal changes were noticed only in the ipsilateral thalamus and the contralateral striatum. Decreases of tracer binding potential and increases of blood flow (fMRI BOLD signal) were significantly correlated in the thalamus, suggesting that in this area of the brain, enhanced opioid

neurotransmission is a major cause of the increases of flow, i.e., of neuronal activation. However, no correlation was observed between the PET and fMRI signal changes in the striatum, suggesting that in this region, additional receptor systems were contributing to the observed flow changes (Wey et al. 2014).

A British study used the PET tracer [ $^{11}\text{C}$ ]DPN to relate opioid receptor availability in the human brain to pain perception in arthritis. Within the patient group with arthritis ( $n = 17$ ), the perception of higher levels of chronic pain in the week before the scan was associated with higher OR availability, particularly in the caudate, nucleus accumbens, and subcallosal area. This observation suggests that opioid receptors are upregulated in individuals experiencing greater chronic pain. Higher OR availability was associated with higher acute thermal pain thresholds, suggesting that OR upregulation is an adaptive response aimed at dampening the pain perception. When the entire patient group was considered, OR availability in the patients caudate nucleus was reduced in comparison to healthy controls ( $n = 9$ ), suggesting a greater release of endogenous opioid ligands and greater occupancy of ORs in the arthritis group (Brown et al. 2015).

Patients with fibromyalgia report widespread pain in muscles and connective tissue, whereas no pathology can be identified in the painful body areas. The central nervous system of these patients may have become overly sensitive to painful stimuli. In order to test this hypothesis, pain-evoked increases of cerebral blood flow were quantified with fMRI and regional  $\mu\text{OR}$  availability was quantified in 18 female patients with fibromyalgia, using PET and [ $^{11}\text{C}$ ]CFN. Patients showed a lower  $\mu\text{OR}$  availability in dorsolateral prefrontal and anterior cingulate cortex than healthy controls, and this was associated with a lower blood flow response to painful stimuli. Thus, the endogenous opioid system appears to be dysregulated in fibromyalgia, resulting in lower activation of antinoceptive brain regions after incoming painful stimuli and abnormal sensitivity to pain (Schrepf et al. 2016).

### 21.5.3 Pain Treatment

Monitoring the relationship between opioid receptor occupancy and pain relief could be useful in the development of improved techniques for the treatment of pain (Sadzot et al. 1990; Sadzot and Frost 1990). This study paradigm has been employed in several reports.

An initial study examined the contribution of the endogenous opioid system to pain relief induced by motor cortex stimulation (MCS). Eight patients with refractory neuropathic pain were scanned three times with [ $^{11}\text{C}$ ]DPN and PET: twice preoperatively with a 2-week interval and once after 7 months of chronic MCS. The preoperative test-retest scans did not show any significant differences, but comparison of the last scan with the two preoperative scans revealed significant decreases of tracer binding in the anterior middle cingulate cortex, periaqueductal gray, prefrontal cortex, and cerebellum. The magnitude of the changes in the first two brain areas was significantly correlated with pain relief. These findings were interpreted as

evidence for the release of endogenous opioids in brain structures involved in pain processing after MCS (Maarrawi et al. 2007b).

The effects of traditional Chinese acupuncture and placebo needle stimulation have been compared in chronic pain patients diagnosed with fibromyalgia. [ $^{11}\text{C}$ ]CFN-PET scans were made at both the beginning and the end of the 4-week treatment period. Acupuncture was shown to result in both short- and long-term increases in [ $^{11}\text{C}$ ]CFN binding potential in multiple pain and sensory processing regions. In the sham group, small reductions of [ $^{11}\text{C}$ ]CFN binding were observed, consistent with previously reported placebo effects. The long-term increases after acupuncture were correlated with reductions in clinical pain. Thus, acupuncture and sham acupuncture may have different effects on cerebral  $\mu\text{OR}$  binding and alterations of  $\mu\text{OR}$  binding potential may be involved in the analgesic effect of acupuncture (Harris et al. 2009).

A case study reported that transcranial direct current stimulation (tDCS) of the motor cortex in a trigeminal neuropathic pain patient considerably decreased the binding potential of [ $^{11}\text{C}$ ]CFN in several brain areas compared to sham tDCS. The effect was particularly striking in the posterior thalamus. tDCS increased the threshold for experimental cold pain in the patient but did not improve her clinical pain. The authors suggested that the release of endogenous opioids is immediately increased by tDCS, but repetitive tDCS sessions are required to revert neuroplastic changes related to the neuropathic pain (DosSantos et al. 2012b). A follow-up study from the same group examined the mechanisms underlying tDCS. Nine healthy volunteers were scanned with [ $^{11}\text{C}$ ]CFN on two different occasions: at baseline and during application of tDCS. The last scan comprised a placebo phase (in which tDCS electrodes were present but no stimulation was applied) and a real stimulation phase. The placebo phase was associated with a decrease of tracer binding potential in the periaqueductal gray, precuneus, and thalamus, indicating release of endogenous opioids in response to the placebo. Real tDCS was associated with similar changes in periaqueductal gray and precuneus and an additional decrease of tracer binding in the left prefrontal cortex. Significant analgesic effects (increases of pain thresholds) were only observed after real tDCS. Thus, the endogenous  $\mu\text{OR}$  system is already activated by a placebo, and the placebo effect seems to be extended and optimized by real tDCS (DosSantos et al. 2014).

A PET study with [ $^{11}\text{C}$ ]DPN examined OR availability in 15 patients suffering from refractory neuropathic pain. The patients were subsequently treated by receiving chronically implanted motor cortex stimulation. Preoperative levels of opioid binding in several brain regions (insula, thalamus, periaqueductal grey, anterior cingulate, and orbitofrontal cortex) were found to be significantly and positively correlated with postoperative pain relief after 7 months. Thus, PET scans with an opioid receptor ligand may help clinicians to select patients with neuropathic pain, who are likely to benefit from motor cortex stimulation (Maarrawi et al. 2013).

A Chinese study evaluated the impact of 30 min of transcutaneous electrical acupoint stimulation (TEAS) on  $\mu\text{OR}$  availability in the brain of anaesthetized rhesus monkeys, using [ $^{11}\text{C}$ ]CFN and PET. The study showed that 2 Hz but not 100 Hz stimulation resulted in a significant increase of tracer binding potential in the

anterior cingulate cortex, caudate nucleus, putamen, temporal lobe, somatosensory cortex, and amygdala (compared to a 0 Hz control scan). These results suggest that not only the release of endogenous opioid peptides, but also an upregulation of  $\mu$ OR plays a role in the mechanism of action of TEAS (Xiang et al. 2014).

A decreased availability of  $\mu$ OR in the human brain (suggesting the release of endogenous opioids) has also been observed in healthy volunteers ( $n = 10$ ) after transcranial magnetic stimulation (TMS). After active but not sham TMS, the binding potential values of [ $^{11}\text{C}$ ]CFN were lowered in the right ventral striatum, medial orbitofrontal, prefrontal and anterior cingulate cortices, left insula, superior temporal gyrus, dorsolateral prefrontal cortex, and precentral gyrus. In contrast to these changes of  $\mu$ OR, dopamine D2R availability was not altered after TMS (Lamusuo et al. 2017).

## 21.5.4 Substance Abuse

### 21.5.4.1 Cocaine Dependence

The endogenous opioid system appears to play a role in the reinforcing actions of non-opioid drugs. American investigators examined  $\mu$ OR binding in the brain of cocaine addicts, 1–4 days after their last use of cocaine, using [ $^{11}\text{C}$ ]CFN and PET. An increase of tracer binding was observed, particularly in the frontal cortex and caudate nucleus, and this increase persisted even after 4 weeks of drug abstinence. The magnitude of [ $^{11}\text{C}$ ]CFN uptake was positively correlated with the severity of cocaine craving experienced by the subjects. These findings confirmed the involvement of the opioid system in cocaine dependence (Zubieta et al. 1996).

Increases of  $\mu$ OR binding potential in the anterior cingulate and anterior frontal cortex of cocaine users during abstinence are not only positively correlated with the intensity of cocaine craving, but the elevation of [ $^{11}\text{C}$ ]CFN binding in these areas is also correlated with the percentage of days with cocaine use and the amount of cocaine used per day before the study period (Gorelick et al. 2005).

Regional increases of  $\mu$ OR binding appear to be related not only to the intensity of cocaine craving but also to the time interval before relapse to cocaine use after discharge from a closed research ward. In an interesting study, 15 cocaine-dependent subjects were housed in this ward for 12 weeks of monitored abstinence. During this period, regional brain  $\mu$ OR binding was measured at 1 and 12 weeks, using [ $^{11}\text{C}$ ]CFN and PET. A shorter time interval before relapse was associated with increased  $\mu$ OR binding in the frontal and temporal cortex in both PET scans and with a lesser decrease in binding between 1 and 12 weeks. The significance of this correlation persisted even after accounting for clinical variables. Thus, increased  $\mu$ OR binding in the frontal and temporal cortex is a predictor of time to relapse to cocaine use (Gorelick et al. 2008).

A PET study examined regional  $\mu$ OR ([ $^{11}\text{C}$ ]CFN) binding in the brain of treatment-seeking cocaine users before the onset of treatment and correlated this parameter with subsequent treatment outcome. Elevated  $\mu$ OR binding in brain regions associated with reward sensitivity correlated with greater cocaine use



during treatment and shorter duration of cocaine abstinence. Thus,  $\mu$ OR binding in these brain areas was a significant independent predictor of treatment outcome (Ghitza et al. 2010).

The mechanism of action of psychostimulant drugs in the mammalian brain appears to involve the release of endogenous opioids (endorphins) in the reward system. Two recent PET studies have attempted to obtain experimental proof for this mechanism by administering d-amphetamine to healthy volunteers (i.v.) and examining the impact of this challenge on [ $^{11}\text{C}$ ]CFN binding in the brain. A Swedish study in which ten subjects were scanned under three conditions (baseline, placebo, amphetamine 0.3 mg/kg) could not detect any significant differences in the binding potential of [ $^{11}\text{C}$ ]CFN after the amphetamine challenge (Guterstam et al. 2013). However, another study in which 12 subjects were scanned before and 3 h after an amphetamine challenge (either a high dose, 0.5 mg/kg, or an ultralow dose 0.017 mg/kg) reported that [ $^{11}\text{C}$ ]CFN binding was reduced after the high but not the ultralow dose in the frontal cortex, putamen, caudate, thalamus, anterior cingulate cortex, and insula (Colasanti et al. 2012).

#### 21.5.4.2 Opioid Dependence

Similar changes of opioid receptor binding as were reported for cocaine addicts have also been observed in early abstinence from opioid (heroin) dependence. Compared with 20 healthy controls, ten subjects with opioid dependence who had completed inpatient detoxification showed increased [ $^{11}\text{C}$ ]DPN binding in the whole brain and in 15 out of 21 studied brain areas (Williams et al. 2007).

#### 21.5.4.3 Alcohol Dependence

Whereas studies in cocaine and heroin addicts suggest increases of opioid receptor binding during abstinence, studies in alcohol dependence have produced conflicting results. In an early PET study with [ $^{11}\text{C}$ ]CFN,  $\mu$ OR binding potential was measured in eight male alcohol-dependent subjects during alcohol withdrawal and eight matched healthy controls. Contrary to the findings in cocaine abusers, alcoholics showed *reductions* of  $\mu$ OR binding potential in the right dorsal lateral prefrontal cortex, right anterior frontal cortex, and right parietal cortex. Lower  $\mu$ OR binding potential was associated with higher craving and with depressive symptoms in the subjects (Bencherif et al. 2004b). Based on the findings in subsequent studies, one could speculate that alcohol abstinence and depression have different effects on  $\mu$ OR binding (an increase and a decrease, respectively), the latter effect predominating in this early investigation.

Opposite results were reported in a German study. The authors examined [ $^{11}\text{C}$ ]CFN binding in the brain of alcoholics after 1–3 weeks of abstinence. Significantly *higher* binding was observed in the ventral striatum of the patients compared with healthy controls. This increase persisted up to 5 weeks, and its magnitude was significantly correlated with the intensity of alcohol craving. Elevated availability of  $\mu$ OR in the striatum was considered as a neuronal correlate of alcohol urge since the ventral striatum is known to be an important part of the reward system in the human brain (Heinz et al. 2005). Similar results were reported in a large study

involving 25 alcohol-dependent subjects who were scanned on day 5 of abstinence and 30 healthy controls. Alcohol-dependent subjects had a significantly higher binding potential of [ $^{11}\text{C}$ ]CFN in many brain regions. There was a significant inverse relationship between [ $^{11}\text{C}$ ]CFN binding in several brain regions and the intensity of craving in the alcohol-dependent group. No significant group differences in binding of the  $\delta$ OR ligand [ $^{11}\text{C}$ ]MeNTI were observed, but the binding potential of this ligand in the caudate nucleus was positively correlated with recent drinking in alcohol-dependent subjects (Weerts et al. 2011).

In alcohol-dependent subjects completing a 19-day inpatient protocol (alcohol abstinence followed by naltrexone treatment, 50 mg daily, on days 15–19), a virtually complete (95%) occupancy of  $\mu$ OR by naltrexone was observed, using [ $^{11}\text{C}$ ]CFN and PET. However, a much lower and highly variable (0–50%) occupancy of  $\delta$ OR was detected in [ $^{11}\text{C}$ ]MeNTI scans. The relationship between  $\delta$ OR occupancy and treatment outcome could be explored in future studies (Weerts et al. 2008).

A British study with the non-subtype-selective opioid ligand [ $^{11}\text{C}$ ]DPN reported a trend toward increased tracer binding in 11 alcohol-dependent subjects during early abstinence, compared to healthy controls, although this trend did not reach statistical significance. A significant correlation between alcohol craving and the global and regional distribution volume of [ $^{11}\text{C}$ ]DPN was noted in the alcohol-dependent subjects (Williams et al. 2009).

A recent study examined [ $^{11}\text{C}$ ]CFN binding in the human brain before and immediately after alcohol consumption in 13 heavy drinkers and an age-matched control group of 12 healthy subjects. Drinking alcohol reduced tracer binding in the nucleus accumbens and orbitofrontal cortex, indicating release of endogenous opioids in brain areas involved in the valuation of reward. The magnitude of change in the orbitofrontal cortex was significantly correlated with problem alcohol use and with subjective high in the drinker group. Thus, the release of endogenous opioids by ethanol promotes further alcohol consumption, and altered function of the opioid system appears to contribute to alcohol abuse (Mitchell et al. 2012).

Increases of the plasma level of the hormone cortisol after administration of an opioid antagonist (such as the drug naloxone) are supposed to provide information about opioid receptor activity. In an American study, naloxone was administered to 18 healthy control and 25 alcohol-dependent subjects (after 5 days of medically supervised alcohol withdrawal). The dose of naloxone required to induce a peak cortisol response was two times higher in the alcohol-dependent group. PET scans with [ $^{11}\text{C}$ ]CFN were made in all subjects, and the relationship between tracer binding potential and rising slope of the cortisol response was examined. Alcohol-dependent subjects showed a significantly higher tracer binding potential in several brain regions than healthy controls, as was previously reported (Weerts et al. 2011). In healthy controls, there was a significant negative correlation between cortisol slope and regional binding potentials of the tracer, but in alcohol-dependent subjects, these parameters were not correlated. Thus, in healthy subjects, the naloxone test can provide information about individual differences in  $\mu$ OR availability, but in shortly abstinent alcohol-dependent subjects, the relationship between naloxone-induced cortisol response and  $\mu$ OR availability is disrupted (Wand et al. 2012).

In a follow-up study from the same group, PET scans with the  $\delta$ OR ligand [ $^{11}\text{C}$ ]methylnantrindole were made in 15 healthy control and 20 shortly abstinent alcohol-dependent subjects, and  $\delta$ OR binding potential in various regions of the brain was correlated to cortisol area under the curve (AUC) after administration of naloxone. Similar to the findings for  $\mu$ OR, significant negative correlations were observed between cortisol AUC and  $\delta$ OR binding potential in several brain regions of healthy subjects. However, in alcohol-dependent subjects, such correlations were not observed. Thus, in healthy subjects, the naloxone test provides information about individual differences in  $\delta$ OR availability, but in alcohol-dependent subjects, the close relationship between opioidergic neurotransmission and hormonal response is lost (Wand et al. 2013).

A PET study with the  $\mu$ OR tracer [ $^{11}\text{C}$ ]CFN examined the relationship between opioid release in the human brain after alcohol consumption and the catechol-O-methyltransferase genotype. Thirteen heavy drinkers and 12 matched healthy control subjects were scanned twice with [ $^{11}\text{C}$ ]CFN: first at baseline and then after the consumption of a standard drink of alcohol. Independent of their drinking history (or gender), individuals with the Val158 variant of the enzyme showed greater opioid release in the right nucleus accumbens and less release in medial orbital frontal cortex. The Val158 variant results in greater activity of catecholamine-O-methyltransferase in the human brain and, consequently, lower dopamine levels in the frontal cortex. Apparently, genetic differences in endogenous dopamine levels can modulate endogenous opioid release in brain regions that contribute to the rewarding effects of alcohol. The Val158 variant could make individuals more prone to impulsive decision-making (Mitchell et al. 2013).

The higher binding potential values of [ $^{11}\text{C}$ ]CFN in the brain of alcohol-dependent subjects that have been reported by various research groups may either reflect increased expression of  $\mu$ OR, or a reduced concentration of endogenous, competing opioid ligands. In order to differentiate between these possible explanations, a German study examined post-mortem brain tissue of 43 alcohol-dependent subjects and 43 healthy controls. The density of  $\mu$ OR in tissue samples was determined by *in vitro* assays, using the ligand [ $^3\text{H}$ ]DAMGO. Tracer binding in the brain of alcohol-dependent subjects was 23–51% lower than in healthy controls. Thus, increased binding potential values of [ $^{11}\text{C}$ ]CFN in the brains of alcoholics seem to reflect reduced competition of endogenous opioids rather than increased receptor expression (Hermann et al. 2017).

Addiction (to alcohol or other substances of abuse) may be considered as a state of “reward deficiency.” Addicts take a rewarding substance to compensate for their deficient reward. If this interpretation of addiction is correct, drug addicts should show a reduced release of endogenous opioids after a release-inducing stimulus compared to healthy controls. In a British PET study, 13 abstinent alcohol-dependent male subjects and 15 healthy volunteers were scanned twice with [ $^{11}\text{C}$ ]CFN, before and 3 h after an oral dose of dexamphetamine. Consistent with the hypothesis, the healthy volunteers showed a significant decline of tracer binding potential in many areas of the brain after the amphetamine challenge, but in alcohol-dependent subjects, this response was reduced, particularly in frontal lobe, insula, thalamus,

anterior cingulate, nucleus accumbens, and putamen (Turton et al. 2018). In pathological gambling, a behavioral addiction with similarities to substance abuse, a similar “blunted” response of the opioid system to amphetamine was noted, although baseline  $\mu$ OR availability was not different in pathological gamblers and healthy volunteers (Mick et al. 2016).

In an extensive PET study with the novel PET tracer [ $^{11}\text{C}$ ]LY2795050,  $\kappa$ OR availability was measured in the brain of 36 alcohol-dependent subjects and 28 healthy controls. Tracer distribution volume in the amygdala and pallidum was significantly lower in the alcohol-dependent group than in the healthy control group. An age effect on tracer binding could not be observed in any brain region (Vijay et al. 2018). These observations for  $\kappa$ OR are in contrast with findings for  $\mu$ OR or total OR that have been reported to be either increased or unchanged in alcohol-dependent subjects. Also, human  $\mu$ OR show an age-related decline, whereas  $\kappa$ OR appear to be preserved in the aging human brain.

#### 21.5.4.4 Nicotine Dependence

Several PET studies have examined the relationship between opioid neurotransmission and smoking. In an initial, groundbreaking publication, six healthy male smokers and six age- and sex-matched nonsmokers were scanned with both [ $^{11}\text{C}$ ]CFN and the dopamine receptor ligand [ $^{11}\text{C}$ ]raclopride. Smokers abstained from smoking for 12 h prior to the study. They first smoked a denicotinized cigarette, 2 and 12 min after scan onset, followed by a normal cigarette after 40 and 50 min. Total scan duration was 90 min. Thus, each scan period consisted of an initial, denicotinized phase and a later, average nicotine phase. Nonsmokers were subjected to the same scanning protocol but were only scanned at baseline (i.e., they did not smoke anything). In the smoker group, significant declines of [ $^{11}\text{C}$ ]CFN binding in the right anterior cingulate cortex were noted during transition from the first to the second phase of the scan, indicating activation of opioid neurotransmission by nicotine. A simultaneous decline of [ $^{11}\text{C}$ ]raclopride binding was noted in the ventral basal ganglia, and the magnitude of this decline correlated with nicotine dependence score. Increases of the release of endogenous opioids and dopamine are probably involved in the rewarding effect of nicotine. Smokers had lower  $\mu$ OR binding potential in the cingulate cortex, thalamus, ventral basal ganglia, and amygdala compared to nonsmokers during the first phase of the scan; these reductions were reversed in the thalamus, ventral basal ganglia, and amygdala after nicotine smoking (Scott et al. 2007b). Greater  $\mu$ OR availability in the basal ganglia and thalamus of nonsmokers compared to overnight abstinent smokers was also observed in a later study (Nuechterlein et al. 2016).

In contrast to the observations of Scott and Nuechterlein et al., another PET study with [ $^{11}\text{C}$ ]CFN did not observe any significant change in tracer binding potential between placebo and active cigarette smoking and also no significant difference in  $\mu$ OR availability between smokers and nonsmokers (Kuwabara et al. 2014). The discrepancy between this and the initial study led to the hypothesis that the expectancy of a smoker, or the sensorimotor effects of denicotinized cigarettes may contribute to the observed release of endogenous opioids. However, a negative

correlation was observed between  $\mu$ OR availability in bilateral superior temporal cortices during placebo smoking and scores for nicotine dependence. Thus,  $\mu$ OR in this area may play a role in nicotine addiction, and stimulation of these receptors by endogenous opioids may be involved in nicotine reward (Kuwabara et al. 2014).

Two recent PET studies showed that psychological and behavioral factors can indeed exert important actions on the  $\mu$ -opioid system. In the first study, smokers showed a reduction of [ $^{11}$ C]CFN binding potential in the thalamus and nucleus accumbens after smoking a denicotinized (placebo) cigarette, indicating  $\mu$ OR activation. No further activation was observed after the smoking of a normal cigarette (Nuechterlein et al. 2016). In the last study, 20 chronic tobacco smokers (males) were asked to abstain for one night from smoking. The next morning, they first smoked a denicotinized cigarette (at 8.00 h) and, subsequently, a normal cigarette with an average dose of nicotine (at 10.00 h). PET scans with [ $^{11}$ C]CFN were made either during the placebo or the nicotine smoking session (on two different days), using a bolus plus infusion protocol. The very low nicotine peak levels in plasma after smoking a denicotinized cigarette were significantly correlated with tracer binding potential after the smoking session, but no correlation was observed between the high peak levels after smoking of the normal cigarette and tracer binding potential (Domino and Hirasawa-Fujita 2019). Thus, “placebo” effects seem to have a major impact on [ $^{11}$ C]CFN binding in the smoker’s brain.

Another study with [ $^{11}$ C]CFN examined the relationship between smoking and the relief of negative affect. Twenty-two smokers were scanned after overnight abstinence, once after smoking a denicotinized cigarette and at another scanning day after smoking a normal cigarette. Higher  $\mu$ OR availability in the amygdala was correlated with greater motivation to smoke in order to relieve negative affect, but not with changes in affect after smoking (Falcone et al. 2012). A positive correlation between the craving to smoke and  $\mu$ OR binding potential in several brain regions (amygdala, hippocampus, insula, nucleus accumbens, putamen, and ventral striatum) was also reported in a later study (Domino and Hirasawa-Fujita 2019).

Polymorphisms of the  $\mu$ OR gene affect both the levels of  $\mu$ OR expression in the human brain and smoking behavior. Smokers homozygous for the wild-type *OPRM1* A allele show significantly higher [ $^{11}$ C]CFN binding in the bilateral amygdala, left thalamus, and left anterior cingulate cortex than smokers carrying the *OPRM1* A118G allele. In bearers of the G polymorphism, the extent of reward difference between smoking a normal and a denicotinized cigarette was significantly associated with the change of [ $^{11}$ C]CFN binding potential in the right amygdala, caudate, anterior cingulate cortex, and thalamus, but in smokers homozygous for the wild-type gene, this association could not be observed (Ray et al. 2011). Later PET studies in  $\mu$ OR-genotyped overnight abstinent smokers confirmed that smokers bearing the G polymorphism have less free  $\mu$ ORs in some brain areas (amygdala and nucleus accumbens) than carriers of the AA wildtype (Nuechterlein et al. 2016; Domino et al. 2015). In one of these studies, GG carriers were found to show also a less extensive decrease of [ $^{11}$ C]CFN binding potential after the smoking of a “real” cigarette than AA carriers, i.e., a blunted response of the  $\mu$ -opioid system (Domino et al. 2015).

Another PET study examined whether individual differences in  $\mu$ OR availability in alcohol-dependent subjects are associated with tobacco use, nicotine dependence, and level of nicotine craving. Subjects had withdrawn from alcohol use under medical supervision. They were not allowed to smoke but received transdermal nicotine maintenance (21 mg/day). Higher scores in a nicotine dependence test were found to be associated with a lower binding potential of [ $^{11}\text{C}$ ]CFN in the amygdala, cingulate, globus pallidus, thalamus, and insula. The number of cigarettes which subjects had used per day prior to the study was also negatively correlated with [ $^{11}\text{C}$ ]CFN binding potential in these areas. Thus, smoking intensity and severity of nicotine dependence appear to be related to reduced  $\mu$ OR binding potential in several brain regions of alcohol-dependent subjects (Weerts et al. 2014).

### 21.5.5 Eating Disorders

Bulimia nervosa, a disorder characterized by cycles of food restriction, binge eating, and vomiting, shares certain phenomena with addiction and substance abuse. Compared with controls, bulimic individuals show significantly decreased  $\mu$ OR binding in the left insular cortex, a brain area involved in taste discrimination and eating reward. This finding may reflect either downregulation of  $\mu$ OR in the bulimic state as a consequence of chronically increased release of opioid peptides or a personality trait that increases the reward value of dieting (Bencherif et al. 2005).

In another PET study, 7 subjects with binge eating disorder, 15 subjects with pathological gambling, and 17 healthy control subjects were scanned with [ $^{11}\text{C}$ ]CFN and [ $^{18}\text{F}$ ]DOPA. Patients with binge eating disorder showed reductions of  $\mu$ OR binding potential in many cortical and subcortical areas of the brain and a significant decrease of the striatal influx constant of [ $^{18}\text{F}$ ]DOPA. However, in subjects with pathological gambling,  $\mu$ OR binding potential was decreased mainly in the anterior cingulate, and the striatal influx of [ $^{18}\text{F}$ ]DOPA was not significantly altered compared to healthy controls. Thus, two forms of addiction (to eating and to gambling) displayed different patterns of neurobiological changes (Majuri et al. 2017). A later study from the same group used [ $^{11}\text{C}$ ]CFN to measure  $\mu$ OR binding potential in the brains of patients with binge eating disorder, patients with morbid obesity and healthy controls. Both eating disorders were associated with similar, widespread reductions of  $\mu$ OR binding potential. Thus, these two eating disorders shared a common opioid abnormality (Joutsa et al. 2018). In a third study from the same group, 13 pathological gamblers and 15 age-, sex-, and weight-matched healthy control subjects were scanned with [ $^{11}\text{C}$ ]CFN and [ $^{18}\text{F}$ ]DOPA. In both groups, a similar correlation was observed between  $\mu$ OR availability in the basal ganglia (putamen, caudate nucleus and globus pallidus) and the presynaptic capacity for dopamine synthesis. Thus, presynaptic dopamine neurotransmission and opioid receptor function in the basal ganglia appear to be linked and this link is not affected by behavioral addiction (Majuri et al. 2018).

### 21.5.6 Obesity

An initial PET study on the involvement of endogenous opioid systems in obesity scanned seven chronically obese men with [ $^{11}\text{C}$ ]CFN in the fasted and fed condition, at the onset of therapy and after 15% weight loss had been achieved by a low-calorie diet. Seven age- and ethnicity-matched lean control subjects were scanned only once, in the fasted and fed condition. Thus, four PET scans were made in the obese subjects and two scans in the healthy controls. In the initial scans, obese subjects showed lower  $\mu\text{OR}$  binding potentials in several brain regions than lean subjects or obese subjects after weight loss. In healthy controls, the observed decline of tracer binding potential in the right temporal pole after feeding was highly correlated to reductions in negative affect, but this correlation was not observed in the patient group. In patients, the magnitude of this decline at the onset of treatment tended to be associated with weight regain, 1 year after treatment. Thus, the magnitude of  $\mu\text{OR}$  activation in the right temporal pole after an acute meal may predict the risk of weight regain of obese subjects after therapy (Burghardt et al. 2015).

Since both the endogenous opioid system and the dopaminergic system are involved in the regulation of food intake and reward processing, a Finnish study examined pathological changes in  $\mu\text{OR}$  and dopamine  $\text{D}_2\text{R}$  availability in obese subjects. Thirteen obese (mean BMI 42 kg/m $^2$ ) and 14 age-matched healthy women were scanned with the PET tracers [ $^{11}\text{C}$ ]CFN and [ $^{11}\text{C}$ ]raclopride. Obese subjects had significantly lower  $\mu\text{OR}$  binding potentials in ventral striatum, orbitofrontal cortex, amygdala, putamen, insula, and anterior cingulate. [ $^{11}\text{C}$ ]CFN binding potential in such areas was negatively correlated with BMI and self-reported food addiction. However, there were no significant differences between healthy and obese subjects concerning the regional binding potentials of [ $^{11}\text{C}$ ]raclopride (Karlsson et al. 2015). These results can be interpreted as evidence for downregulation of  $\mu\text{OR}$  in the human brain as a consequence of excessive stimulation by endogenous opioids, related to persistent overeating. However, a low density of  $\mu\text{OR}$  could also be a neurochemical trait that predisposes subjects to becoming obese.

In a follow-up study from the same year, group sizes were expanded to 25 obese and 20 healthy female subjects, and the correlation between regional  $\mu\text{OR}$  and  $\text{D}_2\text{R}$  binding potentials was examined (Tuominen et al. 2015). In healthy subjects,  $\mu\text{OR}$  and  $\text{D}_2\text{R}$  availabilities in ventral striatum and dorsal caudate nucleus were positively associated, and  $\text{D}_2\text{R}$  availability in ventral striatum was also associated with  $\mu\text{OR}$  availability in the ventral tegmental area. In obese subjects, an association between  $\mu\text{OR}$  and  $\text{D}_2\text{R}$  binding was observed in the caudate nucleus, but the association in the ventral striatum was much weaker than in lean subjects and the association between striatal  $\text{D}_2\text{R}$  and tegmental  $\mu\text{OR}$  binding was abolished. Thus, the interaction between the opioid and dopaminergic systems may be altered in obesity.

A second follow-up study addressed the question whether reduced  $\mu\text{OR}$  availability is a state or a trait in obesity. Sixteen obese women were scanned twice with [ $^{11}\text{C}$ ]CFN and [ $^{11}\text{C}$ ]raclopride, both before and 6 months after bariatric surgery (gastric bypass or sleeve gastrectomy). Measured binding potentials were compared to those of an age-matched healthy control group ( $n = 14$ ). A reduced availability of

$\mu$ OR in ventral striatum, insula, amygdala, and thalamus was initially observed, but the binding potential of [ $^{11}$ C]CFN in these areas normalized after gastric surgery and weight loss ( $26.1 \pm 7.6$  kg). Apparently, the reduced availability of  $\mu$ OR in the patient group was associated with the obese phenotype and was not a persistent neurochemical trait. In contrast to  $\mu$ OR, the binding potential of D<sub>2</sub>R was not altered after surgery (Karlsson et al. 2016).

### 21.5.7 Epilepsy

Opioid receptors and endogenous opioid peptides are known to play a role in the mechanisms underlying seizures. Many PET studies with opioid receptor ligands have been performed in patients with epilepsy. The first of these involved [ $^{11}$ C]CFN and subjects with complex partial seizures due to unilateral temporal seizure foci. [ $^{11}$ C]CFN binding in the temporal neocortex was greater on the side of the focus than on the contralateral side. This increase was interpreted as evidence for involvement of  $\mu$ OR in a tonic anticonvulsant system that limits the spread of electrical activity (Frost et al. 1988). In contrast to [ $^{11}$ C]CFN binding, [ $^{11}$ C]DPN uptake (Mayberg et al. 1991) and [ $^{18}$ F]cyclofoxy (Theodore et al. 1992) uptake in the ipsilateral and contralateral lobes were not significantly different, suggesting that opioid receptor subtypes are differentially regulated in temporal lobe epilepsy. Evidence for differential regulation was obtained in a later study in which subjects with temporal lobe epilepsy were scanned both with the  $\delta$ OR ligand [ $^{11}$ C]methylnaltrindole ([ $^{11}$ C]MeNTI) and with the  $\mu$ OR ligand [ $^{11}$ C]CFN. Binding of both ligands was increased in the ipsilateral temporal cortex, but the regional pattern of the changes was different (more extended for [ $^{11}$ C]MeNTI than for [ $^{11}$ C]CFN). Upregulation of  $\delta$ OR may indicate an anticonvulsant action for this receptor subtype (Madar et al. 1997). A more recent study examined [ $^{11}$ C]DPN binding in the brain of patients with temporal lobe epilepsy both shortly (within a few hours) after spontaneous epileptic seizures and interictally. In 14 healthy controls, no changes of [ $^{11}$ C]DPN binding were observed as a function of time, but in the patients, an increase of tracer binding was noted in the ipsilateral temporal pole and fusiform gyrus, shortly after seizures. The magnitude of this increase was inversely correlated to the interval that had elapsed since the last seizure, suggesting an increase of receptor binding during seizures followed by a gradual return to baseline (Hammers et al. 2007b). A follow-up article that was published after 6 years pointed out that in order to correctly quantify opioid receptor availability after spontaneous epileptic seizures, [ $^{11}$ C]DPN data need to be corrected for the partial volume effect. After such correction, post-ictal increases in tracer distribution volume could also be detected in the hippocampus (McGinnity et al. 2013). Thus, results from PET studies with three different ligands ([ $^{11}$ C]CFN, [ $^{11}$ C]MeNTI, and [ $^{11}$ C]DPN) have supported the idea that the opioid system is involved in seizure control.

The interictal distribution volume of [ $^{11}$ C]DPN in the cortex and thalamus of eight patients with childhood and juvenile absence epilepsy was found to be not significantly different from values in healthy age-matched controls; thus, there



appeared to be no overall abnormality of opioid receptors in this patient group (Prevett et al. 1994). To examine whether absence seizures are associated with release of endogenous opioids, investigators scanned eight patients with primary generalized epilepsy and eight control subjects with [ $^{11}\text{C}$ ]DPN and PET. Serial absences were precipitated in the patients by hyperventilation for 10 min, starting 30–40 min after injection of the tracer. Increased washout of DPN was observed in the association cortex but not in other brain areas during seizures, as compared with control subjects and patients scanned without provocation of absences. Pharmacokinetic modeling suggested that absence seizures resulted in a significant, 15–41% decrease in the rate constant of association ( $k_3$ ) of the tracer. These data were interpreted as evidence for the release of endogenous opioids in the association cortex at the time of absences, leading to increased opioid receptor occupancy (Bartenstein et al. 1993).

A later study involved five patients with reading epilepsy who were scanned both at baseline (reading a string of symbols) and during seizure activation (reading a scientific paper). The latter condition was associated with significantly lower binding of [ $^{11}\text{C}$ ]DPN in the left parietotemporal-occipital cortex of the patients as compared with six healthy controls, suggesting release of endogenous opioids as part of a mechanism to terminate reading-induced seizures (Koepp et al. 1998).

Two patients with mesobasal temporal lobe epilepsy have been scanned with [ $^{11}\text{C}$ ]DPN, before and after selective amygdalohippocampectomy. After removal of the epileptic focus, [ $^{11}\text{C}$ ]DPN binding in the ipsilateral frontal cortex was found to be reduced, suggesting either downregulation of opioid receptors in the absence of seizures or postoperative neuronal dysfunction (Bartenstein et al. 1994).

## 21.5.8 Neurodegenerative Diseases

Losses of opioid receptors with different regional patterns have been observed in several neurodegenerative diseases.

### 21.5.8.1 Huntington's Disease

Significant decreases of [ $^{11}\text{C}$ ]DPN binding (24–40%) were noted in the caudate and putamen of Huntington's disease (HD) patients using striatum-to-occipital cortex uptake ratios, spectral analysis, voxelwise parametric analysis, and statistical parametric mapping (SPM) for quantification (Weeks et al. 1997).

### 21.5.8.2 Alzheimer Disease

Decreases of opioid receptor binding (up to 40%) have been detected in the brain of Alzheimer disease (AD) patients using the tracer [ $^{18}\text{F}$ ]cyclofoxy, and these changes were not correlated with decreases in regional cerebral blood flow (Cohen et al. 1997).

### 21.5.8.3 Parkinson Disease and Related Disorders

An initial study on eight clinically defined Parkinson disease (PD) patients, seven subjects with the striatonigral degeneration type of multiple system atrophy and six subjects with Steele-Richardson-Olszewski (SRO) syndrome examined opioid receptor binding in the human striatum with the tracer [ $^{11}\text{C}$ ]DPN and PET. In the PD patient group, tracer binding was not significantly altered compared to healthy controls. Striatonigral degeneration was associated with reduced binding of [ $^{11}\text{C}$ ]DPN in the putamen, but not in the caudate. In the SRO syndrome group, both caudate and putamen opioid receptor binding were significantly reduced. The binding pattern of [ $^{11}\text{C}$ ]DPN may thus help to differentiate between various akinetic-rigid syndromes (Burn et al. 1995). In ten patients with the olivopontocerebellar variant of multiple system atrophy, a significant 12% reduction of the caudate-occipital ratio and a 15% reduction of the putamen-occipital uptake ratio of [ $^{11}\text{C}$ ]DPN were noted. The latter decline was correlated to loss of [ $^{18}\text{F}$ ]DOPA uptake in the putamen (average reduction 29%) (Rinne et al. 1995).

Using [ $^{11}\text{C}$ ]DPN-PET and either a region-of-interest (ROI) or an SPM approach, a significantly reduced opioid receptor binding was later noted in the striatum and thalamus of PD patients with levodopa-induced dyskinesias but not in non-dyskinetic subjects (Piccini et al. 1997). No difference in striatal dopamine  $\text{D}_1$  or  $\text{D}_2$  receptor binding was found between the two subgroups, and measurements with PET and [ $^{15}\text{O}$ ]water showed that rCBF after oral administration of levodopa was increased during dyskinesias in lentiform nuclei and motor, premotor, and dorsal prefrontal cortex. Thus, dyskinesias appear to arise not from a disturbance of dopamine receptor availability but rather from overactivity of opioid transmission, particularly the basal ganglia-frontal projections (Brooks et al. 2000).

In bilaterally 1-methyl-4-phenyl-1,2,3,6-tetrahydropyridine (MPTP)-lesioned monkeys which had clinically recovered from the acute motor effects of dopaminergic neuron lesioning, a 65% decrease in the accumulation of [ $^{18}\text{F}$ ]DOPA was noted in the basal ganglia. This decrease was associated with a 30–35% decline of [ $^{18}\text{F}$ ]cyclofoxy binding in the caudate, anterior putamen, thalamus, and amygdala. The authors concluded that altered opioid receptor signaling (probably increased levels of Met-enkephalin) had contributed to the behavioral changes which were observed in the animals, i.e., the masking of their motor symptoms (Cohen et al. 1998). A later study from the same group showed that in unilaterally MPTP-lesioned monkeys with parkinsonian symptoms, opioid receptor availability is reduced by 30–35% on both the lesioned and the non-lesioned sides of the brain (Cohen et al. 1999).

In contrast to the decreases of opioid receptor binding which were observed in akinetic-rigid syndromes and PD patients with dyskinesias, no abnormalities of [ $^{11}\text{C}$ ]DPN binding were noted in patients with primary torsion dystonia (carriers of the *DYT1* gene), and no correlation between the severity of dystonia and opioid binding could be detected (Whone et al. 2004).

### 21.5.9 Opioid Receptor Expression in Lung Tumors

A pilot PET study involving seven patients with lung carcinomas examined the feasibility of tumor imaging with the radioligands [ $^{11}\text{C}$ ]MeNTI and [ $^{11}\text{C}$ ]CFN. Four of these patients were also scanned with [ $^{18}\text{F}$ ]FDG for clinical indications. All tumors showed an accumulation of opioid receptor ligands above background. The tumor-to-nontumor binding ratio of the  $\delta\text{OR}$  ligand [ $^{11}\text{C}$ ]MeNTI ( $4.3 \pm 1.3$ ) was greater than that of the  $\mu\text{OR}$  ligand [ $^{11}\text{C}$ ]CFN ( $2.4 \pm 1.2$ ) but lower than that of [ $^{18}\text{F}$ ]FDG ( $7.7 \pm 0.5$ ). About 50% of [ $^{11}\text{C}$ ]MeNTI uptake and 44% of [ $^{11}\text{C}$ ]CFN uptake in the tumors could be blocked by naloxone. Particularly [ $^{11}\text{C}$ ]MeNTI appears suitable for investigation of lung carcinoma biology, since  $\delta\text{OR}$  may be involved in tumor invasion and metastasis (Madar et al. 2007). A ligand targeting  $\mu\text{OR}$  and  $\delta\text{OR}$  was tested in tumor-bearing mice as a strategy for delivering cytotoxic drugs to lung cancer, but unfortunately, this strategy failed since the specificity of payload delivery was inadequate (Li and Low 2017).

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## 21.6 Conclusion

PET imaging has provided insight in the involvement of opioid system in affective and sensory processing. Changes of opioid receptor expression related to increasing age and altered hormonal status have also been reported. Moreover, OR imaging has been successfully applied in the study of the pharmacokinetics and pharmacodynamics of novel and existing drugs.

Regional changes of OR availability have been observed in psychiatric and neurodegenerative disorders, epilepsy, pain, bulimia nervosa, behavioral addiction, and substance abuse. PET imaging may contribute to the development of improved techniques for the treatment of pain (and addiction) by monitoring the relationship between OR occupancy and symptom relief.

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# PET Imaging of ABC Transporters at the Blood-Brain Barrier

# 22

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## Abstract

The function of ATP-binding cassette (ABC) transporters at the blood-brain barrier (BBB) is to protect the brain from toxic compounds. Additionally, they play a crucial role in the onset and progression of several central nervous system (CNS) diseases as well as in drug resistance. Many compounds were identified as substrates, inhibitors, inducers, or activators for ABC transporters, causing important drug-drug interactions. PET imaging represents an excellent tool for assessing the function and expression of ABC transporters. Over the last years, many PET tracers with different characteristics have been developed, mainly for measuring P-glycoprotein (P-gp) function at the BBB. Although (*R*)-[<sup>11</sup>C]verapamil or [<sup>11</sup>C]*N*-desmethyl-loperamide are considered as the “gold standard” P-gp tracers, they have several drawbacks such as its high affinity to P-gp which limits its use for assessing P-gp increased function. Therefore, PET tracers with lower affinity to the transporter have been developed and studied in different species. The assessment of ABC transporters by PET imaging can provide new insight into the physiology and pathophysiology of different CNS diseases and may open new avenues for therapies. Moreover, PET can be used for screening the affinity of new entities toward various ABC transporters and thus enhance the development of CNS drugs.

## 22.1 ABC Transporter Expression at the Blood-Brain Barrier in Neurological Diseases

### 22.1.1 Blood-Brain Barrier

The function of the human brain requires a strictly controlled environment. The blood-brain barrier (BBB) plays an important role in maintaining homeostasis of this micro-environment by protecting the brain from neurotoxic substances since it functions as a biochemical and physical barrier (Hawkins and Davis 2005). This barrier was first described by Paul Ehrlich, who found that trypan blue injections in the peripheral blood circulation did not reach the spinal cord and brain in rabbits (Templeton et al. 1997). Nowadays, we know the BBB consists of a monolayer of non-fenestrated brain microvessel endothelial cells connected by tight junctions, which limits transport by passive diffusion across the BBB. Only small lipophilic molecules and drugs with molecular weights less than 400 Da and less than eight hydrogen bonds can cross the barrier by passive diffusion (Morris et al. 2017).

For compounds that cannot enter the brain by passive diffusion due to their ionization, hydrophilicity, or size, membrane transporters at the endothelial cells can facilitate the transport across the BBB (Zhao et al. 2015). Apart from the limited transport by passive diffusion, the BBB also presents numerous efflux transporters that carry a wide variety of molecules from the brain to the blood. The best-known efflux transporters are the ATP-binding cassette transporters (ABC transporters). These transporters play an important role in protecting the central nervous system (CNS). In this chapter, we will focus on the ABC transporters, with a particular emphasis on P-glycoprotein and its involvement in several neurological disorders, such as Alzheimer's disease, Parkinson's disease, and epilepsy.

### 22.1.2 ABC Transporters

The ABC transporter superfamily represents the largest family of transmembrane proteins and consists of about 50 members classified into 7 different subfamilies. These transporters are ATP-dependent and use this energy to drive the transport of various endogenous and exogenous molecules across cell membranes (Dean and Allikmets 2001; Dean et al. 2001). They play a significant role in various CNS disorders and their pharmaceutical treatments in two different ways: First, many pharmaceuticals used as therapeutic drugs in CNS diseases are substrates of ABC transporters, which limits their access to the brain; and second, the expression and function of ABC transporters may be altered in certain CNS diseases, either as a cause or a consequence of the disease, resulting in altered drug entry in the CNS (see Table 22.1).

P-gp, also known as ABCB1, was the first ABC transporter described and has by far been the most studied. This transporter is found at the luminal side of the endothelial cells at the BBB and acts as an efflux transporter for various structurally

**Table 22.1** CNS diseases involving impairments in ABC transporters

CNS disease	ABC transporter involved	The main change in ABC transporters activity	References
Alzheimer's disease	ABCA1, ABCA2, ABCB1, ABCG2	Decreased P-gp, BCRP, and MRP1 expression and function	Wolf et al. (2012), Macé et al. (2005), Xiong et al. (2009), Lam et al. (2001), Krohn et al. (2011), Kuhnke et al. (2007), Riddell et al. (2007))
Parkinson's disease	ABCB1	Decreased P-gp expression and function	Kortekaas et al. (2005), Westerlund et al. (2008), Zschiedrich et al. (2009), Liu et al. (2013)
Epilepsy	ABCC1, ABCCs, ABCG2	Increased P-gp, BCRP, MRP1, and MRP2 expression and function	Löscher and Potschka (2005a), Wang et al. (2016), Summers et al. (2004), Löscher and Potschka (2005b), Hartz et al. (2017), van Vliet et al. (2005), Syvänen et al. (2013), Feldmann and Koepf (2012), Aronica et al. (2012)
Brain tumors	ABCA1, ABCB1, ABCC1, ACC4, ABCG2	Increased P-gp, BCRP, and MRP1 function and expression	Haber et al. (2006), Agarwal et al. (2011), Porro et al. (2010), Parrish et al. (2015), Liu et al. (2015), Deeken and Loscher (2007), Sane et al. (2014)
Ischemic stroke	ABCB1, ABCG2, ABCC1, ABCC5	Increased P-gp, BCRP, and MRP1 expression	ElAli and Hermann (2010), Kilic et al. (2008), Dazert et al. (2006), Ueno et al. (2009)
Multiple sclerosis	ABCB1, ABCD3, ABCG2	Decreased P-gp expression	Kooij et al. (2011), Kooij et al. (2010), Gray et al. (2014)
Amyotrophic lateral sclerosis	ABCB1, ABCG2	Increased P-gp and BCRP activity	Milane et al. (2010), Jablonski et al. (2012, 2014)
HIV-encephalitis	ABCB1, ABCC1, BCC4, BCC5, ABCG2	Increased BCRP and MRP1 expression	Robillard et al. (2014), Hayashi et al. (2005)
Lysosomal storage diseases	ABCC4	Increased MRP4 function	Schaheen (2006)
Depression	ABCB1	Increased P-gp function	de Klerk et al. (2009), Brzozowska et al. (2017)
Schizophrenia	ABCB1	Decreased P-gp expression	Bernstein et al. (2016)
Creutzfeldt-Jakob disease	ABCB1	Decreased P-gp expression	Vogelgesang et al. (2006)

different substrates (see Table 22.2). Besides P-glycoprotein, other ABC transporters with a similar function exist. ABCG2 (or breast cancer resistance protein BCRP), ABCC1 (or multidrug resistance protein MRP1), ABCC4, and ABCC5 are also localized at the luminal side of the human brain capillary endothelial cells of the

**Table 22.2** Examples of drugs that act as substrates of P-gp, BCRP, and MRP1 transporters (updated from (Mahringer and Fricker 2016; Qosa et al. 1628))

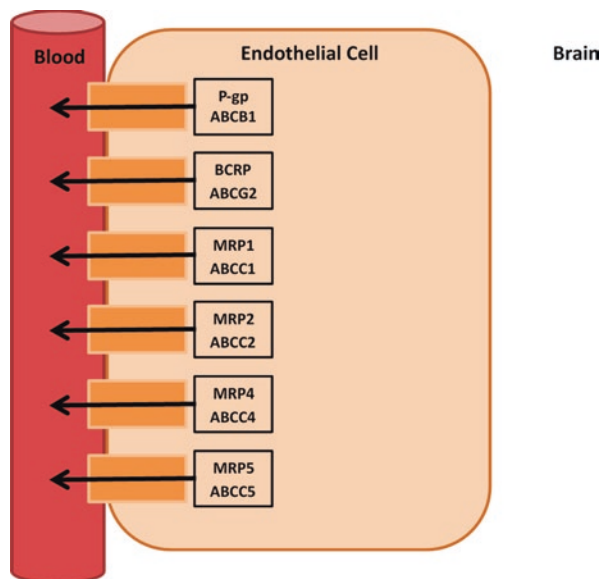
ABC transporter	Drug class	Substrates drugs
P-gp	Cytostatic	Vinblastine, vincristine, doxorubicin, paclitaxel, methotrexate
	Immunosuppressant	Cyclosporine A, tacrolimus
	Corticoids	Dexamethasone, aldosterone, cortisol
	Analgesics	Morphine, methadone, fentanyl
	Antiretroviral HIV	Ritonavir, indinavir
	Anthelmintic	Ivermectin
	Antidiarrheal	Loperamide
	Antiemetic	ondansetron
	Antibiotic	Erythromycin, tetracyclines, rifampicin
	Antiarrhythmic	Amiodarone
	Cardiac glycoside	Digoxin
	Calcium channel blocker	Verapamil, nifedipine, diltiazem
	Beta-blocker	Propranolol
	Statin	Lovastatin, simvastatin
	Antiepileptic	Phenytoin, carbamazepine, phenobarbital
Antipsychotic	Chlorpromazine	
Antidepressant	Paroxetine, amitriptyline, venlafaxine	
BCRP	Cytostatic	Anthracyclines, mitoxantrone, etoposide, erlotinib, gefitinib
	Antiretroviral HIV	Zidovudine, lamivudine
	Antibiotics	Ciprofloxacin, norfloxacin
	Immunosuppressant	Cyclosporine A, tacrolimus
MRP1	Cytostatic	Etoposide, vincristine, doxorubicin, methotrexate
	Immunosuppressant	Tacrolimus, cyclosporine A
	Antiretroviral HIV	Saquinavir, ritonavir
	Antiepileptic	Levetiracetam, phenobarbital
	Antibiotic	Azithromycin, ciprofloxacin

BBB (see Fig. 22.1). ABCC6, ABCC11, and ABCC12 are also present in the microvasculature of the brain, albeit at lower expression levels (Dauchy et al. 2008).

### 22.1.3 P-Glycoprotein

P-glycoprotein (P-gp) or ABCB1, encoded by the MDR1 gene, was the first ABC transporter discovered and remains the most studied one. This 170 kDa membrane protein is located between the cytoplasm and the periplasm and consists of a single chain with two homologous parts. At the intracellular side of the two homologous parts, six transmembrane (TM) helices and two ATP nucleotide-binding domains (NBDs) are present. During the efflux of a substrate, translational and rotational

**Fig. 22.1** ABC transporters in the human endothelial cell at the BBB update from (Mahringer and Fricker 2016; Liu 2019)



motions in P-gp take place: ATP causes a change in the TM helix, which will catalyze the efflux of the P-gp substrate through the transmembrane domain (TMD) and lipid bilayer.

P-gp is involved in the efflux of a broad range of structurally different substrates including analgesics, antihistamines, and antibiotics and is therefore thought to play an important role in drug resistance (Tamaki et al. 2011) (see Table 22.2). Besides the efflux of many pharmaceuticals, P-gp transports also endogenous compounds such as amyloid- $\beta$  and, therefore, may play an important role in the pathophysiology of Alzheimer's disease.

### 22.1.4 BCRP

BCRP is encoded by the ABCG2 gene located on chromosome 4q22. This transporter was first discovered in 1998 by Doyle and colleagues in a multidrug-resistant human breast cancer cell line (Doyle et al. 1998). BCRP consists of a polytopic TM protein with 655 amino acids. It differs from P-gp in its structure since it is a half ABC transporter, consisting of only one NBD and one membrane-spanning domain (MSD) (Ni et al. 2010). BCRP is responsible for the efflux of more than 200 different substrates, including chemotherapeutics, tyrosine kinase inhibitors, antivirals, carcinogens (see Table 22.2) and also physiological substances, such as uric acid and estrone-3-sulfate. This makes BCRP one of the most important efflux transporters at the BBB. Although the substrate and inhibitor specificity of BCRP show some overlap with P-gp and MDR1, they are not completely identical, as can be explained by the different protein structures and working mechanisms (Mao and Unadkat 2015).

#### 22.1.4.1 MRP Family and MRP1

Besides P-gp and BCRP, members of the multidrug resistance protein (MRP) family are efflux transporters at the BBB. The transporters in this family are less studied and characterized than P-gp and BCRP, and their function at the BBB is less clear, although it is known that the MRP family is responsible for the efflux of a diverse range of compounds including xenobiotics and endogenous substrates. Subfamily ABCC, which expresses the MRP transporters, contains 12 proteins, and 9 of these are known to be involved in ATP-dependent transmembrane efflux of drugs in various cells and tissue types in the human body. Proteins in the MRP subfamily share common structural features including multiple TM helices in MSDs and NBDs at the intracellular side for the binding of ATP. MRP1, MRP2, MRP3, MRP6, and MRP7 have an additional NH<sub>2</sub>-proximal MSD, and therefore they are called the “long MRPs” (Haimeur et al. 2004). The first of the nine drug-transporting MRPs is MRP1, which is encoded by the ABCC1 gene on chromosome 16p13.1 and is the most studied and best-known MRP family member. It was discovered by Cole et al. in 1992 as a cause of multidrug resistance in a drug-selected human lung cancer cell line (Cole et al. 1992) and it is because of this broad substrate specificity (Table 22.2) that MRP1 continues to be of considerable interest in preclinical and clinical studies.

#### 22.1.4.2 Alzheimer’s Disease

The incidence of neurodegenerative diseases has increased significantly over the past few decades because the life expectancy of the human population has increased. In 1906 Alzheimer’s disease was first described by Alois Alzheimer as a particularly severe disease process of the cerebral cortex (Alzheimer 1906; Hippus and Neundörfer 2003). Clinically Alzheimer’s disease is characterized by a decline in memory, language, and cognitive skills, which affects the patient’s ability to perform everyday activities (Ballard et al. 2011). Alzheimer’s disease is an irreversible and incurable condition. The underlying cause of these symptoms can be found in the destruction of neurons in brain regions involved in the cognitive function. Although the exact pathogenesis of Alzheimer’s remains unclear, hallmarks of AD are the accumulation of neurotoxic amyloid- $\beta$  ( $A\beta$ ) plaques in the extracellular compartment of the brain, inflammation, accumulation of hyper phosphorylated tau in intracellular neurofibrillary tangles and neuronal loss (Zlokovic 2011). The amyloid hypothesis suggests that this accumulation of  $A\beta$  is caused by an imbalance of overproduction and impaired clearance, and this may be the key element triggering neuronal degeneration (Hardy and Allsop 1991). Although no significant association was found between ABCB1 polymorphisms and the risk of developing AD in humans, P-gp is thought to play an important role in the pathophysiology of AD since this efflux transporter facilitates transport of  $A\beta$  out of the brain (Vogelgesang and Jedlitschky 2014; Cirrito et al. 2005). Various studies have reported an inverse correlation between  $A\beta$  deposition and cerebrovascular ABCB1 expression (Pereira et al. 2017; Zhao et al. 2017).



### 22.1.4.3 Parkinson's Disease

Parkinson's disease (PD) is a neurodegenerative disorder characterized by clinical symptoms of uncontrolled movement and in a later stage by dementia. The symptoms of this disease are caused by the death of dopamine generating cells in the substantia nigra. The reason for this cell death is unknown, although it is known that exposure to neurotoxic pesticides is a risk factor for the development of PD (Steece-Collier et al. 2002; Wirdefeldt et al. 2011; Pezzoli and Cereda 2013). Pesticides are a group of compounds with various mechanisms of action and pharmacokinetic properties, including fungicides, herbicides, and insecticides. In epidemiologic studies in humans, toxicity models in animals, and in vitro studies, exposure to pesticides was shown to be associated with an increased risk of the development of PD. An alteration of the BBB transport of such neurotoxic pesticides has been proposed as an underlying pathophysiological mechanism that may lead to PD. Since P-gp has an extensive range of substrates, P-gp may be involved in the removal of neurotoxic pesticides out of the brain. However, it is still unknown if P-gp is truly involved in this transport, although multiple studies did investigate the association between ABCB1 and PD. In only a few studies, an association between ABCB1 alterations and Parkinson's disease was found, especially in later stages of the disease (Drożdżik et al. 2003; Marzolini et al. 2004; Westerlund et al. 2009; Bartels et al. 2008a, b).

### 22.1.4.4 Epilepsy and Antiepileptic Drugs

Epilepsy is known as a serious neurological disorder characterized by unpredictable seizures. Epilepsy is affecting more than 60 million people worldwide. Intractable epilepsy (IE), in which a patient's seizures fail to come under control with pharmacotherapeutics, is a severe form of epilepsy, occurring in 20–30% of patients diagnosed with epilepsy. The underlying mechanisms of IE are not fully clear, although there are several explanations found in the literature. One of these focuses on overexpression of P-gp transporters, MDR transporters, and other efflux transporters at the BBB in the cerebrovascular endothelium around the epileptic focus in the brain (Wang et al. 2016; Feldmann et al. 2013). However, it is not clear whether this upregulation of P-gp is caused by epilepsy, by uncontrolled recurrent seizures, by drug-associated induction of efflux transporters, by genetic polymorphisms, or by a combination of all of these factors (Tang et al. 2017).

### 22.1.4.5 Ischemic Stroke

During ischemic stroke, the blood flow is locally disrupted leading to deprivation of brain tissue from oxygen and glucose. Subsequently, a lack of sufficient oxygen activates hypoxia-inducible factor 1- $\alpha$  (HIF-1 $\alpha$ ) and nuclear factor  $\kappa$ B, leading to a time-dependent increase in P-gp expression (Wang et al. 1995; Koong et al. 1994; Cen et al. 2013). This increase is thought to be an attempt of the ischemic brain to maintain a functional barrier between the blood and the brain when the BBB is compromised (Spudich et al. 2006; Lazarowski et al. 2007). However, an increase in P-gp function might cause a reduced uptake of therapeutic drugs in the brain after ischemic stroke.

#### 22.1.4.6 Aging

Since aging is the most important risk factor for the development of AD, and P-gp is a known efflux transporter of A $\beta$ , there is an increasing interest in the function of P-gp in normal aging and Alzheimer's disease. A $\beta$  transport at the BBB is significantly altered with increasing age. In 2002, Vogelgesang et al. studied brain tissue samples of 243 non-demented subjects and found there was an age-related decrease in P-gp in association with an increase of amyloid deposition (Vogelgesang et al. 2002a). A later PET study with the tracer (*R*)-[<sup>11</sup>C]verapamil also reported a decrease of P-gp function with increasing aging (Bauer et al. 2009). Chiu et al. showed a trend of reduced P-gp function with advancing age and also a significant reduction in P-gp function in Alzheimer's disease patients compared with healthy elderly controls (Chiu et al. 2015). Amyloid transporter gene expression is age-dependent, which may result in changes in protein expression at the BBB (Osgood et al. 2017). Besides a reduction in P-gp function, a reduction in lipoprotein receptor-related protein 1 (LRP-1) efflux transporter function and an increase in the function of receptors for advanced glycation end products (RAGE) were found in animal studies. The latter changes may result in an increased influx of A $\beta$  into the brain (Church et al. 2014).

#### 22.1.4.7 Polymorphism, Genotyping, and Individualized Medicine

The ABCB1 gene, which codes for P-gp, displays more than 50 single nucleotide polymorphisms (SNPs) (Sharom 2008). Variability in pharmacogenes can influence the metabolism, transport, and elimination of drugs. This makes the variability in pharmacogenes an interesting field of research since it may alter the response of patients to specific pharmaceuticals. Different forms of such genes may be associated with interindividual variations in drug efficacy or the severity of unwanted side effects. Pharmacogenetic data is also of great value in the prediction of neurological diseases. Since ABC transporters influence the uptake of many CNS pharmaceuticals, the question arises whether polymorphisms may predict therapy outcome. The hypothesis that variation in the ABCB1 gene is associated with antidepressant treatment outcome has been tested in various clinical studies, and the results were inconclusive. In a meta-analysis of 16 studies in which the association between ABCB1 polymorphisms and antidepressant treatment outcome was analyzed, Breitenstein et al. found that SNP rs2032583 and rs2235015 have a significant influence on treatment outcome and could possibly be used as predictors of treatment outcome in depression (Breitenstein et al. 2015). However, despite the extensive number of studies performed, no unambiguous answer to the question of the predictive value of ABCB1 polymorphisms was reached. Multiple studies have found an association between MDR1 C3435T polymorphism and antiepileptic drug resistance (Wang 2015; Keangraphun et al. 2015).

#### 22.1.4.8 Drug Resistance and ABC Transporters

Pharmaceuticals can cross the BBB and enter the brain via six major pathways: paracellular transport, passive transcellular diffusion, carrier-mediated transport, receptor-mediated transcytosis, adsorptive-mediated transcytosis, and cell-mediated transport. In various diseases including CNS disorders, oncologic diseases,

and infectious diseases, the effect of treatment is limited because of drug resistance. A prerequisite for adequate distribution of the drug to its target is adequate transport between various tissues and compartments. Distribution of drugs to the brain is dependent on both the properties of the BBB and the properties of the drug (Yamamoto et al. 2018; Elsinga et al. 2004). An efflux transporter as P-glycoprotein can have a great impact on the distribution of various drugs and is thought to be responsible for drug resistance in cancer, epilepsy, Alzheimer's disease, and schizophrenia (Bartels 2011; Loscher et al. 2012; Rosenhagen and Uhr 2011). Drug resistance may be due to two underlying causes: First, the drugs used in the treatment of those diseases are substrates of ABC transporters and second, the expression of the ABC transporters involved in drug efflux can be altered by the disease itself. Studies in the field of drug resistance have led to important new developments, such as specific transport inhibitors, specific antibodies for the transporters, and new *in vivo* imaging techniques (e.g., novel PET tracers). However, drug resistance is still a major therapeutic obstacle, and further studies are required.

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## 22.2 Mechanism of Action of ABC Transporter-Binding Substances

Typically, the ABC transporters consist of a pair of NBDs, present on the membrane side in contact with the cytoplasm and two TMDs. There exist ABC transporters with a different number of TMDs and NBDs. The NBDs are responsible for the ATP binding and hydrolysis, and thus they provide the energy for the active transport of substances. The TMDs are the substrate-binding site which contributes to the transport specificity (Linton and Higgins 2007).

It was initially thought that the P-gp transporter possessed two binding sites (Shapiro and Ling 1997). Then, a third ligand-binding site was discovered (Shapiro et al. 2001). And later on, Martin et al. suggested that the P-gp transporter contained four ligand-binding sites (Martin et al. 2000). Sites I, II, and III work as transport sites because they interact with substrates (such as paclitaxel, vinblastine) and modulators (tariquidar). Site IV, on the other hand, acts as a regulatory site. Since a wide number of compounds can also interact with BCRP and MRP1, these transporters may also have several ligand-binding sites similar to the P-gp transporter (Gameiro et al. 2017; Colabufo et al. 2010).

The compounds that interact with the ABC transporters can behave as substrates, inhibitors, inducers, or activators. Substrates are pumped out of the cell in an ATP-dependent manner, inhibitors reduce the ABC transporter function, inducers increase the transporter expression, and activators enhance the transporter function. Notice that one compound can have different modes of action (depending on the administered dose): drugs that are P-gp substrates can also act as P-gp inhibitors or inducers (Zhou 2008). It is important to mention that induction and activation are not always related. A compound can increase the expression level of the transporter but not increase its functionality and vice versa (Gameiro et al. 2017).

Different hypotheses try to explain how inhibitors interfere with the transporter. The inhibitor may act by blocking the ligand-binding site, by preventing the ATP hydrolysis or by modifying the integrity of the lipid membrane. There are different types of inhibitors, classified by their potency and selectivity (Silva et al. 2015). Some P-gp inhibitors are used as therapeutic agents and are co-administrated with chemotherapeutic agents or antiepileptic drugs to overcome drug resistance in cancer or epilepsy (Zhou 2008).

An inducer promotes upregulation of the protein expression (Gameiro et al. 2017). Many structurally unrelated compounds have been reported as P-gp inducers such as abacavir, aldosterone, hypericin, morphine, and rifampicin (Silva et al. 2015). Numerous routes can regulate the transcription of the MDR1 (ABCB1) gene, which highlights the ability of P-gp to respond to a wide variety of stimuli. Upregulation of the MDR1 gene depends on different transcription factors (pregnane X receptor (PXR), heat-shock transcription factor 1 (HSF-1), nuclear factor Y (NF-Y), and early growth response protein 1 (EGR-1)) that do not bind directly to P-gp (Silva et al. 2015). The nuclear factor PXR can be activated by diverse compounds such as rifampicin, phenobarbital, and mifepristone. PXR upregulates the MDR1 expression, which subsequently leads to an increased expression of P-gp (Zhou 2008; Miller et al. 2008).

Activators bind to the protein and induce a conformational change that enhances the transport of other substrates bound to a different ligand-binding site (Gameiro et al. 2017). These compounds increase the P-gp function without increasing the protein expression. Therefore, activators induce the P-gp function as long as they are present. This is a more rapid mechanism than the action of P-gp inducers (Silva et al. 2015).

Since many compounds can modify the function of the ABC transporters (in particular P-gp and BCRP), the Food and Drug Administration (FDA) and the European Medicine Agency (EMA) developed guidelines for the screening of drugs that cause transporter-mediated drug-drug interactions (DDI) (Colabufo et al. 2010; Clinical Drug Interaction Studies — Study Design, Data Analysis, and Clinical Implications Guidance for Industry 2017; Guideline on the Investigation of Drug Interactions 2012). Concomitant administration of drugs that can interfere with ABC transporters may cause alterations of drug concentrations and, therefore, may lead to decreases in drug efficacy or to toxicity. Appropriate measures should be taken to prevent the clinical implications of transport-mediated DDI. Treatment effects should be monitored, administered dose should be adjusted if necessary, and alternative therapies may have to be sought (Lund et al. 2017).

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### **22.3 Use of P-gp Modulators to Treat Drug Resistance, Neurological Disorders, and Other Conditions**

Since P-gp dysfunctions are encountered in many different diseases of the CNS, P-gp modulators may restore the proper function and may improve the disease condition.

### 22.3.1 P-gp Inhibitors and Drug Resistance

Resistance to drug treatment is a common clinical issue in cancer and infectious diseases, and it also occurs in rheumatoid arthritis, epilepsy, depression, and other psychiatric disorders. The role of P-gp and other ABC transporters in drug resistance has gained increasing attention (Löscher and Potschka 2005c).

P-gp inhibitors have been developed and used to overcome drug resistance in a wide variety of diseases. P-gp inhibitors can be classified into four generations (Silva et al. 2015; Li et al. 2016): The first generation of P-gp inhibitors includes verapamil and cyclosporine A. These inhibitors are characterized by their low therapeutic response and high toxicity (cardiovascular toxicity). The second generation (e.g., valspodar) was developed in order to achieve higher potency and reduced toxicity. However, these compounds also inhibit cytochrome P450, and many DDIs were encountered when the inhibitors were co-administrated with chemotherapeutic agents. Thus, a third generation of P-gp inhibitors was developed. These compounds were 200-fold more potent than the first and second generation of inhibitors (Li et al. 2016). Some of these inhibitors (e.g., tariquidar) were evaluated in clinical trials (Li et al. 2016; Fox and Bates 2007). Recently, natural substances have been proposed to reverse drug resistance. These compounds, which represent the fourth generation of P-gp inhibitors (Silva et al. 2015), are obtained from different natural sources. They include alkaloids, flavonoids, coumarins, resins, etc. and are characterized by low cytotoxicity and increased oral bioavailability (Li et al. 2016; Dewanjee et al. 2017).

In cancer drug resistance, it has been suggested that the overexpression of efflux transporters in tumors reduces the drug concentration in the target tissue and consequently limits the treatment efficacy (Juliano and Ling 1976; Fletcher et al. 2016). P-gp, MRP1, and BCRP are the most relevant efflux transporters related to drug resistance. A wide variety of anticancer drugs such as vinca alkaloids (vincristine), taxanes (docetaxel and paclitaxel), anthracyclines (doxorubicin), topoisomerase inhibitors such as topotecan, and tyrosine kinase inhibitors (gefitinib) are substrates of the P-gp transporter (Waghray and Zhang 2018).

To date, clinical trials with P-gp inhibitors in combination with anticancer drugs were unsuccessful. Several causes may explain the failures. First, the low affinity of early-generation inhibitors required the use of high drug doses which led to severe side effects and toxicity. Moreover, the concomitant administration of P-gp inhibitors and anticancer drugs reduces the metabolism and excretion of the chemotherapeutic agents and increases their concentration in plasma (Waghray and Zhang 2018). Contradictory conclusions were obtained with regard to tariquidar. Some studies identified tariquidar as a P-gp inhibitor and P-gp substrate depending on the used dose. Others reported that tariquidar can also act as a BCRP inhibitor and substrate (Kannan et al. 2011a). The structure of the binding pocket of the P-gp transporter is not well-known, which precludes the development of potent and highly specific P-gp inhibitors (Waghray and Zhang 2018).

The P-gp transporter located at the BBB also decreases the concentration of numerous unrelated drugs in the CNS. For instance, it has been shown that

antiepileptic drugs such as phenytoin, carbamazepine, phenobarbital, lamotrigine, and gabapentin are substrates of P-gp (Wang et al. 2016). Therefore, it has been suggested that P-gp plays an important role in drug-resistant epilepsy or IE (Wang et al. 2016; Schmidt and Schachter 2014). Many scientific publications support the role of P-gp in drug-resistant epilepsy. Preclinical studies have found that phenobarbital-resistant rats showed higher P-gp expression levels at the BBB than phenobarbital responsive rats (Volk and Löscher 2005). Moreover, chronic administration of phenobarbital caused a reduction of phenobarbital concentration in the hippocampus and an increase of P-gp expression (Yang et al. 2008). Despite the unsuccessful strategy of using P-gp inhibitors to overcome cancer drug resistance, it has been proposed that the co-administration of P-gp inhibitors and antiepileptic drugs could restore the antiepileptic efficacy. In support of this idea, the administration of tariquidar together with phenobarbital to phenobarbital-resistant rats counteracted the phenobarbital resistance (Brandt et al. 2006). Moreover, the co-administration of verapamil and carbamazepine to patients with IE improved the efficacy of the antiepileptic drug (Summers et al. 2004). Therefore, modulation of the P-gp function using P-gp inhibitors may be a new approach to overcome drug resistance in epilepsy.

The use of P-gp inhibitors in treatment-resistant depression has also been suggested. Around 60% of patients with depression do not show a satisfactory response to their medication and again the P-gp transporter may play a role in limiting the brain penetration of the applied antidepressant drugs (O'Brien et al. 2012). In vivo studies performed in mice demonstrated that co-administration of the P-gp inhibitor verapamil and the antidepressant escitalopram for 22 days increased the brain concentration of the antidepressant (O'Brien et al. 2015). However, further studies are needed to elucidate the relationship between P-gp function and treatment-resistant depression.

New strategies to overcome drug resistance in cancer and other pathologies have been explored such as the development of drugs that do not show affinity to the ABC transporters and the use of advanced delivery systems such as liposomes or gold nanoparticles as well as the use of antibody-drug conjugates. These new approaches aim to bypass the P-gp efflux action, to target the drug delivery to a specific tissue, and to reduce the toxic side effects of the drug (Löscher and Potschka 2005c; Li et al. 2016; Waghay and Zhang 2018).

### 22.3.2 P-gp Inducers and Activators to Treat Neurodegenerative Disease and Intoxications

P-gp inducers and activators (such as St. John's wort (*Hypericum perforatum*), dexamethasone, vinblastine, and rifampicin (Silva et al. 2015)) increase P-gp function and/or expression, and they may be used to reduce the brain penetration of harmful substances (Gameiro et al. 2017). The use of P-gp inducers or activators can be important in the treatment of intoxications (Silva et al. 2014; Dinis-Oliveira et al. 2006). For instance, the administration of dexamethasone to paraquat-exposed

rats decreased paraquat accumulation in the lungs and reduced the lung damage (Dinis-Oliveira et al. 2006).

Another potential use of inducers and activators was suggested by Haslam et al. based on the finding that the upregulation of ABC transporters in the hair follicle prevents alopecia induced by chemotherapy. This strategy may reduce hair loss in cancer patients (Haslam et al. 2013, 2015).

Additionally, inducers have been proposed as a potential treatment for Alzheimer's and Parkinson's disease. In the case of Alzheimer's disease, A $\beta$  has been identified as a substrate of the P-gp transporter (Lam et al. 2001). Vogelgesang et al. found an inverse correlation between the deposition of A $\beta$  in the brain of elderly humans and the P-gp expression in endothelial cells of the cerebral vasculature (Vogelgesang et al. 2002b). Recent research also highlighted the relationship between the MDR1 gene, which is encoding P-gp, and the risk of developing Parkinson's disease (Drożdżik et al. 2003). It has been suggested that repeated contact with pesticides is associated with the development of Parkinson's disease in patients carrying a c3435T polymorphism of the MDR1 gene (Kortekaas et al. 2005; Lai et al. 2002; Kamel and Hoppin 2004). Furthermore, PET studies have indicated that the P-gp function is decreased in Alzheimer's and Parkinson's disease (Kortekaas et al. 2005; van Assema et al. 2012); thus the use of P-gp inducers may help to treat the disease.

Some preclinical studies have already demonstrated that P-gp inducers such as pregnenolone-16 $\alpha$ -carbonitrile can restore the P-gp function at the BBB and reduce the A $\beta$  deposition in the brain (Hartz et al. 2010). Similar results were obtained in mice models of Alzheimer's disease after treatment with St. John's wort. Animals receiving St. John's wort showed a reduction of A $\beta$  accumulation and an increase in the cerebrovascular expression of P-gp (Brenn et al. 2014).

Many efforts have been made to find new molecules able to induce or activate the P-gp function, which could be applied to treat intoxication or neurodegenerative disease. Recently, MC111 was identified as a promising candidate, due to its capacity to induce P-gp and BCRP expression in SW480 human colon adenocarcinoma cells (Colabufo et al. 2018). In vivo studies using rats and the PET tracer [<sup>18</sup>F]MC225 are ongoing to confirm these findings.

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## 22.4 PET

Positron-emission tomography (PET) and single-photon emission computed tomography (SPECT) together with magnetic resonance imaging (MRI) and computed tomography (CT) are imaging techniques that allow noninvasive tracking of pathophysiological processes in various neuropsychiatric disorders. Molecular imaging has also been used in various aspects of drug development (Willmann et al. 2008). For an efficient visualization, characterization, and measurement of biological processes at the molecular, cellular, whole organ, or body level, specific imaging probes are needed. These probes are also known as tracers and provide functional information, like the magnitude of regional blood flow, glucose metabolism, or receptor

density which can be presented either as a two- or three-dimensional image. PET is the most sensitive technique for *in vivo* imaging and quantification of biological and pathophysiological processes. However, the spatial resolution of PET is relatively low, and the additional use of CT or MRI imaging is required to provide high-resolution structural images that can be fused with the functional PET image. PET/CT or PET/MRI scanners provide both anatomical and functional information in a single scanning session.

The decay of radionuclides for PET imaging occurs by the emission of a positively charged particle called positron. When emitted from the nucleus, the positron travels a very short distance (positron range, 3–5 mm) in the surrounding tissue before it annihilates by combining with an electron. The mass of the positron and electron is then converted to energy resulting in two 511 keV  $\gamma$ -photons which are emitted simultaneously at an angle of approximately 180°. The pair of photons is detected by rings of detectors in the PET scanner. The acquisition of a large number of coincident events provides data that can be reconstructed to an image with information on the spatial distribution of radioactivity as a function of time.

Traditionally, PET tracers for imaging of ABC transporters are evaluated using a dual-scan protocol. This protocol involves a baseline PET scan where the efflux transporters are working adequately and an after-treatment scan where the function of these transporters is either inhibited or induced. An inhibitor is usually used for the challenge, which will reduce the transporter function and, therefore, increase the brain uptake of the tracer (Syvänen and Eriksson 2013; Kannan et al. 2009). By contrast, if an inducer is used, the cerebral uptake of the tracer will decrease. Since the majority of the ABC transporter tracers behave as strong transporter substrates, their baseline uptake of radioactivity in the brain is very low; thus measuring decreases of this uptake is complicated. For this reason, inhibitors are commonly used to validate PET tracers for ABC transporters. Once, an appropriate and specific tracer is found, PET imaging allows to assess the function of that particular ABC transporter in the healthy or diseased human brain and to evaluate the impact of treatments on the function of this transporter.

Recently, the introduction of whole-body PET, which allows simultaneous assessment of tracer uptake in all organs and tissues of the human body, has roused interest in studying the connection among different systems that may play a role in a particular disease (Rahmim et al. 2019). Using this technique, it may, for instance, be possible to study the role of the gut-brain axis in CNS diseases. It has been pointed out that the microbiota present in the gastrointestinal tract has an important impact on the brain. The gut flora not only affects the enteric nervous system but also the CNS via neuroendocrine and metabolic pathways (Carabotti et al. 2009). Changes in the microbiota are related to anxiety and depression (Foster and McVey Neufeld 2013). The onset of other brain disorders such as schizophrenia, autism, and Parkinson's disease has also been related to dysfunction of the gut-brain axis (Cherry et al. 2018). Recent publications suggest that the gut microbiota can affect the integrity of the BBB and may contribute to the development or the progression of neurodegenerative diseases (Parker et al. 2019). The microbiome of Alzheimer's disease patients may also cause a reduction of P-glycoprotein expression in the



intestinal epithelial cells (Haran et al. 2019). Dynamic whole-body PET may bring the opportunity to explore and compare the functionality of the P-gp transporters at the BBB and the intestinal tract. This new approach may broaden insight in the gut-brain axis and its implication in the onset of neurodegenerative diseases.

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## 22.5 PET Tracers for Imaging of ABC Transporter Function and Expression in the BBB: Background

The function of the ABC transporters can be assessed by positron-emission tomography. Since PET is a noninvasive technique, it allows studying the same subject at different time points, e.g., before and after treatment, or at different stages of a disease. Moreover, since longitudinal studies can be performed, the number of subjects required to reach statistical significance is reduced (Willmann et al. 2008). Specific tracers should be designed in order to explore the function of ABC transporters.

In 1998, Hendrikse et al. used [<sup>11</sup>C]verapamil to image the P-gp function for the first time (Hendrikse et al. 1998). Since then, many other compounds have been radiolabeled with carbon-11 or fluorine-18 to explore the function of P-gp and other ABC transporters at the BBB (Saidijam et al. 2018): (*R*)-[<sup>11</sup>C]verapamil (Luurtsema et al. 2005a), [<sup>11</sup>C]-*N*-desmethyl-loperamide ([<sup>11</sup>C]dLop) (Zoghbi et al. 2008), [<sup>11</sup>C]phenytoin (Verbeek et al. 2012a), [<sup>18</sup>F]paclitaxel (Hsueh et al. 2006), [<sup>11</sup>C]laniquidar (Luurtsema et al. 2009), [<sup>11</sup>C]tariquidar (Bauer et al. 2010), [<sup>18</sup>F]MC225 (Savolainen et al. 2015), and [<sup>11</sup>C]metoclopramide (Pottier et al. 2016). Although numerous research studies have been performed with these tracers, none of the tracers was incorporated into the routine medical practice.

The majority of the ABC transporter tracers have been aimed at the imaging of P-gp. Several attempts have been made to develop PET tracers for other ABC transporters, with limited success. Most BCRP and MRP1 substrates proved to show also affinity toward the P-gp transporter. Because of this lack of specificity, the function of a particular transporter cannot be distinguished from the action of other systems (Saidijam et al. 2018; Wanek et al. 2013).

PET studies with “BCRP-selective compounds” did not show differences of uptake in wild-type and BCRP knockout mice or rats (Tournier et al. 2018; Hosten et al. 2013; Mairinger et al. 2010). Since BCRP is the most abundant ABC transporter in the human brain, Wanek et al. designed a protocol for visualizing this transporter at the BBB. The protocol consists of using the dual P-gp/BCRP tracer [<sup>11</sup>C]tariquidar in combination with cold tariquidar or in P-gp knockout mice. P-gp function is then inhibited or absent, and therefore, the function of BCRP can be assessed (Wanek et al. 2012).

Two PET tracers have been developed for the MRP1 transporter: 6-bromo-7-[<sup>11</sup>C]methylpurine (Okamura et al. 2009) and 6-bromo-7-(2-[<sup>18</sup>F]fluoroethyl)purine (Galante et al. 2014). Both are prodrugs that after entering the brain are conjugated with glutathione, and as a result, become substrates for MRP1. In vivo experiments showed significant differences in the brain clearance of both tracers between wild-type and MRP1 knockout mice (Okamura et al. 2009; Galante et al. 2014). Since a relationship between the MRP1 transporter and A $\beta$  clearance from the brain has

been proposed, additional studies were performed in a beta-amyloidosis mouse model (APP/PSI-21) using 6-bromo-7- $^{11}\text{C}$  methylpurine ( $^{11}\text{C}$ BMP). The study evaluated the MRP1 function in APP/PSI-21 mice and in wild-type mice. The study revealed a significantly higher value of the elimination constant ( $k_{\text{elim}}$ ) after the treatment with the MRP1 inhibitor MK571 in the APP/PSI-21 mice compared to the wild-type mice, which represents an increase of MRP1 expression in the astrocytes. Moreover, Western blot results showed an increase in MRP1 levels which was positively correlated with the induction of the MRP1 activity (Zoufal et al. 2020). This study suggests that  $^{11}\text{C}$ BMP may be used to measure the astrocyte MRP1 function (Zoufal et al. 2020). Unfortunately, species differences in the expression levels of MRP1 hamper the translation of preclinical data to humans (Uchida et al. 2011).

To date, the most commonly used PET tracers for imaging P-gp function at the BBB are (*R*)- $^{11}\text{C}$ verapamil and  $^{11}\text{C}$ dLop. Although these tracers have been extensively used in small animals and humans, both tracers have some important drawbacks (Wanek et al. 2013). They act as strong substrates of the P-gp transporter which means that the tracers have a high affinity for the transporter protein. Consequently, the tracers are pumped out from the brain to the blood with very high efficiency and low uptake of the tracer is observed at baseline conditions, when the transporter is fully functional (Langer 2016; Luurtsema et al. 2016). Due to the low uptake at baseline, these PET tracers are not able to measure increases in the P-gp function as occur in drug-resistant epilepsy (Wang et al. 2016). However, (*R*)- $^{11}\text{C}$ verapamil and  $^{11}\text{C}$ dLop can detect decreases in the P-gp function. Such decreases will result in higher uptake of radioactivity in the brain. Thus, these tracers can be used to study diseases characterized by a decline in P-gp function, such as Alzheimer's disease (van Assema and van Berckel 2016).

Another important disadvantage of (*R*)- $^{11}\text{C}$ verapamil is its extensive in vivo metabolism resulting in the formation of radioactive metabolites (dealkylation fragments and polar metabolites) that can cross the BBB. Some of these metabolites are also P-gp substrates (Luurtsema et al. 2005b). A PET study radiolabeled the main metabolite of (*R*)- $^{11}\text{C}$ verapamil, called D617 (Verbeek et al. 2012b). This study evaluated the kinetics of  $^{11}\text{C}$ D617 to investigate whether the kinetics of (*R*)- $^{11}\text{C}$ verapamil and  $^{11}\text{C}$ D617 are similar, and  $^{11}\text{C}$ D617 can be used as an alternative tracer to measure the P-gp function (Verbeek et al. 2012b). However, although the kinetics of  $^{11}\text{C}$ D617 were comparable to that of (*R*)- $^{11}\text{C}$ verapamil at baseline, the increase of the brain uptake of  $^{11}\text{C}$ D617 after P-gp inhibition was lower. A similar approach was more successful in the case of loperamide.  $^{11}\text{C}$ dLop which is the main metabolite of  $^{11}\text{C}$ loperamide became the preferred P-gp tracer, as its imaging data were more easily interpreted than those of the parent compound (Lazarova et al. 2008).

In the case of (*R*)- $^{11}\text{C}$ verapamil, the presence of radioactive metabolites can lead to quantification errors. Even though the radioactivity in blood samples can be corrected for the metabolite fraction, the radioactivity in tissue cannot be corrected, and metabolites will contribute to the tissue signal (Wanek et al. 2013; Tournier et al. 2018). This issue can be solved by using short PET acquisition times because metabolism is negligible shortly after radiotracer injection (Muzi et al. 2009).

Radiolabeling with carbon-11 has the disadvantage of a short physical half-life of the radiotracer (20.4 min) and a low spatial resolution of PET images (because of the long-range of the  $^{11}\text{C}$  positron ( $^{+}\beta$ )). The use of fluorine-18 for radiolabeling allows the distribution of the resulting PET tracer to remote imaging centers and allows using longer scan times because of the longer physical half-life of  $^{18}\text{F}$  (109.8 min) (Luurtsema et al. 2016; Raaphorst et al. 2017a).

Since the perfect tracer to measure the P-gp function does not yet exist, new tracers for ABC transporters are needed. Ideal PET tracers should be produced in high overall yields and with high molar activities. They should display a lipophilicity above 2 and below 3.5 to ensure adequate brain penetration and limited nonspecific binding in the brain (Pike 2009). Moreover, these molecules should show a high specificity for the desired ABC transporter. They should be slowly metabolized, and their metabolites should not cross the BBB and not be substrates for ABC transporters. The parent tracers should show a moderate affinity to the ABC transporter of interest, resulting in a moderate brain uptake at baseline. They would then be able to measure decreases as well as increases of the transporter function at the BBB. Radiolabeling should be done with fluorine-18, to obtain high-quality images (Kannan et al. 2009; Wanek et al. 2013; Luurtsema et al. 2016).

Many strategies have been explored to overcome a low baseline uptake of radioactivity in the brain. The first approach was to develop PET tracers with greater lipophilicity to enhance their passive diffusion to the brain; however, high lipophilicity may lead to increased nonspecific binding (Sylvänen and Eriksson 2013; Toyohara 2016). Next, well-known P-gp inhibitors such as tariquidar, elacridar, and laniquidar were radiolabeled with carbon-11 to obtain radiotracers that would bind to P-gp but not be transported. In theory, this approach would provide a higher brain uptake at baseline and would allow measurement of P-gp expression in the brain. Unfortunately, at low concentrations, these inhibitors were recognized as substrates by P-gp and BCRP at the BBB of rats. Therefore, these tracers were transported out of the brain, resulting in a low baseline uptake. This strategy of inhibitor labeling proved to be unsuccessful (Wanek et al. 2013). A third approach to increase the baseline uptake of PET tracers for P-gp consists of the development of lower avidity substrates. This approach using weak P-gp substrates aimed at higher baseline uptake in the brain which facilitates the measurement of increases in the P-gp function. Several radiotracers have already been identified as weak P-gp substrates and their kinetics are being evaluated (Verbeek et al. 2012a; Pottier et al. 2016; Toyohara et al. 2016; Savolainen et al. 2017a).

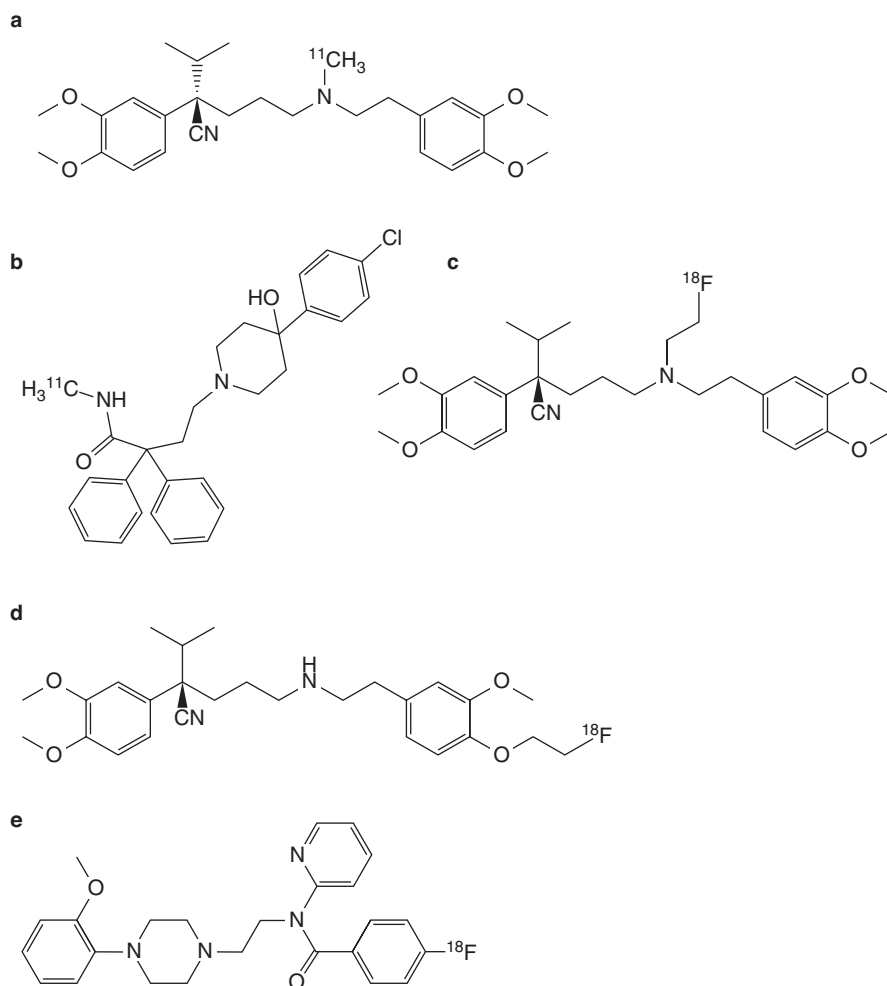
Recently, it has been noticed that (*R*)- $^{[11}\text{C}]$ verapamil and  $^{[11}\text{C}]$ dLop possess limited sensitivity to detect small changes in the P-gp function. The brain distribution of (*R*)- $^{[11}\text{C}]$ verapamil and  $^{[11}\text{C}]$ dLop was not significantly increased in heterozygous mice (+/-), which have 50% lower P-gp levels, compared to wild-type mice. These results indicate that the P-gp function should be reduced by more than 50% to effectively limit the entrance of P-gp substrates in the brain (Wanek et al. 2015). The limited sensitivity of the existing PET tracers may be due to their high affinity to the transporter. It has been proposed that PET imaging with weak P-gp substrates may result in higher sensitivity for detecting small variations in the P-gp function, as may occur in several diseases (Langer 2016; Bauer et al. 2019).

## 22.6 Substrates as Tracers for Measuring P-gp Function

### 22.6.1 Strong Substrates (Fig. 22.2)

#### 22.6.1.1 [ $^{11}\text{C}$ ]Verapamil

The L-type calcium channel blocker verapamil [2,8-bis(3,4-dimethoxyphenyl)-6-methyl-2-isopropyl-6-azaocantirile] is widely used in the treatment of hypertension (McTavish and Sorkin 1989), angina pectoris, and cardiac arrhythmias. Verapamil metabolizes preferentially to norverapamil via *N*-demethylation (Kroemer et al. 1993). After oral administration, the (*S*)-enantiomer of verapamil is



**Fig. 22.2** Structures of strong P-gp substrates: (*R*)-[ $^{11}\text{C}$ ]verapamil (a), [ $^{11}\text{C}$ ]dLop (b), [*R*]-*N*-[ $^{18}\text{F}$ ]fluoroethylverapamil (c), (*R*)-*O*-[ $^{18}\text{F}$ ]fluoroethylnorverapamil (d), and [ $^{18}\text{F}$ ]MPPF (e)

metabolized more actively than (*R*)-verapamil in humans, resulting in a 2.5 times higher concentration of the (*R*)-enantiomer in plasma (Häussermann et al. 1991). (*R*)-verapamil has only 5–10% of the calcium channel blocking activity of the (*S*)-enantiomer. However, the (*R*)-enantiomer is equipotent to racemic verapamil and (*S*)-verapamil in its efficacy to inhibit P-gp. Verapamil is metabolized by cytochrome P450 enzymes, and some of the radioactive metabolites are themselves substrates and inhibitors of P-gp (Pauli-Magnus et al. 2000). In addition, polar radiolabeled metabolites are formed, which result in a non-P-gp-mediated signal (Luurtsema et al. 2005b). Quantification of racemic [<sup>11</sup>C]verapamil is difficult because the enantiomers have different pharmacokinetic properties (Lagerge et al. 2001). Therefore, PET studies need to be performed with an optically pure enantiomer. (*R*)- and (*S*)-[<sup>11</sup>C]verapamil were prepared from (*R*)- and (*S*)-desmethylverapamil by methylation with no-carrier added [<sup>11</sup>C]methyl iodide or [<sup>11</sup>C]methyl triflate (Luurtsema et al. 2002). Although no difference in P-gp-mediated transport selectivity between (*R*)- and (*S*)-verapamil was found (Luurtsema et al. 2003), (*R*)-[<sup>11</sup>C]verapamil was selected as the most appropriate tracer for measuring P-gp function with PET because the (*R*)-enantiomer was less metabolized in humans and showed a lower affinity for calcium channels. This tracer showed a low brain uptake at baseline both in humans and other vertebrate species, but co-administration with P-gp inhibitors as cyclosporine A (CsA) or tariquidar increased its volume of distribution ( $V_T$ ) and kinetic constant  $k_1$  (Muzi et al. 2009; Bankstahl et al. 2008). The increase in brain  $V_T$  after administration of tariquidar was approximately +300% in rats and +24 ± 15% in humans (Kuntner et al. 2010; Wagner et al. 2009), which points to a pronounced species difference regarding the tariquidar effect.

### 22.6.1.2 [<sup>11</sup>C]N-Desmethyl-Loperamide

Loperamide is an antiarrheal agent that acts through agonism at  $\mu$ -opioid receptors in the gut (Awouters et al. 1993). This drug is normally without harmful central effects because P-gp expels it from the brain. [<sup>11</sup>C]loperamide is avidly metabolized to [<sup>11</sup>C]dLop via demethylation (Zoghbi et al. 2008). This metabolite is also a substrate for P-gp, and it has been used as a PET tracer instead of [<sup>11</sup>C]loperamide. Demethylation of [<sup>11</sup>C]dLop may occur, but it will lead only to the formation of single-carbon radiometabolites such as [<sup>11</sup>C]methanol. These metabolites will be oxidized and ultimately expired as [<sup>11</sup>C]carbon dioxide. They should not accumulate in brain tissue and will therefore not cause problems in the kinetic analysis of the acquired PET data. The uptake of [<sup>11</sup>C]dLop 30 min after its injection was about 3.5-fold higher in the forebrain of P-gp knockout mice compared to wild-type mice supporting the results from previous studies which indicate that [<sup>11</sup>C]dLop is a substrate of P-gp (Lazarova et al. 2008). Similar results were acquired in monkeys. [<sup>11</sup>C]dLop was found to be selective for P-gp when tested in human cell lines that overexpress P-gp, MRP1, or BCRP and also in P-gp knockout mice (Kannan et al. 2010). At low concentrations ( $\leq 1$  nM), dLop acts only as a substrate, but at higher concentrations ( $\geq 20$   $\mu$ M), it acts both as a substrate and an inhibitor. Because low concentrations of radiotracer are used in PET imaging, [<sup>11</sup>C]dLop will act only as a selective substrate for P-gp at the BBB. Pretreatment with the inhibitor tariquidar

increases the brain uptake of [ $^{11}\text{C}$ ]dLop markedly in monkeys and humans (Kreisl et al. 2010). After entry of [ $^{11}\text{C}$ ]dLop into the brain, it is trapped and does not wash out. Although dLop is an opiate agonist, its trapping is not a result of high-affinity binding to the opiate receptor. Brain uptake of [ $^{11}\text{C}$ ]dLop cannot be displaced by receptor-saturating doses of an opiate agonist or antagonist (Lazarova et al. 2008). This irreversible trapping amplifies the PET signal, so it is advantageous from an imaging perspective. dLop has a  $\text{p}K_a$  of 7.3 and may be trapped by passive diffusion into acidic organelles, mainly lysosomes, followed by protonation within the organelle (Kaufmann and Krise 2007; Macintyre and Cutler 1988). To test this hypothesis, the accumulation of [ $^3\text{H}$ ]dLop was measured in human cell lines and compared to the uptake of other weak bases (Kannan et al. 2011b). [ $^{11}\text{C}$ ]dLop was found to be ionically trapped in acidic lysosomes and tariquidar competed for lysosomal accumulation, both in vitro and in vivo. This interaction is problematic in peripheral organs but not in the brain. Thus [ $^{11}\text{C}$ ]dLop can still be used, either alone or in combination with tariquidar, to measure the function of P-gp at the BBB. However, the low brain uptake of [ $^{11}\text{C}$ ]dLop at baseline conditions complicates the evaluation of increases in the P-gp function, as is also the case with (*R*)-[ $^{11}\text{C}$ ]verapamil.

### 22.6.1.3 Fluorine-18 Verapamil Analogs

The synthesis of two fluorine-18 (*R*)-verapamil analogs has also been reported: (*R*)-*N*-[ $^{18}\text{F}$ ]fluoroethylverapamil and (*R*)-*O*-[ $^{18}\text{F}$ ]fluoroethylnorverapamil. Since (*R*)-[ $^{11}\text{C}$ ]verapamil is considered as the gold standard for measuring the P-gp function until now, these two candidates were proposed as promising radiofluorinated analogs. (*R*)-*N*-[ $^{18}\text{F}$ ]fluoroethylverapamil was obtained as follows: first, [ $^{18}\text{F}$ ]fluoroethyl triflate was obtained from bromoethyltosylate using a silver triflate column, and secondly, [ $^{18}\text{F}$ ]fluoroethyl triflate was distilled into a vial containing the precursor (*R*)-norverapamil. The synthesis of (*R*)-*O*-[ $^{18}\text{F}$ ]fluoroethylnorverapamil was simpler. The precursor contained a tosyl group which allows direct fluorination, but Boc deprotection was also required. Preclinical studies carried out in rats showed that after administration of the P-gp inhibitor tariquidar, the brain uptake of (*R*)-*N*-[ $^{18}\text{F}$ ]fluoroethylverapamil and (*R*)-*O*-[ $^{18}\text{F}$ ]fluoroethylnorverapamil was increased by 3.6- and 2.4-fold, respectively. A PET study using P-gp knockout and P-gp and BCRP double-knockout mice showed that (*R*)-*O*-[ $^{18}\text{F}$ ]fluoroethylnorverapamil was more specific for P-gp than (*R*)-*N*-[ $^{18}\text{F}$ ]fluoroethylverapamil; thus, this analog was selected as the best tracer candidate (Raaphorst et al. 2017a).

### 22.6.1.4 [ $^{18}\text{F}$ ]MPPF

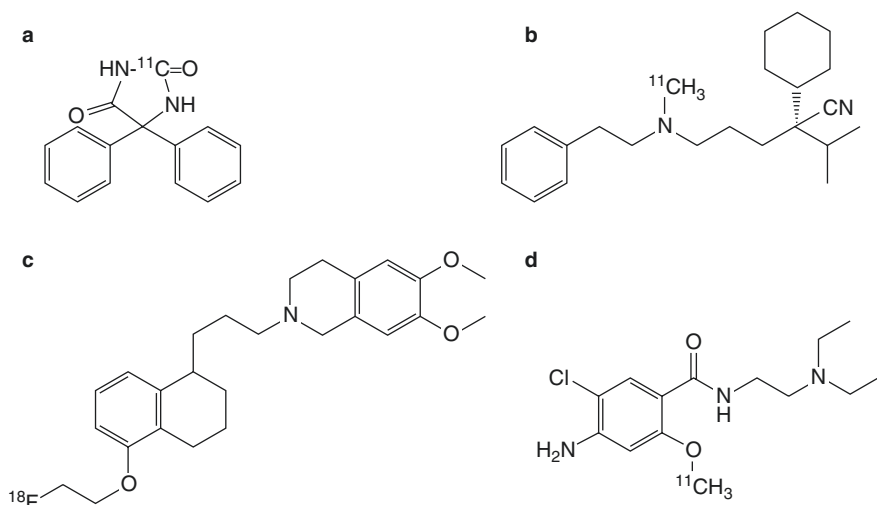
The radiosynthesis of 4-(2'-methoxyphenyl)-1-[2'-(*N*-2''-pyridinyl)-p-[ $^{18}\text{F}$ ]fluorobenzamido] ethylpiperazine ([ $^{18}\text{F}$ ]MPPF) comprises nucleophilic aromatic substitution of a para-nitro compound. The synthesis was performed in the presence of  $\text{K}_2\text{CO}_3$  and Kryptofix<sub>222</sub> and under microwave heating (Le Bars et al. 1998). [ $^{18}\text{F}$ ]MPPF is a PET tracer aimed at the imaging of the serotonin 5-HT<sub>1A</sub> receptor; however, it was discovered to be also a substrate of P-gp. Administration of the P-gp inhibitor cyclosporine A increased the tracer uptake in rat hippocampus (Laćan et al. 2008). This observation was confirmed by biodistribution studies performed in

P-gp knockout mice. The tracer uptake in the brain of P-gp knockout animals was two- to threefold higher than in wild-type mice (Passchier et al. 2000). However, in cells transfected with human P-gp or BCRP genes, [ $^{18}\text{F}$ ]MPPF efflux was not reduced by the P-gp inhibitor valsopodar, and, thus, [ $^{18}\text{F}$ ]MPPF may not be transported by the human P-gp. A study performed in nonhuman primates showed an increase in the uptake of [ $^{18}\text{F}$ ]MPPF in the brain after the treatment with cyclosporine; however, this increase was attributable to an increase of the free fraction of the tracer in plasma. This study noted that cyclosporine A binds to plasma proteins and displaces other compounds (such as [ $^{18}\text{F}$ ]MPPF) from its plasma protein binding sites (Tournier et al. 2012). All these findings restrict the use of [ $^{18}\text{F}$ ]MPPF as a PET tracer for measuring the function of P-gp at the BBB.

## 22.6.2 Weak P-gp Substrates (Fig. 22.3)

### 22.6.2.1 [ $^{11}\text{C}$ ]Phenytoin

In 2010, an in vitro study showed that phenytoin and phenobarbital (antiepileptic drugs) are transported by the human P-gp, which strengthens the hypothesis that P-gp is involved in antiepileptic drug resistance (Zhang et al. 2010). In rhesus monkeys [ $^{11}\text{C}$ ]phenytoin showed a high brain uptake (Stavchansky et al. 1978), which suggests that phenytoin behaves as a weak P-gp substrate (Verbeek et al. 2012a). [ $^{11}\text{C}$ ]Phenytoin was synthesized via [ $^{11}\text{C}$ ]CO using a high-pressure autoclave (Verbeek et al. 2012a). In rats, the baseline uptake of [ $^{11}\text{C}$ ]phenytoin was higher than that of [ $^{11}\text{C}$ ]verapamil. After administration of 15 mg/kg of tariquidar, the brain



**Fig. 22.3** Structure of weak P-gp substrates: [ $^{11}\text{C}$ ]Phenytoin (a), (R)-[ $^{11}\text{C}$ ]emopamil (b), [ $^{18}\text{F}$ ]MC225 (c), and [ $^{11}\text{C}$ ]metoclopramide (d)

uptake of [ $^{11}\text{C}$ ]phenytoin showed a smaller increase compared to avid substrate tracers such as [ $^{11}\text{C}$ ]verapamil. This observation also indicates a moderate affinity of [ $^{11}\text{C}$ ]phenytoin to P-gp (Verbeek et al. 2012a). Quantification of the brain uptake of [ $^{11}\text{C}$ ]phenytoin in humans was assessed by different kinetic models. The results of this study indicated that the tissue-tissue compartment model (1-TCM) is the preferred technique to analyze the data, and  $K_1$  and  $V_T$  can be used for measuring the P-gp function (Mansor et al. 2015). Although [ $^{11}\text{C}$ ]phenytoin seems to be a good tracer candidate to measure increases in the function of P-gp, further studies performed under pathologic conditions are still required.

### 22.6.2.2 [ $^{11}\text{C}$ ]Emopamil

2-Isopropyl-5-[methyl-(2-phenylethyl)amino]-2-phenylpentanenitrile (emopamil) is a calcium channel blocker. Its chemical structure is similar to verapamil. A study evaluated the brain penetration of emopamil, verapamil, and gallopamil in rats and concluded that the cerebral availability and BBB permeability were higher for emopamil than for verapamil and gallopamil (Szabo 1989). The rather high brain penetration of emopamil at baseline condition encouraged the development of [ $^{11}\text{C}$ ]emopamil as a weak substrate tracer for P-gp (Toyohara et al. 2016). (*R*)- and (*S*)-[ $^{11}\text{C}$ ]emopamil were synthesized via methylation of the precursors, (*R*)- and (*S*)-noremopamil, with [ $^{11}\text{C}$ ]methyl triflate. From mice experiments, it was concluded that (*R*)-[ $^{11}\text{C}$ ]emopamil provides higher uptake than (*S*)-[ $^{11}\text{C}$ ]emopamil. (*R*)-[ $^{11}\text{C}$ ]emopamil showed a twofold higher area-under-the-curve (AUC) of the brain radioactivity in baseline scans than (*R*)-[ $^{11}\text{C}$ ]verapamil. However, after the administration of cyclosporine A, the AUCs of the brain radioactivity of both tracers were similar. Unfortunately, both (*R*)- and (*S*)-[ $^{11}\text{C}$ ]emopamil are rapidly metabolized (Toyohara et al. 2016). To verify that (*R*)-[ $^{11}\text{C}$ ]emopamil is suitable for measuring P-gp upregulation, additional experiments in animal models are needed.

### 22.6.2.3 [ $^{18}\text{F}$ ]MC225

Savolainen et al. evaluated three fluorine-18 compounds that shared the same moiety: 6,7-dimethoxytetrahydroisoquinoline, which is also present in tariquidar and elacridar (Savolainen et al. 2015). The compounds were assessed in vitro and in vivo with PET imaging, using P-gp and BCRP knockout mice. The results were compared to those obtained with [ $^{11}\text{C}$ ]verapamil. From this experiment, [ $^{18}\text{F}$ ]MC225 was selected as the most appropriate candidate to measure P-gp function at the BBB. The synthesis of [ $^{18}\text{F}$ ]MC225 consists of two steps: First, [ $^{18}\text{F}$ ]bromomethyl-fluorine is synthesized from 2-bromoethyltosylate by heating at 90 °C for 15 min in dichlorobenzene. Afterward, the distilled [ $^{18}\text{F}$ ]fluorethyl bromide is trapped in a vial containing sodium hydride and the phenol precursor of MC225 in dimethylformamide (DMF), and the vial is heated at 80 °C for 5 min. PET studies demonstrated that the baseline uptake of [ $^{18}\text{F}$ ]MC225 was higher than that of [ $^{11}\text{C}$ ]verapamil in rats. Baseline  $V_T$  values of [ $^{11}\text{C}$ ]Verapamil ranged from 1.1 to 2.3 (Kuntner et al. 2010), whereas values for [ $^{18}\text{F}$ ]MC225 varied from 6.6 to 11 (Savolainen et al. 2017a). The brain uptake of [ $^{18}\text{F}$ ]MC225 was also higher in P-gp and BCRP knockout mice than in the wild-type, and in addition, [ $^{18}\text{F}$ ]MC225 showed greater



metabolic stability (Savolainen et al. 2015). [ $^{18}\text{F}$ ]MC225 showed specificity for P-gp over BCRP or MRP1 (Savolainen et al. 2015; Fusi et al. 2017). Rat experiments confirmed that [ $^{18}\text{F}$ ]MC225 is transported by P-gp and not by BCRP. Inhibition of P-gp increased the brain uptake of the tracer, but inhibition of BCRP did not affect this uptake (Savolainen et al. 2017a). At 30 min after tracer injection, only 24% of the plasma radioactivity represented the parent compound. Different kinetic models were fitted to the data, and it was concluded that a 1-TCM was the preferred model for [ $^{18}\text{F}$ ]MC225 in rats. After inhibition of P-gp, the changes in the influx rate constant  $K_1$  were higher than the changes in  $V_T$ ; for this reason,  $K_1$  was proposed as the best parameter to measure the P-gp function (Savolainen et al. 2017a), as has also been suggested for analysis of the data of (*R*)-[ $^{11}\text{C}$ ]verapamil (Muzi et al. 2009). PET studies with [ $^{18}\text{F}$ ]MC225 are ongoing to demonstrate that this tracer is able to measure increases and decreases of the P-gp function at the BBB.

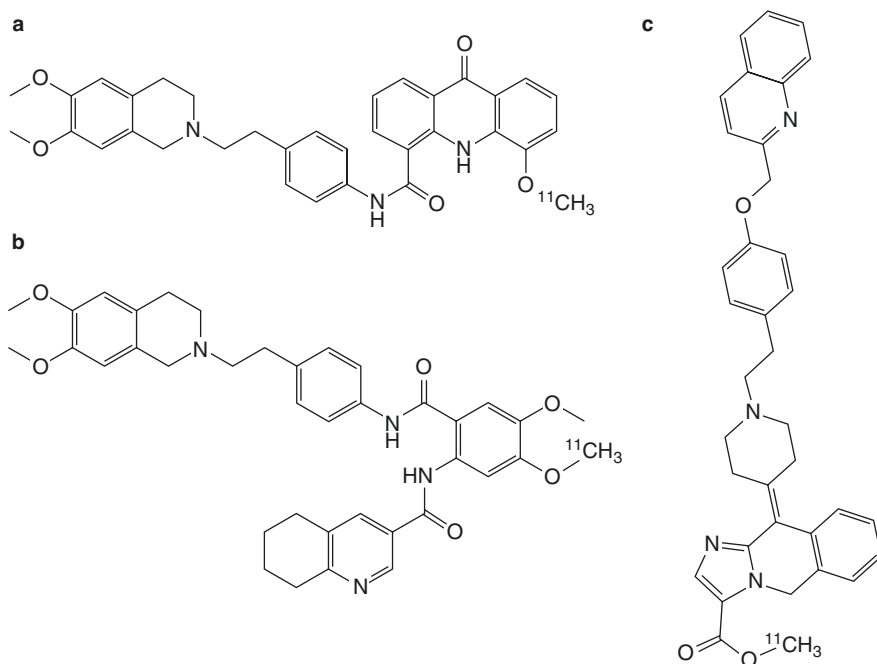
#### 22.6.2.4 [ $^{11}\text{C}$ ]Metoclopramide

Metoclopramide is a dopamine D2 receptor antagonist that is used as antiemetic. The drug was not identified as a P-gp substrate during *in vitro* experiments; however, *in vivo* studies in mice revealed increased (6.6-fold) brain concentration of metoclopramide in P-gp knockout compared to wild-type mice (Feng et al. 2008). Patients treated with metoclopramide reported CNS effects, which suggest that metoclopramide crosses the BBB (Pottier et al. 2016). Therefore, [ $^{11}\text{C}$ ]metoclopramide was synthesized by radiomethylation of the methoxy phenol moiety of nor-metoclopramide using [ $^{11}\text{C}$ ]methyl triflate (Pottier et al. 2016). *In vitro* assays using P-gp and BCRP, overexpressing cells demonstrated that [ $^{11}\text{C}$ ]metoclopramide is specific for P-gp over BCRP. Moreover, the brain uptake of the tracer in rats increases after P-gp inhibition, resulting in an increase of  $K_1$  and the binding potential (BP) in the two-tissue compartment model (2-TCM). At 30 min after tracer injection, only 20% of plasma radioactivity represented the parent compound in rats (Pottier et al. 2016). The kinetics of [ $^{11}\text{C}$ ]metoclopramide were also evaluated in nonhuman primates; P-gp inhibition increased the brain distribution in these animals. In this case, a 1-TCM was used to fit the data, and the most sensitive parameter was the  $V_T$  which increased by 2.02-fold. The inhibition also caused an increase in  $K_1$  (1.28-fold) and a decrease in  $k_2$  (1.64-fold). However, the metabolism of [ $^{11}\text{C}$ ]metoclopramide in nonhuman primates was more rapid than in rats, and 30 min after tracer injection, the fraction of the parent [ $^{11}\text{C}$ ]metoclopramide in plasma was below the limit of detection (Auvity et al. 2018). Recently, the kinetics of this tracer was also assessed in humans. Here, the kinetic analysis showed a significant increase in  $V_T$  by  $29 \pm 17\%$  after P-gp inhibition, mainly due to a significant  $15 \pm 5\%$  decrease in  $k_2$ .  $K_1$  was slightly increased after the treatment' however, this increase was significant in only a few regions compared to the baseline scan (Tournier et al. 2019). Further studies in pharmacoresistant patients are needed to demonstrate that [ $^{11}\text{C}$ ]metoclopramide can visualize the increase of the P-gp function in these patients.

## 22.7 P-gp Inhibitors as Tracers for Measuring P-gp Expression

### 22.7.1 [<sup>11</sup>C]Elacridar, [<sup>11</sup>C]Tariquidar, and [<sup>11</sup>C]Laniquidar

Many well-known P-gp inhibitors such as elacridar, tariquidar, and laniquidar were radiolabeled with carbon-11 (Fig. 22.4) and later also with fluorine-18. Elacridar, tariquidar, and laniquidar belong to the third generation of P-gp inhibitors. Tariquidar is described as a noncompetitive P-gp inhibitor which shows good selectivity toward the P-gp transporter over other efflux transporters, although it also inhibits BCRP transporter but with less potency. Elacridar is considered as a P-gp and BCRP inhibitor. Laniquidar is a noncompetitive inhibitor less potent than tariquidar and elacridar and is highly selective for P-gp (Wanek et al. 2013). Preclinical studies performed in rats showed that elacridar and tariquidar can increase the uptake of [<sup>11</sup>C]verapamil in the brain up to 11-fold, with half-maximum effective doses of  $1.2 \pm 0.1$  and  $3.0 \pm 0.2$  mg/kg, respectively. The results also showed that elacridar is three times more potent than tariquidar (Kuntner et al. 2009).



**Fig. 22.4** Structure of P-gp inhibitors: [<sup>11</sup>C]elacridar (a), [<sup>11</sup>C]tariquidar (b), and [<sup>11</sup>C]laniquidar (c)

For the [ $^{11}\text{C}$ ]elacridar synthesis, the O-desmethyl-elacridar was reacted with [ $^{11}\text{C}$ ]methyl triflate in acetone containing NaOH. [ $^{11}\text{C}$ ]elacridar was administered to mice and rats to evaluate its kinetics. The brain uptake in P-gp knockout mice was 2.5-fold higher, and in BCRP knockout mice was 1.3-fold higher compared to wild-type mice. The co-administration of [ $^{11}\text{C}$ ]elacridar with the unlabeled compound also increased the brain uptake in rats (Dörner et al. 2009). Another PET study found increased  $K_1$  values of [ $^{11}\text{C}$ ]elacridar in P-gp and BCRP knockout mice compared to wild-type mice, suggesting the brain penetration of [ $^{11}\text{C}$ ]elacridar is limited by these two transporters (Kawamura et al. 2010a). In conclusion, [ $^{11}\text{C}$ ]elacridar is not specific for P-gp, and its radioactive signal in baseline scans is very low.

O-desmethyl-tariquidar was reacted with [ $^{11}\text{C}$ ]methyl iodide or [ $^{11}\text{C}$ ]methyl triflate to obtain [ $^{11}\text{C}$ ]tariquidar (Bauer et al. 2010; Kawamura et al. 2010b). In vivo studies performed in mice showed an increase of [ $^{11}\text{C}$ ]tariquidar brain uptake after the administration of unlabeled tariquidar (Kawamura et al. 2010b). Moreover, the brain uptake of [ $^{11}\text{C}$ ]tariquidar was higher in P-gp knockout mice and BCRP knockout mice than in wild-type mice (Bauer et al. 2010; Kawamura et al. 2010b). PET experiments performed in rats also found an increase in the brain uptake of [ $^{11}\text{C}$ ]tariquidar after the administration of tariquidar and elacridar (Bauer et al. 2010). The studies revealed a high metabolic stability of the tracer in the brain and plasma (Kawamura et al. 2010b). In tumor cells, the uptake of [ $^{11}\text{C}$ ]tariquidar and [ $^{11}\text{C}$ ]elacridar increased after the administration of unlabeled tariquidar and this increase was similar to the one observed for [ $^{11}\text{C}$ ]verapamil (Wanek et al. 2011). The kinetics of [ $^{11}\text{C}$ ]tariquidar and [ $^{11}\text{C}$ ]elacridar were also evaluated in healthy human subjects. The brain uptake at baseline condition was very low for both tracers; however, after the administration of unlabeled tariquidar, the brain uptake of [ $^{11}\text{C}$ ]tariquidar increased by  $27 \pm 15\%$  and by  $21 \pm 15\%$  in the case of [ $^{11}\text{C}$ ]elacridar. Both tracers showed a low amount of radiometabolites in plasma, 60 min after injection. The results indicate that [ $^{11}\text{C}$ ]tariquidar and [ $^{11}\text{C}$ ]elacridar behave as P-gp and BCRP substrates in vivo and therefore are unable to visualize P-gp expression at the BBB (Bauer et al. 2013).

The radiosynthesis of [ $^{11}\text{C}$ ]laniquidar involved methylation of a precursor with [ $^{11}\text{C}$ ]methyl iodide in DMSO containing tetrabutylammonium hydroxide (Luurtsema et al. 2009). [ $^{11}\text{C}$ ]laniquidar was injected via the tail vein, and rats were sacrificed at different time points to count radioactivity in various tissues. Moreover, rats were also pretreated with cyclosporine A, and the brain concentration of [ $^{11}\text{C}$ ]laniquidar was determined 30 min after injection. The brain uptake of [ $^{11}\text{C}$ ]laniquidar was low and was increased after treatment with cyclosporine A. The metabolism of the tracer was slow, resulting in a parent fraction of 68% in plasma, 30 min after injection. However, the results suggest that laniquidar may act as a P-gp substrate instead of as a P-gp inhibitor at tracer concentrations (Luurtsema et al. 2009). Co-administration of [ $^{11}\text{C}$ ]laniquidar and unlabeled laniquidar in mice confirmed that laniquidar may act as a P-gp substrate at low doses and as a P-gp inhibitor at high doses (Moerman et al. 2012).

### 22.7.2 [<sup>18</sup>F]Fluoroelacridar, [<sup>18</sup>F]Fluoroethylacridar, and [<sup>18</sup>F]Fluoroethyltariquidar

The synthesis of fluorine-18-labeled analogs of elacridar and tariquidar has been reported. These tracers showed similar properties to their carbon-11 analogs (Dörner et al. 2011; Kawamura et al. 2011). [<sup>18</sup>F]Fluoroelacridar was synthesized by nucleophilic aromatic substitution of the 1-nitro precursor. The reaction was done in the presence of dried kryptofix<sub>222</sub> and K<sub>2</sub>CO<sub>3</sub> complex together with the precursor dissolved in dimethyl sulfoxide (DMSO), and the reaction mixture was heated for 60 min at 175 °C. Preclinical studies performed in rats showed a significant increase in brain uptake of the tracer after the administration of the P-gp and BCRP inhibitor elacridar. Brain uptake was also higher in P-gp and BCRP knockout than in wild-type mice (Dörner et al. 2011). Although [<sup>18</sup>F]fluoroelacridar behaved similarly to [<sup>11</sup>C]elacridar, both tracers are substrates of P-gp and BCRP which hampers the quantification of a specific transporter. [<sup>18</sup>F]Fluoroelacridar also showed an intense defluorination in vivo, which limits its use as a PET tracer (Dörner et al. 2011).

[<sup>18</sup>F]fluoroethylacridar and [<sup>18</sup>F]fluoroethyltariquidar were synthesized by alkylation of its O-desmethyl precursors with [<sup>18</sup>F]2-fluoroethyl bromide. These precursors were heated at 90–120 °C for 10 min in DMF containing tetrabutylammonium hydroxide and [<sup>18</sup>F]2-fluoroethyl bromide. Preclinical studies performed in wild-type and knockout mice revealed that these fluorine-18 tracers present similar kinetics and biodistribution as their carbon-11 analogs. An increase in the brain uptake was observed after the administration of tariquidar or elacridar to wild-type mice, and a higher uptake of the tracer was found in P-gp/BCRP knockout mice compared to wild-type (Kawamura et al. 2011). Both tracers were again substrates of P-gp and BCRP; therefore, they cannot be used to measure only the P-gp function (Raaphorst et al. 2017a; Kawamura et al. 2011).

The radiolabeling of P-gp inhibitors to measure the P-gp expression turned to be an unsuccessful strategy. The results provided by the preclinical and clinical studies were unexpected. First, the baseline brain uptake of these tracers was very low, and, second, the treatment with P-gp inhibitors caused an increase in the brain uptake. At tracer concentrations, these compounds may act as P-gp substrates rather than P-gp inhibitors. (Luurtsema et al. 2016). Therefore, [<sup>11</sup>C]elacridar, [<sup>11</sup>C]tariquidar, and [<sup>11</sup>C]laniquidar as well as their fluorine-18 analogs fail to measure P-gp expression at the BBB (Wanek et al. 2013).

### 22.7.3 Novel P-gp Inhibitors as PET Tracers

Until now, most radiolabeled inhibitors aimed to quantify the expression of the P-gp transporter have not been successful. However, Verbeek and colleges have recently published the synthesis of two novel PET tracers for measuring P-gp expression, [<sup>18</sup>F]2-(4-fluoro-2-oxoindolin-3-ylidene)-*N*-(4-methoxyphenyl)hydrazine-carbothioamide ([<sup>18</sup>F]5) and [<sup>18</sup>F]2-(6-fluoro-2-oxoindolin-3-ylidene)-*N*-(4-methoxyphenyl) hydrazine-carbothioamide ([<sup>18</sup>F]6) (Verbeek et al. 2018). The

kinetics of both tracers were evaluated in P-gp knockout mice and rats before and after administration of the P-gp inhibitor tariquidar. [ $^{18}\text{F}$ ]5 showed greater metabolic stability than [ $^{18}\text{F}$ ]6. PET studies performed in P-gp knockout mice revealed a significantly lower uptake of [ $^{18}\text{F}$ ]5 compared to wild-type mice. Additionally, the administration of tariquidar did not modify the uptake of the tracer in the rat brain. Thus, this study suggested that [ $^{18}\text{F}$ ]5 may be a promising radiotracer that binds to the P-gp transporter and may allow measurement of P-gp expression levels in vivo (Verbeek et al. 2018). However, additional experiments are needed to prove if [ $^{18}\text{F}$ ]5 behaves as a P-gp inhibitor.

## 22.8 Kinetic Modeling of the P-gp Function

As has been explained before, PET is a functional imaging technique that measures the radioactivity concentration of a tracer in a specific target. This technique allows monitoring of the interactions of a radiotracer with its target(s) and quantification of the effects of a drug on physiological processes (Morris et al. 2004). Quantitative analysis of the PET data can involve the normalizing of average values from a region of interest using, for instance, standardized uptake values (SUV) or kinetic modeling methods. Kinetic modeling usually requires the acquisition of dynamic images, which give information about the concentration of the tracer in the target tissue as a function of time, as well as measurements of radioactivity in blood or plasma of the subject at different time points to acquire an arterial input function (Kuntner and Stout 2014; Kuntner 2014; Gunn et al. 2001). Moreover, since radiotracers can be extensively metabolized, the radioactivity concentration in plasma should be corrected for the fraction of metabolites. Reference tissue models have also been developed in order to avoid the need for blood sampling. In this case, the time-activity curve of the radiotracer in the tissue of interest is related to the time-activity curve in a reference tissue that lacks the tracer target (Gunn et al. 2001; Lammertsma and Hume 1996). Compartmental models are frequently used to describe the tracer kinetics, in order to analyze the delivery and elimination of the tracer in the tissue of interest. These methods can be used to calculate the mean transfer rate constants, which describe the movement of the tracer through different compartments, and the volume of distribution ( $V_T$ ) or the binding potential of the tracer to the target (Carson 2003). Kinetic modeling can also be used to determine which parameter describes most adequately the function under study or which parameter is most sensitive to changes in this function (Kuntner 2014).

The P-gp transporter is mainly located at the luminal side of the endothelial cells at the BBB. Thus, the rate constants  $K_1$  and  $k_2$ , which describe the movement of the tracer from the blood to the brain (or first tissue compartment), have been proposed as the best parameters to measure the P-gp function (Tournier et al. 2018). Since the P-gp transporter is an efflux pump, it is expected that the efflux rate constant  $k_2$  would be the parameter which is most affected by a P-gp inhibition, and thus, it would be the best parameter to measure the P-gp function. However, several PET studies with P-gp tracers found a significant increase in  $K_1$

and  $V_T$  after administration of P-gp inhibitors, but no changes in  $k_2$ . Thus, it was suggested that P-gp tracers pass from the blood to the endothelial cells and from there back to the blood by a very rapid P-gp-mediated transport. This transport is much more rapid than the PET acquisition, and therefore, the P-gp function is reflected in the value of  $K_1$ , instead of the value of  $k_2$ . A review of the kinetics of several P-gp PET tracers concluded that  $K_1$  corrected for changes in the blood flow is the preferred parameter to measure P-gp function at the BBB (Lubberink 2016). However, recent studies performed with [ $^{11}\text{C}$ ]metoclopramide in nonhuman primates observed a significant decrease in  $k_2$  after intravenous co-infusion of tariquidar. The results of these studies suggested that the most sensitive parameter to measure P-gp inhibition was  $V_T$  (Auvity et al. 2018). Similar results were obtained in humans where a significant increase in  $V_T$  was found during the infusion of cyclosporine A. In that study,  $K_1$  showed a significant increase in only a few brain regions after P-gp inhibition (Tournier et al. 2019). These publications have reopened the debate about the question which kinetic parameter represents most adequately the P-gp function.

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## 22.9 The Use of P-gp Tracers and PET to Assess Brain Disorders in Humans

The association between Alzheimer's disease and an impaired function of P-gp is an interesting field of research and was studied by various research groups. In the last decade, several radiotracers have been developed to evaluate the function of P-gp in Alzheimer's disease (Leopoldo et al. 2014). Only a few of these tracers survived the in vitro and animal tests and were employed in humans. Until now, (*R*)-[ $^{11}\text{C}$ ]verapamil is the gold standard to measure the function of P-gp in patient brains. This tracer revealed a significant decline of P-gp function in AD patients compared to healthy controls (van Assema et al. 2012). However, (*R*)-[ $^{11}\text{C}$ ]verapamil does not allow assessment of the efficacy of P-gp inducers which are considered as promising drugs for the treatment of AD. For this purpose, weak P-gp substrates are intensively studied and validated in rodents and nonhuman primates. The fact that these compounds have shown a higher uptake at baseline and a higher sensitivity to changes of P-gp function compared to avid substrates is encouraging (Fusi et al. 2017; Raaphorst et al. 2017b; Savolainen et al. 2017b).

In PD, a decreased P-gp function in frontal regions of the brain was seen only in advanced stages of the disease, using (*R*)-[ $^{11}\text{C}$ ]verapamil. In the early stages of PD, no significant reduction of P-gp function was found, which makes P-gp dysfunction at the BBB less likely to be the primary cause of the disease (Bartels et al. 2008a, b).

In epilepsy, the function of P-gp is thought to be increased, which leads to insufficient uptake of antiepileptic drugs in the brain. Although [ $^{11}\text{C}$ ]verapamil alone cannot accurately assess increases of P-gp in patients with epilepsy, [ $^{11}\text{C}$ ]verapamil can be used in combination with tariquidar. This combination indicated that patients suffering from drug-resistant epilepsy show a higher P-gp function in some regions of the brain than drug-sensitive patients or healthy controls (Feldmann et al. 2013).

Additionally, Langer and colleagues studied the P-gp function in a small group of patients suffering from drug-resistant epilepsy, and they found some asymmetries in the kinetic parameters of [ $^{13}\text{C}$ ]verapamil in the parahippocampal and ambient gyrus (PAHG), amygdala, and medial and lateral anterior temporal lobe. However, these regional differences were not statistically significant (Langer et al. 2007). [ $^{13}\text{C}$ ]Phenytoin, a weak substrate for P-gp, is under investigation as a hopefully better technique to evaluate the function of P-gp in patients suffering from drug-resistant epilepsy.

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## 22.10 The Use of P-gp Tracers to Evaluate the Clinical Implications of Drug-Drug Interaction

ABC transporter-mediated drug-drug interactions (DDIs) have gained attention since numerous compounds can interact with these transporters.

On one hand, concomitant administration of drugs can culminate in adverse drug events which can cause patient hospitalization and increased patient morbidity as well as increased treatment costs. Interaction of a drug with ABC transporters can modify the pharmacokinetics of a co-administered drug by changing its absorption, distribution, metabolism, and/or elimination. DDIs can lead to unexpected toxicity or reduced drug effectiveness (Lund et al. 2017; Zhang et al. 2008). For instance, a drug that inhibits the P-gp function can increase the concentration of a concurrently administered drug in the target tissue and can thus increase the risk of an adverse drug event. In the case of drugs with a narrow therapeutic index such as digoxin, which is a P-gp substrate, it is essential to study potential DDIs (Zhang et al. 2008). However, it should be noted that not all DDIs will be clinically relevant (Lund et al. 2017).

On the other hand, transporter-mediated DDIs are crucial in the drug development process. The FDA and EMA have already advised *in vitro* assays to screen whether a new compound is a P-gp substrate, inhibitor, or inducer (Clinical Drug Interaction Studies — Study Design, Data Analysis, and Clinical Implications Guidance for Industry 2017; Guideline on the Investigation of Drug Interactions 2012). However, such *in vitro* methods have disadvantages. They are not yet standardized among laboratories, and P-gp inducers are difficult to recognize *in vitro* (Lund et al. 2017; Zhang et al. 2008). Furthermore, for ABC transporters other than P-gp, *in vitro* methods have not yet been fully developed (Zhang et al. 2008). In addition, if *in vitro* studies show that a new compound may interact with an ABC transporter, *in vivo* studies need to be performed to confirm this finding (Langer 2016; Zhang et al. 2008).

In clinical studies, the drug concentration in the blood of the subjects is usually measured. However, it is known that some DDIs mediated by ABC transporters do not result in a change of drug concentration in the blood, but changes in tissue pharmacokinetics. For example, P-gp inhibition at the BBB is usually not accompanied by changes in the plasma concentration of the drug, but it causes an increase in brain concentration (Langer 2016; Wulkersdorfer et al. 2014).

Since PET allows quantification of the concentration of the drug in the target tissue, imaging represents an interesting technique to study the DDIs mediated by ABC transporters. PET images can be used to quantify several pharmacokinetic parameters of a test drug, such as the tissue and blood concentrations during the scanning time. This usually requires the use of dynamic PET scans and blood sampling. By applying pharmacokinetic models to the data, different kinetic parameters are estimated which may help to interpret some physiological processes (Langer 2016).

PET studies are commonly applied in drug development to measure the binding of a new drug to its target receptor (Matthews et al. 2012; Cunningham et al. 2005; Lee and Farde 2006). Moreover, as many drugs can be affected by the efflux action of ABC transporters and, therefore, do not reach the desired target, PET studies of ABC transporters may be used to predict which drug can reach the CNS in appropriate concentrations.

These PET studies require the administration of a radiotracer to image the function of a specific ABC transporter. Each study consists of a dual-scan protocol where the radiotracer is administered under two different conditions. One scan will be performed without the administration of the test drug. It will provide information about the normal distribution of the tracer. This scan will be referred to as the baseline scan. The second scan will be performed after (or during) administration of the newly developed drug. Changes in the distribution of the radiotracer with regard to the baseline scan will show how the test drug affects the function of the specific ABC transporter. If the drug under study is expected to be a substrate of the ABC transporter, a different study protocol may be used. In this case, the drug under study could be radiolabeled with fluorine-18 or carbon-11. After studying the normal distribution of the test drug, a second PET scan should be performed in combination with a well-known inhibitor or inducer of the specific ABC transporter. This approach will show how changes in ABC transporter function affect the distribution of the test drug (Langer 2016; Wulkersdorfer et al. 2014).

PET studies of ABC transporters play now an important role in the assessment of drug disposition in target tissues, in particular, the CNS. Therefore, the search for improved tracers for PET imaging of ABC transporters has been intensified (Kannan et al. 2009; Wanek et al. 2013; Luurtsema et al. 2016).

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## 22.11 Discussion and Concluding Remarks

The implication of ABC transporters in the onset of neurodegenerative disease as well as in the deposition of CNS drugs makes these efflux transporters an interesting therapeutic target. Since PET imaging allows the *in vivo* assessment of biochemical processes, this technique is well-suited for evaluation of the ABC transporter function under different conditions or at different stages of a disease. Therefore, many research groups are developing novel PET tracers for improved assessment of ABC transporter function *in vivo*. Current research focuses on the development of weak substrates for P-gp which can be used to measure both increases and decreases of the efflux function.



PET imaging of ABC transporters may expand our knowledge regarding their involvement in the onset and progression of several neurodegenerative diseases. Novel treatment approaches for Alzheimer's or Parkinson's disease, such as treatment with P-gp inducers, can also be tested in experimental animals and humans, using PET. Moreover, PET imaging may be a useful tool to evaluate the affinity of novel CNS drugs toward various ABC transporters. This technique may predict the biodistribution and, in particular, the brain penetration of new drug candidates at an early stage in the drug development process, and may thus enhance the success rate of drug development.

It is expected that the use of novel and specific tracers for ABC transporters will increase our understanding regarding the diagnosis and treatment of neurodegenerative disease as well as our insight into potential transporter-mediated DDIs.

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# PET Imaging of Phosphodiesterases in Brain

# 23

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## Abstract

Phosphodiesterases (PDEs) regulate the intracellular concentration of the secondary messengers cGMP and cAMP thereby modulating a variety of cellular activities in brain. Eleven different families of PDEs have been identified of which at least six show significant expression in brain. Phosphodiesterase inhibitors have been evaluated clinically as potential drugs for treatment of different CNS diseases but so far, no PDE inhibitor has been registered as a drug for

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treatment of a CNS disease. PET ligands have been developed and evaluated targeting PDE2,4,5, and 10 and clinical PET studies were conducted with radio-labeled PDE2, 4, and 10 inhibitors. PDEs are interesting targets for in vivo imaging with PET as both their expression level and affinity for inhibitors can be modulated. This chapter provides an overview of the development of PDE PET tracers and their application in preclinical and clinical research.

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## 23.1 Introduction Phosphodiesterases

Phosphodiesterases (PDEs) are a group of 11 families of ubiquitous intracellular enzymes that hydrolyze the cyclic nucleotides 3',5'-cyclic adenosine monophosphate (cAMP) and/or 3',5'-cyclic guanosine monophosphate (cGMP) to the corresponding 5' monophosphates (Maurice et al. 2014).

The different families of PDEs are grouped based on their C-terminal catalytic domain homology. PDEs can be selective for cAMP or cGMP but can also have dual cGMP and cAMP selectivity (Table 23.1). Isoforms within a PDE family differ with regard to their N-terminal regulatory domain which is important for subcellular localization or modulation of catalytic activity.

Both cAMP and cGMP are important intracellular second messenger molecules for G-protein coupled receptors. Both cyclic nucleotides accomplish phosphorylation of a multitude of downstream targets including CREB (cAMP response element binding protein). cAMP activates protein kinase A (PKA), exchange protein activated by cAMP (Epac) (Bos 2006), and popeye domain-containing (POPDC) proteins (Schindler and Brand 2016). cGMP on the other hand stimulates protein kinase G. Both cAMP and cGMP activate cyclic nucleotide gated channels (CNGs).

The simultaneous use of cGMP and/or cAMP as second messengers in various cellular pathways is possible by compartmentalization, i.e., spatial separation of these signalling pathways as was observed in cardiac myocytes (Zaccolo 2011). PDEs can also be largely confined to specific subcellular pools in order to control local cAMP and/or cGMP levels (Mongillo et al. 2004).

The synthesis of cAMP is managed by both transmembrane adenylyl cyclases (tACs) and soluble adenylyl cyclases (sACs). The synthesis of cGMP is controlled by particulate and soluble guanylyl cyclases (pGCs). The concentration and compartmentalization of both cAMP and cGMP is tightly regulated by synthesis and hydrolysis through PDE activity although the hydrolysis outperforms the synthesis capacity (Baillie et al. 2019).

Both the catalytic function ( $V_{\max}$ ) and the affinity of PDEs can be modulated by different mechanisms including oligomerization, modulation by regulatory domains, and post translational modifications such as phosphorylation, ubiquitylation, SUMOylation, S-nitrosylation, and proline hydroxylation (Baillie et al. 2019).

PDEs exist as homodimers except for PDE1 and 6 which occur as heterotetramers (Bender and Beavo 2006). The conformation of the homodimer can change

**Table 23.1** Overview of characteristics of different phosphodiesterase families

PDE family (members)	Substrate	N-terminal regulatory/ localization domain	Relative expression in brain (Lakics et al. 2010) (expression in other organs)	Clinical trial with CNS target	PET tracers for CNS
PDE1 (1A, 1B, 1C)	cGMP, cAMP	Calmodulin	1B caudate and nucleus accumbens, 1C substantia nigra	Schizophrenia, PD, cognition enhancement	–
PDE2 (2A)	cGMP, cAMP	GAFA/B-cGMP stimulated	Hippocampus, cortical regions	Schizophrenia	+
PDE3 (3A, 3B)	cGMP, cAMP	Transmembrane domain	3A cerebellum	Tinnitus, AD, MCI	–
PDE4 (4A, 4B, 4C, 4D)	cAMP	UCR	4A, B, and D all brain regions. 4C cortex, thalamic nuclei and cerebellum	AD, Fragile X, HD, dementia	+
PDE5 (5A)	cGMP	GAFA/B cGMP stimulated	Low levels in vasculature	Migraine, mild TBI	–
PDE6 (6A, 6B, 6C)	cGMP	GAFA/B cGMP stimulated	(PDE6A retina)	–	–
PDE7 (7A, 7B)	cAMP		PDE7B caudate, nucleus accumbens, cortex, hippocampus	–	–
PDE8 (8A, 8B)	cAMP	Per-ARNT-Sim (PAS) domain	(PDE8B thyroid) 8A and 8B low brain levels	–	–
PDE9 (9A)	cGMP	Nuclear localization domain	Dorsal root ganglia, cerebellum	AD, schizophrenia	–
PDE10 (10A)	cGMP, cAMP	GAFA/B cAMP stimulated	Caudate nucleus, nucleus accumbens	Schizophrenia	++
PDE11 (11A)	cGMP, cAMP	GAFA/B cGMP stimulated	Dorsal root ganglia Substantia nigra	–	–

due to activation of the regulatory domain from a low activity conformation with (partially) blocked access to the catalytic site to a highly active conformation with fully accessible catalytic site in a high affinity state.

PDEs have regulatory domains including calcium/calmodulin-binding domains (PDE1), GAF domains (cGMP-binding PDEs, *Anabaena*adenylyl cyclase, and *Escherichia coli* FhlA; PDE2, PDE5, PDE6, PDE10, and PDE11), PAS domains (Per-ARNT-Sim; PDE8), UCR domains (upstream conserved regulatory domains; PDE4), and autoinhibitory domains. The GAF and UCR domains are unique regulatory domains that only occur in PDEs in mammals.

The GAF domains of PDE2,5,6, and 11 allosterically bind cGMP, whereas the GAF domain of PDE10 allosteric binds cAMP. This allosteric binding causes

structural changes relieving autoinhibition of the PDEs (Francis et al. 2011). GAF domains can also provide contacts for dimerization<sup>8</sup>. cGMP binding the GAF-A domain of PDE6 enhances protein–protein interactions that inhibit PDE6 catalytic activity. Allosteric binding of cGMP to the GAF-B domain of PDE2A increases hydrolysis rate of cAMP with a factor of 10 thereby cross-regulating the concentration of both cyclic nucleotides (Wu et al. 2018).

### 23.1.1 Drugs Targeting PDEs

Major marketed PDE inhibitors include non-specific PDE inhibitors used since 1970s for treatment of asthma; PDE3 inhibitors for treatment of heart failure and claudication; PDE4 inhibitors for treatment of chronic obstructive pulmonary disease, psoriasis, dermatitis, and bowel disorders and PDE5 inhibitors for treatment of erectile dysfunction and pulmonary arterial hypertension (Baillie et al. 2019).

Most PDEs are (also) expressed in the brain, and a large number of clinical studies have investigated the use of PDE inhibitors for treatment of various neurological diseases (Table 23.1). Most if not all of these clinical studies however failed due to safety concerns or a lack of efficacy. In some of these clinical trials PET was used to determine the PDE occupancy level of the drug candidate (see below).

Besides PDE inhibitors targeting the catalytic site also allosteric modulators targeting the regulatory domain (which can also activate the PDE activity) are being pursued preclinically for PDE4D (Sutcliffe et al. 2014; Burgin et al. 2010; Omar et al. 2019; Titus et al. 2018), PDE5 (Weeks et al. 2005), PDE10 (García et al. 2017; Jäger et al. 2012), and PDE11 (Weeks et al. 2005).

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## 23.2 PET Studies Targeting Phosphodiesterases

Phosphodiesterases are an interesting target for PET but the binding potential of PET tracers depends both on expression levels and affinity and both parameters can be modulated for PDEs in CNS pathologies. Especially phosphodiesterases with a GAF domain are subject to cGMP and/or cAMP modulation of the affinity of PET tracers directed to the C-terminal catalytic domain. The specific cause for a changed PET binding potential in a studied pathology may therefore be caused by local cGMP and/or cAMP concentration induced PDE affinity modulation vs. PDE expression. The PET binding potential may however provide a more physiologically relevant readout for global PDE activity compared to techniques that only rely on quantifying protein expression levels. On the other hand, PET has only a limited resolution which does not allow to identify potential changes in subcellular compartmentalization of PDEs which have been reported to be important in different CNS pathologies. The existence of 11 different PDE families, each consisting of a number of subtypes requires PET tracers to have high selectivity for the PDE type that is studied as different PDE subtypes will likely be present in the same brain region, cells, and cell compartments.

Different PET tracers for phosphodiesterases have been reviewed extensively (Andrés et al. 2012; Schröder et al. 2016), so in this chapter we will focus on novel developments reported since 2016.

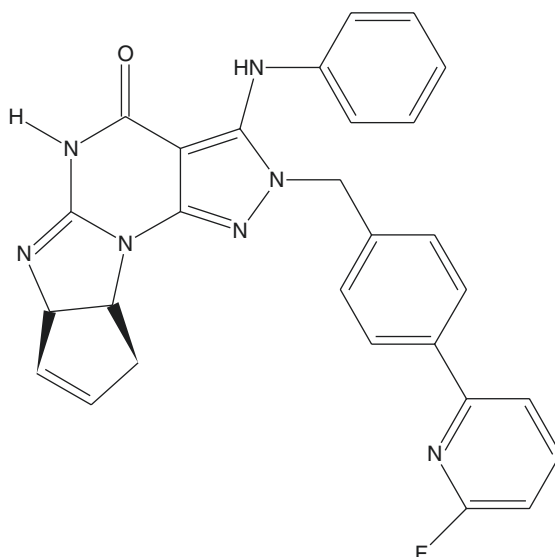
### 23.2.1 PDE1

The PDE1 family consists of PDE1A, PDE1B, and PDE1C isoforms and is the only mammalian PDEs that are transiently activated by the calcium/calmodulin complex. They thus provide intracellular cross-talk between calcium and cyclic nucleotide signaling. PDE1B has the highest expression in brain and co-localizes with dopaminergic D<sub>1</sub> and D<sub>2</sub> receptors in striatal and cortical neurons (Pekcec et al. 2018). Several PDE1 inhibitors have been developed (Chen et al. 2019; Li et al. 2016; Dyck et al. 2017; Khammy et al. 2017; Enomoto et al. 2019) of which ITI-214 (Fig. 23.1) was tested in a clinical study for treatment of schizophrenia (NCT01900522). So far however, no PET tracers targeting PDE1A were reported.

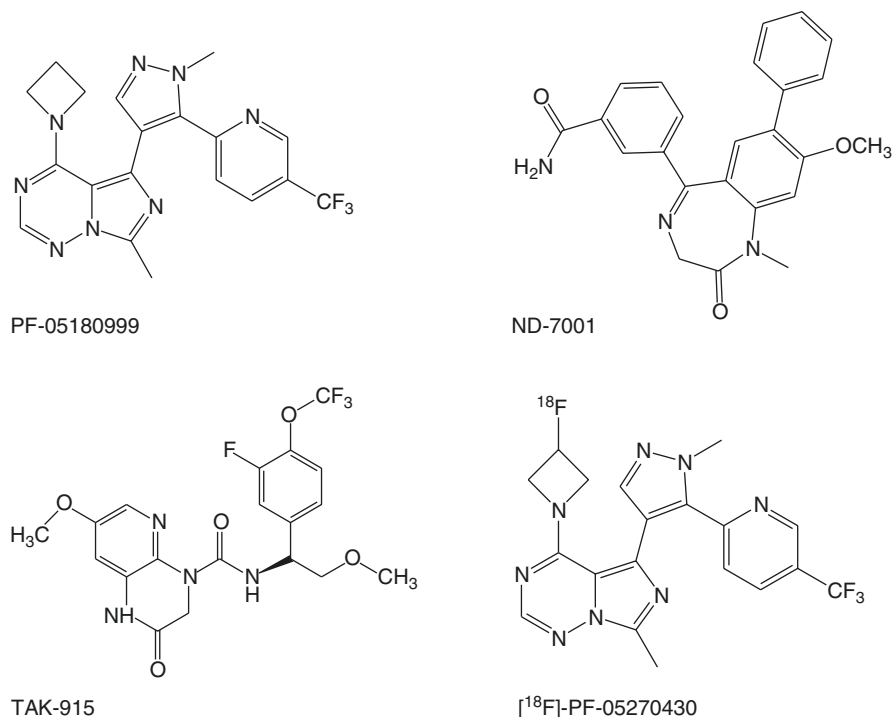
### 23.2.2 PDE2A

PDE2A hydrolyzes both cAMP and cGMP and shows high expression in cortex, amygdala, and hippocampus. Several PDE2A specific inhibitors were developed (Trabanco et al. 2016) and some were evaluated in clinical studies for anxiety, cognition enhancement, schizophrenia, and/or depression including ND-7001 (Gomez and Breitenbucher 2013), PF-05180999 (Fig. 23.2) (Helal et al. 2018) (NCT01981486 and NCT01981499), and TAK-915 (Fig. 23.2) (Mikami et al. 2017) (NCT02461160, NCT02584569).

**Fig. 23.1** Structure of PDE1 inhibitor ITI-214







**Fig. 23.2** Chemical structures of PDE2 inhibitors PF-05180999, ND-7001, TAK-915, and PDE2A PET tracer [<sup>18</sup>F]PF-05270430

Several PDE2A PET tracers have been developed and evaluated preclinically (Chen et al. 2016; Schröder et al. 2015, 2018; Ritawidya et al. 2019) but only two tracers ([<sup>18</sup>F]PF-05270430 and [<sup>18</sup>F]MNI-794, Fig. 23.2) were evaluated clinically. [<sup>18</sup>F]PF-05270430 displayed fast and reversible kinetics in nonhuman primates, as well as PDE2A specific binding blockable by a PDE2A inhibitor. First in human evaluation of [<sup>18</sup>F]PF-05270430 (Naganawa et al. 2016) demonstrated highest uptake in striatum, followed by neocortical regions and white matter, and lowest in the cerebellum. Binding potential relative to the nondisplaceable tracer ( $BP_{ND}$ ) were determined with the simplified reference tissue model and a 70 min scan using cerebellum as reference tissue.  $BP_{ND}$  values ranging from 0.3 (parietal cortex) to 0.8 (putamen) were obtained with a test-retest variability of 16%.

[<sup>18</sup>F]MNI-794 was used to quantify the occupancy of TAK-915 (NCT02584569, Fig. 23.2), but no detailed reports on this PET tracer have been published.

Unfortunately no preclinical nor clinical studies using PET to investigate the expression levels of PDE2A in CNS pathologies have been published so far. Interestingly, at low dose, PDE2A inhibition produced a paradoxical increase of PDE2A autoradiography binding probably due to release of cGMP activating a dormant pool of PDE2A enzymes via binding to the GAF-B domain (Pandit et al. 2009;

Gu et al. 2019). A loss of function mutation in PDE2A was found to be associated with chorea suggesting a role of PDE2A in hyperkinetic movement disorders (Salpietro et al. 2018).

### 23.2.3 PDE4

PDE4 is a highly diverse family of phosphodiesterases which exclusively hydrolyzes cAMP. The family is encoded by four sub-family genes (PDE4A to D) which generate over 16 different splice variants with each a unique N-Terminal regulatory domain. These splice variants can generally be divided into three major categories: long form, short form, and super-short form (Omori and Kotera 2007). The N-Terminal domain of the long form splice variants contains two Upstream Conserved Regions (UCR1 and UCR2) which are conserved in all PDE4 sub-families. Short and super-short isoforms however only express one or even only a part of one UCR (Francis et al. 2011). It is generally accepted that UCR 1 is required for dimerization, while UCR2 plays an important role in the regulation of enzymatic activity and intracellular targeting (Richter and Conti 2002). PDE4 also has a phosphorylation site for PKA that regulates PDE4 activity and activates PDE4 when cAMP levels are high.

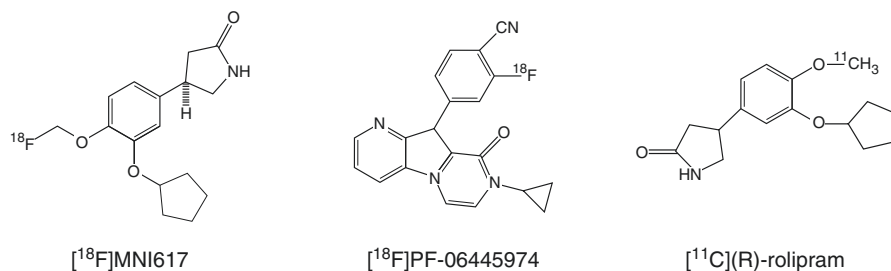
The expression of PDE4 is rather ubiquitous; however high expression in immune cells, brain, and cardiovascular tissues have been reported (Richter et al. 2011; Lakics et al. 2010; Tenor et al. 1996). Among all PDE families, PDE4 is the most abundant in the brain, accounting for 30–70% of all PDE activity in the central nervous system (Lakics et al. 2010). Due to the importance of cAMP signaling in the brain physiology, it is generally accepted that PDE4, as the major cAMP hydrolyzing enzyme in the brain, plays a decisive role in normal functioning of the brain. Alterations in cAMP signaling are believed to be a key element in the pathophysiology of multiple psychiatric and neurological disorders such as depression, Alzheimers Disease, and Parkinson's disease (Missale et al. 1998; Reiach et al. 1999; Martínez et al. 1999; Casacchia et al. 1983). Therefore, it is believed that stabilizing cAMP levels using PDE4 inhibitors can have beneficial effects on emotional and cognitive behaviors in such diseases (Zhu et al. 2001; Li et al. 2009; O'Donnell 1993; O'Donnell and Frith 1999; Zhang and O'Donnell 2000; Zhang 2009). Additionally, PDE4 is highly expressed in inflammatory cells and lung smooth muscle cells. Therefore, it is believed that PDE4 is also an important regulator of the inflammatory response in several lung diseases such as asthma and Chronic Obstructive Pulmonary Disease (COPD). Inhibition of PDE4 could therefore also have beneficial effects on the symptoms of these diseases (Hatzelmann et al. 2010).

Rolipram is a non-subtype selective PDE4 inhibitor which has been used for treatment on regular basis in preclinical and clinical trials (Casacchia et al. 1983; Renau 2004). In a number of single-arm (phase I) trials and in several randomized phase III trials, rolipram demonstrated clinical activity in reducing symptoms of depression. Also in COPD, rolipram was found to be an efficient treatment option.

The application of rolipram was however limited due to dose-limiting side effects including headache, gastric hyper secretion, and emesis (Hebenstreit et al. 1989; Scott et al. 1991; Bobon et al. 1988; Fleischhacker et al. 1992; Krause et al. 1990). Especially the emetic effects have put a serious hold on the further development of rolipram and other related PDE4 inhibitors. It is believed that the lack of selectivity of rolipram toward the different PDE4 subtypes may account for some of the side effects observed. Several studies indicate that the emetic effects of rolipram might be caused by the PDE4D inhibition in the area postrema and nucleus of the solitary tract (Halpin 2008; Lamontagne et al. 2001; Mori et al. 2010; Cherry and Davis 1999; Robichaud et al. 2002), while PDE4B inhibition would be responsible for the anti-depressant effect of rolipram (Zhang et al. 2017).

For the treatment of COPD, the problem of emesis was reduced by the development of a PDE4 inhibitor with limited brain penetration to avoid central inhibition of PDE4D. This PDE4 inhibitor, roflumilast has now been approved for the treatment of COPD (Rabe 2011). When treating neuropsychiatric disorders such as depression, brain penetration is however crucial for its pharmacological activity. Therefore, more efforts are invested on developing subtype specific inhibitors to avoid the emesis problem (Azam and Tripuraneni 2014). Since mice deficient in PDE4B showed anxiogenic and antidepressant effects, PDE4B is believed to be an important target for depression (Zhang et al. 2008). Several PDE4B selective inhibitors have been developed showing similar anti-depressant activities as rolipram (Azam and Tripuraneni 2014). Although PDE4D inhibition has been linked to the emetic effects of non-selective PDE4 inhibitors, it is still believed that PDE4D inhibitors could have therapeutic potential as PDE4D is attributed to play an important role in cognition. Indeed, animal models deficient of the PDE4D gene show increased cognitive performance (Li et al. 2011; Ricciarelli et al. 2017). Therefore, a lot of effort is also put in the development of PDE4D-selective inhibitors as a treatment for Alzheimer's disease and cognitive brain disorders. For example, PDE4D inhibitors such as BPN14770 are currently under development for the treatment of Fragile X Syndrome and other brain disorders (Gurney et al. 2019; Cui et al. 2019).

The majority of the information on PDE4 comes from *ex vivo* data. Enzymatic activity of PDE4 is however highly dependent on the phosphorylation state of the enzyme (MacKenzie et al. 2002). Since dephosphorylation occurs rapidly after death, caution is required when interpreting *ex vivo* measurements of PDE4 activity. Therefore, the availability of PDE4 PET imaging is crucial to quantify PDE4 alterations *in vivo*. The non-PDE4 selective inhibitor Rolipram has 2 enantiomers who have both been radiolabelled with carbon-11 for PET imaging. [<sup>11</sup>C](R)-Rolipram (Fig. 23.3) has a high affinity for PDE4 and PET imaging with [<sup>11</sup>C](R)-Rolipram has been shown to allow monitoring of PDE4 activity and expression *in vivo* (Lourenco et al. 2001, 1999). Since its development, [<sup>11</sup>C](R)-Rolipram has been used to explore PDE4 levels in several preclinical models of disease as well as in patient populations. [<sup>11</sup>C](S)-Rolipram on the other hand has a 20-fold lower affinity toward PDE4 compared to its R-enantiomer. Due to its much lower affinity however, [<sup>11</sup>C](S)-Rolipram can be used to quantify non-displacable binding or changes in tracer brain delivery caused by disease or treatment (Fujita et al. 2005; Itoh et al. 2010).



**Fig. 23.3** Chemical structures of PDE4 tracers  $[^{18}\text{F}]\text{MNI617}$ ,  $[^{18}\text{F}]\text{PF-06445974}$ , and  $[^{11}\text{C}](\text{R})\text{-rolipram}$

A recent study in conscious rats showed that PKA-mediated phosphorylation of PDE4 increased the *in vivo* binding of  $[^{11}\text{C}](\text{R})\text{-Rolipram}$ , consistent with the increased affinity of rolipram toward the phosphorylated PDE4 enzyme (Itoh et al. 2010; Takano et al. 2018). This study was the first to indicate that  $[^{11}\text{C}](\text{R})\text{-Rolipram}$  binding was dependent on the phosphorylation state of PDE4. A further study building upon these results showed that  $[^{11}\text{C}](\text{R})\text{-Rolipram}$  was also sensitive to changes in PDE4 activity due to protein–protein interaction. This study used  $[^{11}\text{C}](\text{R})\text{-Rolipram}$  PET to investigate the interaction between disrupted in schizophrenia protein 1 (DISC1) and PDE4 (Ooms et al. 2019). The fact that PDE4 activity was increased in absence of critical DISC1 protein isoforms suggests a complex regulatory system for PDE4 activity. Additionally, it potentially links PDE4 activity to multiple neurological disorders such as schizophrenia and even Huntington’s disease.

Apart from the preclinical exploratory studies investigating PDE4 activation,  $[^{11}\text{C}](\text{R})\text{-Rolipram}$  PET can also be used to evaluate targeting properties in the development of novel PDE4 inhibitors. For example, brain PDE4 occupancy of Roflumilast was assessed in non-human primates using  $[^{11}\text{C}](\text{R})\text{-Rolipram}$  PET (Takano et al. 2018). Also in human subjects,  $[^{11}\text{C}](\text{R})\text{-Rolipram}$  was used to assess brain occupancy of multiple PDE4 inhibitors such as GSK356278 (van der Aart et al. 2018) and HT-0712 (NCT01215552).

Despite the necessity of preclinical models, they do have their limitations, especially when it comes to evaluating neuropsychiatric disorders. The first-in-man studies using  $[^{11}\text{C}](\text{R})\text{-Rolipram}$  PET imaging in healthy volunteers have been conducted already quite some time ago (DaSilva et al. 2002). Since then,  $[^{11}\text{C}](\text{R})\text{-Rolipram}$  PET has been used in multiple studies to investigate the role of PDE4 in neurological and psychiatric disorders.

Dysregulations in the cAMP pathway have been thought to be an important contributing factor to the symptoms of major depressive disorder (MDD). As the brain’s major cAMP hydrolyzing PDE, PDE4 is therefore hypothesized to be a key player in the pathogenesis of MDD. Initial studies using  $[^{11}\text{C}](\text{R})\text{-Rolipram}$  to investigate PDE4 activity in a small population of unmedicated patients with MDD have found a significant downregulation of PDE4 activity (Fujita et al. 2012). The decline in PDE4 activity could however be reversed by a treatment using selective serotonin reuptake inhibitors. These studies confirm that PDE4 and the cAMP signaling

pathway play an important role in the pathology of MDD. The lack of correlation between [ $^{11}\text{C}$ ](R)-Rolipram binding and symptoms of MDD however indicate a complexity of the disease beyond PDE4 (Fujita et al. 2017).

As stated previously, PDE4 is also believed to be involved in the pathophysiology of several other disorders. Several recent studies have used [ $^{11}\text{C}$ ](R)-Rolipram PET to evaluate PDE4 levels in patient populations with Parkinson's disease. [ $^{11}\text{C}$ ](R)-Rolipram binding was decreased in several brain regions of the striato-thalamo-cortical circuit. The loss of PDE4 expression correlated to the performance in spatial working memory (Niccolini et al. 2017). Additionally, a correlation with excessive daytime sleepiness was found suggesting a role of PDE4 in the pathophysiology of Parkinson's disease (Wilson et al. 2019).

Finally a recent study has been started in which [ $^{11}\text{C}$ ](R)-Rolipram PET is being used to evaluate PDE4 levels in brain and peripheral tissues of patients with McCune-Albright syndrome (NCT02743377). This rare genetic disorder is a mosaic disorder arising from a spontaneous mutation of the GNAS complex locus (Guanine Nucleotide binding protein, Alpha Stimulating complex), which encodes the cAMP pathway-associated G-protein, Gs $\alpha$ . The resulting local dysregulation in cAMP signaling is believed to cause a disperse pattern of fibrous dysplasia. The preliminary data of their study in two patients showed a [ $^{11}\text{C}$ ](R)-Rolipram distribution pattern that correlates with the known locations of fibrous dysplasia in patients with McCune-Albright syndrome (Lora et al. 2019).

In all these clinical trials, determination of [ $^{11}\text{C}$ ](R)-Rolipram distribution volumes is crucial for the correct quantification and interpretation of PET images. Due to the lack of a suitable reference region, quantification of the concentration of [ $^{11}\text{C}$ ](R)-Rolipram in the brain requires arterial sampling. In order to decrease the invasiveness of the procedure and hence increasing clinical applicability, several studies looked for alternative methods for quantification of brain [ $^{11}\text{C}$ ](R)-Rolipram uptake. Both image derived input function using time activity curves from the carotid artery (Lyoo et al. 2014) and a population-based input curve have been shown to be a valuable alternative to avoid arterial sampling (Zanotti-Fregonara et al. 2012).

Despite the success of [ $^{11}\text{C}$ ](R)-Rolipram, the quest for an improved PDE4 PET tracer continues. Recently a fluorine-18-radiolabeled alternative for rolipram, [ $^{18}\text{F}$ ]MNI-617 (Fig. 23.3), was successfully radiolabeled and evaluated in non-human primates. It rapidly entered the brain with brain uptake similar to [ $^{11}\text{C}$ ](R)-Rolipram and a distribution pattern consistent with that of PDE4 expression (Thomae et al. 2016). Together with the increased interest in PDE4 subtype selective inhibitors, also the quest for PDE4 subtype selective PET tracer was engaged. PF-06445974 (Fig. 23.3) is a tricyclic pyrrolopyridine compound with a sub-nanomolar affinity for PDE4B and a good selectivity toward PDE4D (>36 $\times$ ). As a PET tracer, [ $^{18}\text{F}$ ]PF-06445974 showed rapid brain uptake and high target specificity. Clinical trials are currently ongoing to evaluate the feasibility of this radiopharmaceutical for PDE4B visualization in a healthy patient population (NCT03030391).

Parallel to the development of BPN14770 as a PDE4D selective inhibitor for the treatment of several brain disorders including fragile X syndrome and

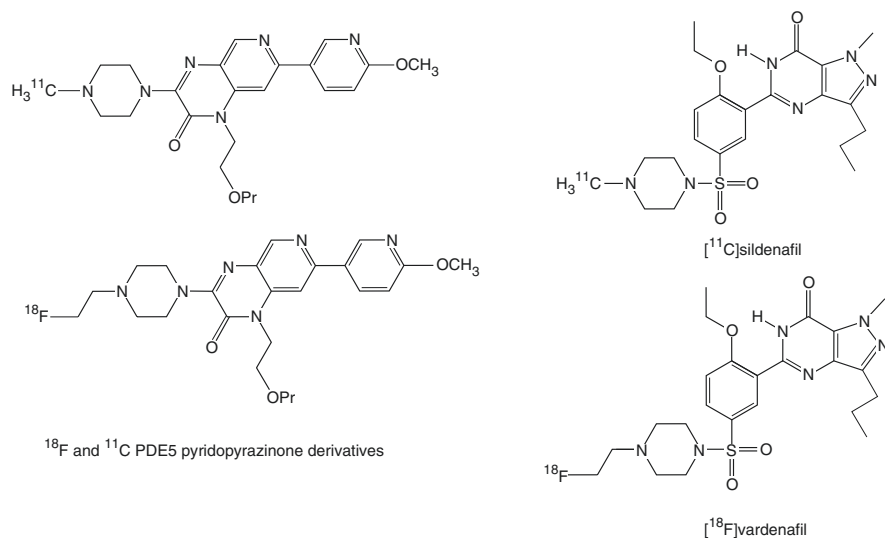
Alzheimer's disease (Gurney et al. 2019), several PDE4D selective PET radioligands were developed. In a recent study, [ $^{11}\text{C}$ ]T1650 was evaluated as a PDE4D specific PET ligand. Studies in non-human primates indicated that [ $^{11}\text{C}$ ]T1650 showed good brain uptake that could be blocked after co-injection with BPN14477056. Currently, clinical trials using [ $^{11}\text{C}$ ]T1650 and [ $^{11}\text{C}$ ]T2310 (undisclosed structures) combined with BPN144770 blocking (NCT03861000, NCT04044781) are ongoing.

### 23.2.4 PDE5

PDE5 is notorious for its regulation of blood flow in the corpus cavernosum through hydrolysis of cGMP. Several PDE5 inhibitors have been marketed for treatment of erectile dysfunction, and some are being investigated for the treatment of pulmonary hypertension (Ahmed 2018).

PDE5 was however also reported to be expressed in the brain, and PDE5 inhibition was suggested to be neuroprotective (Teich et al. 2016; Gulisano et al. 2018; Heckman et al. 2017).

Several PDE5 PET tracers have been evaluated preclinically but so far none of the PET tracers were able to visualize PDE5 in brain. Several vardenafil derivatives, ethoxyethyl pyrazolopyrimidine derivatives, and carbon-11 labelled sildenafil (Fig. 23.4) have relatively high PSA values and showed low brain uptake (Chekol et al. 2014). PDE5-specific retention in the lungs of mice in relation with the PDE5 affinity of the different evaluated tracers was observed.



**Fig. 23.4** Chemical structures of PDE5 PET tracers:  $^{18}\text{F}$  and  $^{11}\text{C}$  PDE5 pyridopyrazinone derivatives, [ $^{11}\text{C}$ ]sildenafil and [ $^{18}\text{F}$ ]vardenafil

Although no PDE5-specific binding was observed for  $^{11}\text{C}$ -sildenafil, therapeutic doses of sildenafil resulted in pharmacologically relevant CSF levels and increased CSF cGMP levels in a non-human primate (Gómez-Vallejo et al. 2016).

A fluorinated quinolone was synthesized and tested *in vitro* and *in vivo* but the tracer was quickly metabolized *in vivo* generating brain permeable radiometabolites (Liu et al. 2016; Wenzel et al. 2019). Other quinolone derivatives were evaluated *in vitro* for their PDE5 affinity and selectivity, but their radiolabelling and *in vivo* evaluation has not yet been reported (Liu et al. 2017).

High affinity fluorine-18- and carbon-11-labelled pyridopyrazinone derivatives (Fig. 23.4) were found to efficiently cross the blood–brain barrier but no PDE5-specific retention was observed in rat brain (Chekol et al. 2017).

### 23.2.5 PDE7

Both PDE7 gene silencing (Morales-Garcia et al. 2015a) and pharmacological PDE7 inhibition (Morales-Garcia et al. 2015b, 2017) were reported to attenuate neurodegeneration in Parkinson model mice.

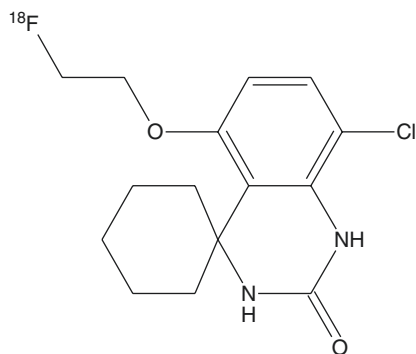
No clinical studies targeting PDE7 have been reported so far, and only few PDE7-specific inhibitors were reported (Safavi et al. 2013; Paterniti et al. 2011).

The spiroquinazolinone [ $^{18}\text{F}$ ]MICA-003 (Fig. 23.5) was evaluated as a PET tracer for PDE7 in the brain, but although the tracer showed good brain entry and fast wash-out, the tracer was found to quickly metabolize generating radiometabolites that accumulated in brain (Thomae et al. 2015). Also, no blocking studies were reported so that is not clear whether brain uptake and retention was (in part) PDE7 specific.

### 23.2.6 PDE9

PDE9 selectively hydrolyzes cGMP with high affinity. *In situ* hybridizations studies indicate that more than 20 different splice variants of PDE9A exist. PDE9 is

**Fig. 23.5** Structure of [ $^{18}\text{F}$ ]MICA-003



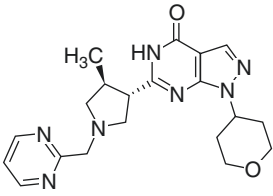
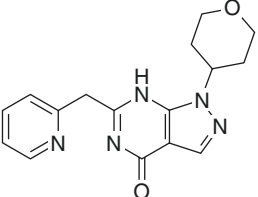
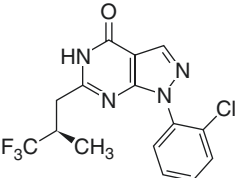
expressed in proximal dendrites and cell bodies of neurons throughout the whole brain (Patel et al. 2018). Different PDE9A inhibitors were advanced to clinical studies for treatment of Alzheimer's disease, schizophrenia and attenuated psychosis syndrome (Boland et al. 2017; Moschetti et al. 2018) (Table 23.2).

No PET radioligands have however been reported so far.

### 23.2.7 PDE10A

PDE10A is a dual substrate (although with a much higher affinity but lower  $V_{max}$  for cAMP than for cGMP) phosphodiesterase with brain expression limited to the basal ganglia and mainly in the axons of the medium spiny neurons in the striatum. PDE10A modulates the cAMP/PKA/DARP-32 signalling cascade and the phosphorylation of many downstream effector molecules. Twelve splice variants of PDE10 were identified but the predominant forms in brain are PDE10A1 that localizes in the cytosol, whereas palmytoylated PDE10A2 is associated with plasma membranes (Jäger et al. 2012). Recently, PDE10A19 was identified as an isoform which is exclusively expressed in primates and also localizes in the cytosol (MacMullen et al. 2017). PDE10 can be modulated via its GAF-B domain that binds

**Table 23.2** Chemical structure of PDE9A inhibitors and pathologies for which they were evaluated clinically

Name	Structure	Clinical trials
PF-04447943		Alzheimer's disease (NCT00930059, NCT00988598), sickle cell disease (NCT02114203)
BI 409306		Schizophrenia (NCT03351244, NCT01892384, NCT02281773), Alzheimers disease (NCT02392468, NCT02337907, NCT02240693), Attenuated Psychosis Syndrome (NCT03230097).
Eisai E2027	Undisclosed structure	Dementia with Lewy bodies (NCT03467152)
BAY 73-6691		–



cAMP with high affinity and effectively stimulates PDE10A hydrolysis of cAMP. GAF-B stimulates cGMP hydrolysing activity at low cAMP concentration but completely inhibits cGMP hydrolysis at high cAMP concentration (Thomae et al. 2015).

PDE10A has been associated with Huntington's disease, Parkinson's disease (García et al. 2014), and schizophrenia (Świerczek et al. 2019). MacMullen et al. also identified an association between PDE10A single-nucleotide polymorphisms within the first large intron and bipolar disorder (Patel et al. 2018).

A clinical study with PF-02545920 (aka MP10) did not reveal significant differences in efficacy compared to placebo, challenging the use of PDE10A inhibitors as antipsychotic agents for schizophrenia (DeMartinis 3rd et al. 2019). Also in the treatment of Huntington's disease PF-02545920 did not perform significantly better than placebo (NCT02342548).

The efficacy and safety of TAK-063 was evaluated in schizophrenic patients. Although this study did not meet the primary endpoint, the effects of TAK-063 may be suggestive of antipsychotic activity (Macek et al. 2019). TAK-063 was also evaluated for prevention of ketamine-induced brain activity changes and psychotic-like symptoms using fMRI BOLD (NCT01892189). In rats and monkeys, TAK-063 increased fMRI BOLD signal in striatum and amygdala and reversed ketamine-induced BOLD signals in the cortex (Tomimatsu et al. 2016), but the full results of this clinical study have not yet been reported.

OMS643762 (NCT01952132) and EVP-6308 (NCT02037074) were also evaluated in schizophrenic patients, but results of these clinical studies have yet to be disclosed.

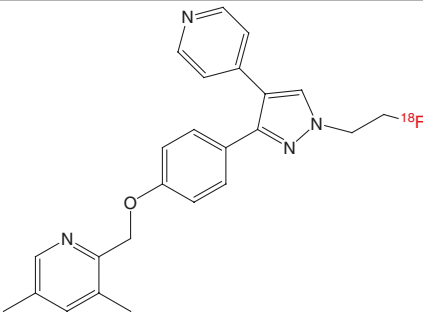
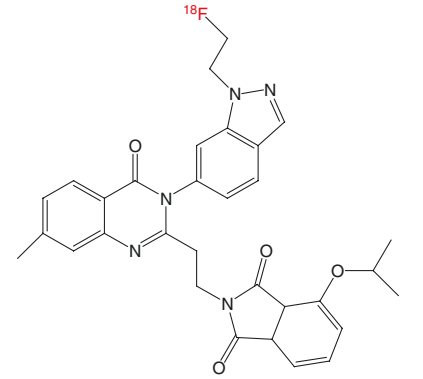
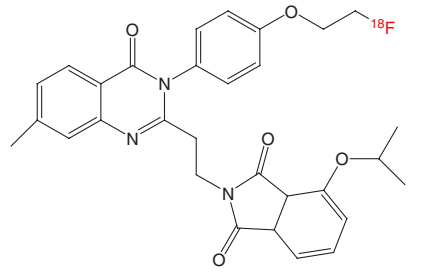
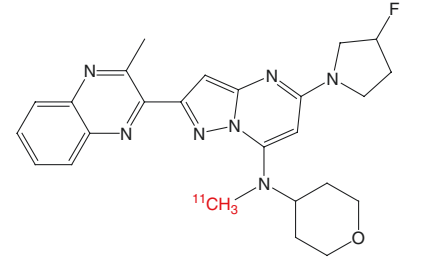
A multitude of PDE10A radioligands have been evaluated preclinically and at least six PDE10A tracers (Table 23.3) were advanced to clinical studies. High specific binding of PDE10A tracers is observed in striatum and globus pallidus and low binding in other brain regions.

In vivo PET studies in rats showed higher striatal binding of [<sup>18</sup>F]JNJ42259152 after pretreatment with the PDE4 inhibitor rolipram or the PDE2 inhibitor JNJ49137530 demonstrating profound interactions and cooperation between different PDEs in brain in vivo (Ooms et al. 2016).

In preclinical PET studies on rodents no modulation of PDE10A binding potential was identified after chronic treatment with haloperidol (Natesan et al. 2014). Repeated D-amphetamine treatment however significantly increases PDE10A binding in rats, but this effect can be abolished by selective D<sub>1</sub> receptor blocking (Ooms et al. 2017). Early decrease of PDE10A binding was observed in R6/2 (Ooms et al. 2014) and Q175 (Bertoglio et al. 2018; Häggkvist et al. 2017) Huntington mice.

After careful comparison, Bioscutti et al. concluded that [<sup>11</sup>C]IMA107, [<sup>18</sup>F]MNI659, and [<sup>11</sup>C]Lu AE92686 have the most appropriate characteristics as PDE10A PET ligands for clinical application, based on brain kinetics, test-retest variability, and specific signal considerations (Boscutti et al. 2019). Since expression of PDE10A is very low in the cerebellum, a dynamic scanning protocol using the simplified reference tissue approach can be used to quantify the BP<sub>ND</sub>.

**Table 23.3** PDE10A PET tracers that were advanced to clinical studies

Name	Structure	Clinical studies
[ <sup>18</sup> F] JNJ42249152		Dosimetry (Van Laere et al. 2013a) test-retest (Van Laere et al. 2013b), Huntington (Ahmad et al. 2014)
[ <sup>18</sup> F]MNI654		First-in-man and dosimetry (Barret et al. 2014)
[ <sup>18</sup> F]MNI659		First-in-man and dosimetry (Häggkvist et al. 2017), Huntington (Russell et al. 2016), age and gender (Fazio et al. 2017a), occupancy of PF-02545920 (DeInommedieu et al. 2017), occupancy of EVP-6308 (NCT02001389)
[ <sup>11</sup> C]IMA107		First in man (Plisson et al. 2014) Huntington (Niccolini et al. 2015a), Parkinson (Niccolini et al. 2015b), schizophrenia (Marques et al. 2016) PDE10A p.Tyr107Cys mutation (Diggle et al. 2016) Cocaine users (Tollefson et al. 2019), premanifest Huntingtons (Wilson et al. 2016), occupancy of RO5545965 (NCT01923025)

(continued)

**Table 23.3** (continued)

Name	Structure	Clinical studies
[ <sup>11</sup> C]T-773		Occupancy TAK-063 (Takano et al. 2016)
[ <sup>11</sup> C]Lu AE92686		First in man (Kehler et al. 2014), schizophrenia (Bodén et al. 2017)

Loss of PDE10A levels (quantified as  $BP_{ND}$ ) in caudate (−28%), putamen (−26%) and globus pallidus (−19%) were found to correlate well with the gradual and progressive increase of motor symptoms in patients with Parkinsons disease, indicating that PDE10A imaging provides a similar potential as dopamine transporter imaging to monitor Parkinsons disease progression (Niccolini et al. 2015b). No difference was found in the  $BP_{ND}$  of patients responding to levodopa and patients with levodopa-induced dyskinesias (Takano et al. 2016).

In line with preclinical findings, significantly lower levels (−50 to −70%) of PDE10A were also detected in striatum and globus pallidus of patients with manifest Huntington's disease (Ahmad et al. 2014; Russell et al. 2016; Wilson et al. 2017). Also in premanifest Huntington's disease gene expansion carriers (HDGECs), a decrease in PDE10A binding was observed correlating with worse UHDRS (Unified Huntington's Disease Rating Scale) motor scores, disease burden pathology, and regional atrophy (Russell et al. 2014). In far-onset premanifest HDGECs (predicted on average 25 years from symptomatic onset), striatal and pallidal PDE10A tracer binding was reduced by about 30%, whereas a 35% increase was observed in the motor thalamic nuclei (Niccolini et al. 2015c). Overall PDE10A alterations seem to be one of the earliest biochemical changes in HD identified to date, and PDE10A PET likely can be used to monitor HD progression and follow up efficacy of novel treatments (Boscutti et al. 2019).

In patients with PDE10A mutations associated with a hyperkinetic movement disorder, PDE10A PET also demonstrated loss of striatal PDE10A tracer binding (Diggle et al. 2016). Interestingly, these mutations are located in the regulatory allosteric GAF-A or GAF-B domains, and the results of this study thus indicate that these patients would benefit from strategies aiming to increase PDE10A function (Whiteley et al. 2019).

Patients with schizophrenia showed in one study similar (Marques et al. 2016) but in another study lower PDE10A BP<sub>ND</sub> in striatum compared to healthy controls. The striatal PDE10A binding potential was found to correlate with the prefrontal cortical thickness (Bodén et al. 2017), and a combined PET/MR study further demonstrated that PDE10A levels correlate with striatal function in schizophrenia (Persson et al. 2019).

Considerable loss (about 8% per decade) of PDE10A binding potential in striatum but less (about 4%) in the globus pallidus as a function of age was found together with a trend of higher binding potential in women compared to man (Fazio et al. 2017b). In studies comparing PDE10A binding in patients and healthy controls, it is therefore important to assure age-matching. PDE10A BP<sub>ND</sub> correlated with D<sub>2/3</sub> receptor BP<sub>ND</sub> in the striatum but not in the globus pallidus (Whiteley et al. 2019).

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### 23.3 Conclusions

Phosphodiesterases are an intriguing family of intracellular enzymes of which several subtypes show high expression in the brain. Not only expression of PDEs can be altered, but also their affinity for inhibitors can be allosterically modulated. This modulation can be quite different in vivo compared to in vitro, e.g., because of post-mortem degradation of allosteric cAMP and/or cGMP or because of dephosphorylation of PDE4. Successful PET radioligands were developed and translated to clinical application for visualization of PDE2A, PDE4, and PDE10A in the brain. Especially for the latter two subtypes, many (pre)clinical studies have been performed to investigate their role in different CNS diseases. It can be expected that novel PET radioligands for so far unexplored PDEs (e.g., PDE1 and PDE7) will be available in the future to enable new clinical studies looking at phosphodiesterase function and interplay in brain in health and disease.

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# PET Imaging of Purinergic Receptors

# 24

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## Abstract

Over the recent years, interest in the purinergic signaling system has sparked in the field of positron emission tomography (PET). Purinergic receptors play key roles in physiological and pathological processes, although the exact role in these processes is not always fully understood. PET provides an excellent imaging technique to study (patho)physiological processes in the living brain, and therefore radiopharmaceuticals (or tracers) targeting different purinergic receptors have been developed recently. This chapter will discuss the state of the art of PET tracer development for the P2 family of purinergic receptors, which is especially of interest for visualization of neuroinflammation, more specifically microglial activation.

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## 24.1 Introduction

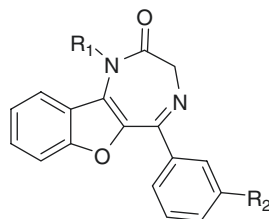
The involvement of purinergic signaling in the modulation of many physiological processes is more and more recognized, although the exact role of some of the purinergic receptors is still not fully understood. Physiological processes involving these receptors include (neuro)inflammation, immune cell recruitment, neurotransmission, and pain perception (Woods et al. 2016). The two families of purinergic receptors are classified in P1 and P2 receptors, natural ligands mainly being adenosine and adenosine 5'-triphosphate (ATP), respectively (Woods et al. 2016). As adenosine (P1) receptor imaging is already described in Chap. 8 (Elmenhorst et al) of this book, this chapter will focus on purinergic P2 receptor imaging. P2 receptors are subdivided in P2X and P2Y receptor families, where the seven P2X receptors (P2X<sub>1-7</sub>) are ATP-gated cation channels and the eight P2Y receptors (P2Y<sub>1</sub>, P2Y<sub>2</sub>, P2Y<sub>4</sub>, P2Y<sub>6</sub>, P2Y<sub>11-14</sub>) are G-protein coupled receptors (Woods et al. 2016; Burnstock 2016). Upon damage to the cell membrane, e.g., caused by pathological insults, the extracellular concentration of ATP significantly increases, activating P2X receptors, while P2Y receptors are activated by ATP but also by adenosine 5'-diphosphate (ADP), uridine triphosphate, uridine diphosphate (UDP), and UDP-glucose (Calovi et al. 2019). In the central nervous system (CNS), ATP degradation and signaling is mainly regulated by microglia, the resident immune cells in the CNS. More specifically, microglia status determines ATP release and degradation and influences microglia behavior, ranging from restoration of cell viability to induction of cell death (Calovi et al. 2019). Hence, tracers targeting purinergic P2 receptors are predominantly developed for imaging of the neuroinflammatory response in CNS pathologies. The most significantly expressed P2 receptors on microglia are P2X<sub>4</sub>, P2X<sub>7</sub>, P2Y<sub>6</sub>, P2Y<sub>12</sub>, and P2Y<sub>13</sub>. Expression of the different receptors varies and is related to factors like biological function, sex, and age (Calovi et al. 2019). This chapter will discuss the P2 targeting tracers developed to date, which are limited to ligands for receptors P2X<sub>4</sub>, P2X<sub>7</sub>, P2Y<sub>1</sub>, and P2Y<sub>12</sub>.

## 24.2 The P2X<sub>4</sub> Receptor

Although the precise role and mechanism of activation of the P2X<sub>4</sub> receptor (P2X<sub>4</sub>R) are largely unknown (Suurväli et al. 2017), it likely plays a role in cellular processes under physiological conditions, since activation of the receptor only requires low concentrations (nM) of extracellular ATP (Calovi et al. 2019), unlike, e.g., the P2X<sub>7</sub> receptor (P2X<sub>7</sub>R). Nevertheless, the current hypothesis is that P2X<sub>4</sub>R primes microglia toward a pro-inflammatory state associated with neuropathic pain and brain injury (Calovi et al. 2019; Suurväli et al. 2017), making it a potential target of interest for PET imaging. The only tracers targeting P2X<sub>4</sub>R to date have been developed by Wang et al., and all are analogues of 5-(3-bromophenyl)-1,3-dihydro-2H-benzofuro[3,2-*e*][1,4]diazepin-2-one (5-BDBD, IC<sub>50</sub> = 1.2 μM (Balázs et al. 2013)) (Wang et al. 2017). Some of the analogues were successfully radiolabeled (Fig. 24.1); however, in vivo evaluation of these ligands has not been reported thus



**Fig. 24.1** Radiolabeled 5-BDBD analogues targeting the P2X<sub>4</sub> receptor



[<sup>11</sup>C]8c R<sub>1</sub> = <sup>11</sup>CH<sub>3</sub>, R<sub>2</sub> = Br

[<sup>11</sup>C]9 R<sub>1</sub> = <sup>11</sup>CH<sub>3</sub>, R<sub>2</sub> = OH

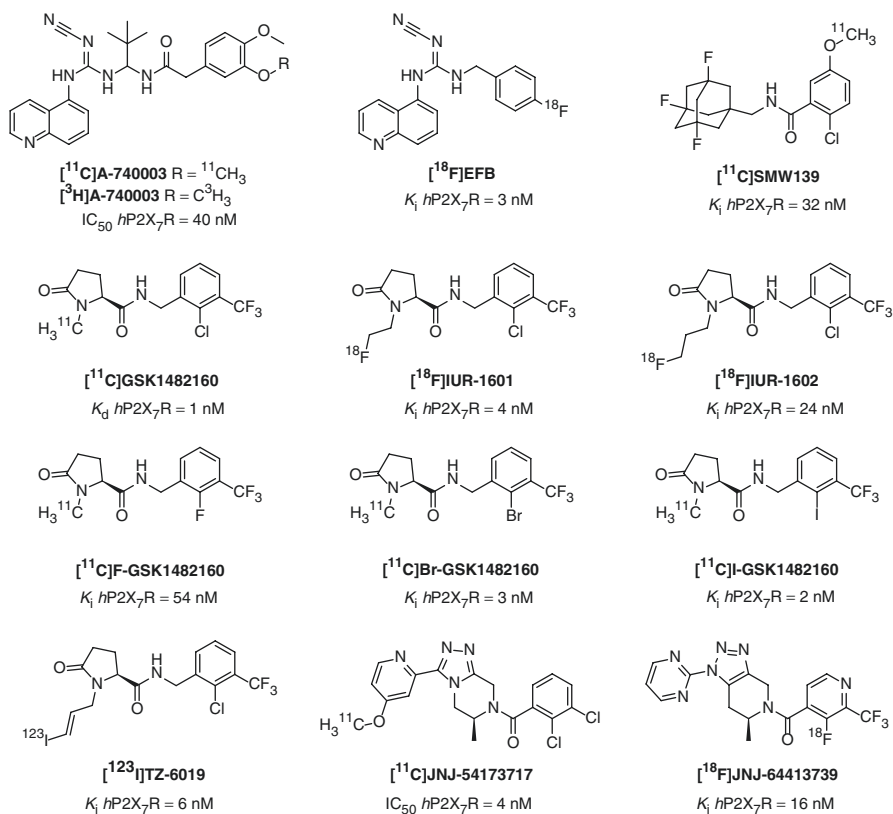
[<sup>18</sup>F]5a R<sub>1</sub> = H, R<sub>2</sub> = <sup>18</sup>F

[<sup>18</sup>F]11 R<sub>1</sub> = H, R<sub>2</sub> = OCH<sub>2</sub>CH<sub>2</sub><sup>18</sup>F

far, nor have in vitro experiments in tissue of animal models of neuroinflammation or postmortem tissue of human patients. All analogues displayed similar affinity toward P2X<sub>4</sub>R as the lead compound 5-BDBD, and so it remains questionable if this affinity is high enough for in vivo imaging of P2X<sub>4</sub>R.

### 24.3 The P2X<sub>7</sub> Receptor

The most studied target in the field of purinergic signaling and thus, not surprisingly, also in PET imaging to date is P2X<sub>7</sub>R. P2X<sub>7</sub>R is expressed in cells of the myeloid lineage, and in CNS P2X<sub>7</sub>R is mostly expressed on microglia cells, but receptor expression has also been shown in other brain cells like astrocytes and sometimes neurons (Bartlett et al. 2014). P2X<sub>7</sub>R is regarded silent in normal physiology, since high (mM) concentrations of its natural agonist ATP are required for activation and functional upregulation of the receptor (Bhattacharya and Biber 2016). The activation of P2X<sub>7</sub>R is associated with multiple signaling pathways, including the activation of the inflammasome leading to release of pro-inflammatory cytokines and reactive oxygen species, and P2X<sub>7</sub>R is therefore strongly linked to neuroinflammation (Bartlett et al. 2013, 2014; Bhattacharya and Biber 2016). Due to this association with the pro-inflammatory activation of microglia, the P2X<sub>7</sub>R is of interest for both PET imaging and drug development in CNS disease. This is exemplified by the fact that quite a number of PET tracers targeting P2X<sub>7</sub>R have been developed in recent years (Fig. 24.2). Two of the reported tracers, [<sup>11</sup>C]A-740003 and [<sup>18</sup>F]EFB, both bearing a cyanoguanidine moiety, have limited application in CNS disease due to poor brain uptake (Fantoni et al. 2017; Janssen et al. 2014). However, by using [<sup>3</sup>H]A-740003 in in vitro autoradiography experiments on brains of rats suffering from experimental autoimmune encephalomyelitis (EAE, an animal model of multiple sclerosis (MS), as well as on human postmortem brain tissue of MS patients, tracking neuroinflammation by targeting P2X<sub>7</sub>R was shown to be feasible, since higher tracer binding was observed at the peak of the disease (EAE)



**Fig. 24.2** Tracers targeting the P2X<sub>7</sub> receptor

and in active MS lesions in human tissue (Beaino et al. 2017). The results in the EAE rat model were further confirmed by a different group using  $[^{11}\text{C}]\text{GSK1482160}$ , with which increased tracer binding was observed in lumbar spinal cord of EAE rats at the peak of the disease (12–14 days post immunization) (Han et al. 2017). Unfortunately, due to too low affinity of  $[^{11}\text{C}]\text{GSK1482160}$  for rat P2X<sub>7</sub>R, this effect could not be observed in vivo. Using the same tracer, Territo et al. demonstrated a 3.6-fold higher uptake in the brain of mice treated with lipopolysaccharide (LPS) 72 h prior to PET scanning, compared with saline-treated animals (Territo et al. 2017). Interestingly, other groups reported peak expression of P2X<sub>7</sub>R and Iba1 12 and 24 h after treatment with LPS (Fantoni et al. 2017; Berdyeva et al. 2019; Choi et al. 2007), a variation that is likely due to high variability observed in LPS-induced neuroinflammation (Catorce and Gevorkian 2016). In addition to the promising results in rodent neuroinflammation models,  $[^{11}\text{C}]\text{GSK1482160}$  also entered brain of nonhuman primates with a maximum standard uptake value (SUV) of 2.7 at 70 min postinjection (Han et al. 2017). Likewise, first-in-human results showed a brain uptake of 2% of the total injected dose (SUV 1.0) in healthy volunteers at

20 min postinjection (Green et al. 2018). Despite the observed high liver uptake in human, clearance of [ $^{11}\text{C}$ ]GSK1482160 seems suited for further clinical evaluation. Attempts to synthesize two radiofluorinated analogues of GSK1482160 ([ $^{18}\text{F}$ ]IUR-1601 (Gao et al. 2018) and [ $^{18}\text{F}$ ]IUR-1602 (Gao et al. 2019a)) were so far unsuccessful; both compounds were found to be unstable; therefore radiochemical yields were too low to enable preclinical evaluation of these analogues. Three more carbon-11-labeled halogenated analogues of GSK1482160 have been developed, but these have not been biologically evaluated yet (Gao et al. 2019b). An iodinated analogue of GSK1482160, [ $^{123}\text{I}$ ]TZ-6019, showed significantly higher binding in the brain of a P301S mouse model of Alzheimer's disease (AD) compared with wild-type mouse brain (1.3-, 1.7- and 2.4-fold increase in the whole brain, hippocampus, and cortex, respectively) (Jin et al. 2018).

A study using the adamantane benzamide [ $^{11}\text{C}$ ]SMW139 showed that tracer binding in postmortem brain tissue of AD patients and healthy controls was not significantly different (Janssen et al. 2018). However, in a rat model locally over-expressing human P2X<sub>7</sub>R unilaterally in striatum (Ory et al. 2016), binding of [ $^{11}\text{C}$ ]SMW139 to P2X<sub>7</sub>R could be demonstrated in vivo (Janssen et al. 2018). In addition, first in vivo results in the EAE rat model using [ $^{11}\text{C}$ ]SMW139 look promising (Beaino et al. 2019), as significantly increased tracer uptake was observed at the peak of the disease, similar to previous in vitro results with [ $^3\text{H}$ ]A-740003 and [ $^{11}\text{C}$ ]GSK1482160 (Beaino et al. 2017; Han et al. 2017). Very recently in a first-in-human study in a small set of MS patients and healthy subjects, [ $^{11}\text{C}$ ]SMW139 showed good pharmacokinetics and quantifiable uptake in the brain (Hagens et al. 2020). Additionally, for all brain regions investigated,  $V_T$  and  $\text{BP}_{\text{ND}}$  were elevated in MS patients compared with healthy controls, with only  $V_T$  differences reaching statistical significance (Hagens et al. 2020). The previously mentioned rat model of local human P2X<sub>7</sub>R expression has also been used to evaluate the imaging potential of [ $^{11}\text{C}$ ]JNJ-54173717, and specific uptake of this tracer in the P2X<sub>7</sub>R-expressing striatum was confirmed (Ory et al. 2016). An in vitro autoradiography study with the same tracer showed a time-dependent increase in tracer binding in the brain of rats treated with 6-hydroxydopamine (6-OHDA), an acute model of Parkinson's disease (PD), with highest P2X<sub>7</sub>R expression 14 days after 6-OHDA injection, which co-localized with Iba1 immunohistochemical staining (Crabbé et al. 2019). On the contrary, in the same study no differences in P2X<sub>7</sub>R binding between the different time points were observed in a chronic A53T viral vector model of PD. [ $^{11}\text{C}$ ]JNJ-54173717 readily crossed the blood-brain barrier in nonhuman primates (Ory et al. 2016), and in a first-in-human study, the tracer was found to be suitable for quantification of P2X<sub>7</sub>R in human brain (Van Weehaeghe et al. 2019a). However, no differences in brain uptake of the tracer were observed between PD patients and healthy volunteers (Van Weehaeghe et al. 2019a), which is in line with the findings in the chronic animal model of PD. In addition, a possible genotype effect (polymorphism rs3751143) on tracer binding was detected, which should be further explored. Similar to the results in PD patients, no difference was observed between healthy

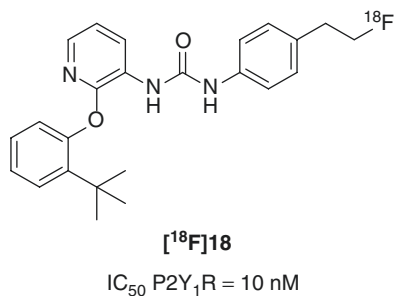
volunteers and patients with amyotrophic lateral sclerosis (ALS) (Van Weehaeghe et al. 2019b). The lack of difference between volunteers and patients in both studies may be due to time dependence of P2X<sub>7</sub>R expression, as it might be most prominent in early stages or acute neuroinflammation, since microglial activation is driven by P2X<sub>7</sub>R activation. Another Janssen R&D compound, [<sup>18</sup>F]JNJ-64413739, showed a significant increase in P2X<sub>7</sub>R binding in the ipsilateral, LPS-treated side of rat brains 2 days posttreatment compared with the contralateral hemisphere, both in vivo with PET and ex vivo with autoradiography (Berdyeva et al. 2019). P2X<sub>7</sub>R binding levels also correlated with Iba1 immunohistochemical staining. Interestingly, this increase was not observed in in vitro autoradiography with [<sup>18</sup>F]JNJ-64413739, which might be indicative of a difference in binding mechanism in vitro and in vivo (Berdyeva et al. 2019). Brain uptake of [<sup>18</sup>F]JNJ-64413739 peaked in nonhuman primates within 10 min postinjection, and a relatively fast metabolic rate (40% intact tracer at 30 min postinjection) but no defluorination was observed (Kolb et al. 2019). A first-in-human study in healthy volunteers showed limited bias, good test-retest variability and excellent reliability with a dynamic 90 min acquisition (Koole et al. 2018). No appropriate reference region could be determined in both nonhuman primates and humans (Kolb et al. 2019; Koole et al. 2018). In human subjects, high interindividual signal variability was observed, which is possibly due to a binding effect of a P2X<sub>7</sub>R polymorphism (Koole et al. 2018), as was also observed for [<sup>11</sup>C]JNJ-54173717 (Van Weehaeghe et al. 2019a).

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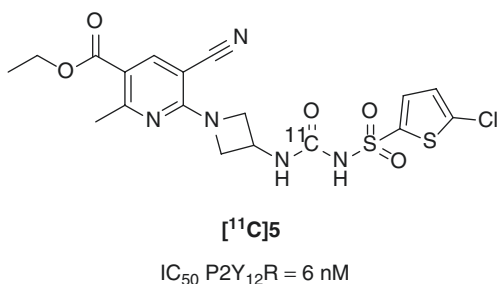
## 24.4 The P2Y<sub>1</sub> Receptor

There is limited knowledge about the expression of the P2Y<sub>1</sub> receptor (P2Y<sub>1</sub>R) in normal and diseased brain; however, alterations in its activity or expression are suggested to play a pathophysiological role in, e.g., neurodegeneration, neuroprotection, and neuroregeneration (Moldovan et al. 2019). P2Y<sub>1</sub>R binds preferentially ADP over ATP and was shown to be involved in microglial migration together with the P2Y<sub>12</sub> receptor (P2Y<sub>12</sub>R) (De Simone et al. 2010). Only a limited number of non-nucleotidic compounds with high affinity for P2Y<sub>1</sub>R have been reported, one of which is a highly selective P2Y<sub>1</sub>R antagonist developed by Bristol-Myers Squibb ( $K_i = 6$  nM) (Chao et al. 2013). In order to make an analogue amenable to radiolabeling, Moldovan et al. pursued the synthesis of fluoro-substituted analogues (Moldovan et al. 2019). In vitro evaluation of their small compound library led to the selection of compound **18** (Fig. 24.3,  $IC_{50} = 10$  nM vs.  $IC_{50} = 180$  nM for the lead compound in the same assay). Metabolism of [<sup>18</sup>F]**18** was studied in CD-1 mice at 30 min postinjection, and only minor amounts of intact tracer were observed, with the main metabolite (85%) being [<sup>18</sup>F]fluoride. The rather unexpected but extensive defluorination of [<sup>18</sup>F]**18** hampers any future use in vivo, and the authors are currently focusing on more metabolically stable analogues of selective P2Y<sub>1</sub>R antagonist **18** (Moldovan et al. 2019).

**Fig. 24.3** P2Y<sub>1</sub> receptor tracer



**Fig. 24.4** Non-brain penetrant P2Y<sub>12</sub> receptor tracer



## 24.5 The P2Y<sub>12</sub> Receptor

P2Y<sub>12</sub>R is involved in chemotaxis and directs the microglia to move toward higher ATP concentrations, e.g., to a site of injury. P2Y<sub>12</sub>R is exclusively expressed by microglia in the CNS and is used as a microglial marker; however, its expression is downregulated in pro-inflammatory conditions (Calovi et al. 2019; Haynes et al. 2006). One potential PET tracer has been developed to date ([<sup>11</sup>C]5, IC<sub>50</sub> = 6 nM (Bach et al. 2013), Fig. 24.4), but unfortunately it did not cross the blood-brain barrier (Villa et al. 2018) and therefore has not been evaluated in vivo in neuroinflammation models. Villa et al. did show that while P2Y<sub>12</sub>R is downregulated in pro-inflammatory microglia, in contrast it is highly upregulated in the anti-inflammatory microglial phenotype (Villa et al. 2018). In addition, [<sup>11</sup>C]5 could be used in in vitro autoradiography studies using brain tissues of both a mouse model of stroke and postmortem human brain tissue of a patient deceased after stroke. Indeed, decreased tracer binding was observed in both human and mouse affected tissue compared with the non-affected hemispheres, and tracer binding corresponded to immunohistochemical staining for P2Y<sub>12</sub>R (Villa et al. 2018). Similarly, in an EAE rat model and human brain tissue of MS patients, [<sup>11</sup>C]5 showed downregulation of P2Y<sub>12</sub>R at the peak of the disease in EAE rats and in human MS lesions (Beaino et al. 2017). In addition, in the EAE rat model, P2Y<sub>12</sub>R expression came back up to baseline levels in the EAE recovery phase and was also shown to be complementary to expression of P2X<sub>7</sub>R, a pro-inflammatory microglial marker. These results show

that P2Y<sub>12</sub>R could be a promising target for PET imaging of the anti-inflammatory microglia, provided that a brain penetrant P2Y<sub>12</sub>R ligand can be identified.

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## 24.6 Concluding Remarks

The interest in purinergic P2 receptor imaging has only relatively recently emerged, and thus there is still much to gain in this field. Thus far, P2X<sub>7</sub>R has been the most investigated target and has been identified as a viable target for imaging of pro-inflammatory microglia. First-in-human studies have been conducted; however, no significant differences in tracer uptake have been found between PD or ALS patients and healthy volunteers. In contrast, albeit only in a small set of subjects, higher tracer uptake was demonstrated in MS patients as compared with healthy volunteers. Taken together, these first-in-human results warrant additional clinical evaluation for P2X<sub>7</sub>R imaging, especially since animal studies in both chronic and acute PD models show that P2X<sub>7</sub>R expression is highly time-dependent and may peak in human disease in earlier stages as well, compared with receptor expression in chronic disease. Currently the only purinergic target under investigation for imaging of the anti-inflammatory microglial phenotype is P2Y<sub>12</sub>R, and, once a brain penetrant tracer can be identified, such a tracer could be used as a complementary method to pro-inflammatory microglia imaging (e.g., TSPO, P2X<sub>4</sub>R, P2X<sub>7</sub>R) to study the dynamics of microglial activation *in vivo*. This could be of incredible importance in a deeper understanding of neurological diseases involving neuroinflammation, as well as be a meaningful aid in drug development and follow-up of anti-inflammatory treatment strategies.

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# Imaging of the Serotonin System: Radiotracers and Applications in Memory Disorders

# 25

Gitte Moos Knudsen and Steen G. Hasselbalch

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**Abstract**

The serotonergic system plays a key modulatory role in the brain and is the target for many drug treatments for brain disorders either through reuptake blockade or via interactions at the 14 subtypes of serotonin (5-HT) receptors. This chapter provides the current status of radioligands used for positron emission tomography (PET) and single-photon emission computerised tomography (SPECT) imaging of the human brain 5-HT receptors and the 5-HT transporter (SERT) with particular emphasis on the applications in Alzheimer's disease (AD).

Currently available radioligands for in vivo brain imaging of the 5-HT system in humans include radiolabelled compounds for the 5-HT<sub>1A</sub>, 5-HT<sub>1B</sub>, 5-HT<sub>2A</sub>, 5-HT<sub>4</sub> and to some extent 5-HT<sub>6</sub> receptors and for SERT. Imaging of serotonergic targets in humans has given invaluable insight into the normal brain function and into brain disorders where the serotonergic system is perturbed.

Imaging studies show that the 5-HT<sub>1A</sub> receptor binding is increased, and 5-HT<sub>2A</sub> receptor binding is decreased in mild cognitive impairment (MCI). In early AD, 5-HT<sub>4</sub> receptor binding is increased, whereas in early and more advanced AD, SERT and the 5-HT<sub>1A</sub> and 5-HT<sub>2A</sub> receptor binding are reduced in a region-specific manner. Future studies should focus on the association between serotonergic dysfunction and symptomatology in order to increase our understanding of the neurobiological background for neuropsychiatric symptoms in neurodegenerative and neuropsychiatric disorders.

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## 25.1 Introduction

The serotonergic (5-HT) system plays a key modulatory role in many brain functions. The serotonin transporter and 14 serotonergic receptor subtypes result in a complex pattern of modulatory control over various physiological, emotional and cognitive processes. These include, e.g. mood, sleep, diurnal rhythms, cognition, learning, memory and appetite. Serotonergic dysfunction has been implicated in the aetiology of many psychiatric and neurological disorders, e.g. affective disorders, anxiety, schizophrenia, Alzheimer's disease, migraine and epilepsy. The development of in vivo brain imaging techniques, such as positron emission tomography (PET) and single-photon emission computerised tomography (SPECT), increasingly allows the study of the serotonergic system in the human brain; for review, see Paterson et al. (2013) and Kumar and Mann (2014).

This review covers the current status of which PET and SPECT radioligands are available for imaging serotonergic targets within the brain. Secondly, as an example of one of the applications of PET imaging, the current knowledge about disturbances of the serotonergic system in Alzheimer's disease is given.

### 25.1.1 5-HT Targets for PET and SPECT

The 5-HT receptors are amongst the most diverse group of neurotransmitter receptors in the human genome, and the 5-HT system is also one of the phylogenetically oldest systems. Currently, 14 structurally and pharmacologically distinct mammalian 5-HT receptor subtypes have been described. Based on their structure, affinity for different ligands and second messenger pathway, they are assigned to one of seven families, 5-HT<sub>1-7</sub> (Hoyer et al. 2002). All 5-HT receptors, except the 5-HT<sub>3</sub> receptor, are G-protein-coupled seven transmembrane-spanning receptors (GPCRs). The 5-HT<sub>3</sub> receptor is a ligand-gated sodium ion channel. In addition, the 5-HT transporters (SERTs) responsible for 5-HT reuptake and 5-HT synthetic enzymes, especially tryptophan hydroxylase, are also targets for tracer development. Significant discoveries of the 5-HT system in the human brain have been made following the development of selective PET and SPECT radioligands; some examples are given in (Jones and Rabiner 2012; Rebolz et al. 2018; Spies et al. 2015).

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## 25.2 Current Radioligands for In Vivo Brain Imaging of the 5-HT System

A multitude of radioligands exist for in vitro studies of serotonergic targets, and over the last decade, we have seen an impressive increase in the number of useful PET and SPECT radioligands. Based on PET neuroimaging studies, a high-resolution human brain atlas of the 5-HT<sub>1A</sub>, 5-HT<sub>1B</sub>, 5-HT<sub>2A</sub> and 5-HT<sub>4</sub> receptors, and the 5-HT transporter has recently been made available (Beliveau et al. 2017), downloadable from <https://nru.dk/index.php/misc/category/90-nru-serotonin-atlas>. Published PET and SPECT radioligands for imaging the serotonergic system are extensively reviewed in (Paterson et al. 2013), but here, we will only summarise some of the to date most utilised radiotracers.

### 25.2.1 5-HT<sub>1A</sub> Receptor

The 5-HT<sub>1A</sub> receptor is one of the best characterised receptors in the serotonergic family; its role as an inhibitory autoreceptor in the raphe nuclei and the possible implications of this role for the treatment of depression and anxiety with serotonin reuptake inhibitors are well-known (King et al. 2008).

Many of the 5-HT<sub>1A</sub> radioligands were based on WAY-100635 (*N*-[2-[4-(2-methoxyphenyl)piperazin-1-yl]ethyl]-*N*-pyridin-2-ylcyclohexanecarboxamide), which in its *carbonyl*-<sup>11</sup>C-labelled form is widely used for 5-HT<sub>1A</sub> receptor imaging. Currently, four radioligands are used for PET studies of the 5-HT<sub>1A</sub> receptor in humans: [*carbonyl*-<sup>11</sup>C]WAY-100635, [<sup>18</sup>F]MPPF, [<sup>18</sup>F]FCWAY and [<sup>11</sup>C]CUMI-101. [*carbonyl*-<sup>11</sup>C]WAY-100635 (Pike et al. 1995) is so far the most widely used 5-HT<sub>1A</sub> receptor radioligand. It has a high target-to-background ratio,

but its fast systemic metabolism makes it difficult to quantify accurately. [ $^{18}\text{F}$ ]MPPF has the advantage of the longer lived  $^{18}\text{F}$ -label, and it also selectively labels the 5-HT<sub>1A</sub> receptors with a low non-specific binding. Its major disadvantage is its low brain uptake. [ $^{18}\text{F}$ ]FCWAY is rarely used, probably because of issues with defluorination of the parent compound which leads to high bone uptake of radioactivity (Ryu et al. 2007). [ $^{11}\text{C}$ ]CUMI-101 is a high-affinity partial agonist 5-HT<sub>1A</sub> radioligand that displays high specific binding and seems suitable for imaging the high-affinity site within the human brain (Milak et al. 2008; Pinborg et al. 2012). More recently, however, it has been found that [ $^{11}\text{C}$ ]CUMI-101 has significant binding to  $\alpha$ 1-adrenoceptors in humans (Shrestha et al. 2014, 2016)

### 25.2.2 5-HT<sub>1B</sub> Receptor

The 5-HT<sub>1B</sub> receptor is of particular interest in relation to obesity (Halford et al. 2007) and migraine (Tfelt-Hansen 2012). No less than two PET radiotracers for imaging the 5-HT<sub>1B</sub> receptor are now available for use in humans: [ $^{11}\text{C}$ ]AZ10419369 and [ $^{11}\text{C}$ ]P943. [ $^{11}\text{C}$ ]AZ10419369 (5-methyl-8-(4-[ $^{11}\text{C}$ ]methyl-piperazin-1-yl)-4-oxo-4*H*-chromene-2-carboxylic acid(4-morpholin-4-yl-phenyl)-amide) is a 5-HT<sub>1B</sub> partial agonist used in humans (Varnas et al. 2011); it has a slow systemic metabolism. The high-affinity 5-HT<sub>1B</sub> antagonist radioligand, [ $^{11}\text{C}$ ]P943 (*R*-1-[4-(2-methoxy-isopropyl)-phenyl]-3-[2-(4-methyl-piperazin-1-yl)benzyl]-pyrrolidin-2-one), also shows good properties for quantification in the human brain (Gallezot et al. 2010). For both radiotracers, the cerebellum constitutes an acceptable reference region, and there is evidence that the radiotracers are sensitive to displacement by endogenous 5-HT in non-human primates (Finnema et al. 2010; Ridler et al. 2011) and in pigs (Jørgensen et al. 2018a, b). Normative data from healthy individuals show an 8% decline in 5-HT<sub>1B</sub> receptor binding per decade (Savli et al. 2012), but no gender-related differences have been reported (Matuskey et al. 2012).

### 25.2.3 5-HT<sub>2A</sub> Receptor

5-HT<sub>2A</sub> receptors are of interest for many reasons: they are a primary target of psychedelic compounds (Lee and Roth 2012), contribute to the efficacy of antipsychotic medications and are involved in the aetiology or treatment of various psychiatric disorders (Leysen 2004).

Several different radioligands for imaging the brain 5-HT<sub>2A</sub> receptor have successfully been used in human PET studies, e.g. [ $^{18}\text{F}$ ]altanserin; [ $^{18}\text{F}$ ]setoperone; [ $^{11}\text{C}$ ]NMSP; [ $^{11}\text{C}$ ]MDL 100,907; and [ $^{11}\text{C}$ ]Cimbi-36. In addition, the SPECT tracer [ $^{123}\text{I}$ ]-R91150 is used in imaging studies, but the radiotracer displays a lower signal-to-noise ratio compared to the available PET radioligands.

Despite its lipophilic radiometabolite, [ $^{18}\text{F}$ ]altanserin has been widely used as a PET radiotracer. Because of its longer half-life, the  $^{18}\text{F}$ -labelling facilitates the

application of a bolus/infusion paradigm that in turn enables subtraction of the lipophilic brain metabolite (Pinborg et al. 2003). Imaging data obtained from [ $^{18}\text{F}$ ]altanserin binding in the human brain are highly reproducible (Haugbol et al. 2007). [ $^{18}\text{F}$ ]altanserin was used to determine changes in 5-HT<sub>2A</sub> receptor density in relation to ageing (Adams et al. 2004), and it has been reported that binding in healthy subjects correlates with the body mass index (Erritzoe et al. 2009) but does not vary with gender (Frokjaer et al. 2009). Furthermore, twin studies have shown that [ $^{18}\text{F}$ ]altanserin binding is strongly genetically determined (Pinborg et al. 2008).

[ $^{11}\text{C}$ ]NMSP (*N*-methylspiperone, 8-[4-(4-fluorophenyl)-4-oxobutyl]-2-methyl-4-phenyl-2,4,8-triazaspiro[4.5]decan-1-one) is a dual D<sub>2</sub>/5-HT<sub>2</sub> receptor ligand (Wong et al. 1984). As [ $^{18}\text{F}$ ]setoperone, NMSP has high affinity for both receptors, but since the density of D<sub>2</sub> receptors is low and that of 5-HT<sub>2</sub> receptors is high in the cortical brain regions, then the majority of specific binding in the neocortex is due to 5-HT<sub>2</sub> receptor binding. For subcortical brain regions, the reverse is true (Lyon et al. 1986). Because of its non-selectivity, [ $^{11}\text{C}$ ]NMSP is no longer in common use. The same applies for the radioligand [ $^{18}\text{F}$ ]setoperone (6-[2-[4-(4-[ $^{18}\text{F}$ ]fluorobenzoyl)piperidin-1-yl]ethyl]-7-methyl-2,3-dihydro-[1,3]thiazolo[3,2-a]pyrimidin-5-one).

The 5-HT<sub>2A</sub> receptor-selective tracer 11C-MDL100907 has been characterised in humans (Talbot et al. 2012). Based on *in vitro* data (Kristiansen et al. 2005), [ $^{11}\text{C}$ ]MDL 100907 is more selective for the 5-HT<sub>2A</sub> receptor than [ $^{18}\text{F}$ ]altanserin, but it is questionable if this has any practical implications because of the scarcity of 5-HT<sub>2B</sub> and to some extent 5-HT<sub>2C</sub> receptors. Another promising radioligand is [ $^{11}\text{C}$ ]Cimbi-36 (Ettrup et al. 2014a, b) which is so far the only 5-HT<sub>2A</sub> receptor agonist radioligand that has proven successful in humans. [ $^{11}\text{C}$ ]Cimbi-36 has excellent test-retest reproducibility properties, and head-to-head comparisons with [ $^{18}\text{F}$ ]altanserin suggest that in brain regions with high 5-HT<sub>2C</sub> receptor density, choroid plexus and hippocampus, [ $^{11}\text{C}$ ]Cimbi-36 likely represents binding to both 5-HT<sub>2A</sub> and 5-HT<sub>2C</sub> receptors (Ettrup et al. 2016). In the pig brain, cerebral 5-HT release correlates with [ $^{11}\text{C}$ ]Cimbi-36 PET measures of 5-HT<sub>2A</sub> receptor occupancy, and likewise, [ $^{11}\text{C}$ ]Cimbi-36 binding is reduced in response to a fenfluramine challenge in non-human primates (Yang et al. 2017). Acute SSRI intervention in human leads to variable changes in the brain [ $^{11}\text{C}$ ]Cimbi-36 binding (da Cunha-Bang et al. 2019), possibly due to interindividual variability of the SSRI 5-HT release response. Preliminary data suggest that a stronger 5-HT challenge, such as dexamphetamine, is associated with a reduction in the brain [ $^{11}\text{C}$ ]Cimbi-36 binding (Erritzoe, *in prep*).

### 25.2.4 5-HT<sub>2B</sub> and 5-HT<sub>2C</sub> Receptors

To date, it is questionable if 5-HT<sub>2B</sub> receptors are expressed in the brain in sufficient amounts to allow for imaging. This is not the case for 5-HT<sub>2C</sub> receptors, but currently, all radiolabelled 5-HT<sub>2C</sub> receptor ligands have shared pharmacology with other receptors. No radiotracers have been developed for SPECT or PET imaging of 5-HT<sub>2B</sub> or 5-HT<sub>2C</sub> receptors.

### 25.2.5 5-HT<sub>3</sub> Receptors

Despite a number of research centres undertaking a concerted effort to develop 5-HT<sub>3</sub>-selective PET and SPECT tracers, it seems that the very discrete localisation and relatively low levels of 5-HT<sub>3</sub> receptors that are localised with highest densities in the brainstem (Parker et al. 1996) make it a very difficult target to image in vivo.

### 25.2.6 5-HT<sub>4</sub> Receptors

5-HT<sub>4</sub> receptors are involved in learning and memory and are potential targets for the treatment of Alzheimer's disease (for a review, see Bockaert et al. (2004)). So far, one PET ligand, [<sup>11</sup>C]SB207145 (8-amino-7-chloro-(*N*-[<sup>11</sup>C]methyl-4-piperidylmethyl)-1,4-benzodioxan-5-carboxylate), has been successfully evaluated and used in humans.

The 5-HT<sub>4</sub> receptor antagonist SB207145 was initially radiolabelled with C-11 (Gee et al. 2008) and evaluated for its potential as a PET radioligand for 5-HT<sub>4</sub> imaging. [<sup>11</sup>C]SB207145 was subsequently successfully quantified for use in human brain studies (Marner et al. 2009). In this study, a comprehensive quantification of the binding of [<sup>11</sup>C]SB207145 to cerebral 5-HT<sub>4</sub> receptors in the human brain in vivo was further provided. Distribution volumes and binding potentials of [<sup>11</sup>C]SB207145 showed good test-retest reproducibility and time stability. The blocking study with piboserod confirmed that the cerebellum is a suitable reference region devoid of specific binding and that reference tissue models apply. Subsequently, it was shown that [<sup>11</sup>C]SB207145 is not to any significant degree displaceable by acutely increased levels of endogenous 5-HT (Marner et al. 2010), but cautions need to be taken to ensure that the injected mass of SB207145 does not exceed 4.5 µg (Madsen et al. 2011c). Clinical data (Fisher et al. 2015a, b; Haahr et al. 2014) suggest, however, that [<sup>11</sup>C]SB207145 serves as a proxy for changes in brain 5-HT levels and this feature needs to be considered when interpreting clinical studies. Normative data on age- and sex-related variations have been published (Madsen et al. 2011b).

### 25.2.7 5-HT<sub>5</sub> Receptors

The 5-HT<sub>5</sub> receptor has two subtypes, the 5-HT<sub>5A</sub> and the 5-HT<sub>5B</sub> receptors. The 5-HT<sub>5A</sub> receptor has been identified in the human brain, but the 5-HT<sub>5B</sub> receptor is not expressed in humans because the coding sequence is interrupted by stop codons (Nelson 2004). The 5-HT<sub>5A</sub> receptor shows a particularly high presence in raphe and other brainstem and pons nuclei (Volk et al. 2010). There are no available radioligands for either of the 5-HT<sub>5</sub> receptors.

### 25.2.8 5-HT<sub>6</sub> Receptors

5-HT<sub>6</sub> receptors are found exclusively in the CNS and are predominantly expressed in the striatum, limbic system and cortex (Woolley et al. 2004). They are of particular interest because of their involvement in learning and memory (King et al. 2008), and the receptor shows a significant age-related decline in healthy subjects, primarily in basal ganglia (Radhakrishnan et al. 2018). So far, only one non-selective radioligand for PET imaging of the 5-HT<sub>6</sub> receptor has made its way into humans, namely, [<sup>11</sup>C]GSK215083 (3-[(3-fluorophenyl)sulfonyl]-8-(4-[<sup>11</sup>C]methyl-1-piperazinyl)quinoline (Parker et al. 2012). GSK215083 has high affinity for 5-HT<sub>6</sub> (in vitro  $K_i$ , 0.16 nM) but also has high 5-HT<sub>2A</sub> affinity (in vitro  $K_i$ , 0.79 nM). However, the differential localisation of 5-HT<sub>2A</sub> and 5-HT<sub>6</sub> receptors (predominantly cortical and striatal, respectively) combined with the ~fivefold difference in affinity means that discrimination between these two receptor types can be done to some extent.

### 25.2.9 5-HT<sub>7</sub> Receptors

In spite of many attempts to develop 5-HT<sub>7</sub>-selective PET and SPECT tracers, none of these have yet made their way into human use. The so far most promising radiotracer is [<sup>11</sup>C]Cimbi-717 which in pigs displayed the most promising results for selective 5-HT<sub>7</sub> receptor PET imaging (Hansen et al. 2014), because it is displaceable with 5-HT<sub>7</sub> receptor selective compounds. See also review by L'Estrade et al. (2020).

### 25.2.10 SERT

SERT is the serotonin transporter, and it receives considerable interest, not the least due to the success of its inhibitors in the treatment of depression and anxiety disorders. Several PET and SPECT ligands have been developed for this purpose, and a detailed review of SERT imaging by PET and SPECT can be found in Huang et al. (2010).

Initially, images of SERT in the human brain came from the non-selective cocaine derivative SPECT ligand [<sup>123</sup>I]β-CIT and later with the selective but kinetically irreversible PET ligand [<sup>11</sup>C](+)McN5652. Today, the most successful line of SERT radioligands is that developed from diarylsulfides such as [<sup>123</sup>I]ADAM and especially [<sup>11</sup>C]MADAM and [<sup>11</sup>C]DASB.

The so far best SPECT radioligand for SERT imaging is [<sup>123</sup>I]ADAM (2-((2-((dimethylamino)methyl)phenyl)thio)-5-iodophenylamine), which is potent and selective and has a high target-to-background ratio in human studies (Newberg et al. 2004). Quantification of the SERT binding with [<sup>123</sup>I]ADAM SPECT is most often done with a ratio method, based on data acquired from 200 to 240 min. This has, however, been shown to overestimate the specific binding by about 10%. The



most reliable outcome is based on a 0–120 min SPECT acquisition followed by Logan non-invasive modelling (Frokjaer et al. 2008).

The first selective PET radioligand for imaging SERT in the human brain was [ $^{11}\text{C}$ ](+) $\text{McN5652}$  (*trans*-1,2,3,5,6,10- $\beta$ -hexahydro-6-[4-(methylthio)phenyl]pyrrolo-[2,1-*a*]isoquinoline) (Szabo et al. 1995). Its use has been limited by a relatively low target-to-background ratio in vivo, as well as a slow brain uptake and irreversible kinetics that complicate quantification in high-binding regions (Frankle et al. 2004).

[ $^{11}\text{C}$ ]DASB (3-amino-4-(2-dimethylamino-methyl-phenylsulfanyl)-benzonitrile) is one of a series of  $^{11}\text{C}$ -labelled arylthiobenzylamines developed by Wilson and Houle (1999). [ $^{11}\text{C}$ ]DASB displays good selectivity and high affinity and is today the most widely used radiotracer for in vivo imaging of SERT. [ $^{11}\text{C}$ ]DASB PET can be quantified with reference tissue models (Ginovart et al. 2001); this is the most frequently used way of quantifying [ $^{11}\text{C}$ ]DASB. Furthermore, test-retest data show high reproducibility and reliability (Kim et al. 2006). [ $^{11}\text{C}$ ]DASB PET normative data are available (Erritzoe et al. 2009, 2010), and apart from a multitude of patient studies, it has also been used to measure SERT occupancy at clinical doses of selective reuptake inhibitors, e.g. citalopram and paroxetine (Meyer et al. 2004).

[ $^{11}\text{C}$ ]MADAM (*N,N*-dimethyl-2-(2-amino-4-methylphenylthio)-benzylamine) is yet another PET radioligand that has made its way into humans (Lundberg et al. 2005). When done several weeks apart, test-retest reproducibility of [ $^{11}\text{C}$ ]MADAM is excellent (Lundberg et al. 2006). As for [ $^{11}\text{C}$ ]DASB, [ $^{11}\text{C}$ ]MADAM has been used to estimate relative SERT occupancy of citalopram and escitalopram (Lundberg et al. 2007) and to investigate the relationship between 5-HT<sub>1A</sub> and SERT binding in healthy young men and women (Henningsson et al. 2009; Jovanovic et al. 2008). [ $^{11}\text{C}$ ]DASB and [ $^{11}\text{C}$ ]MADAM seem comparable in terms of their pharmacological and kinetic profile; the majority of clinical studies have, however, been done with [ $^{11}\text{C}$ ]DASB.

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### 25.3 PET Imaging of the Serotonergic System in Alzheimer's Disease

As an example of the application of PET imaging in neurology and psychiatry, studies of the serotonergic system in Alzheimer's disease (AD) will be reviewed in this section. With increasing incidence and prevalence worldwide, AD presents a unique challenge to researchers in order to provide better diagnostic tools, as well as a better understanding of the pathophysiology of the disease. Biomarkers such as medial temporal lobe atrophy on magnetic resonance imaging, glucose metabolism measured with PET or beta-amyloid and tau in cerebrospinal fluid become increasingly incorporated into the diagnostic process (Dubois et al. 2010; Jack Jr et al. 2018). Therefore, the focus of neuroreceptor imaging with PET has changed in the direction of better understanding the pathophysiology behind the symptomatology of AD (Xu et al. 2012). The ultimate goal of this research is better and more specific treatment options.

Postmortem and clinical studies have shown widespread dysfunction of the serotonergic transmitter system in AD. Thus, postmortem brain studies have consistently found significant loss of serotonin-producing neurons or reductions in the presynaptically located serotonin transporter (SERT) in the raphe nuclei (Aletrino et al. 1992; Halliday et al. 1992; Hendricksen et al. 2004; Tejani-Butt et al. 1995) and in serotonergic neuronal projections (Tejani-Butt et al. 1995; Thomas et al. 2006; Tsang et al. 2003). Also, in several postmortem studies, pre- and postsynaptic receptor subtypes have been found reduced: 5-HT<sub>1A</sub> (Becker et al. 2014; Lai et al. 2003b; Vidal et al. 2016), 5-HT<sub>1B</sub> (Garcia-Alloza et al. 2004), 5-HT<sub>2A</sub> (Bowen et al. 1989; Cheng et al. 1991; Lai et al. 2005; Lorke et al. 2006) and 5-HT<sub>6</sub> receptors (Garcia-Alloza et al. 2004; Lorke et al. 2006). Since serotonergic dysfunction and neuropsychiatric symptoms have been linked to mood disorders, disturbances in the serotonergic system could be related to the presence of neuropsychiatric symptoms in AD. This would be especially relevant for depressive symptoms, which are frequent in AD (Ballard et al. 1996; Lyketsos et al. 2002). Serotonin may, however, also play a role in normal cognitive functions (for review, see Schmitt and co-workers (Schmitt et al. 2006). More recently, the involvement of serotonin in cognition has further been substantiated by studies in healthy young subjects, showing an association between high SERT binding in fronto-striatal regions and better performance on tasks involving executive function and logical reasoning (Madsen et al. 2011a) and an inverse association between 5-HT<sub>4</sub> receptor binding in the hippocampus and memory performance (Haahr et al. 2013). As 5-HT<sub>4</sub> receptor binding is suggested to be a biomarker of serotonergic tone in human (Haahr et al. 2014), low levels of central 5-HT could partly be responsible for not only neuropsychiatric symptomatology but also cognitive impairment in patients with AD.

### 25.3.1 5-HT<sub>1A</sub> Receptor Binding in AD

Using [<sup>18</sup>F]MPPF, Kepe and co-workers found significant reductions in 5-HT<sub>1A</sub> receptor densities in both the hippocampus and raphe nuclei in patients with mild cognitive impairment (MCI) (24%) and AD patients (49%) (Kepe et al. 2006). They interpreted their findings as secondary to neocortical degeneration of synapses and neurons, because a positive association of 5-HT<sub>1A</sub> receptor binding with glucose metabolism was found in the hippocampus. Further, 5-HT<sub>1A</sub> receptor binding was negatively correlated with [<sup>18</sup>F]FDDNP, a marker of tau pathology (Kepe et al. 2006). Interestingly, using the same tracer, Truchot and co-workers found a biphasic change in 5-HT<sub>1A</sub> receptor binding in the hippocampus, parahippocampus and inferior temporal cortex with upregulation in MCI and a marked reduction (approx. 50%) in AD (Truchot et al. 2007) which may suggest a compensatory upregulation due to lower serotonin levels in the pre-dementia stage of AD. Following this upregulation, 5-HT<sub>1A</sub> receptor binding may eventually decrease in early-to-moderate stages of AD due to neurodegeneration, as suggested by Kepe et al. (2006).

### 25.3.2 5-HT<sub>2A</sub> Receptor Binding in AD

Several studies have found widespread reduction in 5-HT<sub>2A</sub> receptor binding in mild-to-moderate AD. Using PET and [<sup>18</sup>F]setoperone in patients with AD, Blin and colleagues found a 35–69% reduction in temporoparietal cortical 5-HT<sub>2</sub> receptor binding (Blin et al. 1993). In a similar patient group (mean MMSE 20) but using [<sup>18</sup>F]altanserin, Meltzer and colleagues found reductions of approximately 30% in the anterior cingulate, prefrontal and sensorimotor cortices (Meltzer et al. 1999). In two consecutive [<sup>18</sup>F]altanserin studies in MCI and AD patients, these findings were corroborated by widespread neocortical reductions in receptor binding (MCI: 22–29% and AD: 28–39%) (Hasselbalch et al. 2008; Marner et al. 2011). Importantly, all three [<sup>18</sup>F]altanserin studies employed correction for partial volume error due to atrophy, and atrophy did not explain the findings.

In contrast to the 5-HT<sub>1A</sub> receptor findings above, the reductions in 5-HT<sub>2A</sub> receptor are widespread and occur early in the course of the disease (Hasselbalch et al. 2008), but the reason for this reduction is largely unexplained. It is unlikely that loss of serotonergic neurons projecting to neocortex has any major impact on the post-synaptic 5-HT<sub>2A</sub> receptor reductions, since reductions in 5-HT<sub>2A</sub> receptors are not accompanied by similar reductions in presynaptic serotonin transporter binding (Marner et al. 2012). One plausible explanation for the diffuse neocortical 5-HT<sub>2A</sub> receptor reductions could be that local AD pathology, especially in the form of widespread beta-amyloid accumulation, is responsible for the reduction in 5-HT<sub>2A</sub> receptors. Beta-amyloid accumulation has a spatial distribution similar to the reduction in 5-HT<sub>2A</sub> receptor binding, whereas neurodegeneration in the form of tau accumulation follows a different pattern (Braak and Braak 1991). As an indirect support of the proposed beta-amyloid/5HT<sub>2A</sub> receptor association, Marner and co-workers found a nonsignificant 5–10% reduction in <sup>18</sup>F-altanserin binding in neocortical regions in a 2-year follow-up study of 15 MCI patients (Marner et al. 2011). This reduction inversely mirrors the small increase in beta-amyloid accumulation found in early AD in most studies (Jack Jr et al. 2010). Further, experimental studies have shown an association between increased beta-amyloid load and decreased 5-HT<sub>2A</sub> receptor binding (Christensen et al. 2008; Holm et al. 2010). Thus, 5-HT<sub>2A</sub> receptor may be especially sensitive to beta-amyloid pathology, but the nature of this interaction needs to be clarified in future studies.

### 25.3.3 5-HT<sub>4</sub> Receptor Binding in AD

The 5-HT<sub>4</sub> receptor is especially interesting in AD as it is both linked to memory function and beta-amyloid accumulation. There is ample evidence for pro-cognitive actions of agonists to the 5-HT<sub>4</sub> receptor. Thus, in animals, pre-task systemic injections of 5-HT<sub>4</sub> receptor agonists have shown to improve performance in a variety of memory tasks such as olfactory associative learning (Marchetti et al. 2000; Marchetti-Gauthier et al. 1997), place and object recognition (Lamirault and Simon 2001), the Morris water maze (Lelong et al. 2001) and matching to sample (Terry Jr et al. 1998). In the only study in healthy humans, using [<sup>11</sup>C]SB207145 and Rey's

Auditory Verbal Learning Test in healthy young subjects, Haahr and co-workers found significant negative associations between the immediate and delayed recall scores and hippocampal 5-HT<sub>4</sub> receptor binding (Haahr et al. 2013). This paradoxical finding was explained by upregulation of 5-HT<sub>4</sub> receptor levels to partially compensate for lower hippocampal 5-HT levels in subjects with poorer memory function (Haahr et al. 2013). This explanation has been corroborated by findings of a reduction in 5-HT<sub>4</sub> receptor binding following a 3-week administration of a selective 5-HT reuptake inhibitor (Haahr et al. 2014). In early AD, Madsen and co-workers found some support for this hypothesis: In mild stage AD patients, beta-amyloid imaging with [<sup>11</sup>C]PiB and 5-HT<sub>4</sub> receptor imaging using [<sup>11</sup>C]SB207145 was performed in the same subjects (Madsen et al. 2011d). PiB-positive individuals had 13% higher 5-HT<sub>4</sub> receptor binding compared to PiB-negative individuals. The 5-HT<sub>4</sub> receptor binding was positively correlated to beta-amyloid burden and negatively to the MMSE score (Mini-Mental State Examination—a measure of global cognitive function) of the AD patients. These findings suggest that cerebral 5-HT<sub>4</sub> receptor upregulation starts at a preclinical stage of AD and continues, while dementia is still at a mild stage, which contrasts other receptor subtypes. As in the study of healthy subjects mentioned above, it was speculated that the upregulation was a compensatory effect of decreased levels of interstitial 5-HT in early AD (Madsen et al. 2011d). Further, agonism of the 5-HT<sub>4</sub> receptor increases alpha-secretase activity and thus promotes non-amyloidogenic degradation of the amyloid precursor protein (APP) (Cachard-Chastel et al. 2007). Therefore, an upregulation of the 5-HT<sub>4</sub> receptor level in early AD may serve to counteract beta-amyloid accumulation, but further studies are needed to elucidate these rather speculative interactions.

### 25.3.4 5-HT<sub>6</sub> Receptor Binding in AD

The 5-HT<sub>6</sub> receptor has been implicated in memory functions, as 5-HT<sub>6</sub> antagonism can reverse scopolamine-induced episodic memory dysfunction (de Bruin et al. 2011), although the effect may be strongest for working memory and aversive learning (Da Silva et al. 2012). The pro-cognitive effects of 5-HT<sub>6</sub> antagonism have been investigated in clinical trials of mild-to-moderate AD but have failed to show clinical effects (Maher-Edwards et al. 2010; Atri et al. 2018). In the postmortem AD brains, reductions in 5-HT<sub>6</sub> binding have been found in frontal as well as temporal cortical areas (Garcia-Alloza et al. 2004; Lorke et al. 2006). As described above, a PET tracer non-selective for the 5-HT<sub>6</sub> receptor is being used, but so far, no studies examining patients with AD have been published.

### 25.3.5 SERT Binding in AD

As described above, several studies in the postmortem AD brains have shown significant losses of serotonin-producing neurons or reductions in the presynaptically located SERT in the raphe nuclei and in serotonergic neuronal projections (Aletrino et al. 1992; Halliday et al. 1992; Hendricksen et al. 2004; Tejani-Butt et al. 1995;

Thomas et al. 2006; Tsang et al. 2003). Three studies have investigated the SERT binding in AD in vivo by PET in AD. Ouchi and co-workers used [ $^{11}\text{C}$ ]DASB in 15 mild to moderately demented AD patients, of whom 7 also had depressive symptoms (Ouchi et al. 2009). They found that [ $^{11}\text{C}$ ]DASB binding in the dense SERT regions (the midbrain, including the raphe nuclei, nucleus accumbens, putamen, and thalamus) of the depressed AD group was significantly lower than that in an age-matched control group (Ouchi et al. 2009). In the midbrain, the reduction amounted to 33%. In the nondepressed group, a significant SERT reduction was found only in the putamen. In contrast, in 12 mild-to-moderate AD patients, Marner and co-workers did not find reductions in [ $^{11}\text{C}$ ]DASB binding in the raphe nuclei (Marner et al. 2012). In this study, SERT binding was reduced in mesial temporal lobe only, whereas 5-HT<sub>2A</sub> receptor binding measured using  $^{18}\text{F}$ -altanserin was globally reduced by 28–39% in the same subjects. Finally, in 28 MCI subjects, Smith et al. found widespread reductions in SERT binding in cortical and limbic regions, as well as in the raphe nuclei, which were associated with both verbal and visual memory scores (Smith et al. 2017). The discrepancies between these studies may arise from methodological differences and also from patient selection, and all three studies document early disruption of serotonergic transmission in AD. In conclusion, in end-stage AD, postmortem studies clearly show degeneration of serotonergic neurons in the raphe and in serotonergic neuronal projections, and PET studies corroborate these findings with serotonergic neurons and projections being affected even in early stages of the disease.

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## 25.4 Can Serotonergic Dysfunction Explain AD Symptomatology?

As described above, dysfunction of the serotonergic transmitter system is well-documented in AD. The important question is whether this dysfunction explains part of the symptomatology of the disorder. In postmortem studies of AD, low densities of neocortical SERT and 5-HT<sub>1A</sub> receptors have been associated with depression and aggression, respectively (Chen et al. 1996; Lai et al. 2003a). Also, Garcia-Alloza et al. found associations between cognition and frontal cortex 5-HT<sub>1B</sub> binding and between behavioural symptoms and temporal 5-HT<sub>6</sub> binding (Garcia-Alloza et al. 2004). Postmortem studies are associated with problems regarding time interval between symptom evaluation and receptor binding analyses, and also, they represent an end stage of the disease. But in vivo studies have so far not been able to give a clear answer to the question. Although Kepe et al. (2006) demonstrated a correlation between hippocampal 5-HT<sub>1A</sub> receptor binding and MMSE, they interpreted this association as secondary to the well-known correlation between neurodegeneration and cognitive function. No correlation between MMSE score and 5-HT<sub>2A</sub> receptor binding was found in two studies of MCI and AD patients (Meltzer et al. 1999; Hasselbalch et al. 2008). However, these studies included rather small sample sizes, and also, the limited range of cognitive dysfunction may have made it difficult to determine correlations between serotonergic and cognitive dysfunction. In a SPECT study of AD patients

(Versijpt et al. 2003), a correlation between MMSE scores and 5-HT<sub>2A</sub> receptor binding in the left orbitofrontal region was found, but the relevance of this finding is not clear. Patients with significant neuropsychiatric symptoms can be very difficult to include in PET studies, and generally, patients covering the spectrum from mild to severe neuropsychiatric symptoms are lacking in *in vivo* studies. Meltzer and co-workers found no difference in 5-HT<sub>2A</sub> receptor binding in AD subjects with depression compared to AD subjects without depression. Further, in the depressed patients, there was no correlation between 5-HT<sub>2A</sub> receptor binding and Hamilton depression scale score (Meltzer et al. 1999). Likewise, an inverse correlation between neuropsychiatric symptoms scored by the Neuropsychiatric Inventory (NPI) and striatal 5-HT<sub>2A</sub> receptor binding observed in MCI (Hasselbalch et al. 2008; Marner et al. 2011) could not be corroborated in with the same methodology in a sample of AD patients (Marner et al. 2012). Similarly, SERT findings have been controversial with regard to neuropsychiatric symptoms: In a postmortem study of depressive versus nondepressive AD patients, Hendricksen et al. (Hendricksen et al. 2004) found no differences in raphe pathology. In contrast, Ouchi and co-workers found significant correlations between SERT high-binding regions (the midbrain, striatum and thalamus) and geriatric depression scores when grouping depressed and nondepressed AD patients (Ouchi et al. 2009). However, Marner and co-workers did not find significant associations between measures of depression and SERT binding in AD patients, but in this study, patients had less cognitive and neuropsychiatric symptoms (Marner et al. 2012) as compared to Ouchi et al. (2009). Thus, postmortem and *in vivo* imaging data are only consistent when it comes to the association between reduced SERT expression in the midbrain and depressive symptoms in AD (Ouchi et al. 2009; Chen et al. 1996). In the study by Smith et al. (Smith et al. 2017), SERT binding was positively correlated with performance on tests of visual and verbal memory and to a greater extent than a reduction in regional cerebral blood flow and grey matter volume. These findings may suggest a specific role of 5-HT transmission in cognitive dysfunction in AD, but future *in vivo* studies need to include larger samples of patients with a spectrum of cognitive and neuropsychiatric symptoms. Even when doing so, the complex neuropathology and the interaction between several affected transmitter systems may make the aim of establishing associations between serotonin receptor subtypes and symptomatology too simplistic. However, once the pathophysiology of AD symptoms is further elucidated, *in vivo* imaging studies may help us to determine available drug targets.

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# Monoamine Oxidase A and Serotonin Transporter Imaging with Positron Emission Tomography

# 26

Jeffrey H. Meyer

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## Abstract

Serotonin has a major role in neural circuits of mood regulation, substance abuse, and neurodevelopment. Although they differ in cellular and anatomical location, the two brain proteins most strongly implicated in influencing serotonin levels are monoamine oxidase A and the serotonin transporter. The advances in positron emission tomography (PET) radioligand development to image these two

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proteins have been applied considerably over the past 15 years, greatly increasing our knowledge about fundamental processes in many important, impactful conditions including major depressive disorder, early postpartum, perimenopause, cigarette smoking, alcohol dependence, aggressive behavior, antidepressant development, ecstasy abuse, obsessive compulsive disorder, and seasonal affective disorder as reviewed in this chapter.

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## 26.1 Why Image Indices of Monoamine Oxidase A Density?

Monoamine oxidase A (MAO-A) is an enzyme that metabolizes three major monoamines (serotonin, norepinephrine, dopamine) in the brain and has a role in creating oxidative stress and predisposition to apoptosis (Youdim et al. 2006). Monoamine levels influence mood (Freis 1954; Ruhe et al. 2007), and oxidative stress and apoptosis influence brain development (Bortolato et al. 2008). Hence MAO-A is a logical target for investigations in mood disorders, substance abuse (during which markers of dysregulated monoamine levels occur (Stockmeier 2003; Meyer 2008)), and aggression for which abnormal neurodevelopment is implicated (Buckholtz and Meyer-Lindenberg 2008). Illnesses and conditions for which dysregulation of monoamine levels is implicated include major depressive disorder, substance abuse, neurodegenerative diseases, and aggression (Youdim et al. 2006; Meyer et al. 2006; Fowler et al. 1996; Bacher et al. 2011). MAO-A is predominantly located on outer mitochondria membranes in neurons (as well as astrocytes and glia) and has a high density in brain regions that influence mood (Saura et al. 1992) although the cellular distribution varies by region (Saura et al. 1992, 1996; Finberg 2012). MAO-A density is highest in the brainstem (within the locus coeruleus); moderately high in the cortex, hippocampus, and striatum; lower in cerebellar cortex; and minimal in white matter tissue (Saura et al. 1992, 1996).

The density of MAO-A correlates highly with the level of its metabolic activity (Saura et al. 1992; Nelson et al. 1979); hence, neuroimaging of measures of MAO-A binding can be interpreted as having functional implications (Saura et al. 1992). This approach represents an optimal practical method for applying positron emission tomography (PET) neuroimaging toward a strongly functional index of MAO-A activity. Another imaging approach would be to measure a radiotracer which is metabolized by MAO-A and trapped to some extent in the brain, but this is extremely difficult because the potential brain radiotracer would require kinetics in which the dominant, separable, measurable effect is its metabolism by MAO-A, rather than its crossing of the blood-brain barrier or impairment of removal of trapped radiometabolites. Most radiotracers that assess metabolism accumulate as a product in the

brain which can be a disadvantageous strategy in radiotracer development because such radiotracers are less likely to exhibit reversible time activity curves, and irreversible time activity curves are usually subject to bias from blood flow. Hence, the most optimal, practical neuroimaging approach toward quantitating MAO-A function with PET or SPECT is radioligands that bind to MAO-A as reversible inhibitors.

## 26.2 Radioligands Available for Neuroimaging Monoamine Oxidase A

There are three PET radiotracers that have been applied for measurement of available monoamine oxidase A in a human study, and a fourth promising radiotracer is in development (see Table 26.1). The first radiotracer developed [ $^{11}\text{C}$ ]clorgyline, demonstrated adequate brain penetration and selectivity, but did not have reversible time activity curves (Fowler et al. 1987). A deuterium-labeled version of [ $^{11}\text{C}$ ]clorgyline was created to attempt to improve reversibility, but it is also not optimally reversible (Fowler et al. 1996). More recently [ $^{11}\text{C}$ ]befloxatone and [ $^{11}\text{C}$ ]harmine were developed. [ $^{11}\text{C}$ ]befloxatone shows promising radiotracer qualities of selectivity and reversibility (Bottlaender et al. 2010; Curet et al. 1998; Dolle et al. 2003) and has been modeled in humans (Bottlaender et al. 2010; Curet et al. 1998; Dolle et al. 2003; Zanotti-Fregonara et al. 2014, 2013). However, test-retest reliability in humans is not known, and it hasn't been studied under blocking conditions in humans.

[ $^{11}\text{C}$ ]Harmine demonstrates the properties of an excellent PET radiotracer for an index of available MAO-A density in humans. Harmine has high affinity for the MAO-A enzyme, ( $K_i = 2 \text{ nM}$ ), has high brain uptake in humans, and is selective for MAO-A (Meyer et al. 2006; Bergstrom et al. 1997a; Ginovart et al. 2006): The affinity of [ $^{11}\text{C}$ ]harmine is three orders of magnitude higher for MAO-A than MAO-B (Bergstrom et al. 1997a). Displacement studies in baboons using MAO-A selective inhibitors show complete displacement of [ $^{11}\text{C}$ ]harmine in regions with high density of MAO-A (Bergstrom et al. 1997b). The uptake is specific to the parent compound as metabolites of harmine are polar and do not cross the blood-brain barrier (Tweedie and Burke 1987). The uptake of [ $^{11}\text{C}$ ]harmine is highest in brain regions with greater MAO-A density such as the cortex and lowest in regions with low MAO-A density such as white matter (Bergstrom et al. 1997c). The MAO-A  $V_T$  (MAO-A distribution volume) is primarily an index of specifically bound [ $^{11}\text{C}$ ]harmine at equilibrium since 80 to 85% represents specific binding and the levels of free and nonspecific binding are similar among different people (Ginovart et al. 2006; Sacher et al. 2011). The modeling of this compound in humans best fits a two tissue compartment model, and negligible bias is found when the Logan model is applied (Ginovart et al. 2006).



**Table 26.1** Comparison of PET radiotracers for monoamine oxidase A (from Meyer (2014))

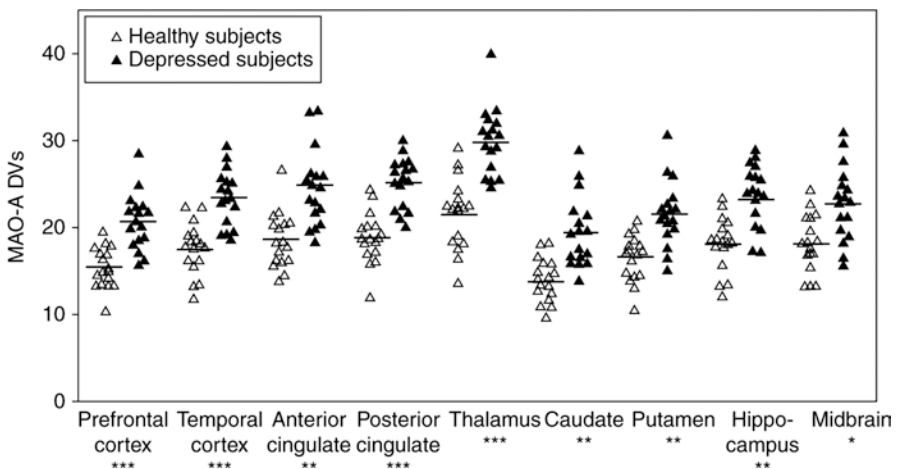
	[ <sup>11</sup> C]Clorgyline	[ <sup>11</sup> C]Harmine	[ <sup>11</sup> C]Befloxadone	[ <sup>18</sup> F] Fluoroethyl-harmol
Selectivity	Excellent (Fowler et al. 1987)	Excellent (Bergstrom et al. 1997a)	Excellent (Dolle et al. 2003)	Excellent (Schieferstein et al. 2015)
Reversibility	Not reversible (Fowler et al. 1987)	Highly reversible (Ginovart et al. 2006)	Highly reversible (Bottlaender et al. 2010)	Moderate in rodent (Schieferstein et al. 2015; Maschauer et al. 2015)
Brain uptake	Very good (Fowler et al. 1987)	High (Ginovart et al. 2006)	High (Zanotti-Fregonara et al. 2013)	Very good (Schieferstein et al. 2015; Maschauer et al. 2015)
Modeling	2 tissue compartment (Fowler et al. 1987)	2 tissue compartment (Ginovart et al. 2006)	2 tissue compartment (Zanotti-Fregonara et al. 2013)	Not yet completed in humans
Specific binding to free and nonspecific binding ratio	Very good but limited assessment (Fowler et al. 1987)	High ~4 (Ginovart et al. 2006; Chiuccariello et al. 2015)	High in baboon (Bottlaender et al. 2010) Unclear in human (Zanotti-Fregonara et al. 2014, 2013)	High in rodent (Schieferstein et al. 2015; Maschauer et al. 2015)
Reliability in humans	Very good (Fowler et al. 1996)	Excellent (Sacher et al. 2012)	Not reported	Not yet completed
Brain-penetrant radioactive metabolites?	Unlikely	No brain-penetrant metabolites (Wilson et al. 2003)	Unlikely	No brain-penetrant metabolites (Maschauer et al. 2015)
Measureable in diverse regions?	Gray matter (Fowler et al. 1987)	Gray matter (Ginovart et al. 2006)	Gray matter (Zanotti-Fregonara et al. 2013)	Gray matter in rodent (Schieferstein et al. 2015; Maschauer et al. 2015)

## 26.3 Major Depressive Disorder

The in vivo aspect of neuroimaging offers the ability to readily sample medication-free subjects and to selectively choose early-onset major depressive disorder (MDD) which is the most common form (World\_Health\_Organization 2008). Late-onset

MDD is often attributable to neurodegenerative diseases such as Parkinson's disease and Alzheimer's disease (Krishnan 2002), which would be expected to have very different pathologies from early-onset MDD. Since postmortem studies often preferentially sample older subjects, sometimes they are not targeted toward early-onset MDD. The first three postmortem studies of MAO-A levels and activity had not selectively targeted early-onset MDD, and the early perspective of the field was that there was no abnormality of MAO-A level or activity in MDD (Mann and Stanley 1984; Ordway et al. 1999; Sherif et al. 1991).

In 2006 MAO-A  $V_s$ , an index of MAO-A density was measured using [ $^{11}\text{C}$ ] harmine PET in medication-free major depressive episodes (MDE) secondary to early-onset MDD (Meyer et al. 2006). Subjects with MDE were drug-free for at least 5 months, and most were antidepressant naive. All MDE subjects and controls were otherwise healthy. The MAO-A  $V_s$  was highly significantly elevated ( $p < 0.001$  each region, average magnitude 34% (or two standard deviations)) during MDE (see Fig. 26.1). This was the first definitive study of MAO-A binding in MDE from early-onset MDD: The samples excluded comorbidity, thereby focusing on the differences between this common, early-onset type of MDD and health. The effect size was large and the radiotracer has outstanding qualities for measuring MAO-A binding. In 2008, Barton et al. reported a consistent finding of elevated brain serotonin turnover in unmedicated depressed patients (Barton et al. 2008). In 2009, the finding of greater MAO-A binding in MDE was replicated with [ $^{11}\text{C}$ ]harmine PET, and in 2011 the finding was replicated in antidepressant-free MDE subjects in postmortem study of orbitofrontal cortex applying Western blot (Meyer et al. 2009a; Johnson et al. 2011). The replication finding of [ $^{11}\text{C}$ ]harmine PET was extended to 42 MDE subjects and 37 healthy controls, demonstrating the greatest levels of MAO-A in those with greater severity and/or reversed neurovegetative symptoms (hypersomnia, hyperphagia, or weight gain); an interesting finding since MDE with these



**Fig. 26.1** On average, MAO-A  $DV_s$  was elevated by 34% or 2 SDs in depressed individuals. The hippocampal region also samples the parahippocampus. Differences between groups were highly statistically significant in each region. \* $p < 1 \times 10^{-5}$ ; † $p < 1 \times 10^{-4}$ ; ‡ $p = 0.001$

particular symptoms is well known to be more likely to respond to MAO inhibitors (Chiucciariello et al. 2014). It was further demonstrated in a comparison among MDE with comorbid severe borderline personality disorder, MDE with moderate borderline personality disorder, and the healthy that MAO- $V_T$  was considerably elevated, particularly in the prefrontal and anterior cingulate cortex in the most severe group (by 43% and 42%, respectively) but also to a lesser extent in the moderate group (by 23 and 21%, respectively). Interestingly, in the overall group of patients in this study, there was also a positive correlation between current suicidality and prefrontal cortex (PFC) MAO-A  $V_T$  (Kolla et al. 2016).

Greater MAO-A binding in affect-modulating regions has important implications for understanding the pathophysiology of MDD. A key longstanding theory concerning MDD is that monoamines were low, but no clear mechanism of monoamine loss had been previously identified, and this finding represents the clearest mechanism to support this theory (Meyer et al. 2006). In addition, since MAO-A has multiple functions, metabolizing serotonin, norepinephrine, and dopamine, oxidizing, and facilitating apoptosis (Youdim et al. 2006), it suggests that new antidepressants should optimally target these additional functions rather than solely raise extracellular serotonin. Such approaches have notable potential to address the issue that at least 40% of MDE does not respond to the common first-line treatment of selective serotonin reuptake inhibitors (Trivedi et al. 2006).

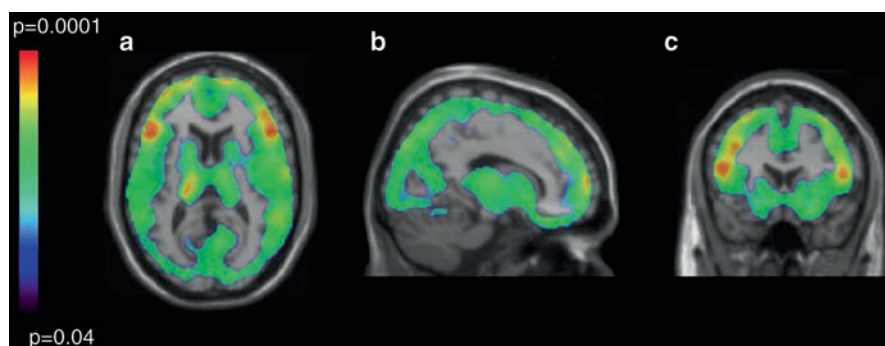
Investigations of MAO-A  $V_T$ , or monoamine oxidase distribution volume, were subsequently extended into different states of MDD, an approach which is readily feasible with positron emission tomography. While it might be expected that MAO-A levels would normalize with euthymic mood, the recovered state of MDD is also a state of high risk for another MDE. The risk for a recurrent MDE over 2 years is 20–50% depending upon treatment conditions (Frank et al. 1990; Fava et al. 2004). Elevated MAO-A  $V_T$  may be considered an index of a monoamine-lowering process, and in the 1950s during treatment with reserpine-based antihypertensives, it was discovered that chronic monoamine lowering is associated with subsequent onset of MDEs which typically occurred 2 weeks to 4 months later (Freis 1954). In recovered MDD, MAO-A  $V_T$  was significantly elevated in the PFC, anterior cingulate cortex (ACC), striatum, hippocampus, thalamus, and midbrain in a sample of 18 medication-free recovered MDD subjects compared to 28 healthy controls (Meyer et al. 2009a). Recovered MDD subjects who had recurrence of their MDE in the subsequent 6 months had the highest levels of MAO-A binding in the prefrontal and anterior cingulate cortex at the time of scanning (Meyer et al. 2009a). The prefrontal cortex and anterior cingulate cortex were of particular interest because these regions (and/or subregions of these structures) are activated in mood induction studies (Liotti et al. 2002) and during cognitive functions like pessimistic perspective that generate sad mood (Sharot et al. 2007; Tom et al. 2007). To address the possibility that other factors related to risk of recurrence did not contribute to the finding, subjects were medication-free for at least a year, had no cognitive behavioral therapy within 3 years, were currently asymptomatic, and had no comorbid medical, psychiatric, or substance abuse illnesses. Given the link between elevated MAO-A binding in prefrontal and anterior cingulate cortex and subsequent MDE, this argues that new interventions are needed to decrease MAO-A levels in these

brain regions even beyond the duration of a therapeutic intervention so as to prevent recurrence.

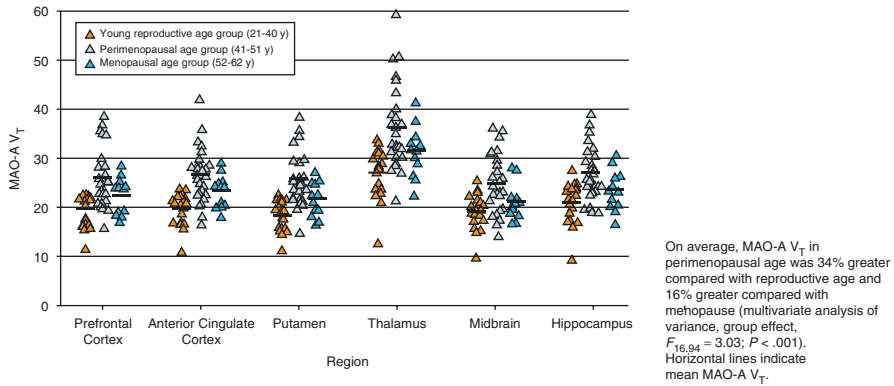
## 26.4 Early Postpartum and Perimenopause

Both early postpartum and perimenopause are high-risk points for MDE with a 13% prevalence for a MDE over the first 6 months postpartum and a 17% prevalence rate for new-onset MDE during perimenopause (reviewed (Mann and Stanley 1984; Ordway et al. 1999)). It is known that estrogen levels decline more than 100-fold over the first several days postpartum in humans (O'Hara et al. 1991), and changes in estrogen levels across the menstrual cycle heighten approximately fivefold during perimenopause (reviewed (Mann and Stanley 1984; Ordway et al. 1999)). While several studies reported an inverse relationship between decline in estrogen and rise in MAO-A levels, mRNA and activity (Luine et al. 1975; Ma et al. 1993, 1995; Holschneider et al. 1998), prior to PET imaging studies in 2010 and 2014, MAO-A binding, and mRNA or activity had never been studied in any species in early postpartum, nor during perimenopause.

A [<sup>11</sup>C]harmine PET study measured MAO-A binding during early postpartum (days 4–6) (Sacher et al. 2010). A highly significant elevation of MAO-A binding was found, which was, on average, 43% greater across the brain regions assayed (prefrontal cortex, anterior cingulate cortex, striatum, thalamus, hippocampus, mid-brain) as compared to women not recently pregnant (Sacher et al. 2010). A voxel-based analysis demonstrated that the elevation in MAO-A binding was present throughout the gray matter of the brain (see Fig. 26.2) (Sacher et al. 2010). A subsequent [<sup>11</sup>C]harmine PET study completed several years later demonstrated a 34% elevation in MAO-A  $V_T$  across gray matter regions sampled in perimenopause, including the PFC and ACC (Sacher et al. 2010; Rekkas et al. 2014) (see Fig. 26.3).



**Fig. 26.2** Parametric maps of elevated monoamine oxidase A binding in the postpartum group versus the control group. Maps are superimposed on a T1-weighted magnetic resonance image that is normalized to the T1-weighted template (SPM2; Department of Cognitive Neurology, Wellcome Trust Centre for Neuroimaging, London, England). (a) Transverse. (b) Sagittal. (c) Coronal. The individual voxel threshold was set at  $p < 0.05$ ; a total of 86,412 voxels comprised a single cluster, which had a cluster-corrected significance of  $p = 0.03$



**Fig. 26.3** Monoamine oxidase A total distribution volume in women of young reproductive age, women of perimenopausal age, and older women in menopause

These studies, which identify elevation in MAO-A  $V_T$  during high-risk states for MDE, have intriguing potential for developing strategies of dietary supplementation to counter the effect of elevated MAO-A level. For example, early postpartum is a time of greater risk for a couple of common, intertwined mood disturbances (O'Hara and Swain 1996). The first is the common, postpartum blues, present in up to 75% of women involving mild sadness, and can be accompanied by some trouble sleeping, irritability, desire for isolation, and difficulty concentrating (O'Hara and Swain 1996). It is considered within the healthy range of experience, typically starting on day 4 postpartum and finishing within the first week. The second is postpartum depression, which is a MDE that occurs within the first 4 weeks after giving birth, although some definitions include onset within the first 3 months after giving birth (O'Hara and Swain 1996; Association AP 1994). When postpartum blues is severe, the risk for MDE is fourfold greater. A dietary supplement strategy composed of blueberry antioxidants (which contain anthocyanins detectable in the brain of rodents), oral tryptophan given the evening of postpartum day 4, and tyrosine given the morning of day 5 was created to counter effects of elevated MAO-A level consisting of greater  $H_2O_2$  production, greater metabolism of serotonin, and greater metabolism of norepinephrine and dopamine. This dietary supplement was shown to prevent depressed mood during early postpartum, assessed with depressed mood induction, in an open trial with a robust effect size of 2.9 (Dowlati et al. 2017) The dietary strategy is also compatible with breastfeeding because tryptophan and tyrosine are already found in breast milk, so the overall concentration of these amino acids in breast milk is not affected by oral administration of these amino acids (Dowlati et al. 2015, 2014).

## 26.5 Cigarette Smoking

The first study of MAO-A binding applying [ $^{11}C$ ] clorgyline PET reported available MAO-A binding is reduced in those who smoke cigarettes in the active smoking state (Fowler et al. 1996). Given that the plasma half-life of the key MAO-A-binding

substances found in cigarette smoke (harman and norharman) is only an hour (Rommelspacher et al. 2002), there was reason to specifically assess MAO-A  $V_T$  during both active smoking and withdrawal conditions. When MAO-A  $V_T$  was assessed in both conditions, it was discovered that prefrontal and anterior cingulate cortex MAO-A  $V_T$  (as well as the other brain regions assessed) rose during withdrawal in those who smoke heavily, that is, more than one pack of cigarettes per day, but not in those who smoke more moderately at less than one pack per day and that during withdrawal from heavy smoking, MAO-A  $V_T$  was greater than in health in all regions assessed (Bacher et al. 2011).

The elevation in MAO-A  $V_T$  has important implications for the problem of sad mood in people who smoke cigarettes. There is a very high comorbidity between cigarette smoking and major depressive disorder with 50% of people with MDD also smoking cigarettes, and the pattern is such that cigarette smoking predisposes to MDD and vice versa (Anda et al. 1990; Breslau et al. 1998). It is the heavy cigarette smoking group who are at much greater risk for MDD (Pratt and Brody 2010), and this group had a 25% elevated MAO-A  $V_T$  in the prefrontal and anterior cingulate cortex during withdrawal as compared to healthy controls, arguing for a process of elevated MAO-A level during withdrawal as a mechanism to create risk for MDD (Bacher et al. 2011). Hence repeated exposure of elevated MAO-A level, in the prefrontal and anterior cingulate cortex, a mechanism associated with onset and maintenance of MDE state, occurring after a short period of 8 hours withdrawal could explain the predisposition to MDE in people who smoke cigarettes heavily.

A second problem with mood in people who smoke cigarettes is the depressed mood of acute withdrawal (Carey et al. 1993; Kenford et al. 2002). In the imaging study of MAO-A during cigarette withdrawal, in those who smoked heavily, the magnitude of rise in MAO-A  $V_T$  in prefrontal and anterior cingulate cortex during withdrawal was significantly correlated with the shift in visual analogue scales toward depressed mood (Bacher et al. 2011). This rise in MAO-A  $V_T$  also correlated with the decline in harman in those who smoke heavily. Harman is the chemical in cigarette smoke that has the highest affinity for MAO-A (Herraiz and Chaparro 2005; Hauptmann and Shih 2001). These results suggest that rapid removal of harman from occupying MAO-A sites leaves a high level of available MAO-A for metabolizing monoamines in prefrontal and anterior cingulate cortex resulting in depressed mood. This argues for testing of MAO-A inhibitor treatments in people who experience sad mood during early cigarette withdrawal as a strategy to assist in quitting, an important issue since 50% of people tend to relapse in the first few days of trying to quit cigarette smoking (Garvey et al. 1992; Law and Tang 1995).

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## 26.6 Alcohol Dependence (AD)

While the mechanism by which MAO-A elevation occurs during early withdrawal from cigarette smoking is not known, mechanisms involving oxidative stress and mitochondrial toxicity are associated with greater expression of MAO-A in glioblastoma and neuroblastoma cell lines (Youdim et al. 2006; Fitzgerald et al.

2007a, b; Ou et al. 2006). The first investigations of MAO-A activity in the post-mortem brain of AD were primarily negative, with no change reported in prefrontal cortex, and a decrease in the hypothalamus and caudate (Oreland et al. 1983; Major et al. 1985). However, none of these studies controlled for factors that influence MAO-A levels, such as cigarette smoking (Fowler et al. 1996; Bacher et al. 2011), exposure to current or past major depressive episodes (Meyer et al. 2006, 2009b; Johnson et al. 2011), and impulsive/aggressive personality traits (Alia-Klein et al. 2008; Soliman et al. 2011). In a [ $^{11}\text{C}$ ] harmine PET study controlling for these factors, MAO-A  $V_T$  was significantly greater in the PFC (37%, independent samples  $t$ -test,  $t_{30} = 3.93$ ,  $p < 0.001$ ) and all brain regions analyzed (mean 32%, multivariate analysis of variance,  $F_{7,24} = 3.67$ ,  $p = 0.008$ ) (Matthews et al. 2014). Moreover, greater duration of heavy drinking correlated positively with greater MAO-A  $V_T$  in the PFC ( $R = 0.67$ ,  $p = 0.005$ ) and all brain regions analyzed ( $R = 0.73$ – $0.57$ ,  $p = 0.001$ – $0.02$ ). A subsequent study of chronic ethanol vapor exposure achieving high levels of ethanol in Sprague Dawley rats comparable to human exposure demonstrated that exposure to alcohol was associated with greater MAO-A activity and protein level in the PFC and ACC (Matthews et al. 2018).

### 26.6.1 Aggression

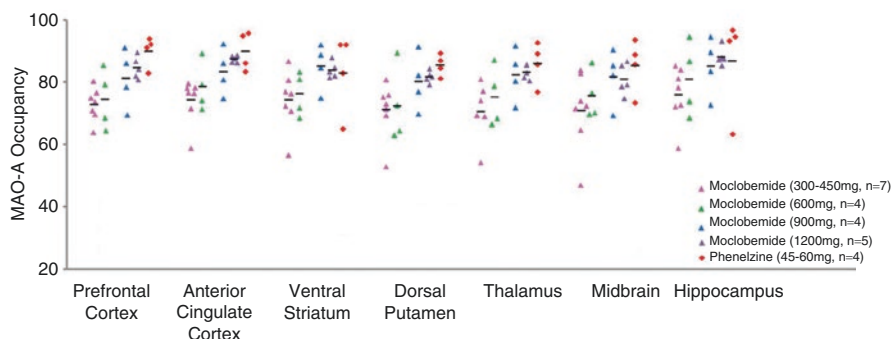
Aggression is associated with major social costs since 50% of prisoners have problems with aggression (Fazel and Danesh 2002). Substantial evidence has accumulated to suggest that inherited levels of MAO-A may also have influence upon neurodevelopment and lead to impulsive behavior: MAO-A knockout (KO) mice and wild-type given MAO-A inhibitors during development demonstrate elevated brain serotonin during development, which disrupts the organization of somatosensory thalamocortical afferents and abnormalities in the somatosensory cortex barrel fields (Cases et al. 1996; Vitalis et al. 1998; Mejia et al. 2002). It has been proposed that impulsive aggression observed in MAO-A KO mice and mice given MAO-A inhibitors during development is consequent to impairment in organizing sensory input secondary to these abnormalities (Bortolato et al. 2008). Men who are deficient in MAO-A, consequent to an X-chromosome mutation that impairs function of the MAO-A gene, demonstrate frequent aggressive behavior (Brunner et al. 1993). Consistent with this, people with a genotype associated with low levels of MAO-A in an in vitro assay have been associated with greater risk of developing antisocial behavior (Caspi et al. 2002; Kim-Cohen et al. 2006).

However, a critical gap in the literature is that until 2015, it was unknown whether brain MAO-A levels are low in more severe, clinical disorders of impulsivity, such as antisocial personality disorder (ASPD). To address this, [ $^{11}\text{C}$ ]harmine PET was applied to measure MAO-A  $V_T$  in 18 male ASPD participants and 18 age- and sex-matched controls. Orbitofrontal cortex (OFC) and ventral striatal (VS) MAO-A  $V_T$  were lower in ASPD compared to controls (MANOVA group effect:  $F_{2,33} = 6.8$ ,  $p = 0.003$ ; OFC and VS MAO-A  $V_T$  each reduced 19%). Similar effects were seen

in other brain regions: anterior cingulate cortex, dorsal putamen, thalamus, hippocampus, and midbrain (MANOVA group effect:  $F_{7,28} = 2.9, p = 0.022$ ). In ASPD, VS MAO-A  $V_T$  was consistently negatively correlated with measures linked to impulsivity including the Iowa gambling task, the psychopathy checklist revised, and the NEO Personality Inventory-Revised ( $r = -0.50$  to  $-0.52$ , all  $p$ -values  $< 0.05$ ) (Kolla et al. 2015).

Two PET imaging studies of MAO-A in relation to aggressive personality traits also show consistent results. First, a [ $^{11}\text{C}$ ]clorgyline PET study reported a positive correlation between reduced MAO-A binding in prefrontal cortex and subregions of the prefrontal cortex such as the orbitofrontal cortex, with self-reported aggression in healthy men (Fig. 26.4) (Alia-Klein et al. 2008). The second study, which applied [ $^{11}\text{C}$ ]-harmine positron emission tomography with the Revised NEO Personality Inventory, found that prefrontal cortex and subregions of the prefrontal cortex (such as the orbitofrontal cortex) correlated strongly with personality facets angry-hostility negatively ( $r = -0.515, p = 0.001$ ) and deliberation facets positively ( $r = 0.514, p = 0.001$ ) (Soliman et al. 2011). In a 2-factor regression model, these facets explained 38% of variance in prefrontal cortex MAO-A binding. In both studies, the MAO-A-binding measures among different regions were intercorrelated, so the correlations of MAO-A binding with aggression-related measures were also present in all brain regions sampled which included both cortical and subcortical regions.

Currently this neuroimaging direction is still progressing toward creating impactful information for society. Given that reduced MAO-A binding is related to pathological aggression, a longer-term strategy could be to identify whether harmful environmental influences predispose to this low MAO-A level phenotype in the prefrontal cortex during neurodevelopment so as to develop novel prevention strategies against harmful neurodevelopment.



**Fig. 26.4** Monoamine oxidase-A occupancy higher in high doses of moclobemide (900–1200 mg) and phenelzine (45–60 mg) than low doses of moclobemide (300–600 mg). There was a significant main effect of doses of moclobemide (1200 and 900 mg) and phenelzine (45 and 60 mg) on occupancy across brain regions sampled when compared to the average clinical doses of moclobemide [300, 450, and 600 mg; MANOVA,  $F(7,16) = 3.94, p = 0.01$ ]



## 26.7 Monoamine Oxidase A Occupancy

Increasingly it is established that clear definitions of target occupancy are key for developing novel therapeutics and occupancy data is now available for two monoamine oxidase inhibitors (MAOI). The irreversible MAOI phenelzine potently reduces MAO-A  $V_T$  by 82–93% after a 6-week clinical trial (Chiucciariello et al. 2015) (see Fig. 26.4). Moclobemide, a selective MAO-A inhibitor with a superior tolerability profile, has a lower occupancy at standard treating doses but can reduce MAO-A  $V_T$  to a similar extent if the dose is double the usual maximum recommended daily dose (Sacher et al. 2011; Chiucciariello et al. 2015) (also see Fig. 26.4). The degree of inhibition associated with therapeutic effect in these trials exceeds the level of disease effect. However, there are several potential explanations for this: It is plausible that the disease effect is present for months to years prior to treatment, and it is possible that downstream consequences of chronically elevated MAO-A are more likely to be targeted after a greater degree of reversal. Furthermore, some antidepressant mechanisms like inducing signal transduction effects and inducing hippocampal neurogenesis are associated with treatments that raise monoamine levels beyond healthy state (Sacher et al. 2011).

## 26.8 Why Image Indices of Serotonin Transporter Density?

The serotonin transporter (5-HTT) is a 630 amino acid long protein with 12 transmembrane domains (Lesch et al. 1993a, b) and is mainly found in the soma, dendrites, and terminals of serotonin-releasing neurons (Zhou et al. 1998; Tao-Cheng and Zhou 1999). Most 5-HTT are located on plasma cell membranes, mainly perisynaptically or along axons (Zhou et al. 1998; Tao-Cheng and Zhou 1999). In the human brain, there is a wide variation in regional 5-HTT density: Superior and inferior raphe nuclei > hypothalamus > thalamus (depending upon nucleus)  $\approx$  amygdala > putamen > caudate  $\approx$  hippocampus > insular cortex > prefrontal cortex > white matter > cerebellar cortex (except vermis) (Cortes et al. 1988; Kish et al. 2005; Laruelle et al. 1988). The serotonin transporter is coupled to sodium, chlorine, and potassium transport (Blakely et al. 1997), but the physiological role of greatest interest of 5-HTT in the brain is its influence upon extracellular serotonin levels. Antidepressants which bind to the serotonin transporter raise extracellular serotonin, and 5-HTT knockout mice have elevated extracellular serotonin, confirming the role of the serotonin transporter in modulating extracellular serotonin levels in vivo (Blier and De Montigny 1983; Bel and Artigas 1992, 1993, 1995; Dreshfield et al. 1996; Moret and Briley 1996; Mathews et al. 2000).

Neuroimaging studies of the 5-HTT offer the opportunity to measure an index of total 5-HTT density, the 5-HTT  $BP_{ND}$ . This marker is useful for determining therapeutic occupancy of antidepressants as well as several different models of 5-HTT dysregulation in disease. Most disease investigations focus upon potential mechanisms for lowering brain serotonin, and there are at least four models to explain how 5-HTT  $BP_{ND}$  could be altered in a disease that lowers brain monoamines (Meyer

2007). The first model is a lesion model in which the density of serotonin-releasing neurons is reduced and reductions in 5-HTT  $BP_{ND}$  occur. The second model is a secondary change in transporter binding consequent to serotonin lowering via a different process. However, available evidence suggests that the different monoamine transporters do not regulate in the same fashion after chronic depletion of their endogenous monoamine. Acute reductions in serotonin lead to reductions in 5-HTT mRNA (Yu et al. 1995), but long-term reductions or elevations in serotonin typically show no effect upon regional 5-HTT density (Benmansour et al. 1999; Dewar et al. 1992; Graham et al. 1987). The third model is increased clearance of extracellular monoamine via greater serotonin transporter density since some radiotracers like [ $^{11}C$ ]DASB preferentially bind to 5-HTT adjacent to extracellular fluid (Quelch et al. 2012). In the third model, greater total available 5-HTT  $BP_{ND}$  is interpreted as representative of greater functional 5-HTT density. The fourth model is endogenous displacement and is dependent upon the properties of the radioligand: Endogenous displacement is the property of a few radioligands to express different bindings after short-term manipulations of their endogenous neurotransmitter. Available evidence suggests that this fourth model is unlikely to be relevant in humans with [ $^{11}C$ ]DASB (Talbot et al. 2005; Praschak-Rieder et al. 2005), but the issue has not been investigated for other radioligands. Abnormalities in 5-HTT  $BP_{ND}$  may be discussed in the context of these models.

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## 26.9 Radioligands Available for Imaging Serotonin Transporters

There are at least six different radiotracer imaging protocols for imaging serotonin transporters, and the quality of technique varies widely. Originally, the only two radiotracer techniques were 2-beta-carbomethoxy-3-beta-(4-iodophenyl)-tropane ( $\beta$ -CIT) single-photon emission tomography (SPECT) and [ $^{11}C$ ](+)McN5652 PET, but both of these methods have significant limitations for quantitation (Kuikka et al. 1993).  $\beta$ -CIT has almost equal affinity for the dopamine transporter as compared to the serotonin transporter (Laruelle et al. 1994). As dopamine transporter density is high in the substantia nigra (Ciliax et al. 1999), the relative contributions of dopamine and serotonin transporter binding to the measure of specific binding cannot be separated in the midbrain, the location where this radiotracer technique is applied. The other early radiotracer, [ $^{11}C$ ](+)McN5652, has a low ratio of specific binding relative to free and nonspecific binding, which, in combination with modest reversibility, makes valid and reliable quantitation difficult in regions other than the thalamus and impossible in the human cortex (Kent et al. 2002; Buck et al. 2000).

The radiotracer [ $^{11}C$ ](DASB, 3-amino-4-(2-dimethylaminomethylphenylsulfanyl)-benzonitrile) was a major advance as a result of its selectivity, reversibility, greater specific binding, and reliability (Praschak-Rieder et al. 2005; Ginovart et al. 2001; Meyer et al. 2001, 2004a, b; Wilson et al. 2000, 2002). This radiotracer was three orders of magnitude more selective for the 5-HTT over other monoamine

transporters and highly selective for the 5-HTT in comparison to a number of other targets screened (Wilson et al. 2000, 2002). Selectivity was also demonstrated when 94% of the specific binding was displaceable by 5-HTT binding medications in animal models (Wilson et al. 2000, 2002). [ $^{11}\text{C}$ ]DASB has very good brain uptake in humans (Ginovart et al. 2001). In humans, its ratio of specific binding relative to free and nonspecific is good, and its free and nonspecific binding has low between-subject variability (Ginovart et al. 2001; Ichise et al. 2003). Multiple brain regions may be assessed (Praschak-Rieder et al. 2005; Ginovart et al. 2001; Meyer et al. 2001, 2004a, b; Wilson et al. 2000, 2002), and reliability of regional 5-HTT  $\text{BP}_{\text{ND}}$  measures is very good (Praschak-Rieder et al. 2005; Meyer et al. 2001, 2004b). The 5-HTT  $\text{BP}_{\text{ND}}$  measures are low in cortex, but with a standardized region of interest methods, very good reliability of 5-HTT  $\text{BP}_{\text{ND}}$  in the human cortex is achievable (Praschak-Rieder et al. 2005; Meyer et al. 2001, 2004b). Thus, the discovery of [ $^{11}\text{C}$ ]DASB PET imaging created a new opportunity for validly quantifying 5-HTT binding in humans.

Subsequent to the development of [ $^{11}\text{C}$ ]DASB, several new similarly structured radiotracers were created for imaging techniques of [ $^{123}\text{I}$ ]ADAM SPECT, [ $^{11}\text{C}$ ]MADAM PET, and [ $^{11}\text{C}$ ]HOMADAM PET (see Table 26.2 for a comparison). [ $^{123}\text{I}$ ]ADAM SPECT is superior to  $\beta$ -CIT SPECT; however, [ $^{123}\text{I}$ ]ADAM SPECT is not as useful a technique as the other methods because of its limited specific binding to free and nonspecific binding signal and lack of formal modeling development. [ $^{11}\text{C}$ ]MADAM PET approaches the quality of [ $^{11}\text{C}$ ]DASB PET, but [ $^{11}\text{C}$ ]MADAM has substantially less brain uptake which would be expected to increase the variability of the measurement, especially in regions of lower 5-HTT density (Lundberg et al. 2006). However, it is clear that [ $^{11}\text{C}$ ]MADAM is superior to [ $^{11}\text{C}$ ](+)McN5652 in terms of the reversibility of time activity curves and specific binding signal relative to free and nonspecific binding signal. [ $^{11}\text{C}$ ]HOMADAM PET shows some advantage over [ $^{11}\text{C}$ ]DASB PET (and the other radiotracers) in that it has a more reversible time activity curve and a somewhat greater specific binding to free and nonspecific binding signal (Ginovart et al. 2001; Nye et al. 2008), but there are still key areas of development for [ $^{11}\text{C}$ ]HOMADAM PET, including a reliability assessment of binding measures and assessment of its sensitivity to endogenous serotonin.

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## 26.10 Developing New Antidepressants with Serotonin Transporter Neuroimaging

In two seminal studies published in 2001 and 2004 applying [ $^{11}\text{C}$ ]DASB PET, it was demonstrated across five different selective serotonin reuptake inhibitors (SSRI) that after 4 weeks of treatment, for doses which clinically distinguish from placebo, an 80% occupancy occurs in most brain regions (Meyer et al. 2001, 2004b) (see Fig. 26.5). These SSRI are defined as such by the property of having at least two orders of magnitude preferential affinity for the serotonin transporter over other target sites. Within the datasets, there was increasing occupancy with increasing dose (and plasma level), with plateauing at the higher doses and higher plasma levels. The dose-occupancy relationships consistently demonstrated that the threshold of reaching 80% occupancy was important. Even though these SSRI have a 100-fold range in

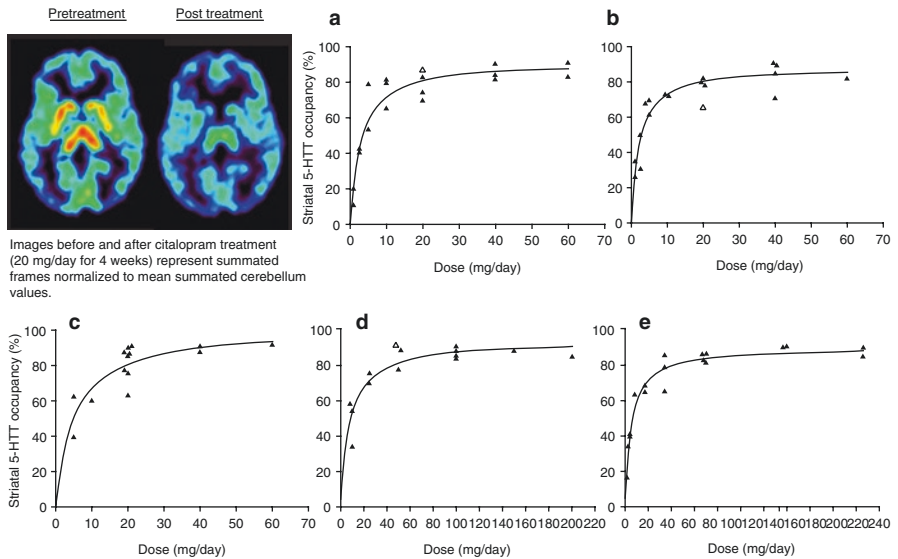
**Table 26.2** Comparison of PET radiotracers for the serotonin transporter (from Meyer (2014))

	[ <sup>123</sup> I]β-CIT SPECT	[ <sup>11</sup> C]MnN5652 PET	[ <sup>11</sup> C]DASB PET	[ <sup>123</sup> I]ADAM SPECT	[ <sup>11</sup> C]MADAM PET	[ <sup>11</sup> C]HOMADAM PET
Selectivity	Non-selective – near 1:1 affinity for 5-HTT to DAT (Laruelle et al. 1994; Carroll et al. 1995)	Likely selective 10:1 to 100:1 affinity for 5-HTT over NET (Shank et al. 1988; Kung et al. 1999)	Highly selective 1000:1 affinity for 5-HTT over NET or DAT; negligible affinity for many other targets (Wilson et al. 2000, 2002)	Highly selective 1000:1 affinity for 5-HTT over NET or DAT (Choi et al. 2000; Oya et al. 2000)	Highly selective (Hallidin et al. 2005; Chalton et al. 2003)	Highly selective (Jarkas et al. 2005)
Displaceability of specific binding	Incomplete (Pirker et al. 1995; Tauscher et al. 1999)	In most, reports (Kent et al. 2002; Parsey et al. 2000; Suhara et al. 2003)	Highly displaceable (Meyer et al. 2001, 2004b; Wilson et al. 2000, 2002)	Highly displaceable (Choi et al. 2000; Oya et al. 2000; Erlandsson et al. 2005)	Highly displaceable (Hallidin et al. 2005; Chalton et al. 2003; Lundberg et al. 2007)	Highly displaceable (Jarkas et al. 2005)
Reversibility (in human imaging)	Good (Kuikka et al. 1993; Brucke et al. 1993)	Not adequate to adequate, depending upon region (Buck et al. 2000; Parsey et al. 2000; Ikoma et al. 2002)	Adequate in the midbrain, good to very good in other regions (Ginovart et al. 2001; Ichise et al. 2003; Houle et al. 2000)	Adequate in the midbrain (Erlandsson et al. 2005; Catafau et al. 2005)	Adequate in the midbrain, good to very good in other regions	Very reversible in all regions (Nye et al. 2008)
Brain uptake	Adequate (Kuikka et al. 1993; Brucke et al. 1993)	Good (Parsey et al. 2000; Ikoma et al. 2002)	Very good (Ginovart et al. 2001; Ichise et al. 2003; Houle et al. 2000)	Adequate (Erlandsson et al. 2005; Catafau et al. 2005)	Good (Lundberg et al. 2006)	Very good (Nye et al. 2008)

(continued)

Table 26.2 (continued)

	[ <sup>123</sup> I]β-CIT SPECT	[ <sup>11</sup> C]MεN5652 PET	[ <sup>11</sup> C]DASB PET	[ <sup>123</sup> I]ADAM SPECT	[ <sup>11</sup> C]MADAM PET	[ <sup>11</sup> C]HOMADAM PET
Sensitivity to endogenous serotonin	Not known	Not known	Negligible (Talbot et al. 2005; Praschak-Rieder et al. 2005)	Not known	Not known	Not known
Specific binding to free and nonspecific binding ratio**	Good	Not adequate in most regions; adequate in the thalamus (Ikoma et al. 2002; Frankle et al. 2005)	Adequate to very good, depending upon region (Ginovart et al. 2001; Ichise et al. 2003)	Not adequate in most regions; adequate in the midbrain (Erlandsson et al. 2005; Catafau et al. 2005)	Adequate to very good (Lundberg et al. 2006, 2005)	Adequate to very good (Nye et al. 2008)
Reliability of 5-HTT BP**	Not measured	Modest (Kent et al. 2002)	Very good to excellent (Praschak-Rieder et al. 2005; Meyer et al. 2001)	Most regions reasonable (Catafau et al. 2005)	Good to very good (Lundberg et al. 2006)	Not yet reported
5-HTT BP measurable in multiple regions?	Brainstem only (Kuikka et al. 1993; Brucke et al. 1993)	Measurable in the thalamus (Ikoma et al. 2002), not measurable in the cortex (Parsey et al. 2000)	Yes (Ginovart et al. 2001; Ichise et al. 2003)	Measurable in the midbrain; unclear for other regions (Erlandsson et al. 2005; Catafau et al. 2005)	Yes (Lundberg et al. 2006)	Yes (Nye et al. 2008)



**Fig. 26.5** The data was fit using an equation of form  $f(x) = a \times x / (b + x)$ . Each fit was highly significant ( $p < 0.0002$  for all). Scanning occurred before treatment after four weeks of the treating dose. Medication type and dosage that differentiates from placebo are listed as follows: A-citalopram (20–40 mg), B-fluoxetine (20 mg), C-paroxetine (20 mg), D-sertraline (50–100 mg), E-venlafaxineXR (75 mg). Occupancy is the per cent reduction in binding potential ( $BP_{ND}$ ) i.e., Occupancy =  $(\text{baseline } BP_{ND} - \text{treatment } BP_{ND}) / \text{baseline } BP_{ND}$

affinity for the serotonin transporter, an 80% striatal 5-HTT occupancy occurs at minimum clinical dose, thereby demonstrating an additional, important predictive value of the 5-HTT occupancy data in relation to clinical trial development. Furthermore, the added information from occupancy measurement is clear because the in vivo  $EC_{50}$  does not correlate with affinity (Meyer et al. 2004b). This association of 80% occupancy with doses of SSRI that distinguish from placebo in clinical trial is consistently reported (Kent et al. 2002; Suhara et al. 2003). Hence, while affinity is an essential piece of information regarding an antidepressant, it cannot predict occupancy, even when plasma levels are known (Meyer et al. 2004b). Given the association between the clinically relevant dose and 5-HTT occupancy for SSRIs, it is now generally believed that an 80% 5-HTT occupancy is a therapeutic threshold for new antidepressants. This technique is frequently applied during phase I trials to assess whether new antidepressants in development are adequately brain penetrant and to guide dosing selection for subsequent phase II clinical trials.

### 26.11 Major Depressive Disorder

Neuroimaging studies of 5-HTT enable measurement of 5-HTT  $BP_{ND}$  in the midst of a MDE, as well as the ability to select more homogeneous subtypes of MDD like early-onset MDD (in contrast to the more heterogenous late-onset MDD). Sampling

early-onset MDD is usually not feasible for postmortem studies of 5-HTT density due to challenges of sampling subjects at older ages. The two postmortem investigations of 5-HTT density in subjects with recent symptoms of MDE evaluated 5-HTT density in the dorsal raphe and locus coeruleus, finding similar levels between MDD and healthy subjects (see review for more details (Stockmeier 2003)).

Neuroimaging studies of 5-HTT binding in untreated MDD exemplify a wide range of sample sizes and quality of techniques. However, there are three studies that focus upon medication-free subjects, with minimum sample sizes of 15 in each group, apply a radioligand of [<sup>11</sup>C]DASB level quality, and tend to focus upon early-onset MDD. For a broader review of these studies, including smaller studies not meeting these criteria, the reader is referred to Meyer et al. (Meyer 2008). The first study applied [<sup>11</sup>C]DASB PET to a sample of 20 subjects with early-onset MDE and 20 healthy controls (Meyer et al. 2004a). Subjects were medication-free for at least 3 months, age matched, had no comorbid axis I illnesses, were nonsmoking, and had early-onset MDD. There was no difference in 5-HTT BP<sub>ND</sub> between MDE and healthy controls, but MDE subjects with severely pessimistic dysfunctional attitudes had significantly higher 5-HTT BP<sub>ND</sub> (on average 21%) compared to healthy volunteers in brain regions sampling serotonin nerve terminals (prefrontal cortex, anterior cingulate, thalamus, bilateral caudate, bilateral putamen). Consistent with the postmortem studies of MDE, there was no difference in midbrain 5-HTT BP<sub>ND</sub>. Within the MDE group, greater 5-HTT BP<sub>ND</sub> was strongly associated with more negativistic dysfunctional attitudes in the same brain regions. An interpretation of the relationship between the functional measure of dysfunctional attitudes and the measure of 5-HTT BP<sub>ND</sub> is that greater regional 5-HTT levels could provide greater vulnerability to low extracellular 5-HT through excessive clearance of extracellular serotonin, leading to subsequent symptoms of extremely negativistic dysfunctional attitudes. There is particular utility in relating a functional symptom measure to 5-HTT binding because not all neuronal 5-HTT are in a functional location (Zhou et al. 1998; Tao-Cheng and Zhou 1999) although total 5-HTT level correlates with extracellular 5-HT in knockout models in vivo (Mathews et al. 2004)). The second study reported elevated 5-HTT binding in several brain regions (thalamus, striatum, insula) during MDE (Cannon et al. 2007), and the third study reports no difference in remitted MDD (Bhagwagar et al. 2007) (dysfunctional attitudes were not assessed in the latter two studies), demonstrating fairly consistent results across groups.

Among the analyses completed in a recent meta-analysis (Spies et al. 2015), there was a subanalysis of 184 participants who had been studied with radioligands selective for 5-HTT for which the striatum was a suitable region of interest. In this subanalysis, which was restricted to the striatal 5-HTT BP<sub>ND</sub> due to sample availability, it was interesting that younger samples had relatively higher striatal 5-HTT BP<sub>ND</sub>, and older samples had relatively lower 5-HTT BP<sub>ND</sub>. One might speculate that with a longer duration of illness, there is a lower density of 5-HTT BP<sub>ND</sub>, but this will take further study to determine. As these measures were relative to age-matched controls, the result was not a consequence of simply reductions in 5-HTT BP<sub>ND</sub> with age (Spies et al. 2015).

## 26.12 Ecstasy Abuse and Serotonin Transporter Imaging

Ecstasy (3,4-methylenedioxymethamphetamine, MDMA) is a substance that is frequently used in certain social contexts as it enhances emotional well-being, feelings of closeness, and desire to be sociable. Its mechanism, based upon the effects of serotonin transporter blockade and serotonin transporter knockout mice, is likely mainly mediated through binding to the serotonin transporter. Animal models suggest that exposure to MDMA is associated with widespread, persistent loss in serotonin transporter density, raising concern about the safety of exposure to this substance.

PET imaging allows for assessment in regard to the effects of MDMA upon 5-HTT BP<sub>ND</sub> in vivo including the regional distribution of this effect and its persistence at amounts typically self-administered by people. Neuroimaging results applying [<sup>11</sup>C]DASB PET have been very consistent across sites with reductions in 5-HTT BP<sub>ND</sub> in prefrontal, anterior cingulate, parietal, temporal, occipital cortex, insula, and hippocampus but no change in subcortical regions including the striatum, thalamus, and midbrain being typical (McCann et al. 2005, 2008; Urban et al. 2012; Erritzoe et al. 2011; Kish et al. 2010). All studies to date report reductions in 5-HTT BP<sub>ND</sub> in cortical regions and four out of five reports sparing of subcortical structures (McCann et al. 2005, 2008; Urban et al. 2012; Erritzoe et al. 2011; Kish et al. 2010). The question of persistence of the reduction in 5-HTT BP<sub>ND</sub> requires further investigations. McCann et al. (McCann et al. 2008) reported no correlation between recency of use and 5-HTT BP<sub>ND</sub> over the initial 7 months, and Selvaraj et al. (Selvaraj et al. 2009) reported no regional difference in 5-HTT BP<sub>ND</sub> in people who had abstained on average for 2.7 years suggesting a normalization window in between these two time frames, which ideally should be followed longitudinally.

## 26.13 Obsessive Compulsive Disorder

Obsessive compulsive disorder is a disabling illness characterized by repeated obsessions and/or compulsions that are distressing and impair function (First et al. 1995). OCD tends to be either relapsing/remitting or chronic (Association AP 1994) affecting 2% of adults (Robins et al. 1984; Weissman et al. 1994). Most likely time of onset is bimodally distributed with an early-onset group in childhood to teenaged years and a late-onset group occurring in the second to third decade. There have been virtually no postmortem investigations into this condition making neuroimaging studies an important source of information toward understanding potential neurochemical abnormalities in this illness.

There have been three investigations of serotonin transporter imaging in obsessive compulsive disorder applying a newer imaging method (all with [<sup>11</sup>C]DASB PET), and although the individual results focused within each study vary, further scrutiny suggests some consistency (Hesse et al. 2011; Matsumoto et al. 2010; Reimold et al. 2007). Heinz et al. applied a voxel-based analysis and reported decreased 5-HTT BP<sub>ND</sub> in the midbrain and thalamus in a sample of nine subjects



with OCD (Reimold et al. 2007). In a sample of ten subjects with OCD, Matsumoto reported significantly decreased 5-HTT BP<sub>ND</sub> within the orbitofrontal and insular cortex (Matsumoto et al. 2010), and in a sample of 19 OCD subjects, divided by early ( $n = 6$ ) and late ( $n = 13$ ) onset, Hesse et al. reported decreased 5-HTT BP<sub>ND</sub> in the medial prefrontal cortex, hippocampus, caudate, putamen, thalamus, occipital cortex, and midbrain, which was primarily driven by reduced 5-HTT BP<sub>ND</sub> in the late-onset OCD (Hesse et al. 2011). What was reasonably consistent across the studies was a decrease in 5-HTT BP<sub>ND</sub>, as Matsumoto noted reductions in 5-HTT BP ranging from 6.3% to 42% across all regions studied. The data of Hesse et al. suggests that late-onset OCD is primarily responsible for the differences in 5-HTT BP<sub>ND</sub> between OCD and health. It is plausible that the studies conducted to date may be underpowered to fully determine the regional extent of the reduction of 5-HTT BP<sub>ND</sub> in late-onset OCD. The main reasons for being underpowered are that the voxel-based analyses tend to be less sensitive than region of interest-based analyses and the sample sizes may not be large enough to achieve results that are regionally consistent even though the direction of effect is consistent. In support of this perspective, the nonsignificant reductions in OCD across studies suggest a global reduction in 5-HTT BP<sub>ND</sub>. Based upon the data from these studies, future serotonin transporter imaging investigations should consider larger sample sizes focused upon late-onset OCD.

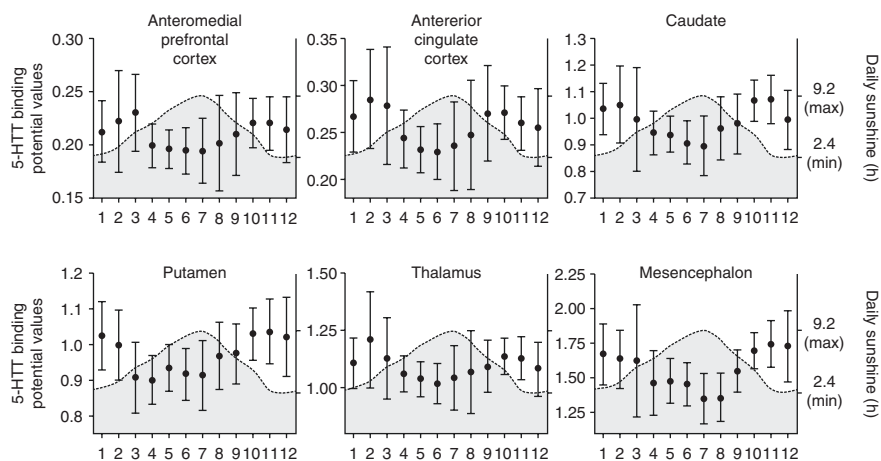
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## 26.14 Season and Serotonin Transporter Imaging

Seasonal affective disorder, that is, major depressive disorder with regular MDE in winter with full remissions in summer, is an important problem, especially in areas of the world at greater extremes of latitude: Prevalence rates of seasonal affective disorder (SAD) are about 1% at 40' latitude and 6% at 60' latitude or greater (Magnusson 2000). The functions of mood, appetite, sleep, and energy which fluctuate with season also are influenced by the serotonin system, and some indicators of serotonin physiology fluctuate with season in humans (Praschak-Rieder et al. 2008).

Prior to serotonin transporter imaging across season, several markers of seasonal fluctuations in serotonin physiology had been reported. In a human postmortem study of serotonin concentrations in the hypothalamus, Carlsson et al. reported lower levels in late winter and higher levels in late summer (Carlsson et al. 1980). Subsequently, Lambert et al. reported seasonal variation in whole brain serotonin turnover in vivo in humans (Lambert et al. 2002). Investigations in rodents find that reduced light exposure is associated with greater 5-HTT density (Rovescalli et al. 1989), lower 5-HT release (Blier et al. 1989), and greater 5-HT clearance (Rovescalli et al. 1989) in the hypothalamus and suprachiasmatic nucleus (two regions with high 5-HT concentration and density) (Tyrer et al. 2016a).

The four largest studies of 5-HTT binding and season have very consistent results. The first study that applied [<sup>11</sup>C]DASB PET, in 88 healthy, nonsmoking



**Fig. 26.6** Serotonin transporter binding potential values were measured using the selective serotonin transporter radioligand carbon 11-labeled 3-amino-4-(2-dimethylaminomethyl-phenylsulfanyl)-benzotriazole and positron emission tomography. The serotonin transporter binding potential values were determined in 6 brain regions [anteromedial prefrontal cortex (a), anterior cingulate cortex (b), caudate (c), putamen (d), thalamus (e), and mesencephalon (f)]. Circles represent bimonthly moving average means of serotonin transporter binding potential values; error bars, 95% confidence intervals of the mean. The shaded areas represent the average duration of sunshine in Toronto, Ontario, Canada (range, 2.4–9.2 h a day). There were 38 study participants in the fall and winter seasons and 50 in the spring and summer seasons. The number of study participants for each bimonthly data point was as follows: January, 11; February, 4; March, 6; April, 14; May, 23; June, 24; July, 14; August, 8; September, 13; October, 28; November, 21; and December, 10

humans in Toronto, Canada, found greater 5-HTT binding in the fall/winter as compared to spring/summer in all the brain regions sampled (medial prefrontal cortex, anterior cingulate cortex, thalamus, striatum, and midbrain) (Praschak-Rieder et al. 2008). See Fig. 26.6. The next [ $^{11}\text{C}$ ]DASB PET study of seasonality evaluated 54 subjects in Copenhagen, Denmark, and replicated the same seasonal variation in 5-HTT  $\text{BP}_{\text{ND}}$  (Kalbitzer et al. 2010). Ruhe et al. reported the same relationship between midbrain binding and season in a sample of 49 healthy and 49 depressed subjects applying [ $^{123}\text{I}$ ]B-CIT SPECT (Ruhe et al. 2009). Buchert et al. reported the same seasonal finding in the midbrain but not in the thalamus in a sample of 39 subjects using [ $^{11}\text{C}$ ]McN5652 PET (Buchert et al. 2006). Thus, studies with a large sample size located in more northern latitudes report greater 5-HTT  $\text{BP}_{\text{ND}}$  in the brain regions sampled in fall/winter compared to spring/summer.

More recently two groups evaluated 5-HTT  $\text{BP}_{\text{ND}}$  in both summer and winter in samples with seasonal affective disorder (SAD) and in health (Tyrer et al. 2016a; Mc Mahon et al. 2016). In SAD, with greater severity of symptoms across seasonal change, 5-HTT  $\text{BP}_{\text{ND}}$  was more elevated as compared to those in health or those with more mild SAD symptoms (Tyrer et al. 2016a; Mc Mahon et al. 2016). These findings tended to be global throughout gray matter brain regions including the PFC and ACC. There was a high consistency of findings across the two studies. In

addition, in a single-blind, placebo-controlled, counterbalanced crossover, controlled trial with light therapy in healthy participants, it was shown that 5-HTT BP<sub>ND</sub> is lowered in the ACC (Harrison et al. 2015). It was also demonstrated that after an open trial of light therapy in SAD, 5-HTT BP<sub>ND</sub> is lowered in both the PFC and ACC in SAD, as well as throughout other brain regions assayed in SAD, such as the hippocampus, ventral striatum, dorsal putamen, thalamus, and midbrain (Tyrer et al. 2016b).

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## 26.15 Conclusions

Neuroimaging studies applying recently optimized radiotracers for MAO-A (Bergstrom et al. 1997a; Ginovart et al. 2006) and serotonin transporter (Ginovart et al. 2001; Wilson et al. 2000) have created new advances in our knowledge of the relationship of these proteins to mood disorder, aggression, substance abuse, anxiety, and antidepressant development.

Prior to 2006, MAO-A had been viewed as a target for treatment of mood disorders, but neuroimaging studies have led to a new view that elevations in MAO-A level, particularly in the prefrontal and anterior cingulate cortex, occur during major depressive episodes; during high-risk states for major depressive episodes in adults such as in remission prior to recurrence, postpartum, and perimenopause; and during certain commonly comorbid illnesses like alcohol and cigarette dependence disorders and borderline personality disorder (Meyer et al. 2006, 2009a; Bacher et al. 2011; Sacher et al. 2010). New advances in understanding substance abuse are also emerging: During active cigarette smoking, available MAO-A binding is lowered, whereas during acute withdrawal, MAO-A binding is elevated (Fowler et al. 1996; Bacher et al. 2011), and MAO-A binding is highly elevated during early withdrawal from alcohol abuse disorder. These findings suggest that changes in MAO-A availability in adulthood predispose to persistent alterations in mood state. Interestingly, pathologically aggressive behavior and the personality traits of aggression are associated with reduced MAO-A binding, particularly in the prefrontal cortex, and in the context of the relationship of MAO-A levels to these traits in other models, we suggest that this may reflect a marker of a neurodevelopmental influence predisposing to a personality characteristic (Alia-Klein et al. 2008; Soliman et al. 2011).

5-HTT imaging now has a major influence upon novel antidepressant development since an 80% occupancy is predictive of a differential response versus placebo (Meyer et al. 2001, 2004b). 5-HTT imaging has verified that the effects of ecstasy upon reducing 5-HTT density in animal models and postmortem humans apply in vivo (McCann et al. 2005, 2008; Urban et al. 2012; Erritzoe et al. 2011; Kish et al. 2010) and are suggesting that diffuse 5-HTT loss occurs in OCD (Meyer et al. 2001, 2004b). In contrast, greater 5-HTT BP<sub>ND</sub> in untreated MDD and winter season suggests that a model of predisposition to serotonin loss may contribute toward a similar predisposition toward sad mood (Meyer et al. 2004a; Cannon et al. 2007;

Bhagwagar et al. 2007; Praschak-Rieder et al. 2008; Kalbitzer et al. 2010; Ruhe et al. 2009; Buchert et al. 2006).

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# PET Imaging of Sigma<sub>1</sub> Receptors

# 27

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## Abstract

Sigma receptors are classified into sigma<sub>1</sub> and sigma<sub>2</sub> subtypes. Sigma<sub>1</sub> receptors are widely distributed in the central nervous system (CNS) and in peripheral tissues. Sigma<sub>1</sub> receptors play a role in a variety of human CNS diseases, including mood disorders, stroke, neurodegenerative disease and drug addiction. Therefore, there is a great deal of interest in imaging of sigma<sub>1</sub> receptors in the living human brain. In contrast, sigma<sub>2</sub> receptors have been the focus of tumour imaging studies. A number of radioligands have been developed for imaging of sigma<sub>1</sub> receptors in the human brain, and a few, including [<sup>11</sup>C]SA4503 and (*S*)-(-)-[<sup>18</sup>F]fluspidine, have been used in clinical studies. Sigma<sub>1</sub> receptors are distributed throughout the grey matter of the human brain. A widespread decrease in [<sup>11</sup>C]SA4503 binding in patients with Alzheimer's disease and a significant decrease in binding on the more affected side of the anterior putamen in patients with Parkinson's disease have been reported. Receptor occupancy studies with [<sup>11</sup>C]SA4503-PET have shown that some antidepressants and antipsychotics have an affinity for sigma<sub>1</sub> receptors in the human brain in addition to their main targets. Recently, it has been reported that the binding of (*S*)-(-)-[<sup>18</sup>F]fluspidine is increased in patients with untreated major depressive disorder.

## Abbreviations

3-PPP	(+)-3-(3-Hydroxyphenyl)- <i>N</i> -(1-propyl)piperidine
AChE	Acetylcholinesterase
AD	Alzheimer's disease
BD-1047	<i>N'</i> -[2-(3,4-dichlorophenyl)ethyl]- <i>N,N,N'</i> -trimethyl-ethane-1,2-diamine
BDNF	Brain-derived neurotrophic factor
BMY14802 (BMS181100)	1-(4-Fluorophenyl)-4-[4-(5-fluoropyrimidin-2-yl)piperazin-1-yl]butan-1-ol
<i>BP</i> <sub>ND</sub>	Specific binding potential relative to non-displaceable binding
CFT	(1 <i>R</i> ,2 <i>S</i> ,3 <i>S</i> ,5 <i>S</i> )-3-(4-Fluorophenyl)-8-methyl-8-azabicyclo[3.2.1]octane-2-carboxylate
CNS	Central nervous system
Cutamesine	SA4503
DHEA	Dehydroepiandrosterone
Donepezil	( <i>RS</i> )-2-[(1-Benzyl-4-piperidyl)methyl]-5,6-dimethoxy-2,3-dihydroindeno[1,2- <i>b</i> ]pyridine-1-one
DTG	1,3-Di(2-tolyl)guanidine
DuP734	2-[1-(Cyclopropylmethyl)piperidin-4-yl]-1-(4-fluorophenyl)ethanonehydrobromide
EBP	Emopamil binding protein (sterol Δ8–Δ7 isomerase)

Eliprodil (SL82.0715)	1-(4-Chlorophenyl)-2-[4-[(4-fluorophenyl) methyl]piperidin-1-yl]ethanol
ER	Endoplasmic reticulum
FDG	2-Deoxy-2-fluoro-D-glucose
Fluspidine	1'-Benzyl-3-(2-fluoroethyl)-3 <i>H</i> -spiro[isobenzofuran-1,4'- piperidine]
Fluvoxiamine	( <i>E</i> )-5-Methoxy-1-[4-(trifluoromethyl)phenyl]pentan-1- one <i>O</i> -2-aminoethyl oxime
FM-SA4503	1-[2-(4-Fluoromethoxy-3-methoxyphenyl)ethyl]-4-(3- phenylpropyl)piperazine
$f_p$	Plasma-free fraction
FPS	1-(3-Fluoropropyl)-4-[(4-cyanophenoxy)methyl]piperidine
Haloperidol	4-[4-(4-Chlorophenyl)-4-hydroxy-1-piperidyl]-1-(4- fluorophenyl)-butan-1-one
HD	Huntington's disease
Ifenprodil	4-[2-(4-Benzylpiperidin-1-yl)-1-hydroxypropyl]phenol
Igmesine (JO1,784)	( <i>E</i> )- <i>N</i> -(Cyclopropylmethyl)- <i>N</i> -ethyl-3,6-diphenylhex-5- en-3-amine
LID	Levodopa-induced dyskinesia
MDD	Major depressive disorder
<i>mI</i> -SA4503	1-[2-(3,4-Dimethoxyphenyl)ethyl]-4-[3-(3-iodophenyl) propyl]piperazine
NE-100	4-[2-(Dipropylamino)ethyl]-2-(2-phenylethoxy)anisole
Nemonapride (YM-09151-2)	<i>N</i> -(1-benzyl-2-methylpyrrolidin-3-yl)-5-chloro-2- methoxy-4-(methylamino)benzamide
NGF	Nerve growth factor
NMDA	<i>N</i> -methyl-D-aspartate
<i>oI</i> -SA4503	1-[2-(3,4-Dimethoxyphenyl)ethyl]-4-[3-(2-iodophenyl) propyl]piperazine
Panamesine	(5 <i>S</i> )-5-[[4-(1,3-Benzodioxol-5-yl)-4-hydroxypiper- idin-1-yl]methyl]-3-(4-methoxyphenyl)-1,3-oxazoli- din-2-one
Paroxetine	(3 <i>S</i> ,4 <i>R</i> )-3-[(2 <i>H</i> -1,3-Benzodioxol-5-yloxy)methyl]-4-(4- fluorophenyl)piperidine
PD	Parkinson's disease
Pentazocine (EMD57445)	(2 <i>RS</i> ,6 <i>RS</i> ,11 <i>RS</i> )-6,11-Dimethyl-3-(3-methylbut-2-en-1- yl)-1,2,3,4,5,6-hexahydro-2,6-methano-3-benzazocin-8-ol
PET	Positron emission tomography
Physostigmine	(3 <i>aR</i> ,8 <i>aS</i> )-1,3 <i>a</i> ,8-Trimethyl-1 <i>H</i> ,2 <i>H</i> ,3 <i>H</i> ,3 <i>aH</i> ,8 <i>H</i> ,8 <i>aH</i> - pyr- rolo [2,3- <i>b</i> ] indol-5-yl <i>N</i> -methylcarbamate
PPBP	4-Phenyl-1-(4-phenylbutyl)piperidine
Raclopride	3,5-Dichloro- <i>N</i> -{[(2 <i>S</i> )-1-ethylpyrrolidin-2-yl]methyl}- 2-hydroxy-6-methoxybenzamide
rCBF	Regional cerebral blood flow



Rimcazole (BW234U)	9-{3-[(3 <i>R</i> ,5 <i>S</i> )-3,5-Dimethylpiperazin-1-yl]propyl}-9 <i>H</i> -carbazole
SFE	WLS1.002 1-(2-fluoroethyl)-4-[(4-cyanophenoxy)methyl]piperidine
SKF 10,047 ( <i>N</i> -allylnormetazocine)	(2 <i>R</i> ,6 <i>R</i> ,11 <i>R</i> )-6,11-dimethyl-3-prop-2-en-1-yl-1,2,3,4,5,6-hexahydro-2,6-methano-3-benzazocin-8-ol
SPECT	Single-photon emission computed tomography
Spiperone	8-[4-(4-Fluorophenyl)-4-oxo-butyl]-1-phenyl-1,3,8-triazaspiro[4.5]decan-4-one
SSRI	Selective serotonin reuptake inhibitor
SUV	Standardised uptake value
TAC	Time-activity curve
Tamoxifen	( <i>Z</i> )-2-[4-(1,2-Diphenylbut-1-enyl)phenoxy]- <i>N,N</i> -dimethylethanamine
TPCNE	1( <i>Trans</i> -iodopropen-2-yl)-4-[(4-cyanophenoxy)methyl]piperidine
Trifluoperazine	10-[3-(4-methylpiperazin-1-yl)propyl]-2-(trifluoromethyl)-10 <i>H</i> -phenothiazine
VACHT	Vesicular acetylcholine transporter
$V_T$	Total volume of distribution

## 27.1 Introduction

Sigma receptors were initially categorised as a subtype of opioid receptors, because the effects of a prototypical opioid agonist, SKF10,047 (*N*-allylnormetazocine), were independent from those of other opioid receptor agonists (Martin et al. 1976). Later, it was determined that sigma receptors are unique binding sites that are different from those of other opioid receptors and at least two subtypes ( $\sigma_1$  and  $\sigma_2$ ) were classified based on pharmacological criteria (Bowen et al. 1989). These subtypes display different tissue distributions and distinct physiological and pharmacological profiles in the central and peripheral nervous systems (Hashimoto and Ishiwata 2006).  $\sigma_1$  receptors are widely distributed in the central nervous system (CNS) and in peripheral tissues (Hayashi and Su 2004). Peripherally, both subtypes are overexpressed in rapidly proliferating cells, such as those in various types of cancer, and the functional roles of the receptors, especially  $\sigma_2$ , have also been the focus of tumour imaging studies (van Waarde et al. 2010). In the CNS, the  $\sigma_1$  receptor may function as a modulator of signal transduction through glutamatergic transmitter systems mediated by the *N*-methyl-D-aspartate (NMDA) receptors.  $\sigma_1$  agonists show anti-amnesic and neuroprotective effects (van

Waarde et al. 2011), and for that reason, they are attractive novel targets for the treatment of neuropsychiatric diseases (schizophrenia, depression and cognition) and neurodegenerative diseases like ischemic stroke and Alzheimer's disease (AD).

The sigma<sub>1</sub> receptor was successfully cloned and characterised as a protein of 233 amino acids with a molecular weight of 25.3 kDa (Hanner et al. 1996). Although the sigma<sub>1</sub> receptor is not homologous to any mammalian protein, it shows 30% overlap with yeast Δ8–Δ7 sterol isomerase, also known as emopamil binding protein (EBP). Only recently, the crystal structure of the human sigma<sub>1</sub> receptor was elucidated and found to possess a trimeric architecture with only a single transmembrane domain in each protomer (Schmidt et al. 2016).

Sigma<sub>1</sub> receptors have been shown to act as ligand-regulated molecular chaperones primarily situated on the mitochondria-associated membrane of the endoplasmic reticulum (ER) (Hayashi and Su 2007). Most recently, the sigma<sub>2</sub> receptor was identified as ER-resident transmembrane protein 97 (Alon et al. 2017), which is a sterol response element binding protein target gene.

Endogenous ligands for the sigma<sub>1</sub> and sigma<sub>2</sub> receptors have not been fully identified, and those for the sigma<sub>1</sub> receptor were first proposed to be neurosteroids, including pregnenolone sulphate and progesterone (Su et al. 1988). Recently, the endogenous hallucinogenic trace amine *N,N*-dimethyltryptamine was found to be a potent endogenous ligand for the sigma<sub>1</sub> receptor (Fontanila et al. 2009).

In this chapter, investigations related to brain imaging of sigma<sub>1</sub> receptors and their role in neurology and psychiatric diseases are reviewed.

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## 27.2 Brain Imaging of Sigma Receptors

### 27.2.1 Post-mortem Studies

Several post-mortem studies on the distribution and densities of sigma receptors in the human brain have been reported. Weissman et al. (1988) reported that the densities of sigma sites in the brain were highest in the cerebellum, nucleus accumbens and cerebral cortex, using a membrane binding assay with [<sup>3</sup>H]haloperidol in the presence of spiperone. The binding is moderate in the caudate nucleus and substantia nigra and low in the thalamic and pontine nuclei. Autoradiographic measurements using [<sup>3</sup>H]DTG showed slightly different results, i.e. high prevalence in the substantia nigra pars compacta and cerebellum with moderate prevalence in the hippocampal dentate granular cells, the striatum and the pineal and pituitary glands (Jansen et al. 1991). In the neocortex, binding was high in the laminae II–IVA and low in the narrow band (5% layer) of midzone. Subtype-selective regional distribution in the brain has been reported in non-human primates but not in humans. Mash and Zabetian (1992) quantified the binding of [<sup>3</sup>H]3-PPP (sigma<sub>1</sub> K<sub>i</sub> = 5.1 nM, sigma<sub>2</sub>/sigma<sub>1</sub> = 87; (Walker et al. 1990)) within 36 brain regions in rhesus macaque monkeys (*Macaca mulatta*) and demonstrated marked enrichment of sigma binding

sites over the paralimbic belt cortices, medial and central nuclei of the amygdala and hippocampal formation. Moderate sigma receptor densities were observed in the caudate and putamen.

Several studies reported on the effects of age on sigma receptor expression. Kornhuber et al. (1996) reported an 8.5% decrease per decade in (+)-[<sup>3</sup>H]pentazocine ( $\sigma_1 K_i = 2.0$  nM,  $\sigma_2/\sigma_1 = 228$ ; (Walker et al. 1990)) in the frontal cortex. Moreover, altered densities of sigma receptors were observed in patients with schizophrenia and AD (see also Sects. 27.3.1 and 27.3.4).

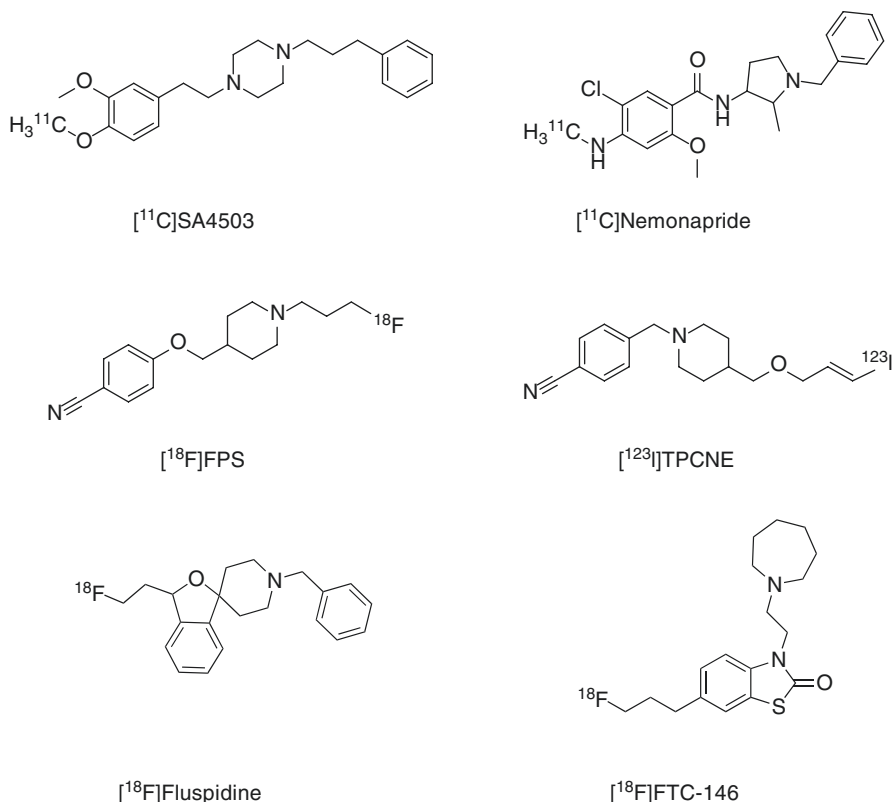
Most post-mortem examinations of the human brain reported to date measured both  $\sigma_1$  and  $\sigma_2$  binding sites using non-selective radioligands. One exception is a previous study of aging effects, which used the  $\sigma_1$  selective ligand (+)-[<sup>3</sup>H]pentazocine (Kornhuber et al. 1996).

## 27.2.2 Radioligands for Imaging of Sigma Receptors

Since the early 1990s, different classes of compounds, such as benzomorphans, phenyl piperidines, *N,N*-disubstituted pyrrolidinyethylamines and *N,N'*-disubstituted piperazines, have been evaluated as radioligands for use in the imaging of sigma receptors by positron emission tomography (PET) and single-photon emission computed tomography (SPECT). A comprehensive overview of the development of radioligands for the different sigma receptor subtypes has been presented in a number of recent reviews (Hashimoto and Ishiwata 2006; Brust et al. 2014a, b; Jia et al. 2019; Mach and Wheeler 2009). The chemical structures of radioligands used clinically for brain imaging of sigma receptors are summarised in Fig. 27.1.

The first radioligand used for imaging  $\sigma_1$  receptors in the CNS was [<sup>11</sup>C]SA4503 (Fig. 27.1), which was developed by Ishiwata and co-workers (Ishii et al. 2001). SA4503 has high and selective affinity for the  $\sigma_1$  receptor ( $\sigma_1$  IC<sub>50</sub> = 17.4 nM,  $\sigma_2/\sigma_1 = 103$ ) but low affinity for 36 neuroreceptors, ion channels and second messenger systems (Matsuno et al. 1996; Matsuno and Mita 1998). Later, four groups reinvestigated the binding properties of SA4503 and reported slightly different affinities and selectivities:  $\sigma_1 K_i = 3.3$ – $4.6$  nM,  $\sigma_2/\sigma_1 = 13.3$ – $55.0$  (Hirata et al. 2006; Lever et al. 2006; Shiba et al. 2006; Wang et al. 2014). Density ( $B_{\max}$ ) of the sigma receptor was estimated to be 30–600 fmol/mg protein (approximately 3–60 nM) in the human brain (Weissman et al. 1988; Kornhuber et al. 1996; Jansen et al. 1993). Theoretically, radioligands with nanomolar affinity (1–6 nM) will be suitable for sigma receptor imaging. From this viewpoint, SA4503 has a suitable range of affinity for quantitative *in vivo* imaging. In human studies, the use of [<sup>11</sup>C]SA4503-PET allowed successful visualisation of the  $\sigma_1$  receptor in the brain as described in Sect. 27.2.3.1, but the affinity of [<sup>11</sup>C]SA4503 was slightly high (i.e. the rate of dissociation of the ligand somewhat low) for measurement of the  $\sigma_1$  receptor on the time scale of the PET scan using a <sup>11</sup>C-labelled tracer with a short half-life (20.4 min) (Sakata et al. 2008).

The second PET ligand that was applied in human studies was [<sup>18</sup>F]FPS, which was developed by Waterhouse and co-workers (Collier et al. 1996). [<sup>18</sup>F]FPS was



**Fig. 27.1** Chemical structures of clinically used sigma ligands for brain imaging

not cleared fast enough from the CNS to reach transient equilibrium by 4 h after administration in healthy volunteers due to its high affinity for the sigma<sub>1</sub> receptor ( $K_d = 0.5$  nM) (Waterhouse et al. 2004; Zhao et al. 2005). In an effort to improve the CNS clearance for in vivo PET studies, Zhao et al. (2005) synthesised and evaluated [<sup>18</sup>F]SFE (previously designated as [<sup>18</sup>F]WLS1.002), a fluoroethyl derivative of [<sup>18</sup>F]FPS that exhibits a lower affinity for sigma<sub>1</sub> receptors ( $K_d = 5$  nM). A preclinical study showed that [<sup>18</sup>F]SFE exhibits very similar regional brain distribution and specific binding in the rodent brain compared to [<sup>18</sup>F]FPS. As expected from its lower affinity, [<sup>18</sup>F]SFE is cleared much faster from the brain than [<sup>18</sup>F]FPS (Waterhouse et al. 2006a). Preclinical toxicity and dosimetry studies of [<sup>18</sup>F]SFE suggested that [<sup>18</sup>F]SFE will be safe for use in human PET studies (Waterhouse et al. 2006b). However, no further clinical imaging studies have been reported to date.

Carbon-11-labelled nemonapride, which was originally developed as a dopamine D<sub>2</sub>-like receptor radioligand (Hatano et al. 1989; Hatazawa et al. 1991), binds not only to D<sub>2</sub>-like receptors in the striatum but also to sigma receptors in other regions, such as the cerebral cortex and cerebellum (Ishiwata and Senda 1999), as it

has rather high affinities for both receptors. Nimura et al. (2004) applied [ $^{11}\text{C}$ ]nemonapride to imaging of sigma receptors in the cerebellum of patients suffering from levodopa-induced dyskinesia (LID), a brain area devoid of  $\text{D}_2$ -like receptors.

More recently, Moussa and co-workers developed a series of *N*-(2-benzofuranylmethyl)-*N'*-(alkoxybenzyl)piperazines as selective sigma<sub>1</sub> receptor ligands. They found high-affinity sigma<sub>1</sub> selective ligands, such as the 4-[ $^{11}\text{C}$ ]methoxy (sigma<sub>1</sub>  $K_i$  = 2.7 nM, sigma<sub>2</sub>/sigma<sub>1</sub> = 38) and 4-(2-[ $^{18}\text{F}$ ]fluoroethoxy) derivatives (sigma<sub>1</sub>  $K_i$  = 2.6, sigma<sub>2</sub>/sigma<sub>1</sub> = 187) (Moussa et al. 2010, 2011). These compounds showed good brain uptake and specific binding to sigma<sub>1</sub> receptors in the *Papio hamadryas* (baboon) brain. However, these compounds showed irreversible binding profiles during the scan periods because of their high affinities.

The sigma<sub>1</sub> receptor-selective ligand [ $^{123}\text{I}$ ]TPCNE ( $K_i$  = 0.67 nM, sigma<sub>2</sub>/sigma<sub>1</sub> = 58) was used in a human SPECT study (Stone et al. 2006; Waterhouse et al. 1997). High brain uptake was reduced by haloperidol pretreatment, suggesting the specific binding of [ $^{123}\text{I}$ ]TPCNE to sigma<sub>1</sub> receptors. However, the time-activity data were best described by an irreversible model. Thus, no further studies for this ligand have been conducted. Radioiodinated analogues of SA4503 were also prepared for SPECT, i.e. *o*I-SA4503 and *m*I-SA4503 (designated as *o*-BON and *m*-BON, respectively) (Hirata et al. 2006). The affinity and selectivity of *m*I-SA4503 for the sigma<sub>1</sub> receptor (sigma<sub>1</sub>  $K_i$  = 8.9 nM, sigma<sub>2</sub>/sigma<sub>1</sub> = 6.1) were approximately half those of SA4503. Although Hirata et al. (2008) used these compounds in a tumour study on rodents, no clinical studies have been conducted to date.

A review of the clinical trials for a few sigma<sub>1</sub> receptor radioligands showed that ligands with affinities ( $K_i$ ) for the sigma<sub>1</sub> receptor of 1–10 nM may be preferable for human studies and that [ $^{18}\text{F}$ ]FM-SA4503 may be more suitable than [ $^{11}\text{C}$ ]SA4503 (Kawamura et al. 2007). However, many sigma receptor ligands have affinity for the vesicular acetylcholine transporter (VACHT) (Efang 2000) and the EBP (Berardi et al. 2001). SA4503 was reported to show affinities for VACHT ( $K_i$  = 50 nM, (Shiba et al. 2006)) and EBP ( $K_i$  = 1.7 nM, (Berardi et al. 2001)), but [ $^{11}\text{C}$ ]SA4503 did not seem to bind to VACHT in the rat brain in vivo (Ishiwata et al. 2006a). Toyohara et al. (2012) confirmed that the brain uptake of [ $^{11}\text{C}$ ]SA4503 in mice was not blocked by high-affinity EBP blockers tamoxifen (EBP  $K_i$  = 2.8 nM, sigma<sub>1</sub>/EBP = 12) and trifluoperazine (EBP  $K_i$  = 3.9 nM, sigma<sub>1</sub>/EBP = 52).

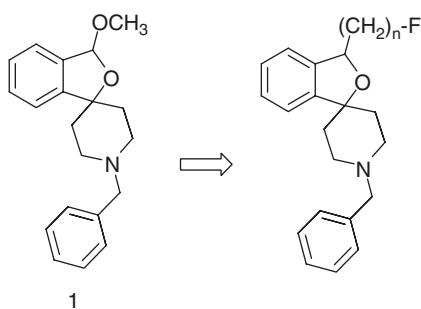
Selectivity toward VACHT and EBP sites should be taken into consideration when designing new selective ligands, although the relationship between the sigma receptor and EBP has not been clearly elucidated (Brust et al. 2014a, b).

A recently developed innovative compound class of spirocyclic benzofuran derivatives showed weak affinity for VACHT and EBP (Fischer et al. 2011). Wunsch and co-workers started the development of spirocyclic sigma receptor ligands with the 2-benzofuran (Fig. 27.2) (Maier and Wunsch 2002a, b). A very high affinity of **1** to sigma<sub>1</sub> receptors ( $K_i$  = 1.1 nM) was found while the sigma<sub>2</sub> affinity ( $K_i$  = 1280 nM) is very low, providing an excellent 1100-fold selectivity for sigma<sub>1</sub> receptors over the sigma<sub>2</sub> subtype. A screening against more than 60 other receptors, ion channels, transporters and enzymes revealed that there are no cross reactions with other

targets including the hERG channel (Wiese et al. 2009). This is a voltage-gated K<sup>+</sup> channel with high expression in the heart, by which potentially new drugs can lead to life-threatening arrhythmia. Generally, during drug development, hERG channel interactions are investigated very early to avoid those problems. Notably, the pharmacophores of sigma<sub>1</sub> receptor ligands and hERG channel blockers are very similar (Recanatini et al. 2005). In the mouse capsaicin assay, **1** showed high analgesic activity, which is in the same range as the analgesic activity of S1RA (E-52862), a prototypical benzomorphan sigma<sub>1</sub> receptor antagonist which was in Phase II clinical trials for pain (EudraCT Number: 2012-000398-21). Therefore, **1** and fluspidine are regarded as sigma<sub>1</sub> receptor antagonists. Incubation with rat liver microsomes provided important information on metabolic transformations. The acetal functional group appeared to be rather sensitive (Wiese et al. 2009).

In order to remove the chemically and metabolically labile acetal functionality and to install a structural element, which allows the introduction of a fluorine atom for potential radiolabelling, the methoxy group of **1** was replaced by homologous fluoroalkyl residues (compounds **2–5** in Fig. 27.2). The homologous fluoroalkyl derivatives **2–5** show low nanomolar up to subnanomolar sigma<sub>1</sub> affinity (see Fig. 27.2). Moreover, all four homologs display very high subtype selectivity (Große Mastrup et al. 2009, 2011; Maisonial et al. 2011, 2012; Wiese et al. 2011). The ethyl derivative with the highest sigma<sub>1</sub> affinity ( $K_i = 0.59$  nM), the highest subtype selectivity (1331-fold), the highest metabolic stability (91% radiotracer remaining in mouse plasma at 30 min p.i.), the highest brain-to-plasma ratio and the highest target-to-nontarget ratio in the brain (4.7 at 45 in p.i.) was termed fluspidine and chosen for further preclinical evaluation.

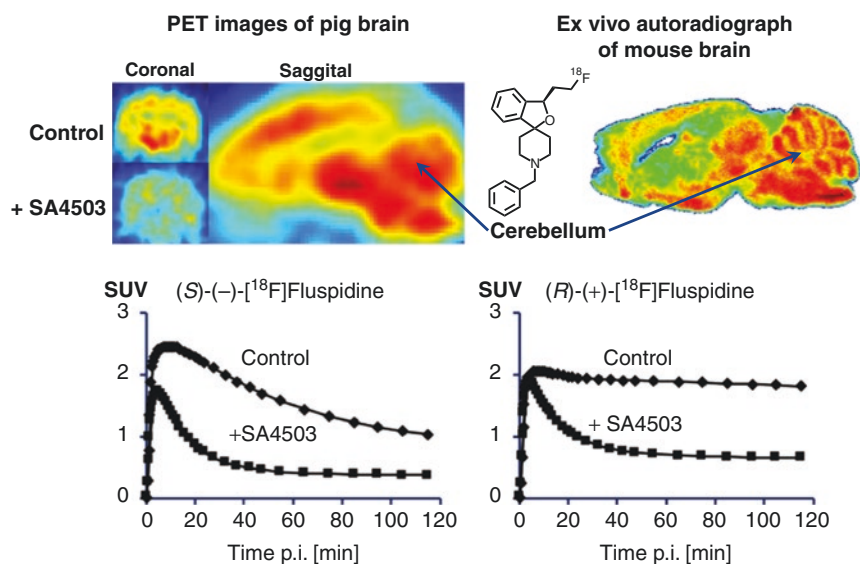
**Fig. 27.2** Development of fluorinated spirocyclic benzofuran starting from methoxy derivative **1**



compd.	n	$K_i(\sigma_1)$ (nM)	$K_i(\sigma_2)$ (nM)	selectivity
<b>2</b>	1	0.74	550	743
<b>3</b>	2	0.59	785	1331
<b>4</b>	3	1.4	837	598
<b>5</b>	4	1.2	489	408

Since fluspidine is a chiral compound, enantiomerically pure (*R*)- and (*S*)-fluspidine were synthesized, and their  $\sigma_1$  affinities, determined in a competitive equilibration binding assay, were 0.57 nM and 2.3 nM, respectively (Holl et al. 2013). Notably, later kinetic analysis revealed an irreversible binding mode of (*R*)-fluspidine which prevented the precise determination of the equilibrium dissociation constant  $K_D$ , while for (*S*)-fluspidine, a  $K_D$  of 0.1 nM was estimated (Baum et al. 2017). Due to the high  $\sigma_1$  affinity of both fluspidine enantiomers (*R*)-**13** and (*S*)-**13**, the radiosynthesis was performed by nucleophilic substitution of the (*R*)- and (*S*)-tosylates with  $K[^{18}\text{F}]\text{F}$  complexed with the cryptand Kryptofix<sup>(R)</sup>, and an automated procedure was developed for further preclinical evaluation (Maisonial-Besset et al. 2014). The uptake of (*S*)-(-)- $[^{18}\text{F}]\text{fluspidine}$  in porcine brain measured with PET was strongly correlated with the uptake in mouse brain measured by ex vivo autoradiography. Furthermore, the PET studies revealed a strong inhibition by SA4503 (see Fig. 27.3).

Both enantiomers of  $[^{18}\text{F}]\text{fluspidine}$  displayed different kinetics in porcine brain (Brust et al. 2014a, b) (Fig. 27.3). Although the initial brain uptake was similar, the accumulation measured at the end of the study was about twofold higher for (*R*)-(+)- $[^{18}\text{F}]\text{fluspidine}$ . Administration of SA4503 reduced the uptake almost equally for both radiotracers by ~65%. The higher affinity of (*R*)-(+)- $[^{18}\text{F}]\text{fluspidine}$  found in vitro was clearly translated into higher  $BP_{\text{ND}}$  values in vivo; however, longer



**Fig. 27.3** Studies with the PET tracer  $[^{18}\text{F}]\text{fluspidine}$  ( $[^{18}\text{F}](S)\text{-13}$ ): Top left: PET image of whole piglet brain and inhibition study with the  $\sigma_1$  ligand SA4503. Top right: ex vivo autoradiography of mouse brain; the region of the cerebellum (highest receptor density) is shown exemplarily. Bottom: kinetics of the uptake of (*S*)-(-)- $[^{18}\text{F}]\text{fluspidine}$  (left) and (*R*)-(+)- $[^{18}\text{F}]\text{fluspidine}$  in porcine brain without and with preinjection of SA4503 (modified according to Brust P, van den Hoff J, Steinbach J (2014) Development of  $^{18}\text{F}$ -labelled radiotracers for neuroreceptor imaging with Positron Emission Tomography. *Neurosci Bull* 30:777–811)

equilibration periods require longer scanning times. The higher metabolic stability of (*S*)-(-)-[<sup>18</sup>F]fluspidine results in a more reliable estimate of the metabolite-corrected plasma input function which is required for kinetic modelling of PET tracers for central sigma<sub>1</sub> receptors due to the lack of a receptor-free reference region. For measuring the sigma<sub>1</sub> receptor occupancy by either administered drugs or endogenous ligands, also the lower-affine and thus faster dissociating (*S*)-(-)-[<sup>18</sup>F]fluspidine seems to be more appropriate due to a higher susceptibility to the competitive compounds (Brust et al. 2014a, b).

Recently, the different brain kinetics of (*R*)-(+)-[<sup>18</sup>F]fluspidine and (*S*)-(-)-[<sup>18</sup>F]fluspidine was confirmed in monkeys (Baum et al. 2017). SA4503 at the dose of 0.5 mg/kg blocked about 85% of the brain uptake of (*S*)-(-)-[<sup>18</sup>F]fluspidine supporting the conclusion that this radiotracer has the kinetic and imaging properties for the investigation of sigma<sub>1</sub> receptors in the human brain with PET.

Accordingly, the preclinical dosimetry of (*S*)-(-)-[<sup>18</sup>F]fluspidine was investigated based on in vivo and ex vivo data from CD-1 mice. The effective dose (ED) estimations for human application were 14.8 μSv/MBq and 17.3 μSv/MBq, respectively (Kranz et al. 2016). Since toxicity results for (*S*)-(-)-fluspidine revealed a no observed effect level of 620 μg/kg, more than 10,000-fold above the injected radiotracer amounts, a dosimetry study in human was performed. An ED of 22.1 μSv/MBq was estimated which is similar to that of [<sup>18</sup>F]FDG (Kranz et al. 2016).

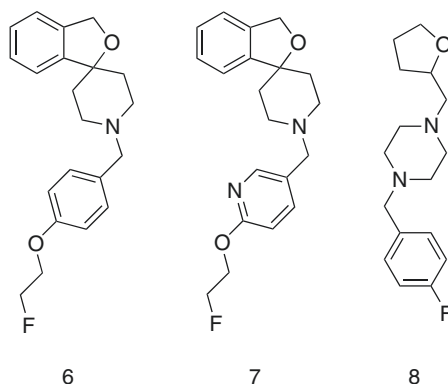
The preclinical studies discussed above provided the scientific data for two ongoing clinical studies using (*S*)-(-)-[<sup>18</sup>F]fluspidine in patients with major depressive disorder (MDD) and Huntington's disease (HD) (EudraCT Numbers: 2014-005427-27, 2016-001757-41).

To further develop PET radioligands from the spirocyclic piperidine series of compounds with suitable properties, Jia and co-workers took another approach, i.e. structural modifications on the *N*-benzyl moiety, while eliminating the substituent on the spirocyclic benzofuran section. Moreover, they replaced the aromatic ring in the *N*-benzyl moiety with a pyridine group to further reduce the lipophilicity of this series of derivatives (Li et al. 2013). Derivatives **6** and **7** (Fig. 27.4) possessed high sigma<sub>1</sub> receptor affinity (subnanomolar and low nanomolar affinity for compounds **6** and **7**, respectively) and excellent sigma<sub>1</sub>/sigma<sub>2</sub> subtype selectivity as well as high sigma<sub>1</sub>/VACHT selectivity. The corresponding radiotracers [<sup>18</sup>F]**6** and [<sup>18</sup>F]**7** showed high initial brain uptakes, high accumulation of the radiotracer in brain areas known to express high levels of sigma<sub>1</sub> receptors and specific binding to sigma<sub>1</sub> receptors in vivo. Interestingly, the radiotracer [<sup>18</sup>F]**7** with lower lipophilicity demonstrated higher specificity than [<sup>18</sup>F]**6**.

Later, four <sup>18</sup>F-labelled radioligands including (*S*)-(-)- and (*R*)-(+)-[<sup>18</sup>F]fluspidine, [<sup>18</sup>F]**6** and [<sup>18</sup>F]**7** from the same spirocyclic piperidine scaffold were evaluated in non-human primates (Baum et al. 2017). (*R*)-(+)-[<sup>18</sup>F]fluspidine and radioligand [<sup>18</sup>F]**6** with subnanomolar affinity for sigma<sub>1</sub> receptors displayed irreversible binding kinetics in the rhesus monkey brain, with no appreciable washout during the 4 h scan sessions. On the other hand, (*S*)-(-)-[<sup>18</sup>F]fluspidine and [<sup>18</sup>F]**7** were found to display fast and reversible kinetics in the monkey brain with good uptake and high specific binding signals. Both are judged to be promising radioligands to image



**Fig. 27.4** Novel *para*-aromatic substituted spirocyclic piperidine and piperazine compounds for imaging  $\sigma_1$  receptors in the brain



compd.	$K_i(\sigma_1)$ (nM)	$K_i(\sigma_2)$ (nM)	selectivity	Log <i>D</i>
6	0.79	277	351	2.55
7	2.3	327	142	2.50
8	3.2	168	53	0.76

$\sigma_1$  receptors in humans. In rhesus monkeys, [ $^{18}\text{F}$ ]7 has a tenfold higher plasma-free fraction ( $f_p$ ), threefold higher  $BP_{\text{ND}}$  and greater  $V_T$  than (*S*)-(-)-[ $^{18}\text{F}$ ]fluspidine.

Most recently, to develop novel radioligands with high specificity, Jia and co-workers eliminated the aromatic ring in [ $^{18}\text{F}$ ]6 and got a selective  $\sigma_1$  ligand as a tumour imaging agent, 8-(4-(2-[ $^{18}\text{F}$ ]fluoroethoxy)benzyl)-1,4-dioxo-8-azaspiro[4.5]decane (Xie et al. 2015). This radiotracer showed twofold lower lipophilicity than lead compound [ $^{18}\text{F}$ ]6. Based on Glennon's pharmacophore model for  $\sigma_1$  receptor ligands, the aromatic ring is required as the primary hydrophobic region (Glennon 2005). However, the above results demonstrated that the 'primary hydrophobic region' in the Glennon's pharmacophore model is more flexible. Similar to the design concept of the above tumour radiotracer with low lipophilicity, Jia and co-workers designed the radiotracer 1-(4-[ $^{18}\text{F}$ ]fluorobenzyl)-4-[(tetrahydrofuran-2-yl)methyl]piperidine ([ $^{18}\text{F}$ ]8) (Fig. 27.4) constructed from a novel scaffold (He et al. 2017). Interestingly, compound 8 possessed low nanomolar affinity for  $\sigma_1$  receptors and high subtype selectivity ( $K_i = 3.2$  nM,  $\sigma_2/\sigma_1 = 52$ ). Moreover, this compound displayed negligible affinity for VACHT, adenosine  $A_{2A}$ , adrenergic  $\alpha_2$ , cannabinoid  $\text{CB}_1$ , dopamine  $\text{D}_1$  and  $\text{D}_2$ ,  $\gamma$ -aminobutyric acid A, NMDA, melatonin  $\text{MT}_1$  and  $\text{MT}_2$  and serotonin 5-HT $_1$  receptors. The corresponding radiotracer [ $^{18}\text{F}$ ]8 showed high stability in the mouse brain, i.e. over 95% of the radioactivity signal represented the intact tracer [ $^{18}\text{F}$ ]8 at 1 h postinjection. Studies in rodents (He et al. 2017) and cynomolgus monkeys (Jia et al. 2018) indicated that this radioligand displayed fast and high brain uptake, suitable kinetics

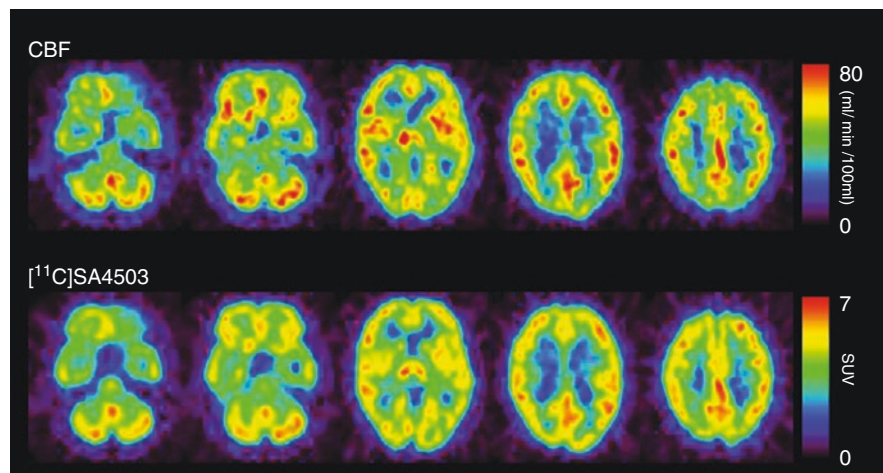
and high specific binding to sigma<sub>1</sub> receptors. Since this tracer possessed low lipophilicity, it showed highest  $f_p$  among the sigma<sub>1</sub> ligands evaluated to date. Most importantly, this radiotracer displayed higher specific signals ( $BP_{ND}$ ) than (*S*)-(-)-[<sup>18</sup>F]fluspidine. Hence, [<sup>18</sup>F]**8** warrants further evaluation in humans.

### 27.2.3 PET Imaging of the Sigma<sub>1</sub> Receptors in the Human Brain

Five radioligands, [<sup>11</sup>C]SA4503 (Ishii et al. 2001; Sakata et al. 2007), [<sup>18</sup>F]FPS (Waterhouse et al. 2004), [<sup>123</sup>I]TPCNE (Stone et al. 2006), (*S*)-(-)-[<sup>18</sup>F]fluspidine (Kranz et al. 2016; Becker et al. 2018; Meyer et al. 2018) and [<sup>18</sup>F]FTC-146 (Hjornevik et al. 2017; Shen et al. 2017), were used for in vivo investigations of the density of the sigma<sub>1</sub> receptor in the human brain by PET or SPECT; however, [<sup>18</sup>F]FPS and [<sup>123</sup>I]TPCNE showed irreversible binding profiles during the scan periods because of their high affinities. Therefore, no further clinical studies were conducted with those radiotracers.

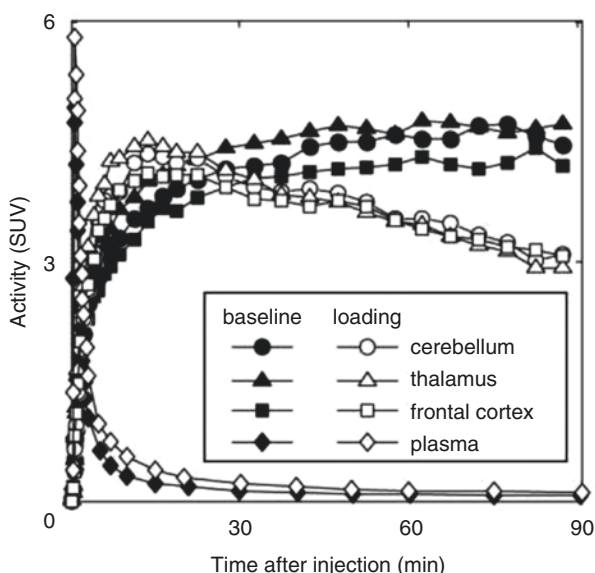
#### 27.2.3.1 Healthy Subjects

[<sup>11</sup>C]SA4503 was distributed in all grey matter regions in the human brain (Fig. 27.5). The distribution patterns did not differ between the early and late phases and were similar to those of regional cerebral blood flow (rCBF). The regional tissue time-activity curves (TACs) of [<sup>11</sup>C]SA4503 in the brain showed a gradual increase in radioactivity over 30 min and then reached a plateau, whereas the radioactivity in plasma decreased very rapidly (Fig. 27.6). These findings suggest that the regional



**Fig. 27.5** rCBF and sigma<sub>1</sub> receptors in the brain of a healthy human subject. rCBF (upper): rCBF was measured using PET and [<sup>15</sup>O]H<sub>2</sub>O and calculated as ml/min/100 ml tissue. [<sup>11</sup>C]SA4503 (lower): static images were acquired 40–60 min after injection of [<sup>11</sup>C]SA4503 and expressed as standardised uptake value (SUV: regional activity divided by administered dose per body weight)

**Fig. 27.6** TACs in the healthy human brain and plasma after intravenous injection of [ $^{11}\text{C}$ ]SA4503 under baseline and haloperidol-loading (3 mg) conditions. Decay-corrected radioactivity was expressed as SUV. Haloperidol was administered orally 18 h before injection of [ $^{11}\text{C}$ ]SA4503



distributions of [ $^{11}\text{C}$ ]SA4503 could depend on the blood flow or permeability rather than receptor density because of its slow dissociation kinetics.

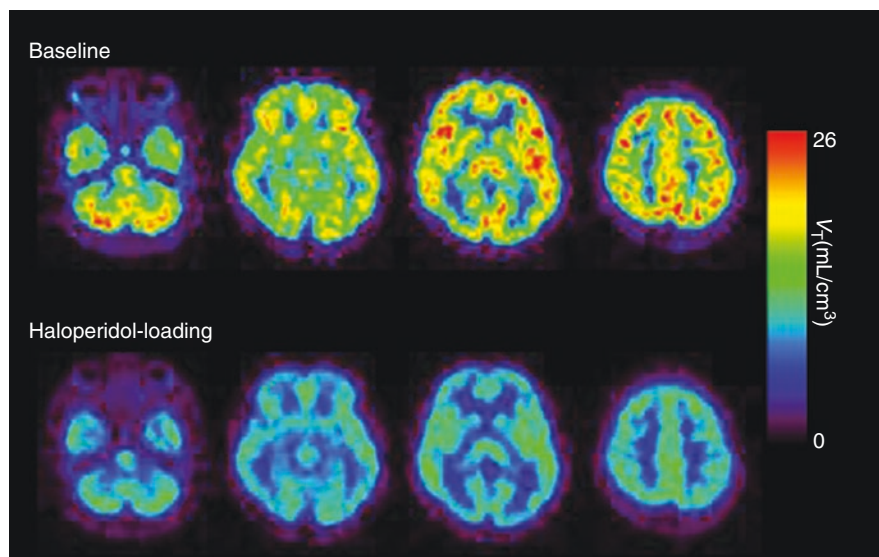
Preclinical studies confirmed age-dependent increases in  $\sigma_1$  receptors in the brains of non-human primates (Kawamura et al. 2003). These results contrast strikingly with the age-dependent decreases in several other receptors in the primate brain. For investigation of the neuroprotective functions of  $\sigma_1$  receptors, a PET study with [ $^{11}\text{C}$ ]SA4503 for evaluation of the aging process in the human brain would be of great interest.

### 27.2.3.2 Kinetic Analysis

In the traditional PET pharmacokinetics analysis for neuroreceptor radioligands, the ratio of  $k_3$  (association constant rate between the free plus nonspecifically and specifically bound compartments) over  $k_4$  (dissociation constant rate between the free plus nonspecifically and specifically bound compartments) in a two-tissue compartmental model (Mintun et al. 1984) is often used as the binding potential, which is currently defined as the specific binding potential relative to non-displaceable binding ( $BP_{\text{ND}}$ ) (Innis et al. 2007). As  $\sigma_1$  receptors are distributed throughout all brain regions, no suitable reference region with negligible specific binding is available. Therefore, in the kinetic analysis of [ $^{11}\text{C}$ ]SA4503 and (*S*)-(-)-[ $^{18}\text{F}$ ]fluspidine, arterial blood sampling measurements corrected by metabolite analysis are essential for obtaining the input function. In general, however, the variability of  $BP_{\text{ND}}$  estimated from direct compartmental analysis without any constraints is clearly larger than that with indirect methods, such as a reference tissue model analysis (Vilkman et al. 2000). In the case of [ $^{11}\text{C}$ ]SA4503, slow kinetics due to its slightly higher

affinity (Fig. 27.6) increased the variability of  $BP_{ND}$  in the time scale of the PET scan using a  $^{11}\text{C}$ -labelled tracer with a short half-life (20.4 min). An alternative parameter for the evaluation of [ $^{11}\text{C}$ ]SA4503 binding is the total volume of distribution ( $V_T$ ) estimated by the linear graphical method (Logan et al. 1990), although  $V_T$  includes both specific and nonspecific binding (Kimura et al. 2007). The estimation of  $V_T$  ( $K_1/k_2 \times (1 + k_3/k_4)$ ) in compartmental analysis tends to be more stable than that of  $BP_{ND}$ . Figure 27.7 shows representative  $V_T$  images of [ $^{11}\text{C}$ ]SA4503 in a healthy brain. The binding of [ $^{11}\text{C}$ ]SA4503 was high in the cerebellar cortex, moderate in the temporal and parietal cortices and low in the caudate and putamen (Sakata et al. 2007). These distribution patterns resemble the distribution patterns of radioligand binding in primate brains, where sigma receptors were noticed to be widely distributed with regionally different densities, using an in vitro binding assay and autoradiography (Weissman et al. 1988; Mash and Zabetian 1992; Shibuya et al. 1992).

An interaction between sigma<sub>1</sub> receptors and steroids has been established (Su et al. 1988; Collier et al. 2007). Currently, no information is available regarding the relationship between plasma hormone levels and [ $^{11}\text{C}$ ]SA4503 binding in the living human brain. If the binding of [ $^{11}\text{C}$ ]SA4503 is sensitive to competition by endogenous steroids, intra-subject variability in  $BP_{ND}$  of [ $^{11}\text{C}$ ]SA4503 in women during the menstrual cycle and inter-subject variability in aged subjects, including postmenopausal subjects, may be found. Therefore, plasma steroid levels should be considered in PET studies of patients and healthy volunteers.

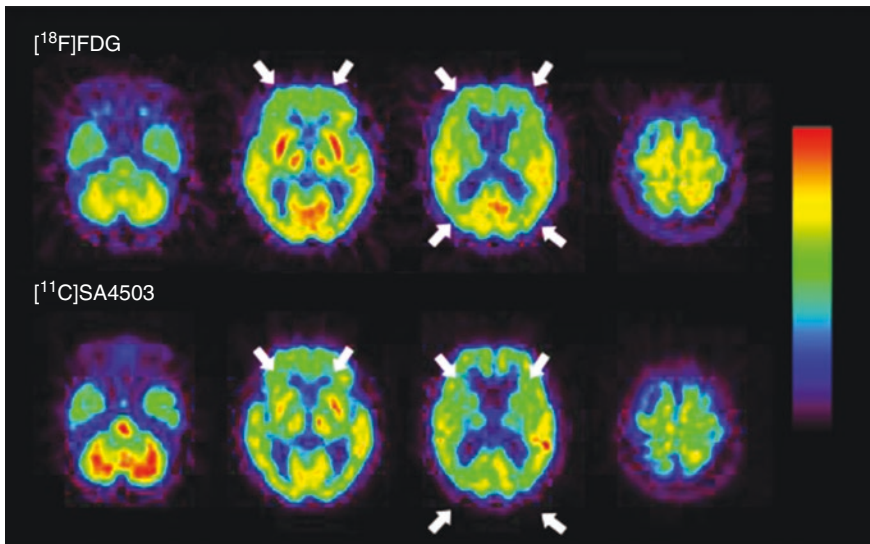


**Fig. 27.7**  $V_T$  images of [ $^{11}\text{C}$ ]SA4503 in the human brain under baseline (upper) and haloperidol-loading conditions (lower). The binding of [ $^{11}\text{C}$ ]SA4503 was considerably reduced after oral administration of haloperidol (3 mg, 18 h prior to the PET scan)

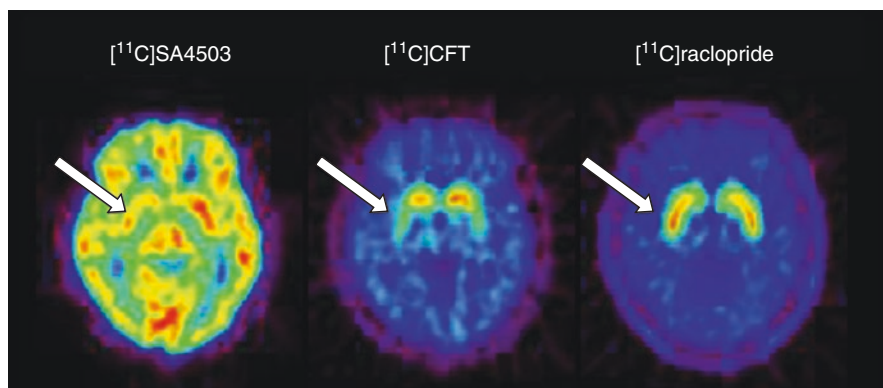
### 27.2.3.3 CNS Disease

Mishina et al. (2008) performed [ $^{11}\text{C}$ ]SA4503-PET in AD patients in the early stages of the disease. Figure 27.8 shows representative [ $^{11}\text{C}$ ]SA4503 images compared with those for glucose metabolism evaluated with [ $^{18}\text{F}$ ]FDG-PET. Although the results may have been influenced by atrophy or changes in regional blood flow, a widespread decrease in [ $^{11}\text{C}$ ]SA4503 binding in the brain was found;  $BP_{\text{ND}}$  for [ $^{11}\text{C}$ ]SA4503 was significantly lower in the cerebellum, thalamus and frontal, temporal and occipital cortices of AD patients than in those of healthy subjects. In other regions of the AD brain,  $BP_{\text{ND}}$  tended to be reduced. However, the data from these PET studies should be interpreted with caution because donepezil taken by some subjects shows a potent  $\sigma_1$  receptor agonist activity (see Sects. 2.3.4 and 3.4).

Parkinson's disease (PD) is also a progressive degenerative disorder of the CNS, characterised clinically by resting tremors, sluggish movements, cogwheel rigidity and postural instability. These symptoms are mainly caused by insufficient dopamine synthesis and death of dopaminergic neurons in the substantia nigra. Some studies have suggested that  $\sigma_1$  receptors are involved in modulating the synthesis and release of dopamine (Chaki et al. 1998). Using [ $^{11}\text{C}$ ]SA4503-PET, Mishina et al. (2005) investigated whether  $\sigma_1$  receptors were involved in the damage to the dopaminergic system in patients with PD, who showed low densities of dopamine transporters and normal or high densities of dopamine  $D_2$  receptors in the putamen using [ $^{11}\text{C}$ ]CFT and [ $^{11}\text{C}$ ]raclopride, respectively. Although the



**Fig. 27.8** Glucose metabolism and densities of  $\sigma_1$  receptors in the brain of a patient with AD. Glucose metabolism (upper): static image acquired 45–51 min after injection of [ $^{18}\text{F}$ ]FDG.  $\sigma_1$  receptors (lower):  $V_T$  derived from Logan graphical analysis of a 90-min dynamic scan of [ $^{11}\text{C}$ ]SA4503. In this patient, uptake of [ $^{18}\text{F}$ ]FDG was primarily decreased in the frontal and parietal cortex, and the distribution of [ $^{11}\text{C}$ ]SA4503 was consistent with this reduction



**Fig. 27.9** Sigma<sub>1</sub> receptors, dopamine transporters and dopamine D<sub>2</sub> receptors in the brain of a patient with PD. [<sup>11</sup>C]SA4503 (left),  $V_T$  derived from the Logan graphical analysis of a 90-min dynamic scan; [<sup>11</sup>C]CFT (middle); and [<sup>11</sup>C]raclopride (right), static image acquired 75–90 min and 40–55 min, respectively, after injection of the radiotracers

differences in  $BP_{ND}$  of [<sup>11</sup>C]SA4503 between normal and PD patients were not clear because of a large inter-subject variability,  $BP_{ND}$  in PD patients was significantly lower on the more affected side of the anterior putamen than on the less affected side (Fig. 27.9).

LID in patients with PD mimics acute dystonic reactions induced by antipsychotic drugs, possibly mediated by sigma receptors. A PET study using [<sup>11</sup>C]nemonapride showed a strong positive correlation between [<sup>11</sup>C]nemonapride binding and the preoperative LID severity score when the patients were receiving medication (Nimura et al. 2004).

Patients with untreated MDD were investigated with (*S*)-(-)-[<sup>18</sup>F]fluspidine in comparison with age-matched healthy controls. Increased cortico-(para-)limbic sigma<sub>1</sub> receptor availability was found in MDD, which was associated with the severity of acute depressive symptoms (Meyer et al. 2018). Furthermore, Teva Pharmaceutical Industries performed a study to evaluate sigma<sub>1</sub> and dopamine D<sub>2</sub> receptor occupancy by pridopidine in the human brain of healthy volunteers and in patients with HD. No results of that study were published so far.

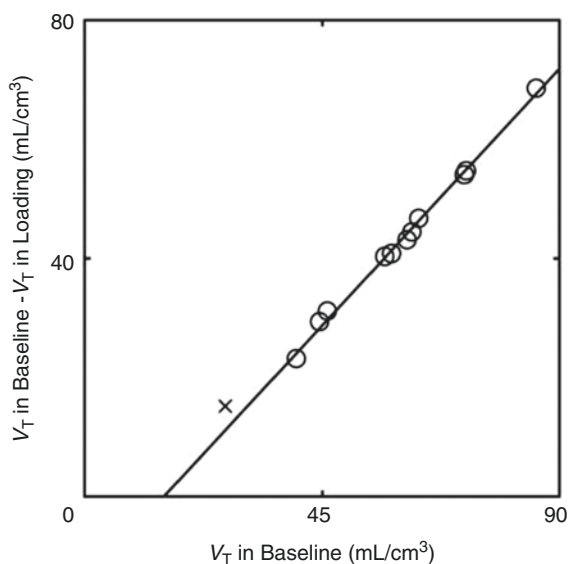
#### 27.2.3.4 Measurement of Sigma<sub>1</sub> Receptor Occupancy in the Human Brain

While the physiological and pathophysiological roles of sigma receptors remain under investigation, a number of neuropsychiatric drugs are known to have moderate to high affinities for sigma receptors. Moreover, sigma<sub>1</sub> receptor ligands represent a new class of therapeutic agents for neuropsychiatric disorders (Hashimoto and Ishiwata 2006; Ishikawa and Hashimoto 2010). Therefore, levels of sigma receptor occupancy of therapeutic drugs in the living human brain are of great interest. In a feasibility study, Ishiwata et al. (2006b) measured sigma<sub>1</sub> receptor and dopamine D<sub>2</sub> receptor occupancy by haloperidol in the human brain by PET using

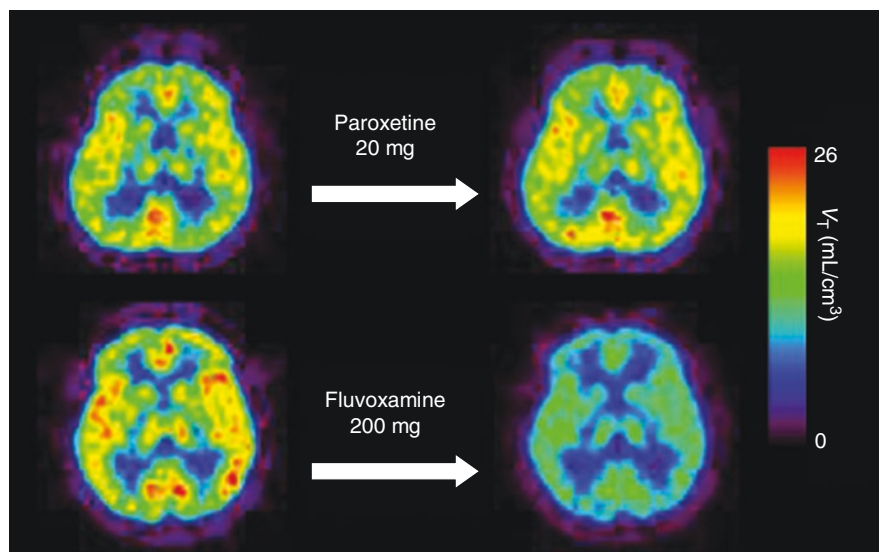
[ $^{11}\text{C}$ ]SA4503 and [ $^{11}\text{C}$ ]raclopride, respectively. Sigma $_1$  receptor occupancy was approximately 80% after oral administration of 3 mg haloperidol, whereas dopamine D $_2$  receptor occupancy was approximately 60%.

With respect to studies involving drugs with good potential for high receptor occupancy, a graphical analysis using a Lassen plot (Cunningham et al. 2010; Lassen et al. 1995) may be applicable for the evaluation of specific binding of the radioligand and receptor occupancy by the therapeutic drug, thus providing more stable estimates. The Lassen plot is based on regional changes in  $V_T$  between baseline and drug-loaded conditions. The assumptions for the analysis are that nonspecific binding is homogeneous, occupancy is the same in all regions of a regression line and there is a steady state of occupancy for the duration of the scan. A typical Lassen plot for haloperidol-loading study is shown in Fig. 27.10, and the parametric  $V_T$  images of [ $^{11}\text{C}$ ]SA4503 for baseline and loading conditions (Fig. 27.7) show that the binding of [ $^{11}\text{C}$ ]SA4503 was globally decreased by haloperidol-loading.

Ishikawa et al. (2007) measured sigma $_1$  receptor occupancy by two therapeutic drugs using [ $^{11}\text{C}$ ]SA4503-PET. Selective serotonin reuptake inhibitor (SSRI) is the



**Fig. 27.10** Typical Lassen plot analysis applied to a haloperidol-loading study. In this plot, the x-axis is  $V_T$  under baseline conditions, and the y-axis is the difference between  $V_T$  under loading and baseline conditions. The circles represent data from the grey matter regions (cerebral cortices, basal ganglia and cerebellar cortex), and the crosses represent data from the white matter region (centrum semiovale). The clear linear relationship suggests homogeneous nonspecific binding across the grey matter regions investigated, and the x-intercept and gradient of the regression line ( $y = 0.96(x - 15.1)$ ) represent the volume of free plus nonspecifically bound radioligand ( $V_{ND}$ ) and the receptor occupancy by nonradioactive haloperidol, respectively. Moreover, the absence of data near the x-axis indicates that no regions, including the white matter, are available as a true reference region devoid of specific binding and that most [ $^{11}\text{C}$ ]SA4503 binding under baseline conditions is specific



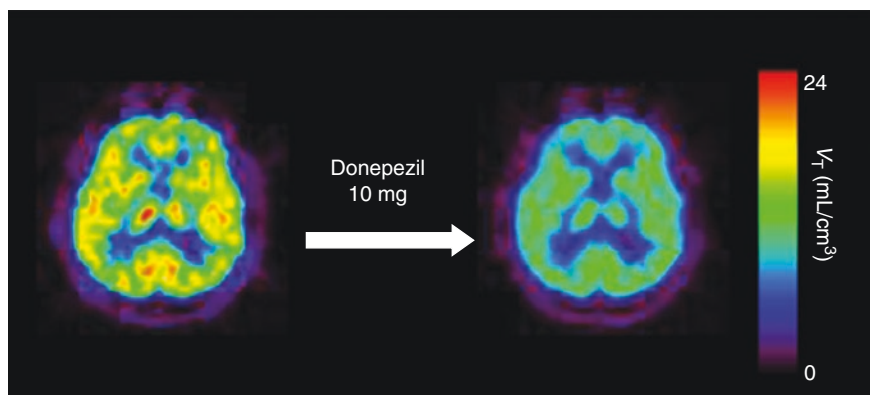
**Fig. 27.11**  $V_T$  images of [<sup>11</sup>C]SA4503-PET before and after a single oral administration of an SSRI. The upper pair represents  $V_T$  images at baseline (left) and at paroxetine (20 mg)-loading (right) in the same subject. The lower pair shows  $V_T$  images at baseline (left) and fluvoxamine (200 mg)-loading (right) in another subject

treatment of choice for many disorders, including major depressive disorder, dysthymia, obsessive–compulsive disorder and obsessive–compulsive spectrum disorders. Fluvoxamine has moderate affinity ( $K_i = 36$  nM) for sigma<sub>1</sub> receptors in addition to the main affinity for serotonin reuptake sites (Narita et al. 1996). A single administration of therapeutic doses of fluvoxamine (50–200 mg) decreased [<sup>11</sup>C]SA4503 binding in the human brain in a dose-dependent manner, whereas that of paroxetine (20 mg), another SSRI with very low affinity for sigma<sub>1</sub> receptors ( $K_i = 1893$  nM), did not (Fig. 27.11). A similar occupancy study of an antipsychotic drug was also preliminarily described (van Waarde et al. 2011). The second example evaluated was donepezil. This drug has high affinity for sigma receptors ( $IC_{50} = 14$  nM measured with [<sup>3</sup>H]DTG) (Kato et al. 1999). Figure 27.12 shows that a single administration of donepezil (5 or 10 mg) decreased [<sup>11</sup>C]SA4503 binding in the human brain in a dose-dependent manner (Ishikawa et al. 2009). The levels of sigma<sub>1</sub> receptor occupancy by fluvoxamine and donepezil were approximately 60% and 75%, respectively. These findings suggest that sigma<sub>1</sub> receptors may be involved in the mechanism of action of fluvoxamine and donepezil.

### 27.3 Sigma Receptors in CNS Diseases

Since the prototypic sigma<sub>1</sub> receptor ligand SKF10,047 (sigma<sub>1</sub>  $K_i = 44.8$  nM, sigma<sub>2</sub>/sigma<sub>1</sub> = 95.1; (Walker et al. 1990)) was found in the 1980s to exhibit psychotomimetic actions, the antipsychotic potential of sigma<sub>1</sub> receptor ligands has





**Fig. 27.12** Representative  $V_T$  images of [ $^{11}\text{C}$ ]SA4503-PET before and after a single oral administration of donepezil (10 mg) in a healthy subject. Left, image at baseline; right, image after donepezil (10 mg)-loading

been examined extensively. During the 1990s,  $\sigma_1$  receptor ligands were synthesised mostly to develop new drugs for the treatment of schizophrenia. However, after cloning of the  $\sigma_1$  receptor gene, molecular biological characterisation indicated relationships between  $\sigma_1$  receptors and a variety of human diseases, including mood disorders, stroke, neurodegenerative diseases and drug addiction.

### 27.3.1 Schizophrenia

Five  $\sigma_1$  receptor ligands, panamesine (EMD57445), eliprodil (SL82.0715), rimcazole (BW234U), BMY14802 (BMS181100) and DuP734, were introduced into clinical trials for the treatment of schizophrenia (Hayashi and Su 2004). The findings of these clinical trials suggest that  $\sigma_1$  receptor ligands may not possess potent antipsychotic actions against the positive symptoms of schizophrenia, but may be useful for ameliorating certain negative symptoms (Hayashi and Su 2004). Recently, it was reported that fluvoxamine, an SSRI possessing potent  $\sigma_1$  receptor agonistic activity, improves negative symptoms and cognitive deficits of schizophrenic patients undergoing antipsychotic treatment (Iyo et al. 2008; Niitsu et al. 2010).

Using [ $^3\text{H}$ ]haloperidol ( $\sigma_1$   $K_i = 0.3$  nM,  $\sigma_2/\sigma_1 = 120$ ; (Walker et al. 1990)) in the presence of spiperone to block binding to dopamine  $D_2$  receptors, Weissman et al. (1991) first reported reductions in the density of  $\sigma$  binding sites in schizophrenic patients on antipsychotic medication; these reductions were most prominent in the temporal cerebral cortex followed by the parietal cortex. Shibuya et al. (1992) measured  $\sigma$  receptor binding with [ $^3\text{H}$ ]DTG ( $\sigma_1$   $K_i = 11.9$  nM,  $\sigma_2/\sigma_1 = 3.2$ ; (Walker et al. 1990)) in 17 areas of the cerebral cortex and

found no significant differences between schizophrenic patients and controls, with exception of the superior parietal cortex in which the binding was significantly increased in the schizophrenic group. In addition, no significant differences were observed between off-drug and on-drug schizophrenic patients in any brain areas. Helmeste et al. (1996) reported no differences in the binding of [<sup>3</sup>H]nemonapride (previously designated as [<sup>3</sup>H]YM-09151-2: sigma<sub>1</sub>  $K_i$  = 8.4 nM, sigma<sub>2</sub>/sigma<sub>1</sub> = 1.1; (Ujike et al. 1996)) in the presence of spiperone in the frontal cortex and cerebellum; however, they observed a decrease in the caudate nucleus. The discrepancies between these results may have been due to differences in schizophrenia types, medication period, quantification methods (receptor binding assay or autoradiography) and radioligands used.

Three out of four case-control association studies failed to find significant associations between alleles of the sigma<sub>1</sub> receptor gene [G-241 T/C-240 T and Gln2Pro (A61C)] and schizophrenia (Uchida et al. 2003).

### 27.3.2 Mood Disorders

Recent animal and human studies demonstrated a tight link between sigma<sub>1</sub> receptors and higher-order brain functions, such as mood and cognition (Hayashi and Su 2004; Fishback et al. 2010). It is noteworthy that some clinically used antidepressants, such as fluvoxamine (sigma<sub>1</sub>  $K_i$  = 36 nM), possess affinities for sigma<sub>1</sub> receptors within the nanomolar concentration range (Narita et al. 1996).

Brain-derived neurotrophic factor (BDNF) plays a critical role in the pathophysiology and in the activity of therapeutic agents in patients with mood disorders (Hashimoto et al. 2004). Furthermore, altered expression of BDNF is considered to be a mechanism of action of antidepressant drugs and mood stabilisers (Hashimoto et al. 2004). Interestingly, cutamesine (SA4503), a selective sigma<sub>1</sub> receptor agonist, was also shown to increase BDNF in the hippocampus of rats (Kikuchi-Utsumi and Nakai 2008). Very recently, Fujimoto et al. (2012) found that cutamesine potentiates the post-translational processing of neurotrophins; in contrast, clinically used antidepressants promote the transcriptional upregulation of BDNF.

The sigma<sub>1</sub> receptor contains binding sites for neurosteroids. It was reported that serum dehydroepiandrosterone (DHEA) and DHEA sulphate levels are altered in patients with depression (Takebayashi et al. 1998). Administration of DHEA is associated with the improvement of symptoms in cases of major depression or dysthymia (van Broekhoven and Verks 2003).

Igmesine (JO1,784) is the first sigma<sub>1</sub> receptor agonist introduced into clinical trials for the treatment of depression. The clinical trial moved into Phase III; however, the antidepressant effect failed to be confirmed in a large sample study (Hayashi et al. 2011). Recently, cutamesine has been introduced into Phase II clinical trials for the treatment of depression (Urfer et al. 2009).

### 27.3.3 Ischemia

There is substantial experimental evidence that sigma receptors play a role in the modulation of ischemic neuronal injury (Maurice and Lockhart 1997; O'Neill et al. 1995). *In vitro* studies demonstrated that sigma<sub>1</sub> agonists exert potent protective effects in a variety of cell types, such as primary cerebral neurons (Yang et al. 2007), retinal ganglion cells (Ha et al. 2011) and lens cells (Wang and Duncan 2006). Similarly, sigma<sub>1</sub> agonists were reported to show robust neuroprotective effects in animal studies. 4-Phenyl-1-(4-phenylbutyl)piperidine (PPBP) prevented early brain injury in rat (Takahashi et al. 1996) and cat (Takahashi et al. 1995) models of transient focal ischemic stroke, suggesting that sigma receptors play an important role in the mechanism of acute injury after transient focal ischemia. Cutamesine has been introduced into Phase II clinical trials for the treatment of post-stroke neurological disturbances (Urfer et al. 2009).

### 27.3.4 Neurodegenerative Diseases

[<sup>3</sup>H]DTG binding was measured in seven hippocampal regions in the brains of Alzheimer's disease (AD) patients by autoradiography; an average reduction of 26% was found, which correlated with an average 29% pyramidal cell loss in the same region (Jansen et al. 1993). One study suggested that haplotype TT-241-240P2 of the sigma<sub>1</sub> receptor gene, which could lead to a reduction of sigma<sub>1</sub> receptor transcription, may be a protective factor against AD (Uchida et al. 2005). However, another study failed to replicate this finding (Maruszak et al. 2007).

Donepezil, an acetylcholinesterase (AChE) inhibitor, is the most widely prescribed drug so far for AD. Donepezil was found to be protective against amyloid β<sub>25–35</sub> peptide-induced neurotoxicity in mice (Meunier et al. 2006). This effect of donepezil was antagonised by the sigma<sub>1</sub> receptor antagonist BD-1047. *In vitro* studies demonstrated that donepezil, but not the AChE inhibitor physostigmine (also known as eserine), significantly potentiated nerve growth factor (NGF)-induced neurite outgrowth in PC12 cells in a concentration-dependent manner and that the effect of donepezil could be antagonised by NE-100 (Ishima et al. 2008). Furthermore, donepezil, but not physostigmine, significantly improved phencyclidine-induced cognitive impairments in mice, and the effect of donepezil could be antagonised by co-administration of NE-100 (Kunitachi et al. 2009). In summary, it is likely that sigma<sub>1</sub> receptors are involved in the mechanism of the neuroprotective pharmacological action of donepezil.

### 27.3.5 Drug Addiction and Alcoholism

Numerous animal studies have demonstrated a relationship between cocaine dependence and sigma<sub>1</sub> receptors (Robson et al. 2012). Many of these studies suggested the potential of sigma<sub>1</sub> antagonists to ameliorate cocaine intoxication, dependence and craving, although these effects have not been tested in clinical studies. Cocaine

at clinically relevant concentrations can interact with sigma<sub>1</sub> receptors and upregulate sigma<sub>1</sub> receptors in animals (Liu and Matsumoto 2008).

Despite the fact that substantial preclinical studies demonstrated roles of sigma receptors in drug abuse, only two clinical studies have been published to date in this regard, both of which were gene-association studies. The association of the sigma<sub>1</sub> receptor polymorphism GC-241-240TT or A61C (Gln2Pro) with methamphetamine dependence was examined in a Japanese cohort study of 143 methamphetamine abusers and 181 control subjects (Inada et al. 2004). This study indicated no significant association between sigma<sub>1</sub> receptor polymorphisms and methamphetamine abuse, although the frequency of the CC genotype of A61C tended to be higher in methamphetamine abusers who had experienced spontaneous relapse of methamphetamine psychosis. One gene-association study indicated a link between sigma<sub>1</sub> receptor polymorphisms and alcoholism (Miyatake et al. 2004). The distribution of sigma<sub>1</sub> receptor gene polymorphisms was analysed in 307 alcoholic and 302 control subjects. The frequencies of the A-485 allele and the TT-241-240/Pro2 haplotype in the 5'-upstream region, which lower transcriptional activity of the sigma<sub>1</sub> receptor gene, were significantly higher in control subjects compared with alcoholic subjects. These results suggest that the A-485 allele or TT-241-240/Pro2 haplotype lowers the expression of sigma<sub>1</sub> receptors and may be a possible protective factor against alcoholism.

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## 27.4 Conclusion

Sigma<sub>1</sub> receptors are implicated in the pathophysiology of some neurological and neuropsychiatric disorders. A number of radioligands have been developed and evaluated for imaging of sigma<sub>1</sub> receptors in the CNS by PET and SPECT. [<sup>11</sup>C]SA4503 was used to generate new findings regarding brain functions in patients with AD and PD, while (*S*)-(-)-[<sup>18</sup>F]fluspidine was used in patients with MDD and HD. Moreover, [<sup>11</sup>C]SA4503-PET demonstrated that some therapeutic drugs for neurological and neuropsychiatric disorders have considerable affinity for sigma<sub>1</sub> receptors in the human brain. Sigma<sub>1</sub> receptor imaging could be useful for pathophysiological studies of the human brain and help in the discovery and validation of potential therapeutic drugs for several CNS disorders. However, clinical sigma<sub>1</sub> receptor imaging is only just beginning to be explored. In view of the recent progress in sigma<sub>1</sub> receptor research and accumulating evidence of the therapeutic potentials of sigma<sub>1</sub> ligands, introduction of new tracers and further clinical studies should be encouraged.

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# Sigma-2 Receptors: An Emerging Target for CNS PET Imaging Studies

# 28

Aladdin Riad, Jinbin Xu, and Robert H. Mach

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## Abstract

The sigma-2 ( $\sigma_2$ ) receptor represents one of the most poorly understood proteins in cell biology. Although this receptor was identified through in vitro binding studies over 30 years ago, the molecular identity of this protein was not unambiguously determined until 2017. It is now known that the  $\sigma_2$  receptor is a pro-

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tein called TMEM97 and that this protein forms a trimeric complex with PGRMC1 and the LDL receptor to increase the rate of internalization of lipoproteins. There is overwhelming data demonstrating that the  $\sigma_2$  receptor is an important biomarker of tumor cell proliferation, which is consistent with the lipoprotein uptake mechanism associated with this protein. In the CNS,  $\sigma_2$  receptor antagonists block A $\beta$ 1-42 oligomer synaptic dysfunction in transgenic mouse models of Alzheimer's disease and have shown promise in animal models of traumatic brain injury and alcohol abuse. These results clearly indicate that this protein is an important therapeutic target for the treatment of a variety of CNS conditions. These observations highlight the importance of having high affinity  $\sigma_2$ -selective radiotracers for imaging this protein in vivo with PET.

## Abbreviations

A $\beta$	Beta amyloid protein
CNS	Central nervous system
LDL	Low density lipoprotein
PET	Positron emission tomography
PGRMC1	Progesterone membrane binding component 1
TMEM97	Transmembrane protein 97
$\sigma$	Sigma

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## 28.1 Introduction

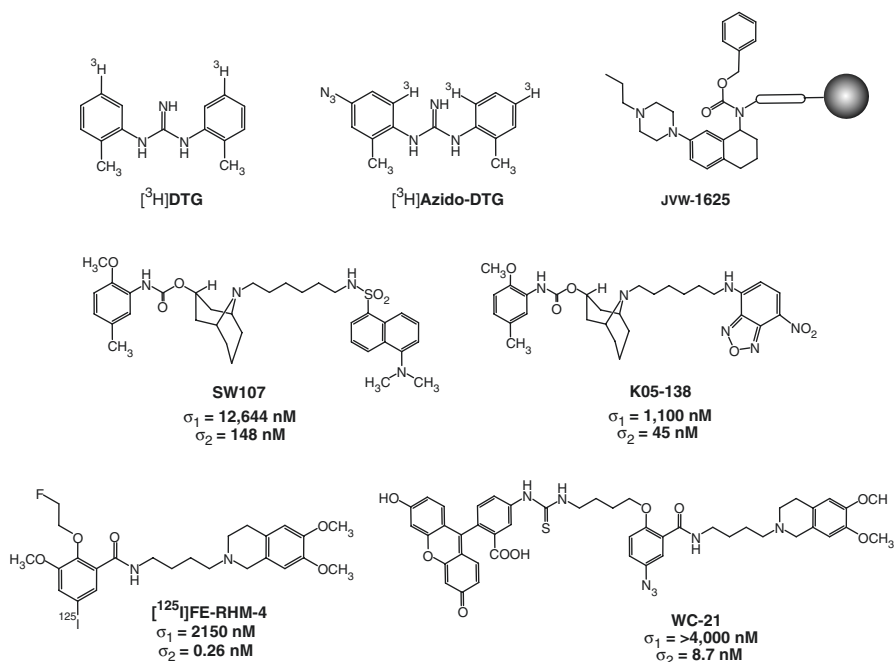
The sigma ( $\sigma$ ) receptors represent one of the most poorly understood protein families in cell biology. Their name was derived from their initial identification as a subtype of the opiate receptors (Martin et al. 1976). This characterization was based on the binding properties and behavioral properties of the benzomorphan analog SKF10047 and that these properties could be blocked by naltrexone, an opioid receptor antagonist. Subsequent studies revealed that the (–)-enantiomer of SKF10047 was a nonselective opioid ligand, acting as a partial agonist at mu and delta receptors, and an antagonist at kappa opioid receptors (Su 1982). On the other hand, (+)-SKF10047 had a high affinity for the  $\sigma$  receptor. By conducting a series of radioligand binding and photoaffinity labeling studies, Hellewell and Bowen were able to demonstrate that there were two subtypes of the  $\sigma$  receptor,  $\sigma_1$  and  $\sigma_2$  (Bowen et al. 1989; Hellewell and Bowen 1990). The  $\sigma_1$  receptor had a molecular weight of ~25 kDa, and [ $^3$ H](+)-SKF10047, [ $^3$ H](+)-pentazocine, and [ $^3$ H]DTG had a high affinity for this binding site. A second  $\sigma$  receptor was found to have a MW of ~21.5 kDa and had a high affinity for [ $^3$ H]DTG but did not bind the benzomorphan analogs [ $^3$ H](+)-pentazocine and [ $^3$ H](+)-SKF10047. This receptor was named the  $\sigma_2$  receptor (Hellewell and Bowen 1990).

Of the two receptor subtypes, much more attention has been placed on the  $\sigma_1$  versus  $\sigma_2$  receptor. This has resulted in publication of a number of key studies including its purification and cloning, resolution of the crystal structure as a homotrimeric complex, subcellular localization, molecular chaperone properties, and its ability to interact with and form heteromeric complexes with G proteins including the opiate and dopamine D2 receptors (Alon et al. 2017a; Beggiato et al. 2017; Borroto-Escuela et al. 2019; Hanner et al. 1996; Hayashi and Su 2007; Navarro et al. 2010; Schmidt et al. 2016). Although less is known about the structure and function of the  $\sigma_2$  receptor, a number of seminal studies have been recently reported that have led to a greater understanding of the function of this receptor in cell biology.

## 28.2 Pharmacological and Molecular Characterization of the $\sigma_2$ Receptor

### 28.2.1 In Vitro Binding Studies

For many years, much of what was known about the  $\sigma_2$  receptor was obtained using radioligand binding studies with tritiated ligands. Photoaffinity labeling studies with [ $^3\text{H}$ ]azido-DTG (Fig. 28.1) were able to determine that the molecular weight of the  $\sigma_2$  receptor was  $\sim 21.5$  kDa, which was smaller than the  $\sigma_1$  receptor (Hellewell



**Fig. 28.1** Structures of the compounds and radioligands used to characterize the  $\sigma_2$  receptor

et al. 1994). In vitro binding studies with [ $^3\text{H}$ ]DTG in the presence of (+)-pentazocine to mask  $\sigma_1$  receptors were useful in studying the density of the protein in various tissues including cancer cells (Hellewell et al. 1994; Vilner et al. 1995). Radioligand binding assays with this ligand have also been instrumental in identifying compounds having a higher affinity and greater selectivity for  $\sigma_2$  versus  $\sigma_1$  receptors, which have served as more optimal ligands for in vitro and in vivo imaging studies.

One of the more important observations made with in vitro binding studies of [ $^3\text{H}$ ]DTG was that  $\sigma_2$  receptors were expressed in high density in a broad panel of cancer cell lines derived from both murine and human tumors (Vilner et al. 1995). This indicated that the  $\sigma_2$  receptor was a potential biomarker for differentiating tumors from the surrounding normal tissue. Another important observation was that the density of  $\sigma_2$  receptors was found to be tenfold higher in mouse mammary adenocarcinoma (66) cells when they were actively progressing through the cell cycle (i.e., proliferating) versus 66 cells that were driven into a state of quiescence (66Q cells) via nutrient deprivation (Mach et al. 1997). The density of  $\sigma_2$  receptors in proliferating 66 (66P) cells was also relatively high, ~one million copies/cell. Furthermore, the time interval for the downregulation of  $\sigma_2$  receptors once the 66 cells entered a quiescent state was ~5 days, which is consistent with the downregulation of a membrane-bound protein (Al-Nabulsi et al. 1999; Wheeler et al. 2000). Finally, the tenfold difference in  $\sigma_2$  receptor density between 66P and 66Q cells was observed in solid tumor xenografts, which indicated that this was not a tumor property limited to cell culture techniques (Shoghi et al. 2013; Wheeler et al. 2000).

### 28.2.2 Microscopy Studies

The next phase in the study of the  $\sigma_2$  receptor involved two-photon and confocal microscopy studies using fluorescently tagged small molecules possessing a high affinity for  $\sigma_2$  versus  $\sigma_1$  receptors. Although a number of these probes have been reported, studies that have yielded the most useful information have utilized the probes **SW-107** and **K05-138** in cancer cells in combination with a series of “tracker” dyes to determine the subcellular localization of the  $\sigma_2$ -targeting fluorescent probes. In these studies, the fluorescent signal of the  $\sigma_2$  receptor ligand was co-localized with tracker dyes for lysosomes/endosomes (LysoTracker), endoplasmic reticulum (ER-Tracker), mitochondria (MitoTracker), and the plasma membrane (FM-1-43FX). These studies demonstrated a co-localization of the  $\sigma_2$  receptor fluorescent ligands with each of the different tracker dyes (Zeng et al. 2007, 2011). In addition, there was a rapid uptake of **K05-138** in time lapsed confocal microscopy studies, and this uptake could be delayed by incubation with the known endocytosis inhibitor phenylarsine oxide. These data indicated that the fluorescent probes undergo receptor-mediated endocytosis into endosomes and that once inside the cell, the fluorescent probes localize to the ER and mitochondria. The receptor-mediated endocytosis of the  $\sigma_2$  receptor fluorescent probes was also consistent with earlier reports of the localization of  $\sigma_2$

receptors in lipid rafts, which play an important role in membrane trafficking (Gebreselassie and Bowen 2004; Zeng et al. 2007).

The observation that the fluorescent probes **SW-107** and **K05-138** undergo receptor-mediated endocytosis and localization in mitochondria led to the use of the [3.3.1] aza-bicyclononane scaffold as a means for delivering anticancer drugs to cancer cells (Hornick et al. 2012; Mach et al. 2013). One of the more interesting results in this series of studies was the observation that this  $\sigma_2$  receptor scaffold increased the rate of uptake of gold nanocages by breast cancer cells; transmission electron microscopy determined that the gold nanocages have a size of ~50 nm and suggested that the “endogenous ligand” binding to the  $\sigma_2$  receptor is likely to have a high molecular weight (Sun et al. 2014). Binding to the  $\sigma_2$  receptor leads to the receptor-mediated internalization of this high molecular weight endogenous ligand into the endosomal compartment. This information was useful in ultimately identifying the structure of the  $\sigma_2$  receptor as discussed in detail below.

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## 28.3 Identification of the $\sigma_2$ Receptor and Its Putative Biological Function

### 28.3.1 Progesterone Receptor Membrane Component 1 (PGRMC1)

Although the above radioligand binding and microscopy studies were able to provide valuable information regarding the molecular weight and subcellular localization of the  $\sigma_2$  receptor, they did not address the most important questions: (1) What is the  $\sigma_2$  receptor? (2) What is its role in cell biology? (3) Why is this protein upregulated in proliferating cancer cells? The only way this could be accomplished was through the purification, sequencing and cloning of the  $\sigma_2$  receptor, and determining the biochemical pathways that utilize this protein.

In order to accomplish this goal, Xu and colleagues (2011) developed a photoaffinity probe based on the conformationally flexible benzamide analogs (Fig. 28.1). These compounds have been shown to have a high affinity for the  $\sigma_2$  versus  $\sigma_1$  receptor, and PET probes based on this class of compounds demonstrated promising results in PET imaging studies of the  $\sigma_2$  receptor (Tu et al. 2005, 2010). The photoaffinity probe also contained a FITC moiety; although it is typically used as a fluorescent probe, this moiety was chosen since there are highly specific antibodies for FITC that could be used to visualize photoaffinity-labeled proteins using classic molecular biology techniques. In this study,  $\sigma_2$  receptors in rat liver membrane homogenates, the tissue used in standard  $\sigma_2$  receptor binding assays, were labeled with the photoaffinity probe under a series of protein enrichment techniques. Western blot analysis using an anti-FITC antibody revealed a protein band having a molecular weight of ~24 kDa, which was larger than the 21.5 kDa band identified by Hellewell and Bowen (Hellewell et al. 1994). This protein was isolated and submitted to two different protein sequencing labs; there was only one protein common to both lists of candidate proteins, the progesterone receptor membrane component 1

or PGRMC1, which has also been termed the 25-Dx protein (Xu et al. 2011). Previous studies have shown that the molecular weight of the PGRMC1 (25-Dx protein) ranged between 22 and 25 kDa in polyacrylamide gels. Interestingly, the PGRMC1 was previously shown to be upregulated in a wide panel of tumors and was proposed as a biomarker for breast cancer (Neubauer et al. 2008). The characterization of protein expression levels of PGRMC1 in tumors on the Human Protein Atlas database paralleled very closely the density of  $\sigma 2$  receptors in cancer cells growing under cell culture techniques reported in the paper by Vilner et al. (1995). In addition, the PGRMC1 was thought to not be functionally active as a monomer but was thought to form either a homodimer in the internalization of heme or a partner protein with a number of other proteins including Insig-1, PAIR-BR1, and EGFR (Ahmed et al. 2010a, b; Cahill 2007).

Since the PGRMC1 was a known protein, it was possible to obtain a number of standard molecular biology reagents to study expression levels under a variety of experimental conditions. This led to a series of studies relating the binding of the radioiodinated  $\sigma 2$  ligand, [ $^{125}$ I]FE-RHM-4, to HeLa cells in which PGRMC1 was either transiently knocked down using siRNA for this protein or transiently upregulated via the transfection of HeLa cells with human PGRMC1 cDNA. Since it was not possible to isolate enough HeLa cells to conduct full Scatchard analysis of the siRNA or overexpression experiments, the binding studies were conducted using a single concentration of [ $^{125}$ I]FE-RHM-4, and the assumption was made that either knocking out or overexpressing PGRMC1 did not alter the Kd value of this radioligand.

In the siRNA experiments, the transient knockdown of PGRMC1 led to a reduction in binding of [ $^{125}$ I]FE-RHM-4. Similarly, transfection of human PGRMC1 cDNA in HeLa cells led to an increase in binding of this radioligand. However, the authors noticed that the level of binding of [ $^{125}$ I]FE-RHM-4 did not match the level of PGRMC1 protein measured via Western blot analysis (Xu et al. 2011). This disconnect between protein levels and radioligand binding data led the authors to conclude that the  $\sigma 2$  receptor was not directly the PGRMC1 but likely represented a binding site on a protein complex involving the PGRMC1 and one of its binding partners. Evidence consistent with the involvement of PGRMC1 with  $\sigma 2$  receptors was reported by Izzo et al. (2014b) and Yi et al. (2017).

In 2015, two papers were published that disputed the results reported by Xu et al. (2011). In the study reported by Abate and colleagues, a stable knockout or overexpression of PGRMC1 in the human breast cancer cell line, MCF-7 cells, did not affect the binding of [ $^3$ H]DTG (Abate et al. 2015). A similar result was reported by Ruoho and colleagues using NSC34 cells using overexpression and CRISPR to knockout the protein (Chu et al. 2015). Both papers stated that the conclusions reached by Xu et al. were incorrect, even though the authors never claimed that the PGRMC1 was the  $\sigma 2$  receptor (Xu et al. 2011). This misquoting of the conclusions stated in the Xu et al. paper has been repeated in subsequent review articles and editorial articles on the  $\sigma 2$  receptor (Blass and Rogers 2018; Hiranita 2016; Schmidt and Kruse 2019).



### 28.3.2 Identification of the Gene Encoding the $\sigma_2$ Receptor

In 2017, Alon et al. published a seminal study that would provide key information to the identity and molecular function of the  $\sigma_2$  receptor (Alon et al. 2017b). In this paper, the authors used affinity chromatography in which the agarose beads were covalently modified with **JVW-1625**, a high affinity  $\sigma_2$  receptor ligand (Fig. 28.1). The authors were able to identify a small panel of candidate proteins from these chromatography experiments, and the protein that displayed high binding of [ $^3$ H]DTG proved to be TMEM97, which has also been named MAC30, a membrane protein previously implicated in a number of cancers. TMEM97 is also a binding partner of the protein NPC1. NPC1 is a lysosomal cholesterol transporter, and loss of this protein leads to the fatal lysosomal storage disorder Niemann-Pick disease type C1.

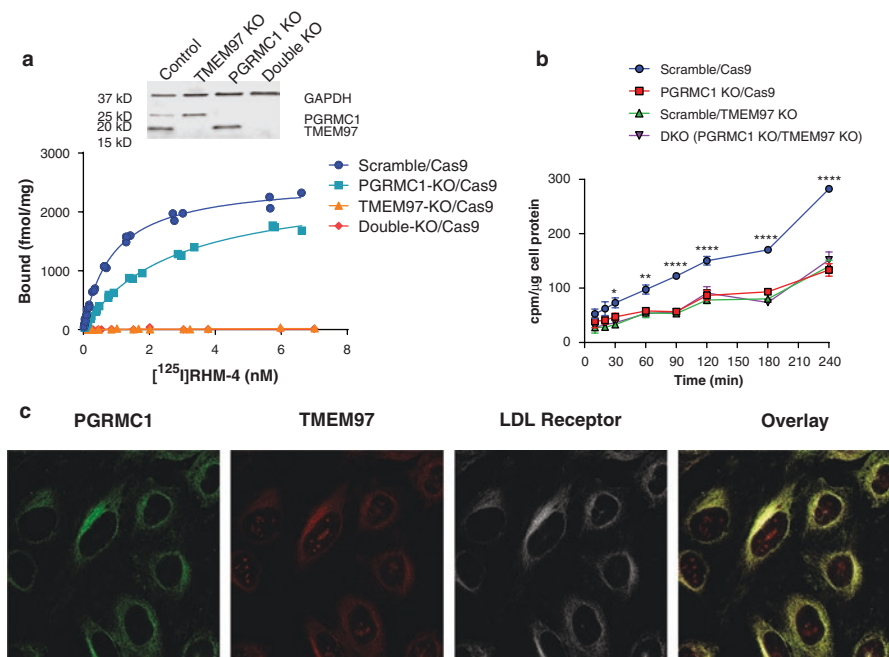
The authors of this study found that siRNA knockdown of TMEM97 in PC-12 cells resulted in a large reduction in binding of [ $^3$ H]DTG. Furthermore, overexpression of TMEM97 in sf9 cells, an insect cell line that does not express TMEM97, resulted in high binding of [ $^3$ H]DTG. Overexpression of PGRMC1 in the sf9 cells did not result in an increase in binding of [ $^3$ H]DTG, which was consistent with the data reported by Abate et al. (2015) and Chu et al. (2015). Finally, introduction of point mutations in several critical residues (Asp56 and Asp29) resulted in a loss of binding of [ $^3$ H]DTG. Taken collectively, these data confirm that the molecular identification of the  $\sigma_2$  receptor protein, and the gene that encodes that protein, is TMEM97. This observation has led to this protein being termed “ $\sigma_2$  receptor/TMEM97” in some literature reports (Sahn et al. 2017; Scott et al. 2018a; Vazquez-Rosa et al. 2019a).

### 28.3.3 The Role of the $\sigma_2$ Receptor in Cholesterol Trafficking

The data described above proved to be instrumental in the identification of the role of the  $\sigma_2$  receptor in cell biology. First, the identification of the  $\sigma_2$  receptor as TMEM97, and the association of this protein with NPC1, strongly implicated this receptor in cholesterol trafficking (Alon et al. 2017b). Second, identification of the  $\sigma_2$  receptor as a binding site in the PGRMC1 protein complex also implicated this protein in cholesterol trafficking since the PGRMC1 has been shown to interact with the LDL receptor, a key process in the cellular uptake of cholesterol (see review by Cahill 2007). Third, the increased rate of internalization of gold nanocages functionalized with the  $\sigma_2$  agonist, **SV119** (Sun et al. 2014), suggested that the endogenous ligand for the  $\sigma_2$  receptor had a high molecular weight; the size of LDL particles range between 22 and 27.5 nm, which is smaller than the size of the gold nanocage described above. Taken collectively, these data suggested that the PGRMC1 protein complex containing the “putative  $\sigma_2$  receptor binding site” consisted of a trimeric complex of  $\sigma_2$  receptor/TMEM97, PGRMC1, and the LDL receptor.

Using the above information, Riad et al. (2018) conducted a series of studies that confirmed the existence of this trimeric complex. Using HeLa cells and the CRISPR/cas 9 gene editing, this group found that a complete knockout of TMEM97 (TMEM97(-) cells) led to a complete loss of binding of [ $^{125}$ I]RHM-4 and a dramatic (but not complete) reduction in the binding of [ $^3$ H]DTG (Fig. 28.2). A knockout of PGRMC1 in the HeLa cells (PGRMC1(-) cells) resulted in no change in the density of  $\sigma 2$  receptors measured with both radioligands but a significant reduction in the *Kd* value of both [ $^{125}$ I]RHM-4 and [ $^3$ H]DTG. Finally, knocking out both TMEM97 and PGRMC1 in HeLa cells (i.e., double knockout or TMEM97(-)/PGRMC1(-) cells) had the same effect as knocking out only TMEM97 (Fig. 28.2).

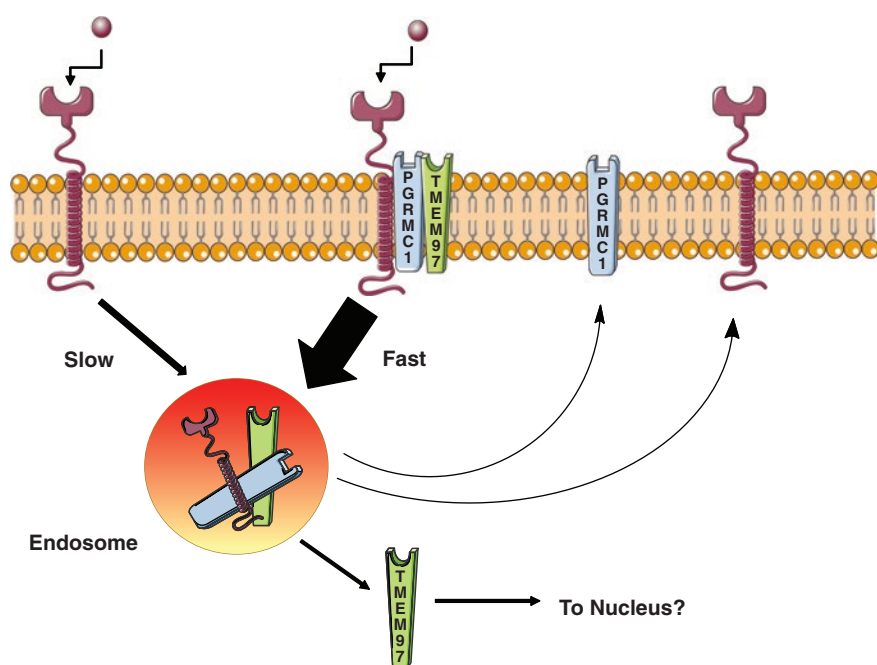
The next series of studies centered on the uptake of LDL particles radiolabeled with [ $^3$ H]cholesterol (i.e., [ $^3$ H]LDL). Knocking out TMEM97 resulted in a dramatic reduction in the rate of uptake of [ $^3$ H]LDL by the HeLa cells. Furthermore, knocking out PGRMC1 resulted in the same reduction in rate of uptake of [ $^3$ H]LDL by HeLa cells. Knocking out both proteins (i.e., double knockout cells) had the same effect as knocking out either TMEM97 or PGRMC1. These data indicated that both TMEM97 and PGRMC1 are integral in the rapid uptake of LDL by the LDL receptor.



**Fig. 28.2** Studies in CRISPR-edited HeLa cells confirming the existence of the TMEM97/ $\sigma 2$  receptor-PGRMC1-LDL receptor complex. (a) In vitro binding studies using [ $^{125}$ I]RHM-4 as the radioligand. (b) Reduction in uptake of [ $^3$ H]LDL in the TMEM97/ $\sigma 2$  receptor, PGRMC1, and double k/o cells. (c) Fluorescence microscopy studies in HeLa cells showing the co-localization of the TMEM97/ $\sigma 2$  receptor-PGRMC1-LDL receptor complex

The third set of studies used a series of antibody labeling studies with confocal microscopy and proximity ligation assays (PLA) to determine the localization of these proteins within the cells (Fig. 28.2). Under serum-starved conditions, it is well established that the LDL receptor localizes on one side of the plasma membrane. A similar observation was made with the localization of both TMEM97 and PGRMC1. Co-localization studies suggested that all three proteins were co-localized. To better visualize the trimeric complex, the authors conducted a series of PLA experiments, a technique that is used to identify protein-protein interactions. A PLA signal was observed when cells were labeled with a pairwise combination of antibodies targeting LDL receptor-TMEM97, LDL receptor-PGRMC1, and TMEM97-PGRMC1. Knocking out TMEM97 eliminated the PLA signal from the LDL-TMEM97 and TMEM97-PGRMC1 labeled cells, but not the LDL receptor-PGRMC1 labeled cells. Similarly, knocking out PGRMC1 eliminated the PLA signal from LDL receptor-PGRMC1 and TMEM97-PGRMC1 cells, but not LDL receptor-TMEM97 labeled cells. Finally, the double knockout cells did not generate a PLA signal with any of the antibody combinations. *These data clearly demonstrate that the TMEM97, PGRMC1, and LDL receptor form a trimeric complex in HeLa cells and that this complex must be intact in order for there to be a rapid uptake of LDL by the cell.*

Figure 28.3 shows a structure of this trimeric complex that is consistent with the results described by Riad et al. (2018) and previous studies demonstrating that the



**Fig. 28.3** Cartoon showing the hypothetical role of the TMEM97/ $\sigma$ 2 receptor-PGRMC1-LDL receptor complex in the internalization of LDL in proliferating versus quiescent tumor cells

$\sigma_2$  receptor is upregulated in proliferating versus quiescent cancer cells (Mach et al. 1997; Wheeler et al. 2000). Cholesterol is a key component of the cell membrane, and the demand for cholesterol is much higher in proliferating than in quiescent cells. Therefore, the “slow” uptake mechanism of LDL uptake is adequate to provide the cholesterol needed when cells are not actively dividing (i.e., quiescent or senescent cells). Consequently, the density of  $\sigma_2$  receptors/TMEM97 (and possibly PGRMC1) is low. However, when cells enter a proliferative state, this slow mechanism is not able to provide the amount of cholesterol needed for synthesis of the cell membrane and other cholesterol-rich organelles. Under these conditions, the “rapid” mechanism of LDL uptake is required, one that involves formation of a trimeric complex consisting of the LDL receptor, TMEM97, and PGRMC1. Therefore, there is an upregulation of the  $\sigma_2$  receptor/TMEM97 (and possibly PGRMC1) in order to increase the level of the trimeric complex in proliferating cells. This hypothesis of “fast” and “slow” uptake mechanisms of LDL is completely consistent with the data reported by Wheeler and colleagues and their characterization of the up- and down-regulation of the  $\sigma_2$  receptor in 66P and 66Q cells (Wheeler et al. 2000).

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## 28.4 Role of the $\sigma_2$ Receptor in the CNS

### 28.4.1 Pharmacological Studies

Although the  $\sigma_2$  receptor has been frequently associated with cancer cell biology because of its high density in proliferating breast cancer cells,  $\sigma_2$  receptor ligands have been shown to have pharmacological properties in the CNS. For example, Lu 28-179 (i.e., Siramesine), a high affinity  $\sigma_2$  receptor agonist having moderate selectivity versus  $\sigma_1$  receptors ( $K_i$  for  $\sigma_2$  receptor = 0.12 nM;  $K_i$  for  $\sigma_1$  receptor = 17 nM), was initially evaluated as an anxiolytic agent. Subsequent studies indicated that siramesine displayed antidepressant properties in rodents.

The  $\sigma_2$  receptor/PGRMC1 complex has been shown to play a role in neurodegenerative diseases, such as Alzheimer’s disease, where  $\sigma_2$  receptor ligands have been found to reduce synaptic loss due to A $\beta$  oligomers and facilitate neuroprotective and anti-inflammatory properties in models of neurodegeneration. For example, Izzo et al. conducted a series of experiments where they discovered that the  $\sigma_2$  receptor mediates A $\beta$ 42 oligomer binding to synaptic puncta on neurons (Izzo et al. 2014a, b). Their work describes that A $\beta$  oligomers act as ligands for the  $\sigma_2$  receptor/PGRMC1 receptor complex and that  $\sigma_2$  expression is upregulated in cultured neurons upon exposure to A $\beta$  oligomers. They reported that their series of compounds, which they described as antagonists for the  $\sigma_2$  receptor/PGRMC1 receptor complex, displaced bound oligomers in neuronal synapses and prevented further binding of A $\beta$ 42 oligomers in neuronal cultures; moreover, their studies showed that  $\sigma_2$  receptor/PGRMC1 complex antagonists inhibited oligomer-induced synaptic loss in neuronal cultures. Due to the ability of their compounds to displace pre-bound oligomers, they conclude that these  $\sigma_2$  ligands function as allosteric antagonists for the  $\sigma_2$  receptor/PGRMC1 receptor complex, rather than direct competitive antagonists. The compounds were able to enter the brain, have an 80% receptor occupancy of the  $\sigma_2$  receptor/PGRMC1 receptor complex, and,

importantly, restored memory deficits in the APP transgenic mouse model of Alzheimer's disease (AD). Phase 1 and 2 clinical trials of a compound designated **CT1812**, which was developed by Cognition Therapeutics, Inc., are currently ongoing in Alzheimer's disease patients.

Yi et al. showed that compounds in the norbenzomorphan class of  $\sigma_2$  receptor ligands modulate intracellular  $\text{Ca}^{2+}$  levels in the human SK-N-SH neuroblastoma cell line, resulting in an increased intracellular concentration (Yi et al. 2017). Dysregulation of  $\text{Ca}^{2+}$  homeostasis has been observed in neurodegenerative diseases such as AD, where A $\beta$  oligomers cause dysregulation of  $\text{Ca}^{2+}$  signaling ultimately resulting in neurodegeneration. Their studies showed that a compound from this class of  $\sigma_2$  receptor ligands, **SAS-0132**, reduced neuroinflammation and reduced behavioral deficits in APP murine models, despite the fact that it had no effect upon A $\beta$  burden or synaptic loss. Vazquez-Rosa et al. found that the  $\sigma_2$  receptor plays a role in other models of neurodegeneration such as traumatic brain injury (Vazquez-Rosa et al. 2019b). Rodents treated with another compound from their series, **DKR-1677**, showed reduced memory loss, axonal degeneration, and loss of neurons and oligodendrocytes.

The  $\sigma_2$  receptor has also been found to be a potential therapeutic target in the context of alcohol addiction. In a study from the same group, Scott et al. characterized a  $\sigma_2$  receptor specific ligand from the same series, **JVW-1034**, that enhanced behavioral performance in a *C. elegans* model of ethanol withdrawal (Scott et al. 2018b). This effect was attenuated when the worm orthologs of  $\sigma_2$  receptor-like protein or PGRMC1, Y38H6C.16, and VEM-1, respectively, were not present. They go on to show that targeting the  $\sigma_2$  receptor ligand with **JVW-1034** caused a reduction in voluntary ethanol intake in ethanol-dependent rats.

Taken together, these studies suggest that the  $\sigma_2$  receptor/PGRMC1 receptor complex represents a promising therapeutic target for ameliorating neurodegeneration, A $\beta$  oligomer-induced synaptotoxicity, and the subsequent neuroinflammatory cascade that ultimately leads to cognitive impairment. The role of the  $\sigma_2$  receptor in facilitating cellular lipoprotein uptake via formation of a trimeric complex with PGRMC1 and the LDL receptor may also prove to be an important consideration in neurodegenerative therapeutics, as the apolipoprotein E4 allele is the greatest risk factor for development of Alzheimer's disease.

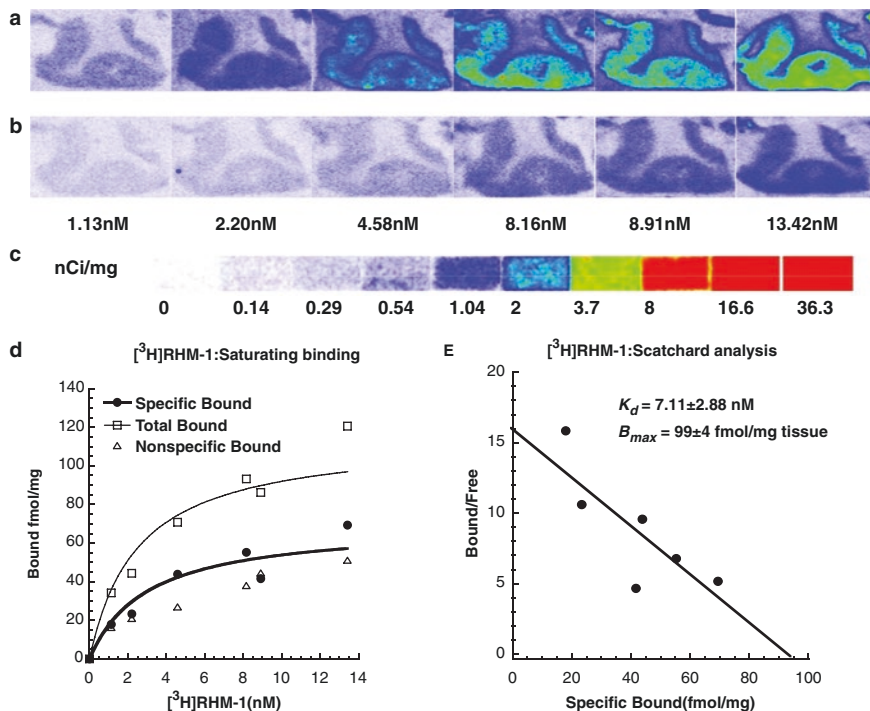
#### 28.4.2 Autoradiography Studies

*N*-[4-(3,4-dihydro-6,7-dimethoxyisoquinolin-2(1H)-yl)butyl]-2-methoxy-5-methyl-benzamide (**RHM-1**), a conformationally flexible benzamide analogue, has been shown to have high affinity and selectivity for  $\sigma_2$  receptor versus  $\sigma_1$  receptor and the dopamine D<sub>2</sub> and D<sub>3</sub> receptors (Mach et al. 2004; Xu et al. 2005). Its high  $\sigma_2$  receptor affinity ( $K_i < 10$  nM) and selectivity ( $\sigma_1:\sigma_2$  ratio  $> 300$ ) indicate that it's a useful radioligand to assess the  $\sigma_2$  receptors in the brain. Therefore, **RHM-1** was radiolabeled with tritium (specific activity = 80 Ci/mmol), and the binding of [<sup>3</sup>H]**RHM-1** to  $\sigma_2$  receptors of aged human postmortem brain tissue (ranging from 84 to 96 years of age) was evaluated in vitro using quantitative autoradiography. Demographic details of the human brains are summarized in Table 28.1.

**Table 28.1** Demographic details of human brains

Case number	Age (year)	Gender	Brain weight (g)	PMI (h)	Braak NFT	Braak amyloid	CDR
1	96	M	1165	12	1	0	0
2	90	M	1150	10	4	A	0
3	84	M	1010	5.5	1	B	0
4	91	M	1170	8.5	1	A	0

PMI postmortem interval, CDR clinical dementia rating

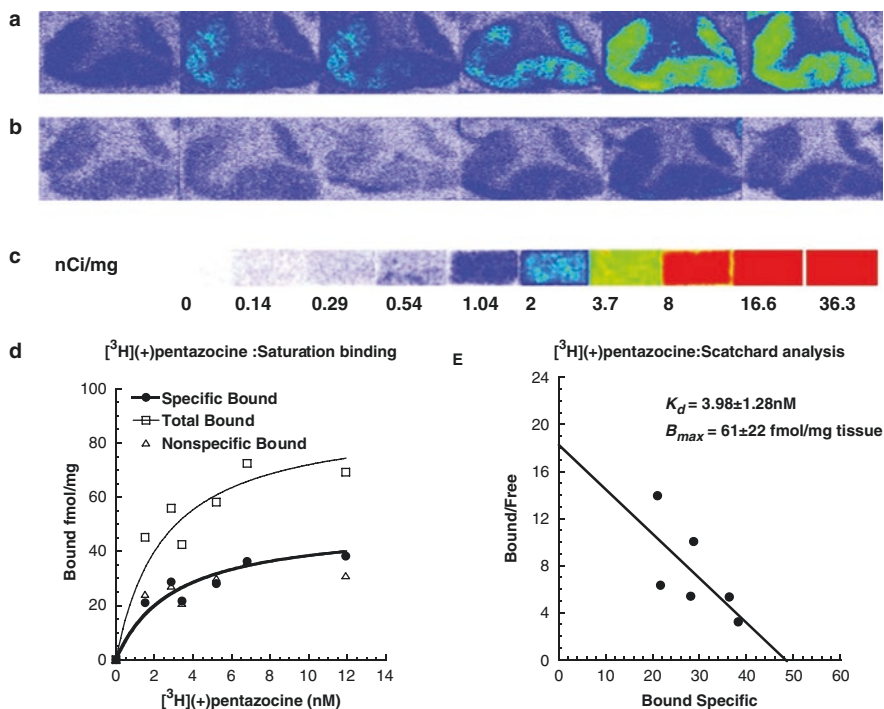


**Fig. 28.4** Representative autoradiography saturation binding of  $[^3\text{H}]\text{RHM-1}$  to  $\sigma_2$  receptors in the frontal cortex of a male cognitively healthy case. Representative autoradiograms show the binding of  $[^3\text{H}]\text{RHM-1}$  to  $\sigma_2$  receptors under different radiotracer concentrations (panel a), Nonspecific binding was determined from adjacent tissue section which contained 1  $\mu\text{M}$  siramesine to mask the  $\sigma_2$  receptor (panel b).  $[^3\text{H}]\text{Microscale}$  standards (ranging from 0 to 36.3 nCi/mg) were also counted for the calibration of radioactivity (panel c). Quantitative autoradiography analysis of the saturation binding (panel d) and Scatchard plots (panel e) was performed to derive the binding densities ( $B_{max}$  values) and the dissociation constants ( $K_d$  values) from three sets of determinations

To measure the binding affinity, direct autoradiography saturation binding studies were conducted using  $[^3\text{H}]\text{RHM-1}$  with a male brain (Fig. 28.4); the  $K_d$  values of the receptor-radioligand binding from Scatchard plots of  $[^3\text{H}]\text{RHM-1}$  to human cortex are shown in Fig. 28.3.  $[^3\text{H}]\text{RHM-1}$  binds with dissociation constant ( $K_d$ ) of 7.11 nM to  $\sigma_2$  receptors in human frontal cortex.

To compare  $\sigma_2$  with  $\sigma_1$  receptors, direct autoradiography saturation binding studies were also conducted using the  $\sigma_1$ -specific radioligand [ $^3\text{H}$ ](+)-pentazocine in the adjacent tissue sections from the same subject. The  $K_d$  values of the receptor-radioligand binding from Scatchard plots of [ $^3\text{H}$ ](+)-pentazocine to human cortex sections are shown in Fig. 28.5. [ $^3\text{H}$ ](+)-pentazocine binds with dissociation constant ( $K_d$ ) of 3.98 nM to  $\sigma_1$  receptors in human frontal cortex.

Autoradiography studies using single concentration radioligands of 8 nM [ $^3\text{H}$ ]RHM-1 and 4 nM [ $^3\text{H}$ ](+)-pentazocine were conducted in three additional male subjects, and the absolute  $\sigma_2$  and  $\sigma_1$  receptor density values ( $B_{\text{max}}$ ) were calculated using the saturation binding isotherm (i.e., Michaelis-Menten equation):  $B_{\text{max}} = B(1 + K_d/L)$ , where  $B_{\text{max}}$  is the receptor density,  $B$  is the apparent binding density measured using single radioligand concentration,  $L$  is the radioligand concentration, and  $K_d$  is the equilibrium dissociation constant. In this study, the  $L$  and



**Fig. 28.5** Binding density and affinity of [ $^3\text{H}$ ](+)-pentazocine to  $\sigma_1$  receptors in the adjacent sections of frontal cortex from the same male subject as shown in Fig. 28.1. Autoradiograms show the binding of [ $^3\text{H}$ ](+)-pentazocine to  $\sigma_1$  receptors under different radiotracer concentrations (panel a). Nonspecific binding was determined from adjacent tissue section which contained 10  $\mu\text{M}$  haloperidol to mask the  $\sigma_1$  receptor (panel b). [ $^3\text{H}$ ]Microscale standards (ranging from 0 to 36.3 nCi/mg) were also counted for the calibration of radioactivity (panel c). Quantitative analysis of the binding was performed to derive the saturation curve (panel d), and Scatchard plots (panel e) were used to determine the binding densities ( $B_{\text{max}}$  values) and dissociation constants ( $K_d$  values) from three sets of determinations

**Table 28.2** Comparison of  $\sigma_1$  and  $\sigma_2$  receptor densities (fmol/mg) in the frontal cortex of aged male human brains

Density (fmol/mg tissue)	$\sigma_2$ receptor	$\sigma_1$ receptor	$\sigma_2/\sigma_1$ ratio	Age (year)
Male ( $n = 4$ )	106.2 $\pm$ 8	70.5 $\pm$ 10.5*	1.6 $\pm$ 0.3	90.3 $\pm$ 2.5

Data were obtained from aged male healthy brains ( $n = 4$ ) and presented as mean value  $\pm$  SEM

\*Denotes  $p < 0.05$ , student t test,  $\sigma_2$  versus  $\sigma_1$

$K_d$  values were 8 and 7.11 nM for  $\sigma_2$  receptors and 4 and 3.98 nM for  $\sigma_1$  receptors, which were determined using saturation binding assay as shown in Figs. 28.4 and 28.5.

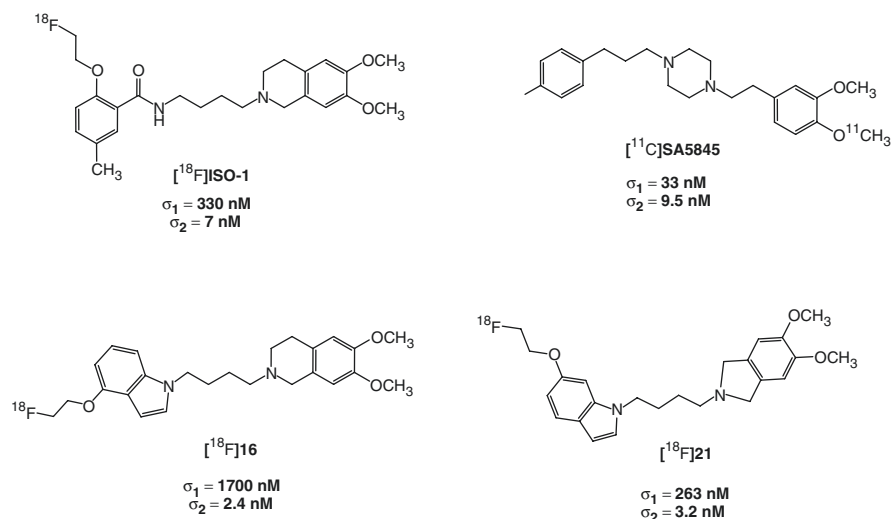
$\sigma_2$  and  $\sigma_1$  receptors are highly expressed in the frontal cortex of healthy control brains.  $\sigma_2$  density (106.2  $\pm$  8 fmol/mg tissue) is higher than  $\sigma_1$  receptor density (70.5  $\pm$  10.5) in the frontal cortex with a ratio of  $\sim$ 1.6-fold, summarized in Table 28.2. The high binding density of  $\sigma_2$  receptor in the aged brains suggests that  $\sigma_2$  receptor may play an important role in the brain disorders, and further studies in both male and female controls, Alzheimer's and Parkinson disease brains, will be important in understanding the regulation of  $\sigma_2$  receptor in central nervous disorders.

### 28.4.3 PET Radiotracers for Imaging the $\sigma_2$ Receptor

To date, a large number of  $^{11}\text{C}$ - and  $^{18}\text{F}$ -labeled radiotracers having a high affinity for  $\sigma_2$  receptors and good selectivity versus the  $\sigma_1$  receptor have been evaluated in in vivo imaging studies. The majority of these studies have focused on tumor imaging studies because of the high correlation of  $\sigma_2$  receptor density and the proliferative status (i.e., P: Q ratio) of the tumor. The only  $\sigma_2$ -selective PET radiotracer translated to human imaging studies is [ $^{18}\text{F}$ ]ISO-1 (Fig. 28.6). A preliminary study of [ $^{18}\text{F}$ ]ISO-1 in patients diagnosed with either lymphoma or breast or head and neck cancer demonstrated that it was possible to separate patients into high and low proliferative status based on a comparison of the PET results with the Ki-67 score of the tumor upon histological analysis (Dehdashti et al. 2013). That is, [ $^{18}\text{F}$ ]ISO-1 had a higher uptake in patients having a Ki-67 score  $> 0.4$  and a lower tracer uptake in tumors below this cutoff value. A follow-up study in breast cancer patients revealed that there was a good correlation of [ $^{18}\text{F}$ ]ISO-1 when compared with Ki-67 score (McDonald et al. 2020). However, the dynamic range for [ $^{18}\text{F}$ ]ISO-1 uptake in tumors having a high versus low Ki-67 was low, which is likely due to the low aggressive nature of breast cancer versus other cancers (McDonald et al. 2020). In addition, [ $^{18}\text{F}$ ]ISO-1 has a very low brain uptake, which has been attributed to it being a substrate for P-glycoprotein (Mach, unpublished data). Therefore, [ $^{18}\text{F}$ ]ISO-1 is not a suitable PET radiotracer for imaging  $\sigma_2$  receptors in the CNS.

A number of promising PET radiotracers for imaging  $\sigma_2$  receptors in the CNS have been reported in recent years. The first PET radiotracer worth noting is [ $^{11}\text{C}$ ]SA5845 (Fig. 28.6), which has a good affinity for  $\sigma_2$  receptors and modest affinity versus  $\sigma_1$  receptors (Kortekaas et al. 2008). PET imaging studies in





**Fig. 28.6** Structures of representative PET radiotracers for imaging the  $\sigma_2$  receptor

nonhuman primates revealed high brain uptake of this radiotracer and kinetically irreversible binding to sigma receptors *in vivo* (Kortekaas et al. 2008). However, a potential limitation of this radiotracer is its relatively low selectivity versus  $\sigma_1$  receptors, which makes it difficult to identify changes in  $\sigma_2$  receptor density in CNS disorders.

Jia et al. (2019) and Wang et al. (2017) have recently reported two  $^{18}\text{F}$ -labeled  $\sigma_2$ -selective PET radiotracers that displayed a high uptake in both rodent and non-human primate brain (Fig. 28.6). Of the two PET radiotracers,  $[^{18}\text{F}]\text{21}$  had the higher brain uptake. *In vivo* blocking studies and *in vivo* metabolite analysis indicated that  $[^{18}\text{F}]\text{21}$  was specific for  $\sigma_2$  receptors; there were no radioactive metabolites of  $[^{18}\text{F}]\text{21}$  observed in rodent brain, indicating that this radiotracer may be useful in quantitative PET imaging studies of  $\sigma_2$  receptors in humans.

## 28.5 Conclusions and Perspectives

The  $\sigma_2$  receptor has undergone a major scientific evolution in recent years. The initial recognition that this protein could serve as a receptor-based marker of the proliferative status of solid tumors resulted in an effort focused on the development of PET radiotracers for oncologic imaging studies. The molecular identification of the  $\sigma_2$  receptor as being a component of the TMEM97/ $\sigma_2$ -PGRMC1-LDL receptor complex suggests that these PET imaging studies will be useful in evaluating the role of lipoprotein uptake in cell proliferation. Although promising results have been obtained in translational imaging studies with  $[^{18}\text{F}]\text{ISO-1}$  in cancer patients, PET radiotracers having a higher affinity for  $\sigma_2$  receptors will likely be needed.

The recent studies showing that  $\sigma_2$  receptor ligands possess a diverse set of pharmacological properties in the CNS have opened a new avenue of research in PET probe development since many of the radiotracers used in the oncological imaging studies lacked either a high brain uptake or suitable  $\sigma_2:\sigma_1$  receptor selectivity required for brain imaging studies. Two promising PET radiotracers were recently reported in imaging studies in rodents and nonhuman primates, but it is not clear if they are suitable for translational imaging studies in humans. The development of  $\sigma_2$  selective radiotracers for brain imaging studies will likely be an active area of research in the years to come.

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# PET Imaging of Synaptic Vesicle Protein 2A

# 29

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## Abstract

The synaptic vesicle protein 2A (SV2A) is an integral 12-transmembrane domain glycoprotein located in the membrane of synaptic vesicles and widely distributed throughout the brain. In 2004, SV2A was shown to be the molecular target of the antiepileptic drug levetiracetam. High-affinity SV2A ligands have been described by the pharmaceutical company UCB more recently, and this enabled the rapid development of SV2A positron emission tomography (PET) radioligands. Because synaptic vesicle proteins, such as SV2A, have previously been used as biomarkers of synaptic density, PET imaging of SV2A may have broad utility across neuropathological diseases. In this chapter we review the current status of SV2A PET imaging.

During the last few years, several SV2A radioligands have been synthesized and evaluated with PET in animals and human. Although [ $^{11}\text{C}$ ]levetiracetam was deemed an unsuitable radioligand, radiolabeled non-acetamide *racetams* (a class of drugs that contain the pyrrolidone moiety) such as [ $^{11}\text{C}$ ]UCB-J and more recently [ $^{18}\text{F}$ ]SynVesT-1 and [ $^{18}\text{F}$ ]SynVesT-2 provided high-quality brain PET imaging results across species. Regional brain SV2A levels were shown to closely correlate with those of synaptophysin, another commonly used marker for synaptic density. PET studies using SV2A-targeting antiepileptic drugs levetiracetam and brivaracetam confirmed that the radioligands bind specifically to SV2A and provided valuable insight into the potential use of a reference region approach for quantification of brain PET signal. Several initial PET studies examining SV2A radioligand binding in patient groups have been reported, and SV2A was found to be reduced in disease-associated brain regions in epilepsy, Alzheimer's disease, major depressive disorder and posttraumatic stress disorder, Parkinson's disease, and schizophrenia.

In conclusion, PET imaging of SV2A has rapidly developed, and qualified radioligands are now available. Initial PET studies in humans indicate that SV2A loss might be specific to disease-associated brain regions and consistent with synaptic density loss. Future studies warrant the use of fluorine-18 labeled radioligands in larger patient cohorts to establish the clinical value of SV2A PET imaging and its potential for diagnosis and progression monitoring of neuropathological diseases, as well as efficacy assessment of disease-modifying therapies.

## 29.1 Synaptic Vesicle Protein 2

The synaptic vesicle protein 2 (SV2) constitutes a conserved protein family in vertebrates (Buckley and Kelly 1985) and is an integral 12-transmembrane domain glycoprotein (Bajjalieh et al. 1992) located on secretory vesicles of neurons and endocrine cells (Buckley and Kelly 1985; Floor and Feist 1989). The SV2 protein sequence has been found to be very similar to sequences of transporter proteins, and SV2 was therefore initially considered to be a vesicular transport protein (Bajjalieh et al. 1992; Feany et al. 1992). SV2 is critical for normal synaptic function and plays a role in vesicle trafficking and exocytosis (Nowack et al. 2010), although its exact functions remain to be fully elucidated (see for reviews Mendoza-Torreblanca et al. 2013; Bartholome et al. 2017).

Further characterization of the SV2 protein has resulted in the identification of three isoforms that share approximately 60% of their sequences (Bajjalieh et al. 1994; Janz and Sudhof 1999). The isoforms have a different regional distribution in the brain; SV2A is ubiquitously expressed in virtually all synapses, while SV2B is more restricted but present throughout the rat brain (Janz and Sudhof 1999; Bajjalieh et al. 1993, 1994). In contrast, SV2C was observed in only a few rat brain areas, including the globus pallidus, substantia nigra, midbrain, brainstem, and olfactory bulb (Janz and Sudhof 1999). On the neuronal level, SV2A is thought to be located in glutamatergic and GABAergic neurons, while SV2B might be more restricted to glutamatergic neurons and SV2C to GABAergic neurons (see for review Bartholome et al. 2017).

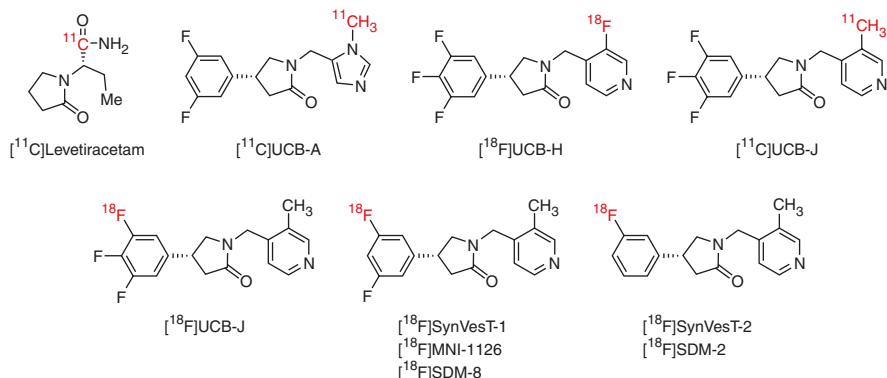
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## 29.2 Development of SV2A PET Radioligands

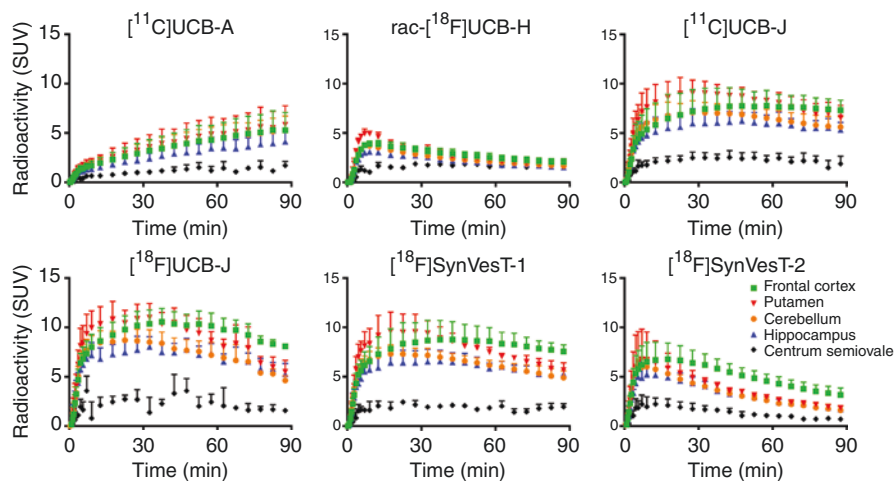
In 2004, Lynch et al. demonstrated that SV2A was the molecular target of the anti-epileptic drug levetiracetam (Lynch et al. 2004; Gillard et al. 2006). Considering the clinical success of levetiracetam as an effective antiepileptic drug, UCB started an extensive medicinal chemistry program to identify a novel generation of antiepileptic drugs with increased SV2A affinity and anti-seizure potency, ultimately resulting in the development of brivaracetam (Gillard et al. 2011; Klein et al. 2018).

As a first attempt to develop a SV2A PET radioligand, levetiracetam was labeled with carbon-11 as [<sup>11</sup>C]levetiracetam (Fig. 29.1) (Cai et al. 2014) but was not evaluated further, probably due to its low binding affinity to SV2A ( $K_i = 1.74 \mu\text{M}$ ) (Danish et al. 2017). In the meanwhile, high binding affinity SV2A-specific ligands with the pyrrolidinone pharmacophore were synthesized and evaluated as candidates for SV2A PET (Mercier et al. 2014, 2017). Three of the SV2A ligands described in the 2014 paper were labeled with carbon-11 or fluorine-18 as [<sup>11</sup>C]UCB-A (Estrada et al. 2016), [<sup>18</sup>F]UCB-H (Warnock et al. 2014; Becker et al. 2017; Warnier et al. 2016; Bretin et al. 2013), and [<sup>11</sup>C]UCB-J (Nabulsi et al. 2016) (Fig. 29.1), and all were initially evaluated in nonhuman primates at the Yale PET Center.





**Fig. 29.1** Chemical structures of SV2A PET radioligands  $[^{11}\text{C}]$ levetiracetam,  $[^{11}\text{C}]$ UCB-A,  $[^{18}\text{F}]$ UCB-H,  $[^{11}\text{C}]$ UCB-J,  $[^{18}\text{F}]$ UCB-J,  $[^{18}\text{F}]$ SynVesT-1, and  $[^{18}\text{F}]$ SynVesT-2. Modified from Cai et al. (2019)



**Fig. 29.2** Regional brain time-activity curves of SV2A PET radioligands  $[^{11}\text{C}]$ UCB-A ( $n = 5$ ),  $\text{rac-}[^{18}\text{F}]$ UCB-H ( $n = 2$ ),  $[^{11}\text{C}]$ UCB-J ( $n = 6$ ),  $[^{18}\text{F}]$ UCB-J ( $n = 4$ ),  $[^{18}\text{F}]$ SynVesT-1 ( $n = 3$ ), and  $[^{18}\text{F}]$ SynVesT-2 ( $n = 3$ ) evaluated in rhesus monkeys. Error bars (up) denote standard deviation. Modified from Cai et al. (2019)

The preclinical results demonstrated that  $[^{11}\text{C}]$ UCB-J possessed the most suitable pharmacokinetic profile as a brain PET radioligand, i.e., rapid and high brain uptake, reversible binding kinetics, and relatively low nonspecific binding in white matter (Nabulsi et al. 2016), while  $[^{11}\text{C}]$ UCB-A showed slower brain penetration and apparently irreversible kinetics in the brains of nonhuman primates, making quantitative analysis more challenging (Fig. 29.2). Furthermore,  $\text{rac-}[^{18}\text{F}]$ UCB-H showed relatively low specific binding in the nonhuman primate brain, probably due to its lower binding affinity to SV2A ( $pI_{C_{50}} = 7.8$  for UCB-H vs. 8.2 for UCB-J)

(Mercier et al. 2014; Nabulsi et al. 2016). After translation to clinical studies, the human imaging data were highly consistent with preclinical data, reaffirming [ $^{11}\text{C}$ ]UCB-J as an ideal imaging radioligand for SV2A in the brain (Finnema et al. 2016), with [ $^{11}\text{C}$ ]UCB-A possessing slow binding kinetics (Lubberink et al. 2017) and [ $^{18}\text{F}$ ]UCB-H low specific binding in the brain (Bahri et al. 2017), limiting both [ $^{11}\text{C}$ ]UCB-A and [ $^{18}\text{F}$ ]UCB-H as PET radioligands for quantifying SV2A in the brain.

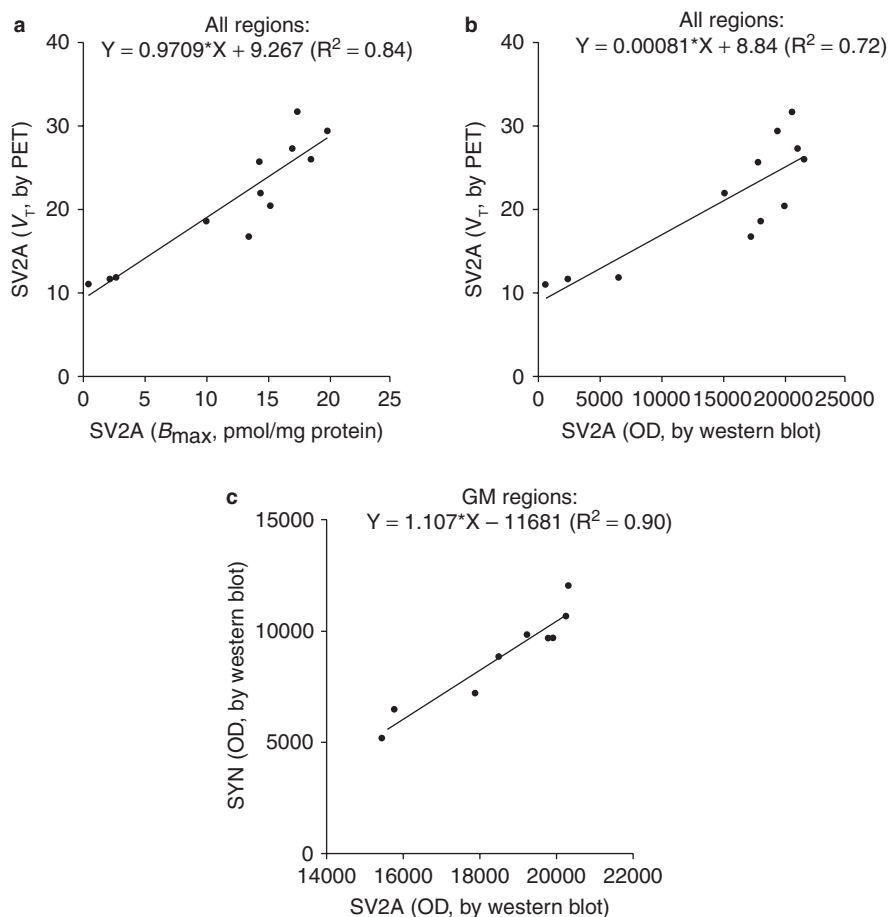
As preliminary data of [ $^{11}\text{C}$ ]UCB-J in preclinical and clinical PET studies were promising, fluorine-18 labeled SV2A radioligands became of interest as the short half-life of carbon-11 (20.4 min) limits the production of [ $^{11}\text{C}$ ]UCB-J to PET centers with cyclotrons. Initial work focused on the synthesis of fluorine-18 labeled counterparts of UCB-J, i.e., [ $^{18}\text{F}$ ]UCB-J (Fig. 29.1), and indeed [ $^{18}\text{F}$ ]UCB-J provided very similar results as [ $^{11}\text{C}$ ]UCB-J in nonhuman primates (Li et al. 2019b). However, due to the harsh labeling conditions and the need for chiral HPLC separation in the production of [ $^{18}\text{F}$ ]UCB-J, the process was considered unsuitable for routine production supporting clinical studies. A second approach focused on the synthesis of mono- and di-fluorinated UCB-J analogs, i.e., [ $^{18}\text{F}$ ]SynVesT-1 ([ $^{18}\text{F}$ ]SDM-8, [ $^{18}\text{F}$ ]MNI-1126) (Li et al. 2019a; Constantinescu et al. 2019) and [ $^{18}\text{F}$ ]SynVesT-2 ([ $^{18}\text{F}$ ]SDM-2) (Cai et al. 2020) (Fig. 29.1). In nonhuman primates, the brain pharmacokinetics of [ $^{18}\text{F}$ ]SynVesT-2 were faster than [ $^{11}\text{C}$ ]UCB-J (Fig. 29.2), while regional binding potential ( $BP_{\text{ND}}$ ) values of the two radioligands were very similar. [ $^{18}\text{F}$ ]SynVesT-1 possessed similar pharmacokinetic profile and higher  $BP_{\text{ND}}$  values as [ $^{11}\text{C}$ ]UCB-J (Fig. 29.2). [ $^{18}\text{F}$ ]SynVesT-1 was evaluated in humans and displayed outstanding imaging characteristics, i.e., very high brain uptake, fast and reversible kinetics, and excellent test-retest reproducibility and binding specificity to SV2A (Li et al. 2019c). Compared to [ $^{11}\text{C}$ ]UCB-J, [ $^{18}\text{F}$ ]SynVesT-1 displayed higher specific binding signals (i.e.,  $BP_{\text{ND}}$ ). Recent human studies with [ $^{18}\text{F}$ ]SynVesT-2 showed slightly lower  $BP_{\text{ND}}$  but exhibited faster kinetics, as in nonhuman primates. The availability of these fluorine-18 labeled SV2A PET radioligands provides the opportunity for multicenter clinical validation of SV2A PET in a variety of neuropsychiatric disorders.

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### 29.3 SV2A as a Biomarker of Synaptic Density

The human brain is estimated to contain close to 100 billion neurons (Azevedo et al. 2009), and with estimates of each neuron potentially contacting >1000 synapses, there might be >100 trillion synapses. Structural disturbances or reductions in synapses can result in dysfunctional neuronal signaling and in network dysfunctions. Synaptic vesicle proteins, such as SV2A, have previously been established as histological markers of synaptic density (De Camilli et al. 1983; Goelz et al. 1981; Masliah et al. 1990) due to their highly restricted localization to synaptic boutons. SV2A is a promising biomarker of synaptic density as it is ubiquitously and homogeneously present in synaptic vesicles (Bajjalieh et al. 1994) and has relative low variation in copy number per vesicle, i.e., SV2 was most “monodispersed” (Mutch et al. 2011).

To examine if [ $^{11}\text{C}$ ]UCB-J binding to SV2A could be used as an index of synaptic density, a baboon (*Papio anubis*) underwent a [ $^{11}\text{C}$ ]UCB-J PET measurement after which the animal was euthanized and the brain harvested for post-mortem tissue studies. Analysis of regional brain tissues confirmed that the PET-measured [ $^{11}\text{C}$ ]UCB-J distribution volume ( $V_T$ ) correlated well with the regional distribution of SV2A measured by radioligand homogenate binding assay (Fig. 29.3a) and western blot (Fig. 29.3b). Importantly, there was also good correlation between SV2A and the “gold standard” synaptic density marker synaptophysin in western blot (Fig. 29.3c) and confocal microscopy experiments, indicating that SV2A can be



**Fig. 29.3** (a) Correlation between regional in vitro SV2A ( $B_{\max}$ ) measured using tissue homogenate binding and in vivo SV2A ( $V_T$  by PET). Data are 12 brain regions. (b) Correlation between regional in vitro SV2A (optical density (OD) by western blot) and in vivo SV2A ( $V_T$  by PET). Data are 12 brain regions. (c) Correlation between in vitro SV2A and in vitro synaptophysin density in gray matter regions determined using western blot analyses. Data are nine brain regions. Modified from Finnema et al. (2016)

used as an alternative to synaptophysin for accurate quantification of synaptic density (Finnema et al. 2016). Analogous biochemical studies with resected human tissue following epilepsy surgery show promising results. Further studies with larger sample sets from post-mortem human tissue samples are required to more firmly establish the value of SV2A as a biomarker of synaptic density.

## 29.4 Quantification of SV2A PET Radioligands

Evaluation of the pharmacokinetic properties of SV2A PET radioligands in humans revealed that the best models to quantify regional  $V_T$  values require measurements of arterial input function: the one-tissue compartment model (1TC) for [ $^{11}\text{C}$ ]UCB-J (Finnema et al. 2018; Koole et al. 2019; Mansur et al. 2020) and the two-tissue compartment model (2TC) or Logan graphical analysis for [ $^{18}\text{F}$ ]UCB-H (Bahri et al. 2017). Furthermore, excellent test-retest reproducibility and low noise  $V_T$  parametric images can be obtained with the 1TC model for [ $^{11}\text{C}$ ]UCB-J (Finnema et al. 2018).

Noninvasive imaging with [ $^{11}\text{C}$ ]UCB-J would not only make blood sampling redundant but is also desired to increase subject comfort and simplify imaging protocols. It is also of interest to conduct parametric imaging to explore SV2A changes in the whole brain using statistical mapping in larger study cohorts. There are two promising kinetic analysis methods which could be used for noninvasive parametric imaging of [ $^{11}\text{C}$ ]UCB-J: using the reference tissue model or the standard uptake value ratio (SUV<sub>R</sub>) approach. Before implementing these methods, identification of a reference tissue was needed first.

For noninvasive quantification of [ $^{11}\text{C}$ ]UCB-J specific binding, the centrum semiovale (CS) has been proposed as a reference region based on *in vitro* data (Finnema et al. 2016), which showed negligible concentrations of SV2A in CS in baboons by biochemical methods. However, a reduction in [ $^{11}\text{C}$ ]UCB-J uptake was observed in CS  $V_T$  in displacement studies with levetiracetam and brivaracetam in humans (Finnema et al. 2016, 2019). Further assessment of CS as a reference region in drug occupancy studies revealed that CS  $V_T$  overestimated the non-displaceable distribution volume ( $V_{ND}$ ) (Rossano et al. 2019; Koole et al. 2019) but showed a significant correlation between CS  $V_T$  and gray matter  $V_{ND}$ , which suggests that the CS remains a promising proxy reference region (Rossano et al. 2019). To avoid a spill-in from surrounding gray matter uptake, a core area of the CS (volume, 2 mL) was determined based on a probability map. Increasing the number of ordered subset expectation maximization (OSEM) iterations decreased the CS  $V_T$  values (35–40% overestimation at 4 iterations of 30 subsets) (Rossano et al. 2019).

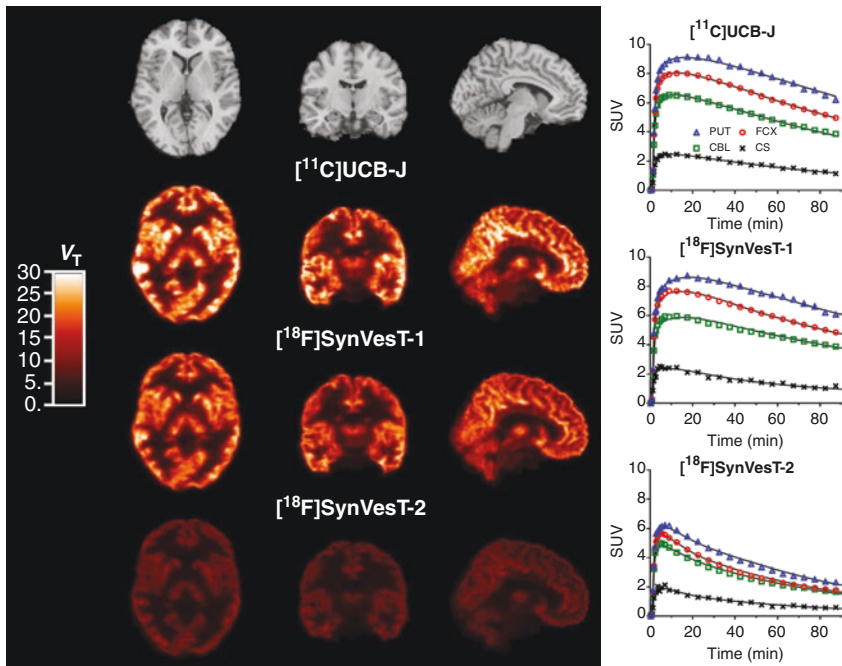
Since 1TC is the model of choice with arterial blood sampling, simplified reference tissue model (SRTM) and SRTM2 were expected to work well as reference tissue models, and SRTM2 would improve the statistical quality of parametric imaging over SRTM. It has been reported that SRTM2 worked well with a sufficiently long acquisition (80 or 90 min) (Mertens et al. 2020), on a GE Signa PET/MR system. For shorter acquisition duration, the variance of  $k'_2$  estimates (i.e., the efflux rate from the CS region of interest) in SRTM2 increased, and SRTM2 failed

more often (11% of the scans at 50 min acquisition). However, the use of higher-resolution PET system (HRRT, high-resolution research tomograph), combined with the use of a small core CS region (2 mL), produced a noisy CS time-activity curve (TAC). This affected the estimation performance of  $k'_2$ . However, the 1TC  $k_2$  value of CS was similar between healthy controls and patients (healthy control,  $0.029 \pm 0.003$ ; Alzheimer's disease (AD),  $0.028 \pm 0.005$ ; all participants,  $0.028 \pm 0.004$  [1/min]) (Naganawa et al. 2018b). Therefore, SRTM2 with population  $k'_2$  is a promising method to generate high-resolution parametric  $BP_{ND}$  images, although this approach will require validation in other populations.

The ratio of tissue activity to plasma activity (i.e., apparent  $V_T$ ) overestimated the 1TC  $V_T$  at any time point in a 90 min scan after bolus injection (Naganawa et al. 2018a). This was the case in both healthy controls and patients. However, the overestimation in the target regions and reference region (i.e., CS) tended to cancel out while computing the SUVR value (i.e., ratio of target tissue activity to CS activity). The choice of the acquisition time window is critical to compute SUVR values, since SUVR values continued to increase monotonically over time throughout a 2 h scan. When comparing SUVR-1 values from several 30 min acquisition time windows with the 1TC  $BP_{ND}$  values, the best match was observed at 60–90 min post-injection in healthy controls (Mertens et al. 2020; Koole et al. 2019) and patients with various diseases (Naganawa et al. 2018a). Similar results have recently been found for [ $^{18}\text{F}$ ]SynVesT-1, which should allow the use of SUVR for application in large studies evaluating the effect of drug treatment in neurodegenerative disorders.

Evaluation of the other radioligands has also been conducted for the imaging and quantification of SV2A in humans. As expected, [ $^{11}\text{C}$ ]UCB-A displayed slow kinetics leading to unreliable estimates of binding parameters (Lubberink et al. 2017), while the specific binding of [ $^{18}\text{F}$ ]UCB-H was low, with  $BP_{ND}$  below 1.0 for all regions (Bahri et al. 2017). For [ $^{18}\text{F}$ ]SynVesT-1, evaluation in humans reveals tissue kinetics similar to that of [ $^{11}\text{C}$ ]UCB-J (Li et al. 2019c) (Fig. 29.4). The 1TC model fits the regional time-activity curves well and provides reliable estimates of regional  $V_T$  values, which are lower than those of [ $^{11}\text{C}$ ]UCB-J. However, regional  $BP_{ND}$  values of [ $^{18}\text{F}$ ]SynVesT-1 estimated with CS as reference region are higher than those of [ $^{11}\text{C}$ ]UCB-J (Table 29.1). Also similar to [ $^{11}\text{C}$ ]UCB-J, 60 min of acquisition time was sufficient to estimate 1TC  $V_T$  values. As for [ $^{18}\text{F}$ ]SynVesT-2, its tissue kinetics in humans are substantially faster than those of [ $^{11}\text{C}$ ]UCB-J and [ $^{18}\text{F}$ ]SynVesT-1 (Fig. 29.4). Regional  $V_T$  and  $BP_{ND}$  values are also lower. However, due to its fast kinetics, a relative short acquisition (i.e., 30 min) is sufficient for stable estimates of regional  $V_T$  values.

In summary, there are to date five radioligands available for imaging and quantification of SV2A in humans, with three of them ([ $^{11}\text{C}$ ]UCB-J, [ $^{18}\text{F}$ ]SynVesT-1, and [ $^{18}\text{F}$ ]SynVesT-2) displaying excellent imaging characteristics of high and reversible brain uptake, excellent test-retest reproducibility, and high specific binding. Application of radiolabeled SV2A radioligands, especially [ $^{18}\text{F}$ ]SynVesT-1 and [ $^{18}\text{F}$ ]SynVesT-2, will advance our understanding of neurologic and psychiatric



**Fig. 29.4** Comparison of  $[^{11}\text{C}]\text{UCB-J}$ ,  $[^{18}\text{F}]\text{SynVesT-1}$ , and  $[^{18}\text{F}]\text{SynVesT-2}$  in one human subject. Left: MR and  $V_T$  images from PET measurements following injection of  $[^{11}\text{C}]\text{UCB-J}$ ,  $[^{18}\text{F}]\text{SynVesT-1}$ , and  $[^{18}\text{F}]\text{SynVesT-2}$ . Right: Regional time-activity curves following injection of  $[^{11}\text{C}]\text{UCB-J}$ ,  $[^{18}\text{F}]\text{SynVesT-1}$ , and  $[^{18}\text{F}]\text{SynVesT-2}$ . Solid lines represent the ITC fit

disorders in large-scale longitudinal and cross-sectional studies in patient populations and facilitate evaluation of disease-modifying therapeutics.

## 29.5 PET Studies in Disease Populations

### 29.5.1 Epilepsy

Epilepsy is one of the most common neurological conditions with a prevalence of 0.5–1.0% (Bell and Sander 2001; Picot et al. 2008). It is characterized by spontaneous and recurrent seizures typically starting in a focal brain area referred to as the seizure onset zone (SOZ). Approximately 20–40% of patients develop drug-resistant epilepsy (Picot et al. 2008; Kwan and Brodie 2000; Cockerell et al. 1995) with a high incidence (50–70%) in temporal lobe epilepsy (TLE), the most common type of epilepsy in adults (Schmidt and Loscher 2005). In patients with medically refractory epilepsy, surgical resection of the epileptogenic tissue may provide high quality-of-life benefit (Jette et al. 2016). Identification of the SOZ can however be challenging and is typically guided by multi-modality data, including magnetic

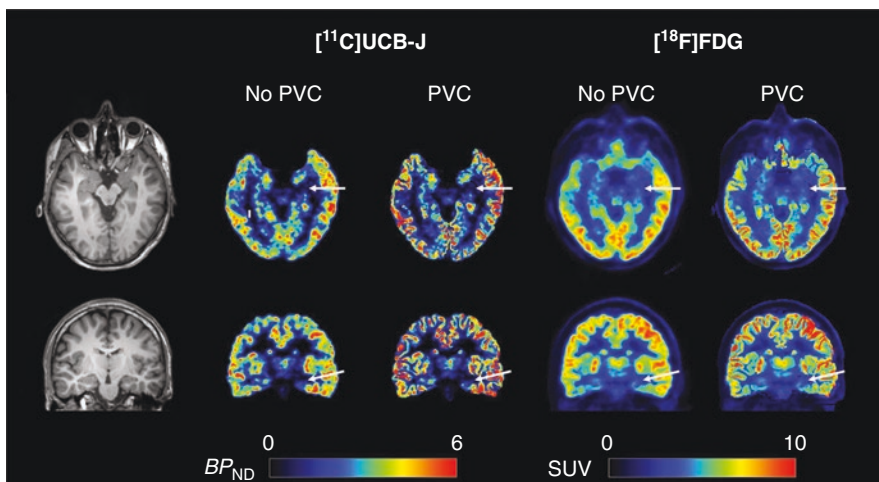
**Table 29.1** Comparison of  $BF_{ND}$  values of [ $^{11}C$ ]UCB-J, [ $^{18}F$ ]SynVesT-1, and [ $^{18}F$ ]SynVesT-2 in humans

Radioligand	Thalamus	Putamen	Frontal	Parietal	Occipital	Temporal	Amygdala	Hippocampus	Caudate	Subjects
[ $^{11}C$ ]UCB-J	2.22	3.69	3.15	3.37	3.37	3.61	2.95	2.18	2.58	$n = 3$
[ $^{18}F$ ]SynVesT-1	2.92	4.48	4.01	4.38	4.34	4.63	4.06	2.84	3.39	$n = 6$
[ $^{18}F$ ]SynVesT-2	1.94	3.02	2.40	2.48	2.43	2.71	2.31	1.59	1.59	$n = 3$

resonance (MRI) and PET imaging (Martinkovic et al. 2014). 2-Deoxy-2- $^{18}\text{F}$ -fluoro-D-glucose ( $^{18}\text{F}$ ]FDG) is the clinical standard PET radioligand, but other radioligands have shown promise for improved identification of the SOZ when compared to anatomical MRI (Theodore 2017).

SV2A is the binding site for the antiepileptic drugs levetiracetam and brivaracetam (Gillard et al. 2006; Lynch et al. 2004; Gillard et al. 2011). In vitro studies using resected brain tissue have shown that SV2A expression is reduced by 30–50% in the epileptogenic anterior temporal neocortex of patients with TLE (Feng et al. 2009). Similar reductions in SV2A have been observed in the hippocampus of TLE patients with hippocampal sclerosis using western blot analysis (Van Vliet et al. 2009) and quantitative mRNA expression (Crevecoeur et al. 2014). Furthermore, in rat models of epilepsy, SV2A loss has been shown to contribute to epileptogenesis and pharmacoresistance (Van Vliet et al. 2009; Hanaya et al. 2012; Wang et al. 2014). For instance, SV2A knockout mice manifest a severe seizure phenotype (Crowder et al. 1999; Janz et al. 1999).

In the first pilot study in subjects with TLE ( $n = 3$ ),  $^{11}\text{C}$ ]UCB-J binding was asymmetric in the mesial temporal lobe, consistent with SV2A loss in the ipsilateral mesial temporal lobe (Finnema et al. 2016). In a follow-up study, 11 subjects with TLE and medial temporal lobe sclerosis (MTS) and 11 control subjects were investigated (Finnema et al. 2017). In the TLE MTS subjects with hippocampal volumetric asymmetry of  $25 \pm 11\%$ ,  $^{11}\text{C}$ ]UCB-J  $BP_{\text{ND}}$  values were lower in the sclerotic hippocampus when compared to control subjects (Fig. 29.5). Following partial volume correction, hippocampal  $^{11}\text{C}$ ]UCB-J  $BP_{\text{ND}}$  asymmetry indices were  $37 \pm 19\%$  in the hippocampus with very limited asymmetry in other brain regions. The corresponding asymmetry in hippocampal  $^{18}\text{F}$ ]FDG uptake was  $22 \pm 7\%$  and correlated



**Fig. 29.5** Parametric images without and with partial volume correction (PVC) of  $^{11}\text{C}$ ]UCB-J  $BP_{\text{ND}}$  (left) and  $^{18}\text{F}$ ]FDG standard uptake values (SUV, right) reveal unilateral mesial temporal sclerosis (MTS) in a representative subject with temporal lobe epilepsy (TLE)



with that of [ $^{11}\text{C}$ ]UCB-J  $BP_{\text{ND}}$  across subjects. Notably, hippocampal asymmetries in [ $^{11}\text{C}$ ]UCB-J binding were 1.7-fold larger than those of [ $^{18}\text{F}$ ]FDG uptake. In summary, [ $^{11}\text{C}$ ]UCB-J binding was reduced in the seizure onset zone of TLE subjects with MTS. PET imaging of SV2A holds promise as a biomarker approach in the presurgical selection and evaluation of TLE patients and may improve the sensitivity of molecular imaging for SOZ detection. Future studies should evaluate SV2A imaging in patients with TLE without MTS and in extra temporal non-lesional epilepsy patients.

## 29.5.2 Alzheimer's Disease and Other Dementias

Aging is a normal physiologic process, which includes gradual decrease in cognitive function as compared to healthy young adults. Cognitive function refers to complex cortical functions predominantly relating to memory, language, judgment, reasoning, planning, behavior, and executional abilities. Dementia is abnormal cognitive decline that is greater than expected as compared to age-matched controls and often causes disability or affects the ability to independently perform daily living activities. Dementia has become a global public health issue with an increasingly aging population. Dementia can be attributed to many causes, the most common and devastating one is AD accounting for more than 50–60% of dementia, while 15–25% of cases are due to frontotemporal dementia (FTD) and dementia with Lewy bodies (DLB), respectively (Brown et al. 2014). In 2016, approximately 5.4 million Americans were estimated to have AD, and it was the sixth leading cause of death in the United States (Scheltens et al. 2016). The number of patients is expected to double by 2050 if no cure is identified (Alzheimer's-Association 2017).

From a diagnostic perspective, AD is increasingly viewed along a continuum from preclinical AD to mild cognitive impairment (MCI) and to AD dementia (Jack et al. 2011). The clinical dementia stage of AD is coupled to a distinct pathology with formation of plaques composed of  $\beta$ -amyloid, neurofibrillary tangles composed of hyperphosphorylated tau protein, and synaptic density loss (Overk and Masliah 2014). Synapses are crucial for cognitive function, and synaptic density loss is a robust and consistent pathology in AD (Selkoe 2002). Cognitive impairment in AD is closely associated with synaptic density loss in the cortex and limbic system (Terry et al. 1991; Dekosky and Scheff 1990; Dekosky et al. 1996; Hamos et al. 1989). Synaptic density damage induced by toxic A $\beta$  oligomers is observed in the earliest stages of clinical AD with loss of synapses and several presynaptic proteins (Pham et al. 2010; Robinson et al. 2014; Beerli et al. 2012). However, the assessment of synapse loss and quantification of presynaptic proteins in AD could previously only be performed in post-mortem brain tissues. Thus, the ability to assess synaptic density in vivo would be extremely valuable in studies of AD as well as in monitoring the efficacy of potential therapies.

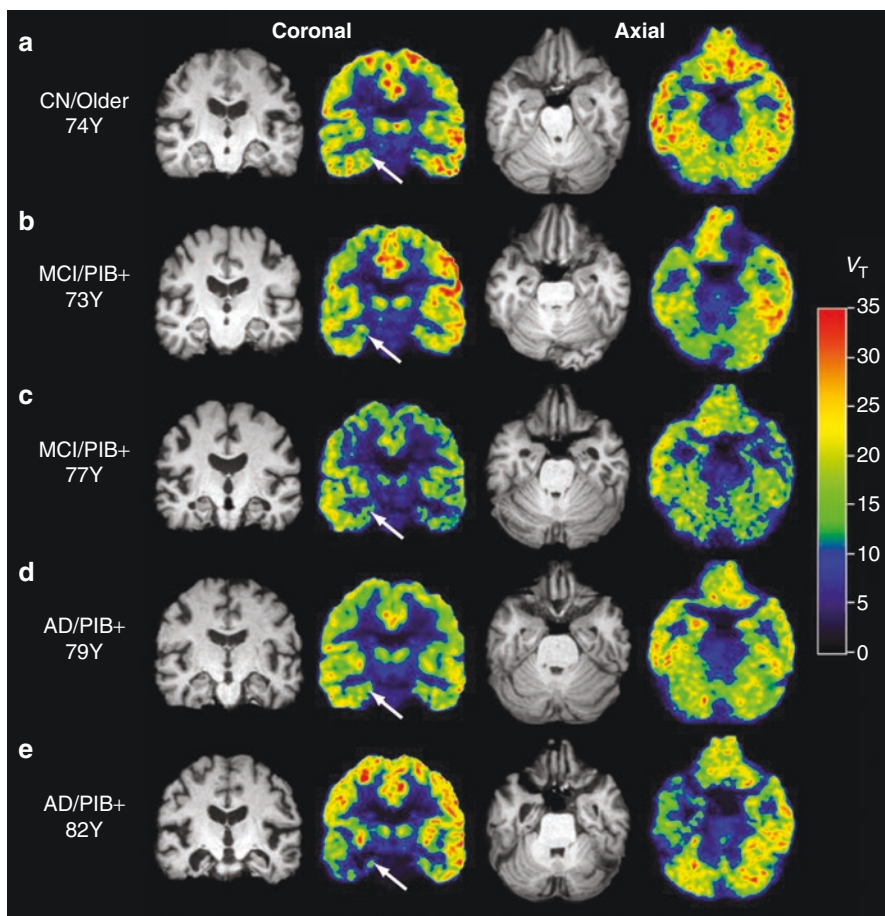
PET imaging is increasingly employed in AD studies to measure glucose metabolism (i.e., [ $^{18}\text{F}$ ]FDG),  $\beta$ -amyloid, and neurofibrillary tangles (Mckhann et al. 2011).

[<sup>18</sup>F]FDG PET is widely used clinically to differentiate AD from FTD and to track disease progression by measuring neuronal activity (Landau et al. 2011). However, [<sup>18</sup>F]FDG is metabolized by both neurons and glial cells (astrocytes, microglia, and oligodendrocytes) and is not a direct biomarker of synaptic density. [<sup>18</sup>F]FDG PET also has its limitations, e.g., brain [<sup>18</sup>F]FDG uptake is affected by stimulation, medication, and blood glucose level (Ishibashi et al. 2015; Burns et al. 2013). Inadequate fasting in healthy subjects can result in false-positive findings of [<sup>18</sup>F]FDG PET, similar to a pattern of AD (Ishibashi et al. 2015; Burns et al. 2013). SV2A imaging has potential to provide a direct indicator of synaptic density in AD.

In the first SV2A PET imaging study on quantitative assessment of synaptic density in AD using high-resolution PET, [<sup>11</sup>C]UCB-J binding was compared in 10 AD and 11 cognitively normal (CN) subjects (Chen et al. 2018). All AD subjects were confirmed  $\beta$ -amyloid positive (A $\beta$ +) by carbon-11 labeled Pittsburgh compound B ([<sup>11</sup>C]PiB) PET and spanned the disease stages from amnesic mild cognitive impairment (MCI,  $n = 5$ ) to mild dementia ( $n = 5$ ). All CN subjects were confirmed A $\beta$ –, and the two diagnostic groups were well balanced for age, sex, and education. The AD subjects had clinical characteristics typical of amnesic MCI and mild AD dementia with MMSE =  $24.1 \pm 4.8$  and CDR-global =  $0.75 \pm 0.26$  (Chen et al. 2018). Reduced [<sup>11</sup>C]UCB-J binding in the hippocampus of AD was hypothesized based on early degeneration of entorhinal cortical cell projections to the hippocampus via the perforant pathway (Gomez-Isla et al. 1996; Braak et al. 2011) and hippocampal SV2A reductions observed in AD post-mortem studies (Robinson et al. 2014). A statistically significant reduction of  $BP_{ND}$  was found by 41% in the hippocampus of AD as compared to CN ( $p = 0.011$ ) (Fig. 29.6).

The reduction of  $BP_{ND}$  in the hippocampus of AD was larger than the volume loss (22%) measured by MRI (Chen et al. 2018). Statistically significant correlations were found between synaptic density as measured by hippocampus  $BP_{ND}$  and the clinical cognitive tests including an episodic memory score (mean  $Z$  scores from LMII and RAVLT,  $R = 0.53$ ,  $p < 0.05$ ) and the clinical dementia rating sum of boxes (CDR-sb,  $R = -0.70$ ,  $p < 0.05$ ) (Chen et al. 2018). This study was the first demonstration of reduced synaptic density in the hippocampus of living AD subjects by SV2A PET imaging with [<sup>11</sup>C]UCB-J.

More recently, Bastin and coworkers reported the evaluation of [<sup>18</sup>F]UCB-H in 24 patients with MCI or AD (all A $\beta$ +) and 19 healthy controls (Bastin et al. 2020). The distribution volume ( $V_T$ ) of [<sup>18</sup>F]UCB-H was lower in the hippocampus (31%,  $d = 1.24$ ), cortical regions (11–18%,  $d = 0.49$ – $0.69$ ), and the thalamus (16%,  $d = 0.73$ ). The difference in hippocampal binding was directly related to patients' cognitive decline and unawareness of memory problems. A similar significant reduction in [<sup>11</sup>C]UCB-J binding (38%) in the hippocampus of AD ( $n = 12$ ) as compared to CN ( $n = 12$ ) was reported by the MINDMAPS consortium (Molecular Imaging of Neurodegeneration Mitochondria, Associated Proteins, Synapses) at the BrainPET 2019 conference (Venkataraman et al. 2019). They also presented a statistically significant reduction in other brain regions such as amygdala and associated cortical regions (~20%). However, the authors did indicate significant reduction



**Fig. 29.6** Representative coronal images of [ $^{11}\text{C}$ ]UCB-J parametric PET ( $V_T$ ) in CN, MCI, and mild AD dementia. The CN participant had a negative [ $^{11}\text{C}$ ]PiB scan and MCI/AD participants had positive [ $^{11}\text{C}$ ]PiB scans. The pseudo-color in PET images represents the intensity of [ $^{11}\text{C}$ ]UCB-J binding ( $V_T$ ). Evident reduction of [ $^{11}\text{C}$ ]UCB-J binding in the hippocampus of AD was noted as compared to CN (arrow denotes the right hippocampus). Adapted from Chen et al. (2018)

of [ $^{11}\text{C}$ ]UCB-J binding (18%) in the cerebellum of AD, which is usually considered spared in AD. Therefore, more studies are needed to further establish the utility of SV2A PET in AD.

In addition to studying pathophysiology in AD, preclinical studies were conducted to explore the potential of SV2A PET imaging for evaluating AD treatment effects. In a recent PET study using an AD mouse model, longitudinal [ $^{11}\text{C}$ ]UCB-J PET was performed to measure disease progression and treatment effects of saracatinib (Toyonaga et al. 2019). Saracatinib is a Fyn kinase inhibitor and has been investigated as potential therapeutic drug for AD. Saracatinib has been previously demonstrated to restore synaptic density loss in AD mice by post-mortem

immunohistochemistry for SV2A (Kaufman et al. 2015; Nygaard et al. 2014). In this study, nine wild-type (WT) mice and nine amyloid precursor protein and presenilin 1 double transgenic (APPswe/PS1 $\Delta$ E9:APP/PS1) mice underwent [ $^{11}$ C]UCB-J PET at baseline, following treatment with saracatinib and during drug washout phase. At first the authors confirmed that [ $^{11}$ C]UCB-J uptake in hippocampus with normalized SUVR values at baseline was significantly lower in AD than WT mice (AD,  $1.11 \pm 0.04$ ; WT,  $1.15 \pm 0.02$ ,  $p = 0.033$ ) (Toyonaga et al. 2019). Using SUVR-1 for the hippocampus, there was a significant difference at baseline (AD,  $0.48 \pm 0.13$ ; WT,  $0.65 \pm 0.10$ ,  $p = 0.017$ ). Following 1 month of treatment with saracatinib, hippocampal SUVR values in AD mice were significantly increased ( $p = 0.037$ ) (Toyonaga et al. 2019). Treatment effects from saracatinib may persist, as there were no significant differences between WT and AD mice after drug washout (Toyonaga et al. 2019). These results support the use of [ $^{11}$ C]UCB-J PET as a potential biomarker for detecting disease-specific synaptic deficits and to monitor treatment effects in AD.

In summary, [ $^{11}$ C]UCB-J PET provides a direct measure of SV2A and a proxy of synaptic density and therefore holds promise as a novel *in vivo* biomarker for AD. The potential clinical applications of SV2A PET imaging with [ $^{11}$ C]UCB-J in AD include the early detection of synaptic density loss as demonstrated by our pilot PET study in human and the monitoring of disease progression and the evaluation of synaptic rescue and recovery through therapeutic interventions as demonstrated by the PET study in an AD mouse model. SV2A PET could also be utilized as an outcome measure for trials of disease-modifying therapies, particularly those that target the preservation and restoration of synapses. For example, the effect of CT1812 (Elayta<sup>TM</sup>, a sigma-2 receptor antagonist) treatment on brain synaptic density in participants with mild to moderate AD is currently being evaluated by CogRx (Cognition Therapeutics Inc.) in collaboration with Yale School of Medicine using [ $^{11}$ C]UCB-J-PET (NCT03493282).

### 29.5.3 Parkinson's Disease

Parkinson's disease (PD) is characterized as a neurodegenerative disease with predominant motor and non-motor symptoms. Despite well-documented dopamine neuron decreases, the etiology of PD is only partially understood. While studies have found cortical and limbic Lewy bodies or amyloid plaque pathology (Sabbagh et al. 2009), the wide range of clinical difficulties in PD have been proposed to represent varying rates and regional patterns of neurodegeneration. Furthermore, there is evidence of widespread brain pathology involving multiple transmitter systems and pathways, including the cholinergic, noradrenergic, and serotonergic systems and not just a single neural structure or neurotransmitter (i.e., substantia nigra and dopamine) (Todorova et al. 2014; Jellinger 2012; Picconi et al. 2012; Sauerbier et al. 2016).

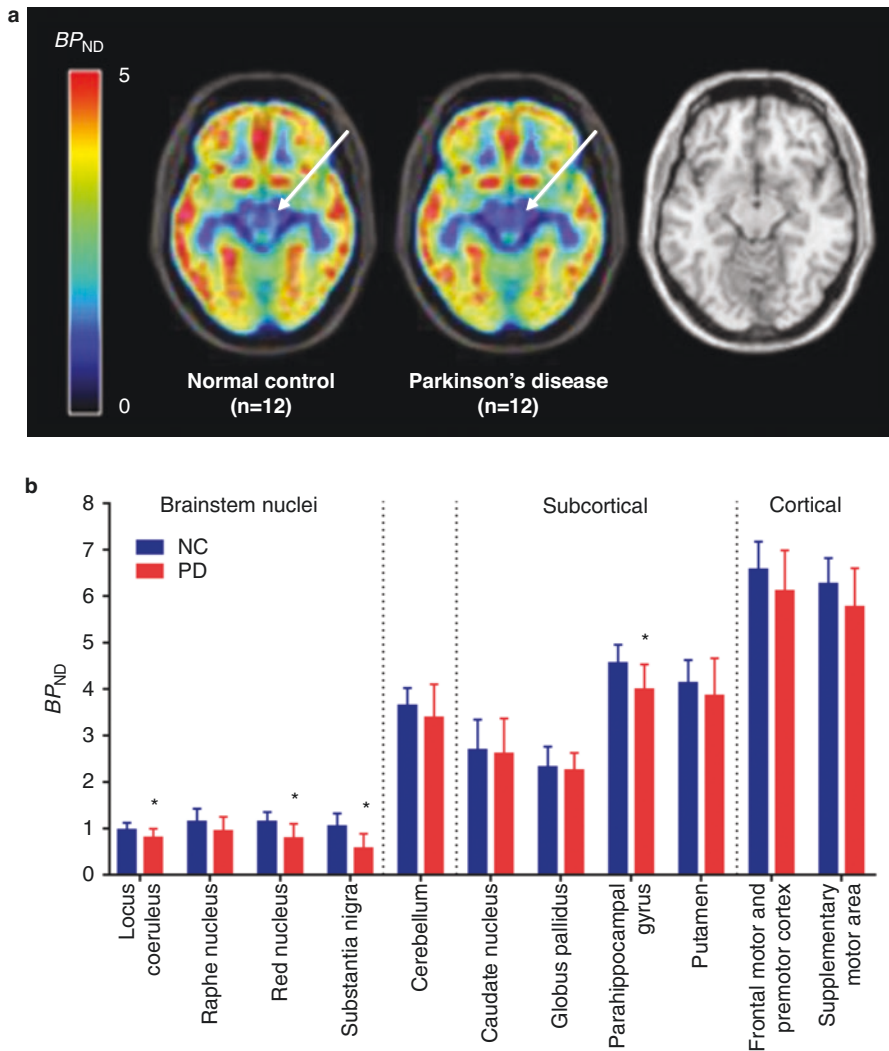
Thus, examining synaptic density is exceedingly relevant in PD as current evidence has demonstrated synaptic changes in both animal models and post-mortem

patients with the disease, particularly in the later stages of PD when clinical manifestations develop and decreases in synaptic density are overwhelmingly prevalent (Picconi et al. 2012; Mathai et al. 2015; Hallett et al. 2005; Villalba and Smith 2013; Zaja-Milatovic et al. 2005). These synaptic losses have also been found outside of the nigrostriatal system and include non-dopamine neurons in the cortex, implicating widespread synaptic damage throughout the brain (Hou et al. 2010). In addition, accumulating evidence implicating gene mutations (e.g.,  $\alpha$ -synuclein and LRRK2) suggests that synaptic dysfunction is centrally involved in PD and that synaptic loss may be a characteristic in the pathogenesis of the disease (Picconi et al. 2012).

The Yale PET Center has recently reported a study with [ $^{11}\text{C}$ ]UCB-J examining 12 PD subjects with mild bilateral disease compared to 12 demographically matched healthy controls (Matuskey et al. 2020). Lower SV2A  $BP_{\text{ND}}$  values were observed in all brain regions examined, with the largest differences in the brainstem regions including the substantia nigra ( $-45\%$ ;  $p < 0.001$ ), red nucleus ( $-31\%$ ;  $p = 0.03$ ), and locus coeruleus ( $-17\%$ ;  $p = 0.03$ ) (Fig. 29.7a,b). Cortical areas also exhibited significantly lower  $BP_{\text{ND}}$  in PD subjects in the posterior cingulate cortex ( $-15\%$ ;  $p = 0.01$ ), parahippocampal gyrus ( $-12\%$ ;  $p < 0.01$ ), orbitofrontal cortex ( $-11\%$ ;  $p = 0.01$ ), and ventromedial prefrontal cortex ( $-11\%$ ;  $p = 0.02$ ; all  $p$  values are uncorrected) (Matuskey et al. 2020). This study provided the first in vivo evidence of decreased synaptic density in PD, and ongoing follow-up work focuses on the sensitivity and longitudinal dimensions of these findings.

### 29.5.4 Major Depressive Disorder

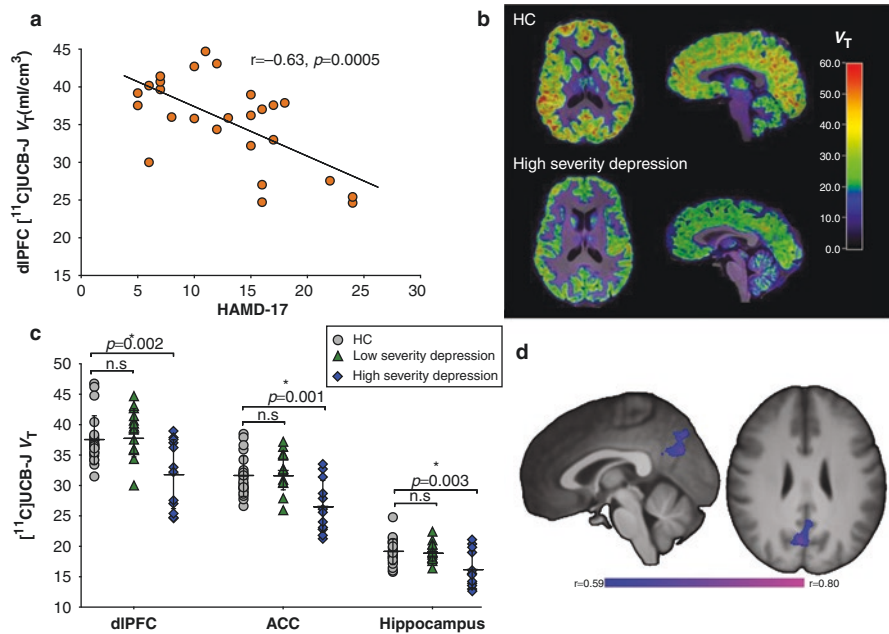
Stress has a profound impact on the human brain, affecting both its structure and function (Duman and Aghajanian 2012). Indeed, persisting levels of stress can lead to a loss of synapses in circuits underlying affective and cognitive processing, which is thought to contribute to the symptoms of major depressive disorder (MDD) and posttraumatic stress disorder (PTSD) (Duman et al. 2016) – disorders which are highly prevalent and debilitating, yet whose treatment is currently inadequate. Convergent research indicates that synaptic loss contributes to stress-related disorders such as MDD and PTSD, and lower synaptic density is consistently observed in preclinical models of chronic stress and depression (Duman and Aghajanian 2012; Liu and Aghajanian 2008; Kassem et al. 2013). A lower number of synaptic-function-related genes and synaptic signaling proteins have been demonstrated in post-mortem brain tissue of individuals with MDD (Kang et al. 2012; Feyissa et al. 2009; Duric et al. 2013), meanwhile MRI studies have consistently shown lower brain volume in the prefrontal cortex (PFC), anterior cingulate cortex (ACC), and hippocampus in both MDD (Price and Drevets 2010; Bora et al. 2012) and PTSD (Kühn and Gallinat 2013), likely in part due to synaptic and neuronal loss (Duman et al. 2016). Stress-related alterations in synaptic connections are also thought to contribute to the network dysfunction observed in both MDD (Scheinost et al. 2018; Kaiser et al. 2015; Murrough et al. 2016) and PTSD (Akiki et al. 2017). Increasing



**Fig. 29.7** (a) The midbrain area with arrows denoting the substantia nigra (SN) is shown on the averaged group  $BP_{ND}$  [11C]UCB-J parametric images for normal control (left) and Parkinson's disease (center) subjects. Magnetic resonance imaging template is also shown for comparison purposes (right). Color scale denotes binding potential ( $BP_{ND}$ ). (b) Region of interest analysis of binding potential ( $BP_{ND}$ ) values for normal control (NC) and Parkinson's disease (PD) subjects.  $BP_{ND}$  values were lower across all primary brain regions between the NC and PD subjects. \*Statistical significance at  $p = 0.05$  or better on post hoc tests. Error bars denote standard deviation. Modified after Matuskey et al. (2020)

synaptic connections, therefore, appears to be a promising approach for the treatment of symptoms of MDD and PTSD which are often resistant to conventional antidepressants. In line with this, ketamine (an NMDA receptor antagonist with rapid antidepressant effect and recently approved by the FDA for treatment-resistant depression) is thought to exert its therapeutic effects in MDD (McGirr et al. 2015; Krystal et al. 2013; Esterlis et al. 2018) and PTSD (Feder et al. 2014) through restoration of synaptic connections (Li et al. 2010; Duman and Aghajanian 2012; Hare et al. 2017), thus adding further support to the notion that synaptic loss contributes to the neurobiology underlying depressive symptoms in these disorders. However, evaluation of synaptic density in MDD and PTSD has until now been confined to post-mortem and preclinical quantification.

In the first in vivo study to investigate synaptic density in depression, the relationship between [ $^{11}\text{C}$ ]UCB-J binding and depression severity was evaluated in a transdiagnostic sample of 26 individuals with MDD and/or PTSD and compared to 21 healthy comparison (HC) subjects (Holmes et al. 2019). The primary outcome measure was  $V_T$ , calculated using a 1TC model with a metabolite-corrected arterial input function, in the dorsolateral PFC (dlPFC), ACC, and hippocampus. Loss of synaptic connections is likely to have consequences for network organization and function. Therefore, a secondary analysis was applied to examine intrinsic connectivity distribution (ICD) (Scheinost et al. 2012), a measure of whole-brain connectivity obtained via functional MRI (fMRI), and a follow-up seed connectivity was used to explore network alterations in relation to synaptic density and symptom severity. The primary finding was a negative correlation between [ $^{11}\text{C}$ ]UCB-J  $V_T$  and symptoms of depression (measured using the Hamilton Depression Rating Scale; HAMD-17 (Hamilton 1967)) across individuals with MDD and PTSD in dlPFC ( $r = -0.63$ ,  $p = 0.001$ ), ACC ( $r = -0.63$ ,  $p = 0.001$ ) and hippocampus ( $r = -0.49$ ,  $p = 0.01$ ), such that lower synaptic density was associated with more severe depression (Fig. 29.8a). There were no significant differences in [ $^{11}\text{C}$ ]UCB-J  $V_T$  between the full clinical sample and HCs. However, after stratifying clinical subjects by severity of depressive symptoms, a significantly lower [ $^{11}\text{C}$ ]UCB-J  $V_T$  was observed in individuals with higher depression severity vs. HCs across ROIs ( $p = 0.005$ ; Fig. 29.8b,c), but no difference between mild depression and HC groups. Exploratory whole-brain functional connectivity analyses revealed significantly lower whole-brain functional connectivity in the dlPFC ( $p < 0.05$ , corrected) in the full clinical sample. Using the dlPFC as seed, a significant negative correlation was observed between dlPFC-posterior cingulate cortex (PCC) connectivity and SV2A density in the dlPFC ( $r = -0.60$ ,  $p = 0.002$ ), as well as with depression severity ( $r = 0.41$ ,  $p = 0.042$ ) (Fig. 29.8d), suggesting that lower synaptic density in the dlPFC is associated with downstream network dysfunction. Specifically, it was demonstrated that higher connectivity between central nodes of the default mode network (PCC) and central executive network (dlPFC) is associated with higher depression severity. Importantly, these two networks are typically anticorrelated (Fox et al. 2005) and are thought to be “blurred” in depression, possibly reflecting a bias toward internal



**Fig. 29.8** (a) Correlation between dIPFC  $[^{11}\text{C}]\text{UCB-J } V_T$  and depression severity (as measured using the HAMD-17) across MDD and PTSD subjects ( $n = 26$ ). Correlations were also significant in ACC and hippocampus. (b) Representative parametric  $V_T$  images registered to MR images from a HC subject (top) and a subject with high severity depression (bottom). (c)  $[^{11}\text{C}]\text{UCB-J } V_T$  in the dIPFC, ACC, and hippocampus across groups. The low severity group (in green;  $n = 14$ ) consisted of participants with HAMD-17 scores  $<14$  and the high severity group (in blue;  $n = 12$ ) consisted of participants with HAMD-17 scores  $\geq 14$ . (d) Negative correlation between dIPFC-PCC connectivity and dIPFC SV2A density in clinical subjects. Significant voxels represent those voxels whose connectivity with the dIPFC negatively correlates with  $[^{11}\text{C}]\text{UCB-J } V_T$  in the dIPFC. Adapted from Holmes et al. (2019)

thoughts at the expense of engagement with the external world (Anticevic et al. 2012). Taken together, the findings suggest that lower synaptic density in the dIPFC may underlie “blurring” of networks in depression, possibly reflecting reduced top-down control over network function.

In summary, the use of  $[^{11}\text{C}]\text{UCB-J}$  PET provided the first in vivo evidence of lower synaptic density in association with depressive severity in MDD and PTSD. Importantly, these findings provide further incentive to discover and evaluate treatment interventions that increase synaptic connections and reverse the loss of synapses caused by stress. Hopefully, SV2A PET imaging will aid in the search for targeted treatments that are critically needed to reduce the suffering and burden caused by depression.



## 29.5.5 Schizophrenia

Schizophrenia (SCZ) is a debilitating psychiatric disorder, of as yet unknown etiology, that affects about 1% of the general population and is associated with significant morbidity, health expenditure, and disability (Owen et al. 2016). Reduced dendritic spine density in several brain regions (Glantz and Lewis 2000; Garey 2010; Roberts et al. 2015; Bakhshi and Chance 2015; Osimo et al. 2019) is one of the most consistent neuropathological findings seen in post-mortem studies of SCZ. Additionally, post-mortem studies show lower expression of region-specific presynaptic transcripts/proteins (e.g., synaptophysin, SNAP-25, and rab3a) (Faludi and Mirnics 2011). Moreover, genetic studies in SCZ have implicated genes encoding presynaptic proteins (including syntaxin1a, SNAP-25) (Dyck et al. 2011; Ayalew et al. 2012) and relevant to synaptic integrity and function (e.g., ZNF804A, complement C4 gene) (Deans et al. 2017; Sekar et al. 2016). Thus, converging evidences implicate synaptic pathology in SCZ. In the meantime, a recent post-mortem study showed that lower synaptic spine density in SCZ was found specifically in immature spines rather than mature spines, suggesting that the synaptic density differences are not secondary to synaptic pruning (Macdonald et al. 2017), thereby raising the possibility that this finding is a result of disease-specific factors such as age at onset of illness, duration of illness, and illness severity.

Post-mortem studies are, however, confounded by a number of methodological limitations related to factors such as the mode of collection of specimens, post-mortem interval, storage, and nature of death (Harrison 2011). Additionally, it is challenging to address the confounding effects of disease-specific factors such as age at onset of illness, duration of illness, illness severity, effects of antipsychotic treatment, drugs/alcohol abuse, and tobacco use, all of which are common in SCZ (Harrison 2011). Furthermore, post-mortem findings cannot be related to concurrent measures of the illness, e.g., symptom severity or cognitive function.

Onwordi et al. recently reported the evaluation of [ $^{11}\text{C}$ ]UCB-J in control subjects ( $n = 18$ ) and SCZ patients on antipsychotic medication ( $n = 18$ ) (Onwordi et al. 2020). A significant group difference and group-by-region interaction effect on  $V_T$  were found, with lower  $V_T$  values in the frontal ( $-13\%$ ,  $p = 0.03$ ) and anterior cingulate ( $-13\%$ ,  $p = 0.02$ ) cortices. Evaluation of distribution volume ratio (DVR) using the CS as a reference region resulted in significantly lower SV2A in frontal and anterior cingulate cortices and the hippocampus in SCZ. Exploratory analysis of additional brain regions also showed lower SV2A in the dorsolateral prefrontal and temporal cortices and occipital lobe in SCZ. Since SCZ patients were not drug-naïve, the authors examined the effect of antipsychotic drug administration on SV2A levels in rats using western blot, SV2A autoradiography, and confocal microscopy and found no significant effect of drug administration on SV2A outcome measures. This was the first PET study to demonstrate lower levels of a presynaptic protein marker in SCZ and confirmed previous post-mortem evidence. These preliminary findings demonstrate the utility of SV2A PET imaging to further understand the neurobiology of SCZ and warrant further evaluation in larger study cohorts.

## 29.6 Future Considerations

In summary, PET imaging of SV2A has been rapidly established during the last decade. High-quality PET radioligands labeled with carbon-11 and fluorine-18 have been developed and validated for application in human studies. The utility of SV2A as a marker of synaptic density is promising and warrants formal validation by comparison of in vivo SV2A PET signal to post-mortem evaluation of SV2A and synapse levels in the human brain. The first PET studies in patient groups indicate SV2A loss specific to disease-associated brain regions and consistent with synaptic density loss. Future studies may apply SV2A PET imaging to study neuronal development and neurodegeneration in the healthy and diseased brain. Further studies also warrant use of fluorine-18 labeled radioligands in larger patient cohorts to establish the clinical value of SV2A PET imaging and its potential for diagnosis and progression monitoring of neuropathological diseases, as well as assessment of drug treatment effects.

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# PET Imaging of Translocator Protein Expression in Neurological Disorders

# 30

David J. Brooks

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## Abstract

Microglia provide the intrinsic immune defence of the brain and are activated by any injurious process. As such they provide a non-specific marker of disease activity. Their function can be both detrimental and beneficial as they release cytokines which drive disease progression but also release restorative growth factors, help clear cellular debris and abnormal protein aggregations and can remodel connections as an adaptive response to brain damage. Activated microglia express translocator protein (TSPO), and this allows them to be imaged in vivo with positron emission tomography (PET) radioligands which are

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substrates. In this review the role of TSPO imaging with PET is discussed in neurodegenerative and inflammatory brain diseases and in focal brain injury due to trauma or stroke.

### 30.1 Introduction

The 18 kDa translocator protein (TSPO), previously known as the peripheral benzodiazepine receptor (PBR), is present in many peripheral organs including the liver, spleen, adrenals and myocardium (Banati 2002b). It is expressed only at low levels in a normal brain, but if microglial cells become activated, TSPO can be detected in their outer mitochondrial membrane. The TSPO serves a number of functions including transport of cholesterol, anions, nucleosides and porphyrins; maintenance of mitochondrial membrane potential; regulation of cell apoptosis and proliferation and immunomodulation (Papadopoulos et al. 2006). The protein expresses a binding site for benzodiazepines and isoquinolines such as PK11195, and this has allowed the detection of TSPO expression in vivo with (*R*)-[<sup>11</sup>C]PK11195 PET. Newer PET biomarkers are now available to image TSPO expression which include phenoxyarylacetamides such as [<sup>11</sup>C]/[<sup>18</sup>F]DAA1106, [<sup>11</sup>C]PBR28, [<sup>18</sup>F]PBR06, [<sup>18</sup>F]PBR111 and [<sup>18</sup>F]FEPPA; pyrazolo-[1,5-a]-pyrimidines such as [<sup>11</sup>C]DPA713 and [<sup>18</sup>F]DPA714; the vinca alkaloid [<sup>11</sup>C]vinpocetine and the imidazo-[1,2a]-pyridines [<sup>11</sup>C]/[<sup>18</sup>F]CLINDE (Dolle et al. 2009; Doorduyn et al. 2008).

Microglia account for around 15% of the cerebral white cell population. They are normally in a resting state sending out long ramified processes which contact neighbouring neurones and astrocytes (Kreutzberg 1996; Ransohoff and Perry 2009). It is, therefore, likely that the function of resting microglia is to monitor changes in the local brain milieu—for a review see (Hanisch and Kettenmann 2007). Microglia are cells of monocyte lineage and are protected from antigens in the plasma by the blood-brain barrier. They form the natural immune defence of the brain, and exposure to plasma proteins such as fibrinogen following blood-brain barrier disruption, or to intrinsic excitotoxic agents such as raised glutamate, nitric oxide or cytokines, causes them to become activated taking on amoeboid- or rod-like morphology. When activated, the microglia express MHC class 1 and 2 antigens and can release cytokines such as TNF $\alpha$ , IL1 $\beta$  and IL6, or growth factors such as TGF- $\beta$ 1. They may become phagocytic clearing cellular debris and stripping and remodelling synapses. It is now considered that activated microglia may exist as two primary phenotypes: M1 which is associated with the release of cidal cytokines so promoting cell damage and M2 which is associated with phagocytosis of dead tissue, synaptic remodelling and growth factor release promoting neurogenesis (Boche et al. 2013; Varnum and Ikezu 2012). These two primary phenotypes are thought to be interconvertible, have overlapping functions and predominate at different disease phases. It has been suggested that substrates binding to the TREM2 receptor on microglia promote activation of the M2 phenotype (Li and Zhang 2018). After an acute stroke, activated microglia act locally to wall off, remove dead tissue and remodel connections, while

in chronic neurodegenerative diseases, they may release cytokines where disease is locally active but remodel distant connections in the brainstem and thalamus. The advent of TSPO PET agents has allowed us to image in life the distribution of activated microglia in the brain. Current TSPO PET ligands are unable to discriminate between M1 and M2 phenotypes and provide a measure of total activated microglia load rather than indicating whether their function is protective or toxic in nature.

## 30.2 Imaging TSPO with PET

The PET ligand that was initially used for imaging TSPO expression in the brain and has seen the greatest use is the isoquinoline (*R*)-[<sup>11</sup>C]-PK11195. The rat unilateral facial nerve crush model results in activated microglia in the disconnected ipsilateral facial nucleus. Autoradiography studies have shown that [<sup>3</sup>H]PK11195 binds selectively to these activated microglia which are involved in remodelling connections to restore facial muscle function (Banati et al. 1997). A human equivalent of this rodent model is Bell's palsy where the facial nerve becomes unilaterally compressed due to local inflammation and swelling as it passes through the auditory canal. (*R*)-[<sup>11</sup>C]PK11195 PET studies have demonstrated tracer uptake in the facial nerve nucleus ipsilateral to the paralysed facial muscles (Banati 2002b). Human subjects with acquired upper limb amputations develop phantom limb phenomena where the absent limb still feels present but telescoped into the stump. In these subjects one can detect thalamic inflammation contralateral to the missing limb with (*R*)-[<sup>11</sup>C]PK11195 PET (Banati 2002a). These studies, therefore, reveal the microglial activation resulting from the disconnection of brain nuclei due to peripheral injury, presumably playing an active role in the remodelling of connections. It remains to be determined whether these cells are primarily expressing an M2 phenotype.

In the majority of brain disorders, (*R*)-[<sup>11</sup>C]PK11195 PET reveals microglial activation due to both local disease activity and the effects of downstream disconnection. Additionally, as endothelial cells also express TSPO, tracer binding is seen in the lateral and sagittal venous sinuses, and this signal can spill over into adjacent brain tissue—particularly the cerebellum—due to the 4–5 mm spatial resolution of most commercial PET cameras. This makes quantitative modelling of brain (*R*)-[<sup>11</sup>C]PK11195 PET problematic as there is no anatomical region that provides a pure tissue reference for non-specific tracer uptake though cerebellar grey matter has been used. The use of an arterial plasma input function with this tracer is also problematic as (*R*)-[<sup>11</sup>C]PK11195 sticks to plastic tubing, making it difficult to obtain blood time-activity curves which allow an accurate measurement of peak height, delay and dispersion. For these reasons, a modelling approach has been developed that uses supervised cluster analysis to identify clusters of voxels that fall into six classes of brain uptake kinetics. One of these clusters represents a collection of grey matter voxels in the subject's brain that have a time-activity curve (TAC) similar to that of a normal grey matter TAC in a population of healthy controls, while another represents vascular binding (Anderson et al. 2007). The normal grey

matter reference cluster can be used to compute non-specific (*R*)-[<sup>11</sup>C]PK11195 uptake in other brain clusters where specific tracer retention is occurring. At the same time, the vascular signal due to tracer binding, which shows the most rapid uptake, can be separated from adjacent brain parenchymal signal. The (*R*)-[<sup>11</sup>C]PK11195 binding potentials (BPs) are then computed using a standard simplified reference tissue model (SRTM) with brain-specific and non-specific compartments. As (*R*)-[<sup>11</sup>C]PK11195 binding increases in a normal thalamus and, to a lesser degree, cortex with age, it is important to age match healthy controls to patients when assessing levels of inflammation in disease states (Cagnin et al. 2001).

Recently, second-generation TSPO tracers have been developed which can have higher affinity and/or a lower non-specific signal in order to provide a more sensitive detection of microglial activation and facilitate modelling approaches. However, with these new tracers has come the realisation that TSPO PET imaging is influenced by TSPO gene polymorphisms expressed by individuals (Owen et al. 2012). The most influential of these is the rs6971 polymorphism where Ala147Thr substitutions result in homozygous subjects becoming low affinity binders of these newer TSPO ligands while heterozygotes express a mixture of TSPO with high and low affinity for these ligands. In Caucasian populations, around 60% of individuals are high, 10% low and 30% mixed affinity binders for the newer TSPO ligands. (*R*)-[<sup>11</sup>C]PK11195 affinity for TSPO, however, appears to be less influenced by genotype (Owen et al. 2012), while [<sup>11</sup>C]PBR28 shows a 75-fold difference in affinity for TSPO between high (K<sub>d</sub> 4 nM) and low (K<sub>d</sub> 300 nM) affinity binders. The PET tracers [<sup>11</sup>C]DAA1106, [<sup>11</sup>C]DPA713 and [<sup>18</sup>F]PBR111 show 4–5-fold differences in affinity for TSPO between high and low binders. Mixed affinity binders express high and low affinity TSPO binding sites in equal proportions. If one is to use these newer TSPO PET markers, then low affinity binders will need to be excluded by prior genetic screening and patient and control populations matched for prevalence of mixed and high affinity binders. When interrogating associations between TSPO PET signals and pathology or clinical parameters, the variance due to TSPO polymorphisms may need to be corrected by ANCOVA.

Modelling the brain uptake kinetics of second-generation TSPO PET tracers has similar problems to those described previously for (*R*)-[<sup>11</sup>C]PK11195. In practice cerebellar grey matter has often been employed as a tissue reference for non-specific binding for tracers such as [<sup>18</sup>F]DPA714 and [<sup>11</sup>C]PBR28 although this can lead to underestimations of specific binding in regions of interest. Supervised cluster analysis has also been applied to define reference clusters for individual patients and correct spill over of vascular signals. Arterial input functions have been also used in some series (Schain et al. 2018).

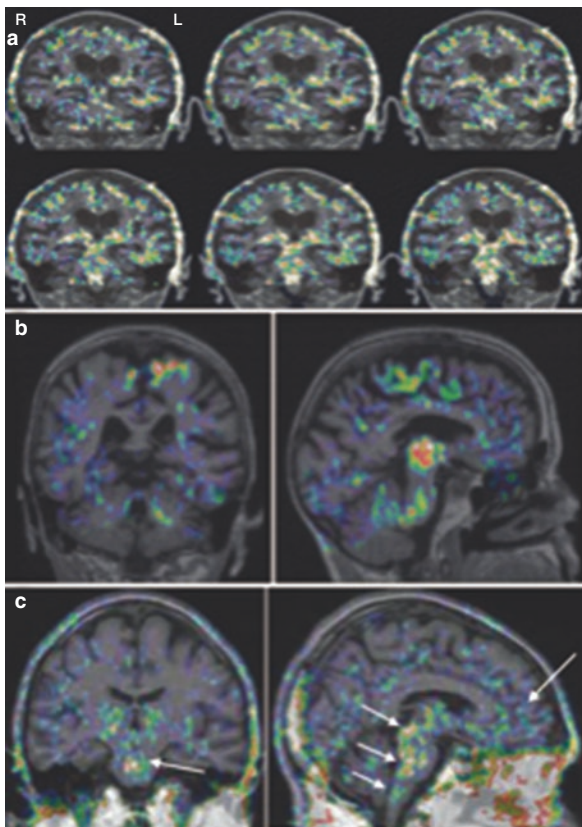
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### 30.3 TSPO Imaging in Alzheimer's Disease

Dementia affects 10% of the over 60s, the prevalence rising to 30% by the ninth decade. It is characterised clinically by progressive impairment of memory, speech and perception along with personality change in the absence of altered conscious

level. Sixty percent of dementia cases have Alzheimer pathology characterised by extracellular fibrillar  $\beta$ -amyloid plaques and intra-neuronal neurofibrillary tangles of hyper-phosphorylated tau at post-mortem. Activated microglia are seen surrounding the neuritic amyloid plaques which target association cortex and cingulate but are less evident in the striatum where plaques are diffuse (Braak and Braak 1997; Dickson 1997). Activated microglia are also found in brain areas surrounding neurites with tau tangle pathology such as the entorhinal cortex and hippocampus.

Raised levels of microglial activation were first imaged *in vivo* in clinically diagnosed Alzheimer's disease (AD) patients with (*R*)-[ $^{11}\text{C}$ ]PK11195 PET, binding potentials being raised by up to 50% in association cortex (Cagnin et al. 2001; Schuitmaker et al. 2013) (Fig. 30.1a). Uptake of the TSPO ligand [ $^{11}\text{C}$ ]DAA1106 has also been reported to be elevated by up to 33% in AD (Yasuno et al. 2008). The cortical distribution of raised (*R*)-[ $^{11}\text{C}$ ]PK11195 uptake parallels that of  $\beta$ -amyloid plaque deposition revealed with [ $^{11}\text{C}$ ]PIB PET, a marker of fibrillar amyloid load



**Fig. 30.1** (*R*)-[ $^{11}\text{C}$ ]PK11195 PET images of microglial activation in patients with (a) Alzheimer's disease, (b) motor neuron disease, (c) Parkinson's disease, (d) multiple sclerosis



**Fig. 30.1** (continued)

(Edison et al. 2007a), targeting association rather than primary areas. Levels of cortical (*R*)-[<sup>11</sup>C]PK11195 and [<sup>11</sup>C]PBR28 uptake in AD have been reported to correlate with cognitive impairment rated with the mini-mental state examination (MMSE) and CDR-SOB, but there was no correlation between amyloid load and cognitive status (Edison et al. 2007b; Kreisl et al. 2016; Yokokura et al. 2011). In established Alzheimer's disease, amyloid plaque load plateaus while cognitive ability declines, and these findings suggests that it may be the cortical microglial activation rather than amyloid plaques in AD that are detrimental to cognitive function, possibly due to cytokine release by cells exhibiting a cidal M1 phenotype. Along with the raised cortical TSPO signal, activated microglia can also be detected in the thalamus, cerebellum and brainstem of Alzheimer patients. As these subcortical areas are only targeted in late disease by plaques, the (*R*)-[<sup>11</sup>C]PK11195 signal there is likely to reflect microglial activation either due to local tau tangle disease or in response to cortical disconnection, the cells acting to remodel connections.

(*R*)-[<sup>11</sup>C]PK11195, [<sup>11</sup>C]DAA1106 and [<sup>18</sup>F]DPA714 PET have all been reported to detect the presence of microglial activation in a proportion of amnesic mild cognitive impairment (MCI) cases. These subjects have isolated progressive recall problems that are not severe enough to interfere with activities of daily living, but around 60% of these cases progress over 5 years to develop Alzheimer's disease (Petersen et al. 2001). In an initial small series, 60% of MCI cases showed evidence of amyloid plaque deposition with [<sup>11</sup>C]PIB PET and so represented prodromal Alzheimer's disease (pAD). (*R*)-[<sup>11</sup>C]PK11195 PET detected microglial activation in 40% of these amnesic MCI cases; that is two thirds of the pAD subjects (Okello et al. 2009). However, another series failed to detect microglial activation in amnesic MCI with (*R*)-[<sup>11</sup>C]PK11195 PET—possibly because these workers used a cerebellar reference rather than defining normal voxels with supervised cluster analysis (Schuitemaker et al. 2013). A group of seven MCI cases showed a significant mean of 27% increase in tracer uptake across brain regions when their TSPO was imaged with [<sup>11</sup>C]DAA1106 PET, and two cases individually had microglial activation elevated more than 2 SD above the control mean (Yasuno et al. 2012). Five of these seven MCI subjects with [<sup>11</sup>C]DAA1106 uptake raised more than 0.5 standard deviations above the normal mean progressed to develop dementia over a 2-year follow-up period.

More recently, Parbo and colleagues scanned 42 MCI cases with [ $^{11}\text{C}$ ]PiB and (R)-[ $^{11}\text{C}$ ]PK11195 PET (Parbo et al. 2017). Twenty-six (62%) of the cases had raised cortical [ $^{11}\text{C}$ ]PiB uptake, and 17 (65%) of these pAD cases had evidence of cortical inflammation. Clusters of correlated levels of amyloid and inflammation were found in the frontal, temporal and parietal cortex. In a follow-up study, tau tangle load was imaged with [ $^{18}\text{F}$ ]flortaucipir PET in a subgroup of 20 MCI cases and 6 AD patients (Parbo et al. 2018). While amyloid load and inflammation levels were again correlated in cortical clusters of voxels, there was surprisingly no correlation evident between tau tangle load and inflammation. However, a 2-year follow-up study on these subjects has now detected a correlation between cortical tau and inflammation levels in the MCI and early AD cohort though this was not evident at baseline (R Ismail, Brooks DJ. *Neurobiology of Disease* 2020;17(1):151).

Using [ $^{18}\text{F}$ ]DPA714 PET, Hamelin and colleagues have studied the levels of microglial activation in prodromal (38) and early clinical Alzheimer's disease (26) and its relationship with clinical progression (Hamelin et al. 2016). The presence of raised amyloid load was confirmed in all cases with [ $^{11}\text{C}$ ]PiB PET, and, based on their MMSE score changes over 2 years, the AD group was divided into fast and slow decliners. The levels of TSPO binding were determined as target region:cerebellar ratios, and findings for high and mixed affinity binders were combined. TSPO binding in AD was raised relative to healthy controls, and cortical levels correlated negatively with MMSE scores and hippocampal volume and positively with CDR-SOB ratings and [ $^{11}\text{C}$ ]PiB uptake. The slow and fast decliners had similar baseline levels of amyloid load, but baseline TSPO uptake was higher in the slow decliners. The authors suggested that microglial activation may be playing a protective role in early AD. In a follow-up study, these workers repeated [ $^{18}\text{F}$ ]DPA714 PET in their AD patients and control subjects and found a greater increase in inflammation in AD (Hamelin et al. 2018). However, the increase in [ $^{18}\text{F}$ ]DPA714 uptake was lower in those cases with higher baseline inflammation. The authors suggested that inflammation has two distinct profiles in AD, an initial protective profile followed by a detrimental effect on disease progression.

In favour of a protective role of inflammation in early AD, an [ $^{11}\text{C}$ ]PBR28 PET series in 37 MCI amyloid-positive cases has reported that levels of TSPO binding correlated positively with cortical and hippocampal volumes (Femminella et al. 2019). Here an arterial input function with Logan graphical analysis was employed to measure tracer volumes of distribution. Longitudinal data is not yet available for this cohort.

Recently, our group has followed 38 MCI cases over 2 years with [ $^{11}\text{C}$ ]PK11195 and [ $^{11}\text{C}$ ]PiB PET (Ismail R and Brooks DJ. *Neurobiology of Disease* 2020;17(1):151). Twenty-two of these MCI subjects also had [ $^{18}\text{F}$ ]flortaucipir PET to determine their tau tangle load. At baseline 23 MCI cases had raised and 15 cases had low/normal levels of cortical PiB uptake. Over 2 years the combined group of 38 MCI cases showed a rise in mean amyloid load but a decrease in mean cortical level of inflammation. In those 22 cases who had [ $^{18}\text{F}$ ]flortaucipir PET, their cortical tau tangle load increased over time. Seven of the low PiB cohort showed increasing tracer uptake over 2 years, three crossing into the abnormal range. These cases

showed a correlation between their cortical PiB uptake and inflammation levels in the absence of tau tangle deposition. In the MCI group with raised PiB, the level of inflammation correlated with tau tangle load. These findings suggest that inflammation plays a role both in early MCI when amyloid load is rising—possibly protective—and in later MCI when tau tangles are forming, possibly toxic. The relationship between pathology and cognitive status was also followed in this MCI cohort. Amyloid plaque load and levels of inflammation across cortical regions at baseline correlated with cognitive deficit, while tau load correlated most strongly at the 2-year follow-up time point.

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### 30.4 Imaging Inflammation in Frontotemporal Dementia and ALS

Dementia with a frontotemporal phenotype comprises around 10% of later onset dementia cases, and clinically it presents as personality change, praxis and language difficulties with memory becoming impaired subsequently. Clinical phenotypes include behavioural variant frontotemporal dementia (bvFTD), semantic dementia (SD) where subjects cannot recall the features and uses of objects and progressive non-fluent aphasia and apraxia (PNFA). The pathology targets the frontal and inferior temporal lobes and most commonly involves TDP-43 protein inclusions when it can also be associated with ALS. Less commonly Pick bodies containing three-repeat tau isoforms or spongiform degeneration are present.

25–50% of FTD cases are genetic in origin, and multiple gene mutations have been implicated. Repeat expansions of the C9ORF72 gene are the most common genetic cause and are associated with TDP-43 inclusions as are mutations of the progranulin (GRN) gene. MAPT gene mutations can cause Pick body tau inclusion disease. Mutations of TREM2 have been associated with FTD implicating a role of inflammation (Zhang 2015). Presenilin gene mutations also can rarely cause FTD as can mutations of the mitochondrial CHCHD10 gene. FTD is usually younger onset than AD and rarely associated with amyloid plaque formation.

The first PET series to study inflammation reported raised frontotemporal uptake of (*R*)-[<sup>11</sup>C]PK11195 in four idiopathic PFNA and a fifth bvFTD case (Cagnin et al. 2004). In some but not all regions, the inflammation was associated with cortical atrophy. More recently, increased [<sup>11</sup>C]PBR28 uptake has been documented in bvFTD cortex targeting association areas in patterns matching the clinical phenotypes of the patients (Kim et al. 2019). Larger series including presymptomatic susceptibility gene carriers are required to determine whether inflammation precedes neuronal dysfunction and aberrant protein aggregation in FTD.

There is both a clinical and pathological overlap between frontotemporal dementia and motor neuron disease (amyotrophic lateral sclerosis). The neuropathology of both can be associated with cortical TDP-43 inclusions and spongiform degeneration, and neuropsychological and imaging studies have indicated that dysfunction extends beyond the motor system in ALS. In an early series, ten probable or definite (El Escorial criteria) ALS patients without dementia had (*R*)-[<sup>11</sup>C]PK11195 PET (Turner



et al. 2004). Significantly increased microglial activation was seen in the motor cortex, brainstem, thalamus, dorsolateral prefrontal cortex and occipital lobes of the ALS patients compared with 14 healthy controls (Fig. 30.1b). There was a significant correlation between the severity of upper motor neuron signs on examination and levels of motor cortex (*R*)-[<sup>11</sup>C]PK11195 uptake. More recently, ten non-demented patients with probable or definite ALS were enrolled prospectively and eight healthy controls matched for age had [<sup>18</sup>F]DPA714 PET (Corcia et al. 2012). A significant increase of microglial activation was found in the ALS sample in primary motor, supplementary motor and temporal cortex. Longitudinal follow-up is required in these series to determine the predictive power of TSPO PET for dementia in ALS cases.

Alshikho and colleagues have scanned 53 ALS and 11 primary lateral sclerosis (PLS) cases with [<sup>11</sup>C]PBR28 PET and compared their findings with those of 21 healthy controls (Alshikho et al. 2018). Inflammation was measured as cortical grey matter SUVRs with a cerebellar reference for non-specific signal. ALS and PLS cases both showed increased inflammation in pre- and post-central gyri along with cortical thinning. Cortical uptake of [<sup>11</sup>C]PBR28 correlated negatively with ALSFRS-R disability ratings. No significant increase in pre- and post-central cortical [<sup>11</sup>C]PBR28 was apparent over a 6-month follow-up although ALSFRS-R ratings continued to decline. The authors suggested [<sup>11</sup>C]PBR28 PET could provide a useful biomarker of the efficacy of neuroprotective strategies in ALS.

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## 30.5 Parkinson's Disease and TSPO Imaging

Parkinson's disease (PD) is the second most common neurodegenerative disorder after dementia, affecting around 1% of the over 60s. It manifests as asymmetrical limb bradykinesia, rigidity and tremor, and these symptoms are usually responsive to oral levodopa. The pathology is characterised by Lewy body inclusions in neurons and Lewy neurites which contain aggregated alpha-synuclein and neurofilaments.

While the dopamine neurons of the substantia nigra pars compacta are targeted by the Lewy body pathology, it is now thought that the pathology can start peripherally in the skin and gut and then track via the vagus nerve to affect the dorsal motor nucleus in the medulla and then ascend through the brainstem to the cortex in stages (Braak et al. 2004). In stage 2 the locus ceruleus, pedunculopontine nucleus and median raphe in the pons become involved, while the dopamine neurons of the substantia nigra pars compacta in the midbrain only become targeted in stage 3 along with the cholinergic nucleus basalis. The limbic cortex and cingulate are involved by stage 4 and the association and primary neocortex in stages 5 and 6.

At post-mortem microglial activation has been reported to accompany Lewy body pathology in affected subcortical and cortical regions (Imamura et al. 2003; McGeer et al. 1993). Aggregations of alpha-synuclein as oligomers and fibrils can stimulate microglial activation *in vitro*, but the relationship between inflammation and disease progression remains uncertain. In an initial PET study, it was reported that levels of increased midbrain (*R*)-[<sup>11</sup>C]PK11195 uptake in PD patients correlated

with reductions in putamen dopamine transporter (DAT) binding measured with [ $^{11}\text{C}$ ]CFT PET (Ouchi et al. 2005). The authors suggested that this provided evidence for involvement of microglia in the dopamine loss that characterises PD.

Later (*R*)-[ $^{11}\text{C}$ ]PK11195 PET series, however, have failed to detect consistent microglial activation in the substantia nigra of PD cases or to replicate an association between midbrain (*R*)-[ $^{11}\text{C}$ ]PK11195 signals and loss of dopaminergic function in PD. Gerhard and colleagues noted that PD patients showed significant striatal and frontotemporal (*R*)-[ $^{11}\text{C}$ ]PK11195 uptake though this did not correlate with disease severity (Gerhard et al. 2006a) (Fig. 30.1c). Iannaccone and co-workers have reported increased (*R*)-[ $^{11}\text{C}$ ]PK11195 binding in the putamen and the substantia nigra of early Parkinson's disease but did not detect inflammation in the cortex (Iannaccone et al. 2013). Measuring [ $^{11}\text{C}$ ]PBR28  $V_T$  with Logan graphical analysis using an arterial plasma input function, Varnas and co-workers (Varnas et al. 2019) were unable to detect any increase in microglial activation in their group of 16 PD patients (8 HABs and 8 MABs).

In contrast, using the second-generation TSPO marker [ $^{11}\text{C}$ ]DPA713 PET, increased signal was detected in the temporal, parietal and occipital cortex of PD cases which increased over 1 year (Terada et al. 2016). This finding supports the report from Gerhard and suggests that microglia activation can involve the substantia nigra, striatum and association cortex and be present in the early stages of the disease even in the absence of dementia.

It is conceivable that the dopaminergic drugs used to treat PD may have an influence on levels of microglial activation. Although none of these dopaminergic agents directly act to suppress microglial activation, it has been suggested that both dopamine agonists and monoamine oxidase B inhibitors have neuroprotective properties (Schapira and Olanow 2004). This claim is, however, controversial, and currently there is no hard evidence that therapy for PD influences levels of the inflammatory response.

The widespread microglial activation seen in PD at post-mortem and the raised (*R*)-[ $^{11}\text{C}$ ]PK11195 uptake seen in the brainstem, basal ganglia and cortical regions of non-demented cases with PET are all regions which are targeted by Lewy body pathology in later Braak stages. More recently we have noted that levels of striatal (*R*)-[ $^{11}\text{C}$ ]PK11195 uptake in PD correlated with disability rated with the Unified Parkinson's Disease Rating Scale (UPDRS), while impaired verbal fluency was associated with raised frontal and insular (*R*)-[ $^{11}\text{C}$ ]PK11195 activity (Simpson et al. 2012). This supports a role for microglial activation in driving disease activity and is in line with the dying back theory of PD pathology which suggests that nigrostriatal terminals may become dysfunctional ahead of nigral cell body loss.

Those few longitudinal studies that have reported (*R*)-[ $^{11}\text{C}$ ]PK11195 uptake in PD suggest levels remain static over a 1–2-year follow-up period despite on-going clinical deterioration and loss of putamen [ $^{18}\text{F}$ ]DOPA storage of PD cases (Gerhard et al. 2004; Terada et al. 2016). It has been suggested that cortical microglial activation is an early phenomenon in PD and may act to promote the superadded dementia that usually follows although levels of inflammation remain fixed. In favour of this viewpoint, it was noted that PD cases with late dementia had similar cortical levels of microglial activation to similarly disabled non-demented cases.

### 30.6 Atypical Parkinsonian Syndromes and TSPO Imaging

Neurodegenerative disorders that cause atypical parkinsonian disorders include multiple system atrophy (MSA) and progressive supranuclear palsy (PSP). MSA is associated with asymmetric parkinsonism along with early autonomic dysfunction, postural instability and ataxia. The pathology is characterised by argyrophilic glial cytoplasmic inclusions (GCIs) that contain alpha-synuclein and targets the substantia nigra, putamen, ponto-cerebellar connections and the lateral columns of the spinal cord. Microglial activation is known to be associated with MSA pathology, and (*R*)-[<sup>11</sup>C]PK11195 PET has revealed raised TSPO binding in the putamen, pallidum, pons, substantia nigra and the dorsolateral prefrontal cortex at higher levels than that associated with PD (Gerhard et al. 2003). In a follow-up series of 14 MSA-P cases, Gerhard and colleagues again reported significant increases in microglial activation in the putamen ( $p = 0.001$ ), pallidum ( $p = 0.002$ ), precentral gyrus ( $p = 0.004$ ), orbitofrontal cortex ( $p = 0.006$ ), presubgenual anterior cingulate cortex ( $p = 0.006$ ) and the superior parietal gyrus ( $p = 0.007$ ) but now also the caudate nucleus ( $p = 0.002$ ) (Kubler et al. 2019). The increases in (*R*)-[<sup>11</sup>C]PK11195 binding potentials correlated with decreases in glucose utilisation in the putamen ( $r = -0.78$ ,  $p = 0.003$ ) and pallidum ( $r = -0.77$ ,  $p = 0.003$ ). No correlations between regional (*R*)-[<sup>11</sup>C]PK11195 BPs and clinical parameters were found.

A proteolipid protein-alpha-synuclein overexpression (PLP- $\alpha$ SYN) transgenic mouse model of MSA showed a correlation between levels of nigral microglial activation and dopamine neurone loss (Stefanova et al. 2007). Minocycline suppressed the inflammation and preserved dopamine neurones in the mouse. A human trial was subsequently set up to determine whether minocycline had neuroprotective efficacy in MSA. The drug failed to alter disease progression at clinically licensed doses though MSA cases receiving active medication showed a 30% reduction in brain (*R*)-[<sup>11</sup>C]PK11195 uptake relative to placebo-treated cases (Dodel et al. 2010).

Three PSP patients with Richardson syndrome who were studied with (*R*)-[<sup>11</sup>C]PK11195 PET showed significantly increased signal in their caudate nucleus, putamen, pallidum, midbrain, substantia nigra, the frontal lobe and the cerebellum (Gerhard et al. 2006b). One of these patients was rescanned after 10 months, and the level of microglial activation had remained stable.

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### 30.7 Detecting Preclinical Huntington's Disease Activity with TSPO PET

Huntington's disease is an autosomal dominant inherited progressive neurodegenerative disorder associated with motor, cognitive and psychiatric symptoms. It arises from an abnormal CAG triplet repeat expansion of the HTT gene on chromosome 4 which leads to an elongated polyglutamine chain at the terminus of the huntingtin protein. This results in cytoplasmic and intranuclear aggregations of huntingtin and progressive loss of medium spiny striatal GABA-ergic and cortical interneurons

(Sapp et al. 1997). The striatal output neurons express either dopamine D1 or D2 receptors depending on whether they are part of the direct or indirect pathway to the internal pallidum. Striatal dysfunction can be detected prior to symptom onset as a loss of availability of D1 and D2 sites for ligand binding (Andrews et al. 1999).

The loss of striatal and cortical neurons in HD is associated with microglial activation (Sapp et al. 2001) which can be detected in vivo with (*R*)-[<sup>11</sup>C]PK11195 PET. Levels of striatal microglial activation correlate with loss of dopamine D2 receptor binding measured with [<sup>11</sup>C]raclopride PET and with locomotor disability rated with the Unified Huntington's Disease Rating Scale (UHDRS) (Pavese et al. 2006). This suggests that microglial activation plays a role in driving the disease process. Supporting this viewpoint, (*R*)-[<sup>11</sup>C]PK11195 PET studies have detected raised microglial activation in a majority of asymptomatic adult HD gene carriers in their fourth decade (Tai et al. 2007). Those asymptomatic carriers with raised TSPO expression also showed reduced dopamine D2 availability with [<sup>11</sup>C]raclopride PET. Cortical microglial activation was also evident in pre-manifesting HD carriers confirming that this is not purely a basal ganglia disorder (Politis et al. 2011). Levels of striatal (*R*)-[<sup>11</sup>C]PK11195 uptake have been shown to correlate with the predicted time of clinical disease onset (Politis et al. 2011; Tai et al. 2007).

Lois and colleagues have measured translocator protein (TSPO) expression in HD using [<sup>11</sup>C]PBR28 PET. Regional signals were quantitated using SUVRs normalised to whole brain uptake with emission data acquired 60–90 min after intravenous radiotracer administration (Lois et al. 2018). Significant TSPO overexpression was detected in the putamen and pallidum of all the individual HD patients. Additionally, some HD patients showed elevated [<sup>11</sup>C]PBR28 uptake in thalamic nuclei, the brainstem and red nuclei. Increased cortical inflammation was not reported in this series. The authors suggested that [<sup>11</sup>C]PBR28 PET might prove a useful biomarker in clinical trials evaluating therapies targeting neuroinflammation.

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## 30.8 Measuring Inflammation in Multiple Sclerosis

Multiple sclerosis (MS) is a disease characterised clinically by relapsing and remitting neurological episodes followed by a progressive phase of disability. Pathologically central inflammatory demyelination is seen as white matter plaques associated with axonal degeneration. MS targets young adults and is their most common cause of non-traumatic disability in the Western world (Compston and Coles 2008). Active plaques of demyelination contain blood-borne B and T lymphocytes and macrophages that have invaded via the disrupted blood-brain barrier. However, there is also an intrinsic immune reaction to the disease process manifested as involvement of activated microglia (Benveniste 2007). Post-mortem investigations have detected activated microglia not just in white matter plaques but also in cortex and apparently normal-appearing brain areas (De Groot et al. 2001; Peterson et al. 2001). In progressive MS activated microglia are seen surrounding the degenerating myelinated axons (Magliozzi et al. 2010).

(*R*)-[<sup>11</sup>C]PK11195 PET has been used to demonstrate the extent of microglial activation in MS patients (Banati et al. 2000; Debruyne et al. 2003; Politis et al. 2012) (Fig. 30.1d). PET findings have been validated with [<sup>3</sup>H]PK11195 autoradiographic studies on human brain slices which showed that this tracer binds to activated microglia rather than activated astrocytes or lymphocytes in plaques of demyelination (Banati et al. 2000). PET reveals that not only can raised (*R*)-[<sup>11</sup>C]PK11195 uptake be seen in active plaques which show raised T2 signal and gadolinium enhancement on MRI but also in normal-appearing white and grey matter (Banati et al. 2000; Debruyne et al. 2003; Versijpt et al. 2005; Politis et al. 2012). This finding supports the hypothesis that in early MS microglial activation may initiate the inflammatory process prior to invasion of the damaged blood-brain barrier by lymphocytes. The activated microglial load at baseline may well be the critical factor when predicting outcome in MS patients (Confavreux et al. 2000). (*R*)-[<sup>11</sup>C]PK11195 uptake has been reported to be higher in secondary progressive than relapsing-remitting MS. Additionally, a significant correlation was reported between levels of (*R*)-[<sup>11</sup>C]PK11195 binding in cortical grey matter and locomotor disability in patients with secondary progressive MS (Politis et al. 2012).

TSPO expression in MS has also been examined with [<sup>11</sup>C]PBR28, [<sup>18</sup>F]DPA714 and GE180 PET. Oh and colleagues reported that global brain [<sup>11</sup>C]PBR28 uptake was increased in their MS cohort and focal increases in binding, corresponding to areas of active inflammation and blood-brain barrier breakdown on gadolinium contrast-enhanced magnetic resonance imaging (MRI), were evident (Oh et al. 2010). Increases in [<sup>11</sup>C]PBR28 binding preceded the appearance of contrast enhancement on magnetic resonance imaging in some lesions, suggesting that glial activation is an early phenomenon in MS lesion formation. In this series global levels of brain [<sup>11</sup>C]PBR28 binding correlated with disease duration but not with locomotor disability. Herranz and colleagues have imaged nine relapsing-remitting multiple sclerosis (RRMS) and ten secondary progressive multiple sclerosis (SPMS) patients with [<sup>11</sup>C]PBR28 PET and 7-Tesla (7 T) MRI (Herranz et al. 2019). Normal white matter signal was used as a reference tissue to compute SUVr<sub>s</sub>. Raised [<sup>11</sup>C]PBR28 uptake was found in cortical lesions in both RRMS and SPMS but in the latter inflammation was also evident in normal-appearing cortex. It was noted that patients with higher levels of cortical inflammation had a worse clinical outcome. The same group also used [<sup>11</sup>C]PBR28 PET to demonstrate significant cerebellar inflammation in RRMS and SPMS patients (Barletta et al. 2019). Levels of cerebellar [<sup>11</sup>C]PBR28 uptake correlated with both physical disability and cognitive deficit.

The utility of the TSPO ligand, [<sup>18</sup>F]DPA-714, has been evaluated in primary and secondary progressive multiple sclerosis (PMS) (Hagens et al. 2018). Eight PMS cases and seven healthy controls had [<sup>18</sup>F]DPA-714 PET, and non-binders were excluded by prior genotyping for the presence of the rs6971 polymorphism. Regional time-activity curves (TACs) were kinetically modelled using a brain two-compartment model and a metabolite corrected plasma input function. Regional  $V_T$ s were higher for high affinity (HABs) than mixed affinity binders (MABs) in both patient and normal groups, but [<sup>18</sup>F]DPA-714 PET was not able to discriminate

patients from healthy controls. The patients, however, only showed focally increased tracer uptake where T2 white matter lesions were seen on MRI. A drawback with [ $^{18}\text{F}$ ]DPA-714 as a TSPO marker was felt to be its large non-displaceable background signal, rather similar to the situation seen with (*R*)-[ $^{11}\text{C}$ ]PK11195.

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### 30.9 Traumatic Brain Injury

Recovery from concussive traumatic brain injury (TBI) is highly variable. (*R*)-[ $^{11}\text{C}$ ]PK11195 PET can detect the presence of microglial activation in many of these cases years after the original injury. In a recent series, ten patients had (*R*)-[ $^{11}\text{C}$ ]PK11195 PET at least 11 months after moderate to severe TBI, and binding was calculated in and around the site of focal brain damage, and in selected distant and subcortical brain regions (Ramlackhansingh et al. 2011). Microglial activation was significantly raised in the thalami, putamen, occipital cortices and the posterior limb of the internal capsules after TBI though there was no increase in binding at the original site of the focal brain injury. Levels of thalamic (*R*)-[ $^{11}\text{C}$ ]PK11195 uptake correlated with impairment on executive tasks but not with either time since the injury or the amount of structural brain injury. The authors concluded that increased microglial activation can persist up to 17 years after TBI in distant brain areas from the original focal lesion, particularly in subcortical regions. They suggested that a chronic inflammatory response to TBI develops over time and that anti-microglial strategies may still be beneficial months to years after the original insult. In a follow-up article, these workers showed that levels of thalamic inflammation correlated with levels of thalamo-cortical fibre damage measured with diffusion tensor imaging rather than cortical or thalamic lesions (Scott et al. 2015). Whether the persistent inflammation was beneficial to recovery or toxic remained uncertain.

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### 30.10 Stroke and Microglial Activation

Stroke is associated with microglial activation both around the infarct, which develops within hours of the event, but also later in disconnected brain regions including the thalamus and brainstem. In an early series, six patients were examined with (*R*)-[ $^{11}\text{C}$ ]PK11195 PET between 3 and 150 days after their infarct, and increased microglial activation was present in all the patients examined (Gerhard et al. 2005). In the first 6 days following the stroke, the focus of inflammation was generally smaller and found adjacent to the MRI lesion with little spatial overlap. The size of the local area of microglial activation then increased over the next months overlapping with the MRI lesion as the blood-brain barrier became disrupted and invasion by macrophages binding (*R*)-[ $^{11}\text{C}$ ]PK11195 alongside local microglial activation occurred. By 6 months after the initial stroke, local microglial activation was subsiding, whereas distant areas of (*R*)-[ $^{11}\text{C}$ ]PK11195 uptake could be seen beyond the

primary infarct site involving disconnected areas of the ipsilateral hemisphere including the thalamus and brainstem. The lesioned cortical area with high initial (*R*)-[<sup>11</sup>C]PK11195 binding became atrophic during that 6-month interval as local inflammation regressed.

Thiel and colleagues used (*R*)-[<sup>11</sup>C]PK11195 PET to prospectively image activated microglia in vivo 2 weeks and 6 months after acute subcortical stroke in humans to investigate their temporal dynamics and relate local and remote inflammation to pyramidal tract (PT) damage detected using diffusion tensor imaging (DTI) (Thiel et al. 2010). As reported previously, microglial activation increased and then regressed around the local lesion. In contrast, brainstem inflammation was only seen in cases with pyramidal tract damage on DTI and occurred later. The brainstem (*R*)-[<sup>11</sup>C]PK11195 signal correlated positively with clinical outcome.

Recently Morris and co-workers used diffusion and perfusion MRI to define the ischemic penumbra around acute stroke and examined neuronal integrity with the GABA marker [<sup>11</sup>C]flumazenil PET and microglial activation with (*R*)-[<sup>11</sup>C]PK11195 PET (Morris et al. 2018). They found evidence of viable neurons in the penumbra in the absence of microglial activation. These workers concluded that viable penumbra may not require the presence of activated microglia during the stroke recovery phase.

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### 30.11 Psychosis

Lewy body dementia and Parkinson's disease are associated with secondary psychosis related to their cortical pathology and possibly the presence of the cortical and limbic inflammation that can be detected in these disorders. Schizophrenia is a primary psychosis but has also been reported to be associated with brain inflammation at post-mortem. In an early series, seven schizophrenic patients in the recovery phase from a psychotic episode were studied with (*R*)-[<sup>11</sup>C]PK11195 PET (Doorduyn et al. 2009). Conventional T1- and T2-weighted MRI showed no significant abnormalities. PET revealed a 51% increase in mean hippocampal and a 30% increase in global cortical (*R*)-[<sup>11</sup>C]PK11195 binding in the schizophrenic subjects compared with age-matched cortical controls. Three of the subjects individually showed significantly raised levels of cortical inflammation. Low-level inflammation in the cortex of schizophrenic cases in remission has also been reported (van Berckel et al. 2008). Doorduyn and co-workers suggested that active episodes of psychosis may be related to increases in background inflammatory activity.

In contrast, using the second-generation TSPO marker [<sup>11</sup>C]PBR28 with PET, Collste and colleagues detected reduced levels of binding ( $V_T$ ) in 16 untreated schizophrenics recovering from their first psychotic episode (Collste et al. 2017). Brain tracer uptake was kinetically modelled using a two-compartmental approach with an arterial plasma input function. These workers suggested that the function of immune cells may be depressed in schizophrenia.

## 30.12 Conclusions

Microglial activation is a non-specific reaction to all forms of brain injury and can be detected with PET in inflammatory, vascular, traumatic, degenerative and, possibly, psychotic brain disorders. The role of activated microglia after stroke and trauma appears to be primarily restorative, the cells adopting a phagocytic phenotype removing debris and remodelling connections to disconnected regions. However, the ischemic penumbra surrounding infarcted tissue may not require activated microglia for its revival. In neurodegenerative diseases, the role of activated microglia is less certain. Initially these cells could act to ingest misfolded proteins and release growth factors, but in later disease this action could fail and local release of toxic cytokines predominate leading to neuronal death and disease progression. In downstream disconnected areas, microglia could be beneficial leading to remodelling of connections as an adaptive response.

Currently PET imaging of activated microglia relies on the use of TSPO radioligands. While these provide a valuable *in vivo* tool for detecting disease activity and tracking the progression of neuroinflammation, binding of second-generation ligands is influenced by the TSPO polymorphisms expressed by subjects—10% of Caucasians are non-affinity binder, 30% mixed affinity binders and 60% high affinity binders. As a consequence, despite its higher non-specific signal and rapid wash-out, (*R*)-[<sup>11</sup>C]PK11195 PET still provides a reasonable approach for measuring brain inflammation as it is little affected by the TSPO polymorphisms expressed. While TSPO imaging reflects disease activity, it does not have diagnostic potential as raised microglial activation is not specific to any one neurological disorder. However, the early detection of microglia with PET provides a potential biomarker for establishing the presence of disease activity in at risk subjects and testing the efficacy of neuroprotective strategies designed to suppress the inflammatory response to local injury or neurodegenerative processes. In the future, hopefully, there will be development of both better TSPO tracers uninfluenced by the genotype along with other markers of microglial activation, such as cannabinoid CB2 expression, allowing us to improve our understanding of the role of activated microglia in CNS disease.

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# Toward Imaging Tropomyosin Receptor Kinase (Trk) with Positron Emission Tomography

# 31

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## Abstract

The three tropomyosin receptor kinases (TrkA, B, and C) fulfill important roles within the central nervous system facilitating neuronal growth, neuronal survival, and neuronal differentiation during cell development at all ages of neuronal

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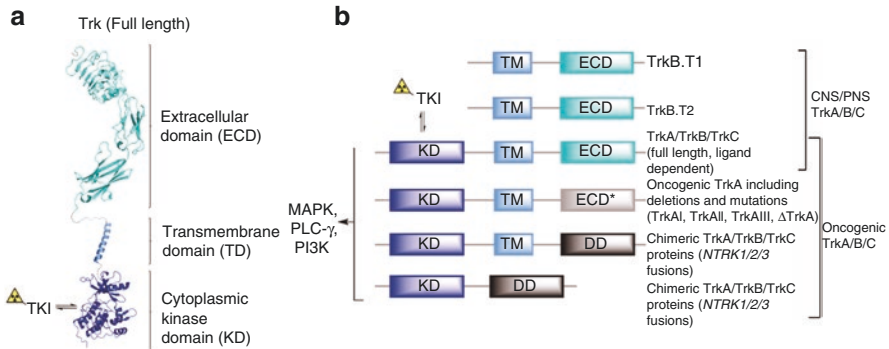
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development. The downregulation of TrkA, B, and C has been identified as an important hallmark of a plethora of neurological diseases such as Alzheimer's disease (AD) and Parkinson's disease (PD). In oncology, Trks have been recognized as tumorigenic drivers. Full-length Trk as well as fusion proteins composed of the kinase portion of the full-length receptor and various other cancer-related proteins has been reported, and the targeting of these receptors for a contingent Trk-based therapy has recently caught momentum in the wake of precision medicine evolution. Both in neurology and oncology, the spatiotemporal changes in Trk expression can only be reviewed via destructive and invasive methods such as taking a tumor biopsy. The quantification of Trk density in neurodegeneration (e.g., AD and PD) as well as cancer treatment, where therapeutic Trk target engagement of antineoplastic Trk inhibitors with Trk fusion proteins is the site of therapeutic action, has triggered the need for a noninvasive methodology to quantify Trk *in vivo* in these Trk-altering conditions. Positron emission tomography (PET), a noninvasive imaging methodology, relying on the highly sensitive detection of radiation, has the potential to take advantage of radioactively labeled probes binding to the kinase domain of Trk to not only reveal the location of Trk but allow for the quantification of receptor density. This article covers the most recent developments in Trk tracer evolution for PET imaging and their first human *in vivo* application.

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### 31.1 The Role of Trk in Neurology and Oncology

The tropomyosin receptor kinases (TrkA, B, and C) constitute transmembrane glycoproteins which are encoded by their individual genes (NTRK1, 2, and 3). These kinases are composed of an extracellular domain (ECD) which is the target for the endogenous neurotrophins such as the nerve growth factor (NGF), binding to TrkA; the brain-derived neurotrophic factor (BDNF) and neurotrophin-4, binding to TrkB; and neurotrophin-3 engaging TrkC (Lessmann et al. 2003; Chao 1992, 2003; Kaplan and Miller 2000; Binder and Scharfman 2004). In addition to the ECD, all Trks possess an intracellular binding domain which encodes a tyrosine kinase binding site which primarily interacts with ATP as a binding substrate (Fig. 31.1a) (Reichardt 2006; Huang and Reichardt 2003). Ligand-induced dimerization of Trk activates the tyrosine kinase and mediates receptor transphosphorylation. In this activated state, the Trks immediately phosphorylate their endogenous substrates: Shc, phospholipase C $\gamma$ 1 (PLC- $\gamma$  1), and the fibroblast growth factor receptor substrate 2 (FRS2/SNT1) (Kaplan and Miller 2000; Huang and Reichardt 2003; Yan et al. 2019; Poo 2001). In addition to the full-length Trks, truncated variants can be formed by alternative splicing which lack the intracellular kinase domain (Fig. 31.1b) (Fenner et al. 2014; Fenner 2012; Luberg et al. 2010). In the case of TrkB, the 90 kDa truncated isoform TrkB.t1, comprised of only the extracellular domain, is the most prominent differentially spliced variant. When ATP binds to the intracellular kinase domain, downstream signaling pathways are activated which control neuronal survival and



**Fig. 31.1** Tertiary and domain structural representations of native and aberrantly expressed Trk proteins. **(a)** Structure (Massa et al. 2010) overview of the representative native TrkA receptor (TKI: tyrosine kinase inhibitor). **(b)** Domain representation of diverse oncogenic Trk proteins, including Trk splice variants and Trk fusion proteins. Dimerization and transautophosphorylation of Trk kinase domains leads to the activation of the downstream signaling pathways, including MAPK1, PI3-K/Akt, and PLC- $\gamma$ 1 (DD: dimerization domain). Reproduced with permission from R. Schirmmacher et al.; Pharmaceuticals (Basel), 2019, 12(1), 7–21

mediate embryonic, postnatal, and mature differentiation of the developing central and peripheral nervous system.

Aberrant or impaired expression/signaling of Trks within the central nervous system is connected to a variety of neuropathological abnormalities encompassing Alzheimer's disease (AD), Parkinson's disease (PD), and other neurodegenerative diseases (Zhang et al. 2012; Song et al. 2015; Murer et al. 2001; Reinhart et al. 2015; Oyesiku et al. 1999; Deng et al. 2007; Gupta et al. 2013). Convincing evidence points to a direct involvement of impaired TrkB and C signaling in AD. Confirmative studies gained through ex vivo analysis of brain tissue from AD patients demonstrated that TrkB levels are significantly reduced in the hippocampus and the frontal and temporal cortex of AD individuals (Ferrer et al. 1999). In addition, the progressive loss of TrkA, B, and C in basal forebrain cholinergic nuclei has been linked to the progression of AD (Allen et al. 1999; Savaskan et al. 2000). There is further evidence from transgenic mouse models of AD that treatment with Trk agonists increases the formation of dendritic spines in hippocampal and cortical regions (Castello et al. 2014; Li and Liu 2010; Nagahara and Tuszyński 2011; Devi and Ohno 2015). During agonist treatment, inhibition of neuronal apoptosis in ongoing neurodegeneration as a result of plaque deposition and neurofibrillary tangle formation results in improved spatial memory performance (Géral et al. 2013). However, there is currently no noninvasive way to follow the spatiotemporal changes of Trk densities over time. PET would provide a unique opportunity to track these dynamic changes without perturbing the underlying biochemistry and would connect disease progression or recession to a quantifiable visual signal provided by the PET camera. Besides an obvious interest in Trk radioligands for neurological imaging, the continuously growing interest in anti-Trk therapy in fusion cancers provides further incentive to develop Trk-radioligands for cancer staging and subsequent

therapeutic investigations after agonist therapy (Cocco et al. 2018). Such radioligands targeting Trks would have an important impact on patient care for individuals enrolled in NTRK fusion-positive cancer trial where pan-Trk inhibitors are administered as part of the treatment protocol (Lange and Lo 2018). The current practice is to acquire tumor biopsies and assay them for Trk occurrence, followed by next-generation sequencing or fluorescence in situ hybridization. This procedure is highly invasive and puts significant stress on the cancer patient. Radioactively labeled Trks have the potential to noninvasively quantify Trk present in tumor tissue and would alleviate an investigative burden from the patient. In addition, Trk PET imaging would significantly expedite Trk status stratification and patient sorting (Hsiao et al. 2019; Khotskaya et al. 2017).

NTRK fusions are generally a low-frequency occurrence in a variety of commonly known cancers, although they occur at a very high frequency in some rare neoplasms (Märkl et al. 2019). A prerequisite for all aforementioned applications is the development of selective pan-Trk inhibitors that can either be used in therapeutic scenarios or be translated into a functioning Trk radiotracer. Fortunately, recent research efforts from academia and industry have provided a great plethora of new Trk lead structures covering selective pan-Trk as well as TrkA subtype selective inhibitors. These inhibitors lay the foundation for efficient radiotracer development where medicinal chemistry provides the scaffold for further translational work into clinical PET tracer advancement. Trk radiotracers would be especially useful in the assessment of Trk status in the early clinical stages of cancer where the treatment regimen is set for the individual patient. Only patients who possess NTRK fusions are responsive to a Trk-inhibitor-driven therapy (Lange and Lo 2018). Patients undergoing Trk inhibitory therapy could be monitored for treatment response including possible downregulation of Trk receptors hampering a continuous therapy. Omitting sequential tumor biopsies as part of the treatment monitoring would benefit the patient by removing invasive procedures from the therapy plan. Radiolabeled Trk inhibitors for in vivo imaging would thus have a profound impact on patient care and disease control.

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## 31.2 The History of Trk Radioligand Development

### 31.2.1 Positron Emission Tomography

The PET detection principle is based on the simultaneous detection of two gamma rays of 511 keV stemming from the annihilation of a positron and an electron (masses are converted into energy) using a PET camera. The positron originates from the radioactive decay of a positron-emitting radionuclide, which is integrated into the structure of the radioligand, and the electron is provided by the surrounding body tissue. The PET camera consists of multilayered ring-shaped arranged detectors surrounding the patient's body. Following the rule of conservation of momentum, the two gamma rays originating from the annihilation are leaving their point of origin in the opposite direction. Traveling with the speed of light, opposite detectors



can detect these two gamma rays in coincidence, drawing a virtual line of response between the two detectors activated. The origin of the positron annihilation consequently must be somewhere on that line of response, narrowed down by the time resolution of the used detector material. Accumulated information from all communicating detector pairs and applying mathematical reconstruction algorithms (e.g., filtered back projection) allow for the accurate reconstruction of the radioactivity distribution within the region of interest. PET's resolution depends primarily on the positron energy but also on the size of the detectors and the photomultipliers intensifying the relatively weak signal from the gamma ray interacting with the detector material. Preclinically, submillimeter resolution is achieved in animal PET imaging, whereas in a human clinical setting, millimeter resolution is routinely accomplished.

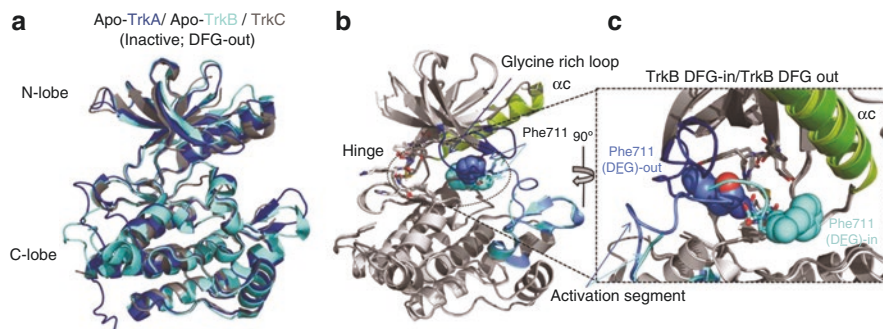
In order to gain the maximum information about a disease state from a radiotracer's binding characteristics, the compound used must comply with a few general criteria. Foremost, and this is an independent biological parameter, the molecular target concentration (e.g., TrkB) must be sufficiently high to generate a quantifiable radioactivity signal upon radiotracer binding. If the concentration of the receptor ( $B_{\max}$ ) is low, only tracers with a very high binding affinity will yield a detectable signal. The radiotracer concentration in the tissue of interest (e.g., the brain) must also be sufficiently high to guarantee binding, and radiotracer equilibrium conditions have to be established where the tracers  $k_{\text{on}}$  and  $k_{\text{off}}$  are the same. Tracer stability is also of utmost importance to preclude bio-active metabolite binding interference, complicating the kinetic analysis of the tracer binding pattern. Another important point is a tracer's on-target specificity. If the tracer binds to more than one target, reconstruction and image interpretation are very difficult to achieve since the only signal that is recorded is the radioactivity concentration in the tissue of interest. The PET camera does not distinguish between radioactivity signals coming from different target interactions. After careful consideration of all the above-mentioned points, the consequent question in the development of potential Trk radiotracers is which domain of the Trk receptor is suitable for targeting in order to answer a clinical question related to a specific disease?

### **31.2.2 The Tropomyosin Receptor and Its Targets: Extra- Vs. Intracellular Target Engagement**

Trk possesses two very distinct binding sites: the extracellular binding domain (ECD), which is engaged by endogenous ligands, and an intracellular kinase binding site which binds ATP and significantly impacts tracer development. Furthermore, the TrkA/B/C isoforms and splice variants/fusion proteins found in certain cancers may complicate the *in vivo* PET imaging depending on the choice of target, extra- vs. intracellular. To illustrate this further, besides the full-length TrkA/B/C containing a fully functional extra- and intracellular domain, a variety of truncated isoforms missing the intracellular kinase domain have been identified for TrkB (TrkB.T1 and TrkB.T2). Their biological function is unclear and still under debate, but it is

accepted that these splice variants are not associated with neurodegeneration or function as oncogenic drivers. However, a most recent study revealed that astrocyte truncated-TrkB exerts a BDNF anti-apoptotic effect that leads to neuroprotection (Saba et al. 2018). Especially within the brain, the expression of the truncated isoforms on the protein level is much higher than that of the full-length, biologically active, TrkB. A radiotracer targeting the ECD of TrkB (e.g., radiolabeled BDNF) would not be able to distinguish between the full-length Trk protein and the truncated isoforms, and consequentially would overestimate the number of biologically active TrkB (Tejeda and Díaz-Guerra 2017). The combined PET signal resulting from the catalytically incompetent truncated isoforms, lacking the kinase domain, and the full-length TrkB would therefore erroneously overestimate the number of catalytically active Trk responsible for an array of downstream signaling on a molecular level. Such a tracer would be useless to investigate TrkB's role in neurodegeneration. In a cancer imaging situation where NTRK fusion proteins are present, an ECD radiotracer would be unable to engage this protein target because it lacks the cognate ectodomain. Besides these obvious hurdles, there are currently no compounds available which warrant a translation into a radiotracer as a result of their poor and challenging *in vivo* properties.

Our research group and collaborators recently demonstrated that radiolabeled fluorine-containing 7,8-dihydroxyflavon (7,8-DHF), which was reported to bind with high affinity to the ECD (Jang et al. 2010), was unsuitable as PET imaging agent for TrkB (Bernard-Gauthier et al. 2013). None of the reported 7,8-DHFs' basic biological properties could be reproduced (unpublished). In one of our pre-clinical PET imaging studies in healthy rats using 2-(4-[<sup>18</sup>F]fluorophenyl)-7,8-dihydroxy-4*H*-chromen-4-one, we observed a low overall radioactivity uptake in the brain (SUV of 0.64) that was followed by a fast elimination, proving the inability of the DHF compounds to serve as a TrkB radioprobe. The plasma half-life was suboptimal at 4 min and suggests an extensive phase II metabolic breakdown accounting for the poor radiotracer properties. In support of our findings, Boltaev et al. most recently incontrovertibly demonstrated the absence of an interaction of these and other compounds with the ECD of TrkB, or any TrkB-related effects attributed to these compounds (Boltaev et al. 2017). Going from putative ECD binding drugs to inhibitors targeting the ATP binding site of the cytoplasmic kinase domains offers several advantages and at the same time adds some challenges to a PET imaging approach. Unfortunately, all Trks display a high homology within their kinome as well as within their Trk family sequence identity, adding an intrinsic challenge to any imaging regime. To achieve selectivity over at least other unrelated kinases, our group always tested the most promising compounds for PET tracer translation in a comprehensive kinase screening to ensure near exclusive binding to TrkA/B/C. All radiolabeled compounds reported so far are pan-Trk radioligands which insufficiently distinguish between the three subtypes. Fortunately, the local deposition of TrkB/C in comparison to TrkA in the human brain is very prominent, in theory yielding a target density-driven PET image of predominantly TrkB/C (Altar et al. 1991a, b; Merlio et al. 1992; Anderson et al. 1995). This would translate into an almost exclusive visualization and quantification of TrkB/C in the human



**Fig. 31.2** Trk kinase domain. (a) Overlap of TrkA, TrkB, and TrkC kinase domains in the inactive DFG-out conformation (PDB: 4F0I, 4ASZ, 3V5Q). (b, c) Views of the conformational differences between “Asp-Phe-Gly” DFG-in and DFG-out TrkB. The Phe residues of the DFG triad are shown in spheres (PDB: 4AT3, 4AT5). Reproduced with permission from R. Schirmacher et al.; Pharmaceuticals (Basel), 2019, 12(1), 7–21

brain. The imaging situation is further complicated by the fact that Trk kinases can exist in two distinct conformations, DFG-in, wherein the active kinase conformation is targeted by type-1 inhibitors, and DFG-out, which is the inactive conformation being engaged by type-2 inhibitors (Fig. 31.2a). Conformational rearrangements within the DFG motive grant access to a deep binding pocket that is favorably targeted by lipophilic moieties inherent of type-2 inhibitors (Fig. 31.2b, c). This extended binding mode for type-2 inhibitors enhances the residence time of the inhibitor at the kinase binding site via cooperative binding in comparison to type-1 inhibitors. Ultimately, only the final in vivo imaging study can reveal what type of radiolabeled inhibitor will be the most suitable for human Trk PET imaging. To investigate the relation of each binding mode to PET image quality, as well as the tracers’ residence time and overall kinetic profile, we developed both type-1 and type-2 radiotracers. The following sections will give an overview of our efforts to preclinically, as well as clinically, introduce these novel Trk radiotracers to the imaging community, culminating in the two first human in vivo studies of a type-1 Trk radioligand in a healthy male subject.

### 31.2.2.1 Radioligands for Brain Imaging: General Considerations

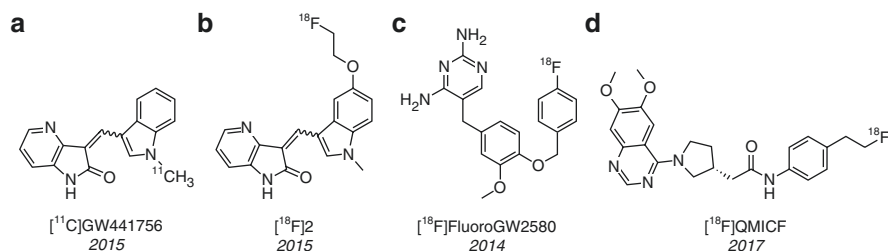
When developing radiotracers for brain imaging, the chance of successful neurological PET imaging is significantly elevated by adhering to a few general principles guiding the structural composition of a potential radiotracer. The following parameters are cardinal points that should help streamlining the tracer design process (Van de Bittner et al. 2014). First and foremost, a radiotracer intended for neurologic PET imaging must cross the highly restricted blood-brain barrier (BBB) to bind to its target. As for Trk target binding, the tracer also has to overcome the neuron cell membrane to bind to the cytoplasmic binding site of the Trk kinase domain. To get a high contrast image, it is advantageous that the radioligand binds with high specificity to the target and only displays minor, or ideally no, nonspecific binding (a

feature rarely seen). A common denominator determining nonspecific binding is the tracer's lipophilicity, or  $\text{LogD}_{7.4}$ , granting passive diffusion through the cell membrane and if too high simultaneously increasing nonspecific binding and influencing the tracer's receptor binding potential ( $\text{BP} = B_{\text{max}}/K_d > 10$ ). Further parameters, like susceptibility to active efflux from the brain compartment, the topological surface area (TSPA), and hydrogen bond donor characteristics, directly influence BBB penetration and binding specificity. Taken together, these considerations and parameters directly determine the quality of a potential PET tracer for neuroimaging.

### 31.2.2.2 Radioligands Based on the 4-Aza-2-Oxindole Scaffold

Following the above-outlined considerations, the first pan-Trk radiotracer developed was structurally based on reported 3-arylideneindolin-2-on Trk inhibitors belonging to the 4-aza-2-oxindole family. The well-characterized type-I inhibitor GW441756 was found to theoretically fulfill most of the parameters accounting for a viable PET neuro-tracer (Zhang et al. 2013; Pike 2009; Rankovic 2015) and was consequently translated into its  $^{11}\text{C}$ -isotopologue and applied in autoradiography ex vivo binding studies as well as preclinical animal PET imaging. GW441756 was found to possess a half maximal inhibitory concentration ( $\text{IC}_{50}$ ) of 29.6 nM for TrkA, and 6.7 nM and 4.6 nM for TrkB and TrkC, respectively (Wood et al. 2004). The  $^{11}\text{C}$ -radiolabeling of GW441756 (Fig. 31.3a) was easily accomplished in high radiochemical yields (RCYs) by using standard  $^{11}\text{C}$ -labeling methodology, although the labeled compound turned out to be a mixture of two distinct isomers, the E and Z isomer, of which only the Z isomer actively binds to the kinase domain (Bernard-Gauthier et al. 2015a). The E/Z ratio after synthesis was 1:1, accounting to 50% of the radioactivity signal stemming from the wrong isomer. In addition, the clinical translation of an impure tracer is very difficult from an image reconstruction and modeling point of view, not to mention the ethical challenge of injecting a useless impurity into a human subject. However, autoradiographic ex vivo imaging using the mixed isomers demonstrated competitive and replaceable binding, matching the almost ubiquitous distribution of TrkB/C throughout the brain.

Following these successful ex vivo investigations, E/Z- $^{11}\text{C}$ -GW441756 was injected into healthy Sprague-Dawley rats to investigate general tracer parameters



**Fig. 31.3** Chemical structures of early Trk-targeted PET radioligands. (a, b)  $^{11}\text{C}$ - and  $^{18}\text{F}$ -labeled analogues of type-I inhibitor GW441756. (c, d) Type-II binding mode  $^{18}\text{F}$ -labeled 2,4-diaminopyridines

such as BBB penetration and general full body biodistribution. Elucidated in PET baseline scans, it was observed that the tracer quickly entered the brain with an SUV peaking at 2 after 30 s, but was rapidly eliminated from the brain over the following 30 min. The distribution of the mixed tracer was uniform throughout the rodent brain generally matching the distribution of TrkB/C expression. A replacement study using a non-radioactive GW441756 yielded ambiguous results, although an indication of specific binding interaction was deduced from successful blocking of pulmonary radioactivity uptake. The attempted derivatization of the accessible 5-hydroxy moiety present in GW441756 with a  $^{18}\text{F}$ -fluoroethyl appendix (Fig. 31.3b) delivered a highly kinase selective compound with generally favorable in vivo properties such as a low nano-molar binding affinity for all Trk subtypes. Unfortunately, the in vitro evaluation in rat plasma revealed a pronounced CYP450 metabolic susceptibility in contrast to the original unmodified compound precluding any further efforts to translate this  $^{18}\text{F}$ -derivative into a PET tracer. An additional in vivo animal experiment confirmed a lack of in vivo blocking after pre-treatment with GW441756 and the presence of unidentified brain-penetrating metabolites preventing proper image interpretation. These accumulated data thwarted any further ambitions to translate compounds from the 3-indolydene 4-aza-2-oxindole scaffold into clinical PET tracers. Even though the labeled compounds showed a high selectivity and potency toward Trk, the unavoidable presence of the E isomer, formed by reaching a stable photostationary state, unwarranted any further research into potential PET tracer development.

### 31.2.2.3 Type-II Trk Inhibitors: 2,4-Diamionopyrimidine and Quinazoline-Derived Radiotracers

As mentioned above, type-II inhibitors targeting the DFG-out conformation are further capitalizing on the deep binding pocket made accessible by the DFG-in to DFG-out switch, leading to a longer residence time and improved overall kinetics. Compared to prototypical type-I inhibitors, type-II binding compounds are typically characterized by a linear, elongated structure which makes it possible to penetrate the lipophilic binding pocket residing alongside the DFG-out kinase binding domain. Type-II inhibitors thus display a higher average molecular weight and a higher hydrogen bond donor ability in comparison to type-I inhibitors which might translate into significantly different in vivo behavior, especially regarding brain imaging. Two structurally diverging type-II inhibitors appertaining to the class of 2,4-diaminopyrimidines (Bernard-Gauthier and Schirmmacher 2014) and quinazoline (Bernard-Gauthier et al. 2017a) have been converted into potential PET radiotracers, [ $^{18}\text{F}$ ]FluoroGW2580 and [ $^{18}\text{F}$ ]QMICF (Fig. 31.3c,d).

Extensive kinome vetting identified GW2580 as a dually active Trk as well as colony-stimulating factor-1 receptor (CSF-1R) inhibitor with a  $K_d$  of 630 nM for TrkA, 36 nM for TrkB, 120 nM for TrkC, and 2 nM for CSF-1R, representing one of the most kinome selective compounds known to date. In addition to this promising selectivity profile, preclinically acquired data on absorption, distribution, metabolism, and excretion (ADME) prompted us to approach the conversion of GW2580 into a PET radiotracer for both Trk and CSF-1R in vivo imaging. Imaging

of CSF-1R might be helpful in imaging tumor-infiltrating macrophages in cancer. We omitted the conversion of GW2580 into a structurally identical  $^{11}\text{C}$ -isotopologue and instead opted for the introduction of  $^{18}\text{F}$  to take full advantage of  $^{18}\text{F}$ 's superior PET imaging properties. A library of fluorinated GW2580 derivatives was synthesized and tested for their kinase selectivity and Trk binding affinities. Among these compounds, the most promising lead was converted into the  $^{18}\text{F}$ -radiotracer, 5-(4-((4- $^{18}\text{F}$ fluorobenzyl)oxy)-3-methoxybenzyl)pyrimidine-2,4-diamine ( $^{18}\text{F}$ FluoroGW2580) (Fig. 31.3c). However, the radiosynthesis of ( $^{18}\text{F}$ )FluoroGW2580 proved to be challenging, involving several consecutive steps which included the synthesis of the 4- $^{18}\text{F}$ fluorobenzylbromide prosthetic group for reaction with a phenolic precursor molecule (Lemaire et al. 2012). Naturally, this complex multistep synthesis provided the radiotracer in only limited amounts, unsuitable for any future clinical application. More recent  $^{18}\text{F}$ -labeling efforts however introduced a simple one-step labeling procedure based on Cu-catalyzed late-state fluorination of a GW2580 boronic ester precursor, providing the radiotracer in just two steps in high RCYs (unpublished results).

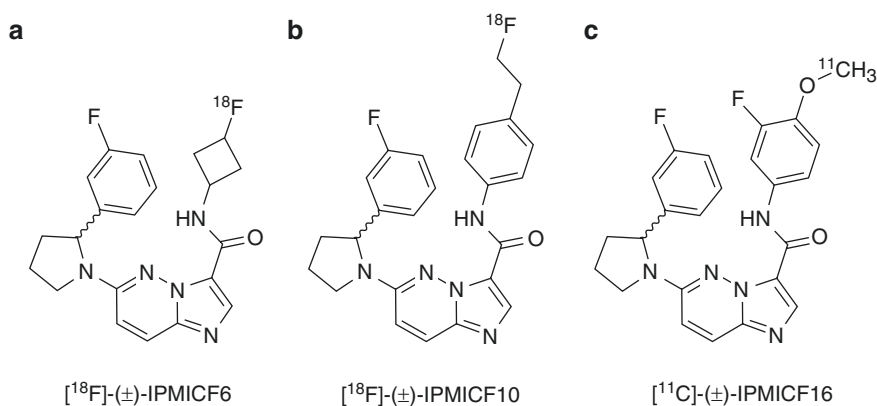
Another more recent addition to the Trk tracer portfolio is the quinazoline motif-bearing type-II pan-Trk inhibitor [ $^{18}\text{F}$ ]QMICF (Fig. 31.3d) constituting a dual Trk and FLT3 inhibitor. The compound exists in two different stereoisomers and extensive structure-activity studies identified the R isomer being the active TrkB inhibitor. Before engaging into radiolabeling, the actual positioning of the fluorine atom within the molecular structure had to be considered carefully to avoid a reduction in kinase inhibitory activity. Molecular modeling revealed that a derivatization of the allosteric pocket interacting portion of the molecule lends itself favorably to  $^{18}\text{F}$ -labeling without compromising the binding interaction. The original lead structure that was chosen as a starting point for tracer development of [ $^{18}\text{F}$ ]QMICF contained an isopropyl moiety connected to the 4-position of a benzene moiety. The isopropyl unit was replaced by a 2-fluoroethyl group of highly adaptable flexibility and able to protrude inside the allosteric binding pocket. The 2-fluoroethyl group provided a perfect opportunity to introduce the  $^{18}\text{F}$  isotope by the simple nucleophilic substitution of a mesyl leaving group of the appropriate precursor molecule in 18% RCY. Yet, even the minor structural change from an isopropyl group to a 2-fluoroethyl moiety reduced the potency of Trk binding by a factor 10. A full kinase profiling of [ $^{19}\text{F}$ ]QMICF demonstrated exceptional selectivity for Trk. In parallel, a Calcein-AM assay in P-gp-overexpressing MDCKII cells demonstrated only minor interaction with P-gp (P-glycoprotein), an important parameter for brain PET imaging. The tracer is currently being evaluated as a peripheral tumor imaging agent visualizing NTKR fusion proteins in cancer. Preclinical PET imaging in mice bearing TrkA-TPM3 overexpressing colon carcinoma KM12 tumor xenografts are ongoing in our lab and will soon be reported.

#### 31.2.2.4 Toward Clinical Application: Synthesis of Imidazo[1,2-*b*]Pyridazine-Related Radioligands

The advent of imidazo[1,2-*b*]pyridazine-related inhibitors happened coincidentally with the increased clinical interest in Trk inhibitor therapy of cancers comprising

NTRK fusions. The existing portfolio of mainly nonselective pan-Trk inhibitors was not ideally suited to address the current need for a Trk inhibitor-based therapeutic regimen. Owing to their improved pan-Trk selectivity profile, and therefore enhanced potential for a specific Trk therapy without engaging other kinases, imidazo[1,2-*b*]pyridazine- and pyrazolo[1,5-*a*]pyrimidine-based inhibitors received increased attention from medicinal chemistry and industry Trk inhibitor programs. In 2014, our group and collaborators became aware of this trend, and we chose to focus our developmental efforts on these promising structural scaffolds for radiotracer development (Bernard-Gauthier et al. 2015b). When compared side by side with all other Trk inhibitors described so far (*vide supra*), the type-I pan-Trk inhibitors originating from the (2-pyrrolidin-1-yl)imidazo[1,2-*b*]pyridazine structural motif displayed the highest affinity with all three Trk subtypes and provided an advantageous starting point for PET probe development.

Our first step in the development of a new library of potential PET-suited compounds was to engage into extensive molecular modeling studies to reveal the best position for  $^{18}\text{F}$ -introduction or  $^{11}\text{C}$ -labeling into this chemical scaffold. These modeling efforts translated into a comprehensive library of compounds suitable for radiotracer development yielding compounds that showed the highest reported potency against TrkB/C to date (<200 pm). Among these compounds, two structures caught our attention, IPMIC6 and IPMICF10. Both compounds demonstrated favorable binding to all Trk subtypes, a high kinome selectivity, and their chemical nature allowed for an easy introduction of  $^{18}\text{F}$  via nucleophilic substitution (Fig. 31.4a, b). The translation of both compounds into experimental radiotracers was straightforward, although the RCYs of the final products were rather low (3–8% RCY). Subjected to autoradiography using coronal rat brain slices, both radiotracers were able to accurately map the established brain distribution of TrkB/C receptors, in congruence with known mRNA and protein level expressions most prominent in cortical areas, the striatum, thalamus, and cerebellum.

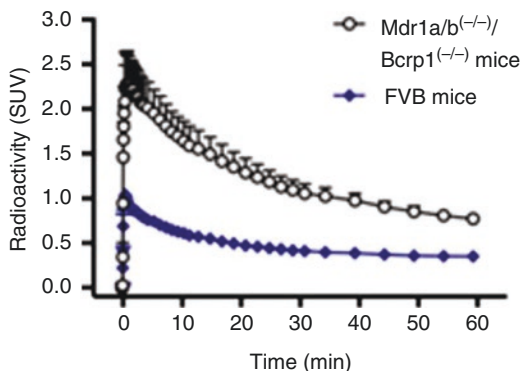


**Fig. 31.4** Chemical structures of preclinical imidazo[1,2-*b*]pyridazine-based Trk-targeting PET radioligands. (a, b)  $^{18}\text{F}$ -labeled racemic tracers. (c)  $^{11}\text{C}$ -labeled racemic lead

Despite these encouraging preliminary results, we decided to focus our attention to another member of that library, IPMICF16, based on its highest scores in binding affinity and general physicochemical properties. The structure of IPMICF16 is perfectly suited for a simple  $^{11}\text{C}$ -methylation using the most common  $^{11}\text{C}$ -labeling synthon,  $[^{11}\text{C}]$ methyl iodide (Fig. 31.4c) (Bernard-Gauthier et al. 2017b). Initially, only the racemic IPMICF16 compound was evaluated, but based on structure/activity studies using computational chemistry, it became clear that only one isomer, namely, the *R* isomer, would be responsible for the high binding affinities observed. To confirm these computational results experimentally, both isomers of IPMICF16 were synthesized and evaluated for their binding potential. (*R*)-IPMICF16 demonstrated an exceptionally high binding affinity toward all Trk subtypes as well as showing an unexpected intra-Trk selectivity.  $\text{IC}_{50}$ s of 4.0, 0.2, and 0.1 nM for human TrkA, B, and C were experimentally determined in addition to observed inhibitory constants ( $K_i$ ) of 2.80, 0.05, and 0.021 nM, respectively. Experimental studies confirmed our docking studies that only the *R* isomer possesses a high affinity to Trks. A full kinome analysis revealed an exceptional selectivity of (*R*)-IPMICF16 toward TrkB/C over 369 other targeted kinases prominent in the human brain. When reacting the desmethyl precursor of *R*-IPMICF16 with  $[^{11}\text{C}]$ methyl iodide in a single one-step reaction, RCYs of >10% were routinely achieved, enough for a non-human primate or even human PET study. Before planning non-human primate PET studies and a first-in-human study with  $[^{11}\text{C}]$ -(*R*)-IPMICF16, the radiosynthesis was established according to good manufacturing practices at four different production sites. Consequently, (*R*)-IPMICF16 was evaluated in vivo first in mice, then non-human primates, and finally in a healthy human subject.

When performing in vivo PET experiments in double knockout *Mdr1a/b*-*Bcrp* mice, it became obvious that  $[^{11}\text{C}]$ -(*R*)-IPMICF16 is a substrate for active efflux transporters such as P-gp and breast cancer resistance protein (Fig. 31.5). However, even after 60-min PET scanning time, the SUV in the rodent brain was still 0.4, pointing toward only a moderate efflux liability. In vivo PET blocking studies irrefutably confirmed specific binding of  $[^{11}\text{C}]$ -(*R*)-IPMICF16 to TrkB/C after challenging receptor binding by adding the clinically approved pan-Trk inhibitor

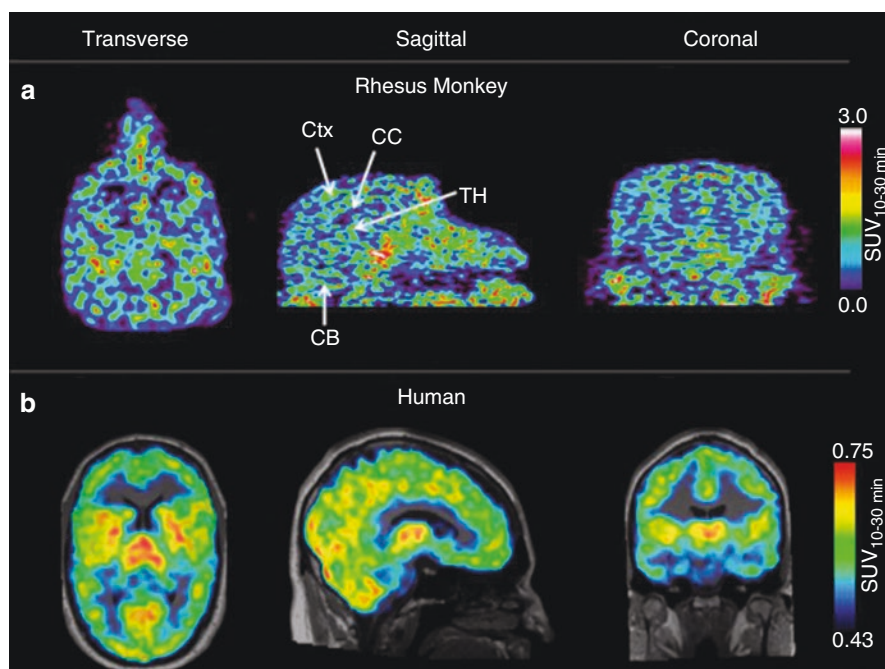
**Fig. 31.5** Time-activity curve for  $[^{11}\text{C}]$ -(*R*)-IPMICF16 brain uptake in *Mdr1a/b*<sup>(-/-)</sup>*Bcrp*<sup>(-/-)</sup> and FVB mice





entrectinib. The blocking agent was injected intraperitoneally in high dose (350 mg/kg) as a result of its low BBB penetrating properties that were reported previously. Eighty-eight percent of the brain radioactivity signal could be blocked during these *in vivo* experiments.

These positive data in rodents prompted us to further evaluate this tracer in non-human primates (NHP). Interestingly, but not entirely unexpected due to known interspecies differences in biodistribution, the brain kinetics in NHP turned out to be significantly different than what we observed in mice. Even though the overall brain uptake was clearly slower than in mice, the overall tracer uptake in the brain was continuously rising without any detectable washout during the 60-min PET scan. Analysis of the regional tracer distribution in the NHP brain showed a high uptake in TrkB/C-rich gray matter areas of the brain, reflecting the known distribution of TrkB/C in the NHP brain (highest uptake in the thalamus was SUV 0.8 (Fig. 31.6a)). In contrast to gray matter uptake, the white matter tracer accumulation was reduced in accordance with the near absence of Trk receptors in these regions. In comparison to the PET data obtained in rodents and the obvious efflux liability observed in this species, the PET data in NHP did not indicate a liability concern in higher species as a result of interspecies discrepancies.



**Fig. 31.6** In vivo PET imaging of  $[^{11}\text{C}]$ -(R)-IPMICF16. (a) Representative in vivo PET images in NHP. SUV summed over 30–60 min p.i. CB cerebellum, CC corpus callosum, Ctx cortex, TH thalamus. (b) Representative in vivo PET images in a healthy human brain with MRI overlay. Static SUV images averaged over 10–30 min p.i.

Motivated by these outcomes, the decision was made to evaluate (*R*)-[<sup>11</sup>C]IPMICF16 for human brain PET imaging. This move was further substantiated by some very encouraging autoradiography results demonstrating that (*R*)-[<sup>11</sup>C]IPMICF16 reflects the regionally specific and significant reduction of hippocampal TrkB/C concentrations in AD brain slices compared to healthy, age-matched control brains. The accumulation of [<sup>11</sup>C]-(*R*)-IPMICF16 could be challenged in both AD brains and control brains using a structurally distinct pan-Trk inhibitor evidencing specific binding of [<sup>11</sup>C]-(*R*)-IPMICF16 in the hippocampus and other TrkB/C-rich brain compartments. [<sup>11</sup>C]-(*R*)-IPMICF16 was consequently injected into a 41-year-old healthy male subject (Bernard-Gauthier et al. 2017b). The radioactivity distribution principally matched the PET data acquired in NHP with a peak SUV of 1.5 after in the thalamus after a short tracer delivery phase of 25 s to the brain (Fig. 31.6b). After 60 min, equilibrium was reached, and a SUV in the thalamus of stable 0.7 was observed followed by a slow washout. The human PET data corresponded well to the previously acquired NHP PET data, reflecting equivalent topographical distribution of TrkB/C in all gray matter regions, as well as in white matter areas of low Trk occurrence. More importantly, the general brain distribution of [<sup>11</sup>C]-(*R*)-IPMICF16 extracted from both in vivo studies corroborated the known biodistribution of TrkB/C allocated from ex vivo studies. The most pronounced uptake was found in the thalamus, followed by cerebral uptake and cortical accumulation of the tracer. No adverse effects after the injection of [<sup>11</sup>C]-(*R*)-IPMICF16 were recorded, and the tracer is now ready for further clinical application. A larger follow-up study to fully characterize [<sup>11</sup>C]-(*R*)-IPMICF16 in vivo is currently underway.

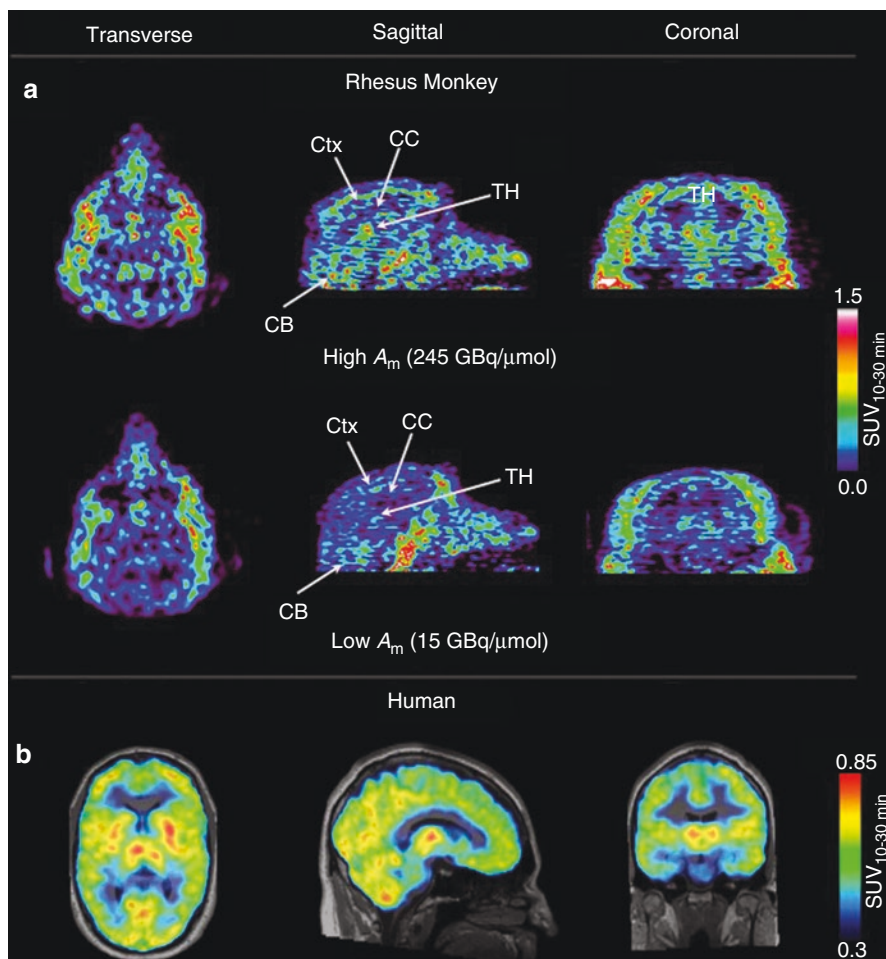
Based on our extensive previous computational studies which revealed an accurate picture of the structure-activity relationship of the Trk inhibitor binding site interaction, we aimed to modify [<sup>11</sup>C]-(*R*)-IPMICF16 toward a possible <sup>18</sup>F-radiolabeled analogue, taking full advantage of the longer half-life and more optimal decay properties (e.g., lower positron energy yielding PET images of higher resolution) of <sup>18</sup>F. The structure of [<sup>11</sup>C]-(*R*)-IPMICF16 lends itself perfectly to a <sup>18</sup>F-introduction via the so-called late-stage copper-catalyzed fluorination of aromatic non-activated boronic esters, a technique pioneered most recently by different radiochemistry groups (Preshlock et al. 2016; Brooks et al. 2014; Zlatopolskiy et al. 2015; Tredwell et al. 2014). Even so, this radiochemical methodology still has to cope with unwanted side reactions of boronic esters (e.g., protodeboronation during the radiofluorination contaminating the <sup>18</sup>F-tracer and reducing its molar activity ( $A_m$ )); it is one of the most valuable new additions to the growing portfolio of <sup>18</sup>F-labeling procedures and enables the <sup>18</sup>F-introduction into organic compounds that were previously regarded as challenging. However, the presence of a protodeboronated side product, having very similar physicochemical properties as the <sup>18</sup>F-fluorinated tracer, can significantly and undesirably compete for the ATP kinase binding site, reducing the quality of the PET image via receptor challenge.

Another main motivation to structurally amend IPMICF16 was the intention to reduce its efflux liability. The methoxy group close to the fluorine atom at the

aromatic amide fragment was clearly determined to be a major contributor to P-gp and BCRP efflux of the compound. The methoxy group was therefore removed yielding a pan-Trk inhibitor with increased subtype selectivity and a slightly decreased affinity for all Trk subtypes. The latter feature was welcome since a reduced, yet still high, affinity will more quickly achieve an equilibrium binding state. The resulting compound was named TRACK (the *R* isomer), and the  $^{18}\text{F}$ -labeled isotopologue was obtained by late-stage copper-catalyzed fluorination of the corresponding boronic ester precursor in good RCY for clinical translation. The formation of a side product stemming from protodeboronation could not be separated during the radiofluorination reaction, yielding the radiotracer [ $^{18}\text{F}$ ]TRACK with a low effective  $A_m$  of 15 GBq/ $\mu\text{mol}$ —too low for a clinical application and Trk visualization with PET. Fortunately, the use of a specialized HPLC column material, a pentafluorophenyl-modified stationary phase, achieved the baseline separation of [ $^{18}\text{F}$ ]TRACK and the protodeboronation side product, resulting in high  $A_m$  of >100 GBq/ $\mu\text{mol}$ —sufficient for human application and complying to the tracer principle (Bernard-Gauthier et al. 2018).

During the planning phase of the human study and the acquisition of ethic approval, [ $^{18}\text{F}$ ]TRACK was evaluated in the first study in a rhesus monkey affirming the uptake of [ $^{18}\text{F}$ ]TRACK in TrkB/C-rich regions of the brain (Fig. 31.7a), reminiscent of the brain distribution profile of [ $^{11}\text{C}$ ]-(*R*)-IPMICF16 in the same species yet with a slightly higher uptake in the cerebellum (SUV 0.9), thalamus (SUV 0.8) and cortex (SUV 0.6). The radioactivity uptake in white matter areas was comparable to [ $^{11}\text{C}$ ]-(*R*)-IPMICF16. Most interestingly, the overall uptake kinetics of [ $^{18}\text{F}$ ]TRACK were distinguishably different, hinting at a more reversibly binding radiotracer. As expected, injection of [ $^{18}\text{F}$ ]TRACK of different  $A_m$  into the same NHP revealed the paramount importance of a high  $A_m$  for optimal PET image quality (Fig. 31.7a).

Most recently we reported on the first-in-human evaluation of [ $^{18}\text{F}$ ]TRACK alongside with a thorough investigation of its CNS off-target engagement (Fig. 31.7b) (Bailey et al. 2019). TRACK demonstrated excellent kinome selectivity and high on-target in vitro potency for all three Trk subtypes (4.21, 0.15, and 0.31 nM for human TrkA, B, and C, respectively). The basic PET scan was performed in the same human subject that previously took part in the [ $^{11}\text{C}$ ]-(*R*)-IPMICF16 PET study, enabling the best possible and accurate comparison between the two tracers. After injection of a bolus radioactivity of 185 MBq, the radioactivity concentration due to flow-related first pass peaked after 30 s and steadily increased over the first 20 min, after which the concentration was slowly decreasing. The tracer distribution in the human brain matched the regional distribution of TrkB/C accurately. In stark difference to [ $^{11}\text{C}$ ]-(*R*)-IPMICF16, which demonstrated a steadily increasing accumulation of radioactivity in Trk-rich areas, [ $^{18}\text{F}$ ]TRACK contrarily reached a plateau with no discernable washout between 5 and 60 min scan time. Human plasma analysis revealed only a minor metabolic fraction in addition to the unaltered tracer. [ $^{18}\text{F}$ ]TRACK was 80% intact after 90 min post injection, rendering this tracer highly suitable for human brain PET imaging. In addition, no defluorination was observed attesting to the tracer's



**Fig. 31.7** In vivo PET imaging of  $[^{18}\text{F}]$ TRACK (Bernard-Gauthier et al. 2018). (a) Representative in vivo PET images in an NHP brain at high and low effective  $A_m$ . SUV summed over 10–30 min p.i. CB cerebellum, CC corpus callosum, Ctx cortex, TH thalamus. (b) PET/TIMP-RAGE MRT in vivo overlay in a healthy human brain. Static SUV images averaged over 10–30 min p.i.

high in vivo stability and robustness. Several studies in human AD and MCI (mild cognitive impairment) patients are currently planned as well as a full characterization study of the whole body dosimetry of  $[^{18}\text{F}]$ TRACK in healthy control subjects. As the contingent of newly disclosed Trk inhibitors grows over time (Bailey et al. 2017a, b), new lead compounds will be translated into potential radiotracers with a higher overall brain uptake (SUV >1) and longer residence time at the Trk receptor (type-II inhibitors).

### 31.3 Conclusion

The visualization and quantification of the Trk receptor system with potent and reversibly binding Trk radioligands is of undeniable importance to the investigation of neurodegenerative diseases such as AD and may become a diagnostic tool to assess and characterize Trk fusion proteins in cancer to support individualized Trk inhibitor-based therapies. The so far disclosed prototypical imaging probes [<sup>11</sup>C]-(R)-IPMICF16 and [<sup>18</sup>F]TRACK, although displaying a suboptimal brain uptake in humans (SUV brain <1), are certainly suitable to start the investigation whether Trk PET imaging constitutes a valuable tool for neuroimaging in the context of neurodegeneration and cancer Trk-fusion research. The most important goal is to find Trk radiotracers that display an improved general brain uptake surpassing the current generation of type-I ATP challenging probes. Macrocyclic pyrazolo[1,5-*a*]pyrimidines, such as LOXO-195, could delineate a seminal starting point for the development of a third generation of Trk imaging probes with improved physicochemical properties.

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# Radioligand Development for PET Imaging of the Vesicular Acetylcholine Transporter (VACHT) in the Brain

# 32

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## Abstract

Nowadays, it is general consensus that the cholinergic transmission system in the brain is heavily involved in the development, progress, and therapy of certain neurodegenerative diseases. In particular cholinergic presynaptic components such as the acetylcholinesterase (AChE) or the vesicular acetylcholine transporter (VACHT) are considered to be affected by early changes in neuropathological processes as observed, e.g., in Alzheimer's disease (AD). The VACHT is a transmembrane protein located at synaptic vesicles and responsible for the

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transport and storage of the neurotransmitter acetylcholine (ACh) in the vesicles. Therefore, the VAcHT is regarded as a potential target for neuroimaging of cholinergic alterations with positron emission tomography. To date, the development of PET radioligands for this transporter is based on a single known lead compound named vesamicol. A challenge was arising due to the finding that vesamicol also binds to the sigma receptors which are partly co-localized with the VAcHT in several brain regions. In the last three decades, a multitude of structural diverse vesamicol analogs has been designed resulting in a considerable number of  $^{11}\text{C}$ - and  $^{18}\text{F}$ -labeled PET as well as a few  $^{123/125}\text{I}$ -labeled SPECT tracers which were mainly preclinically evaluated. However, only very few of them had the potential for translation to human studies. Therefore, a routinely used VAcHT PET imaging method could not be established in the clinics so far. However, first results of recently published studies using the potent candidate [ $^{18}\text{F}$ ]FEOBV in patients with neuropathologies are very promising and possibly a breakthrough in this field.

This review addresses the efforts in ligand design and PET radioligand development for the VAcHT with a special view on the difficulties arising from the lead compound vesamicol and its low selectivity.

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## 32.1 Introduction

### 32.1.1 The Vesicular Acetylcholine Transporter

The vesicular acetylcholine transporter, coded by the gene *SLC18A3* (solute carrier family 18 member A3), belongs to the major facilitator superfamily (MFS). These transmembrane proteins are expressed in all organisms and transport dissolved small organic molecules across the cell membrane along a chemiosmotic gradient of either the solute itself (passive diffusion) or a second substrate (co-transport). In principle, all MFS transporters consist of 12 transmembrane helical sequences separated in two bundles of six N- and six C-terminal helices forming the cavity for the respective substrate (Bai et al. 2017).

Mainly related to neurotransmission in vertebrates, the transport of acetylcholine and other positively charged amine neurotransmitters from the cytoplasm into secretory vesicles is mediated by a specific subgroup of the MFS, the SLC18 transporters (Alexander et al. 2017). Multiple aspects of the history, function, and structure of these vesicular neurotransmitter transporters have recently been reviewed (Lawal and Krantz 2013; Anne and Gasnier 2014). The neurotransmitter ACh is the endogenous substrate of the VAcHT which utilizes a proton gradient to mediate the vesicular storage of one molecule ACh by the antiport of two protons (Alexander et al. 2017; Eiden et al. 2004). First reports, indicating an active proton-driven uptake of

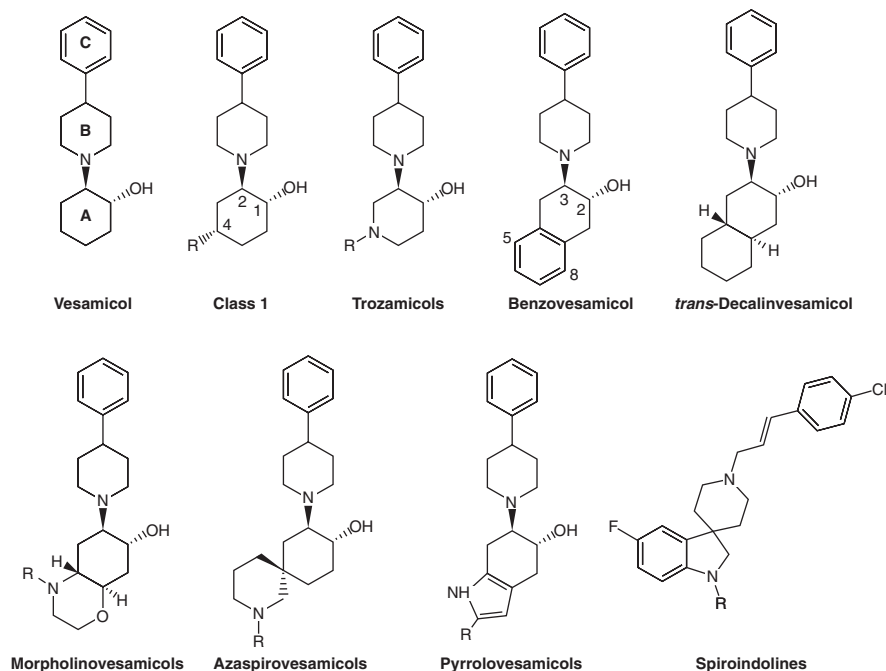
ACh into synaptic vesicles, date back to the 1980s (Koenigsberger and Parsons 1980; Michaelson and Angel 1981), followed soon by the identification of AH1853 (vesamicol) as potent inhibitor of this vesicular uptake of acetylcholine (Anderson et al. 1983; Carroll 1985) postulated by I. G. Marshall (1970) already in 1970.

With this tool in hands, biochemical research on structure-function aspects of the assumed VAcHT, at first named “synaptic vesicle vesamicol receptor” (Bahr and Parsons 1986), was strongly stimulated. The cDNA encoding VAcHT has been cloned in the early 1990s, initially in worms (Alfonso et al. 1993) and shortly after in mouse, rat, and humans (Bejanin et al. 1994; Erickson et al. 1994; Roghani et al. 1994) allowing detailed analyses of the binding sites of protons, ACh, and vesamicol (Song et al. 1997; Kim et al. 1999, 2000; Keller et al. 2000; Zhu et al. 2001; Khare et al. 2010). As one result it has been shown that vesamicol binds to an allosteric binding site of the VAcHT and inhibits the ACh transport non-competitively (Marshall and Parsons 1987; Prior et al. 1992). Although no crystal structure has been reported so far, the group of S. Schuldiner created a homology model of the human VAcHT (see Fig. 1 in Khare et al. 2010) based on the principle structure of MFS proteins, in particular the neurotransmitter transporters of the SLC18 family (Vardy et al. 2004; Elbaz et al. 2010).

As already mentioned, the VAcHT couples the transport of acetylcholine into the synaptic vesicle with an efflux of protons. By this energy-dependent process, a remarkably high gradient of ACh can be obtained—from 1–4 mM in the cytoplasm up to 600 mM in the cytoplasmic vesicle (Parsons et al. 1993)—which is of crucial importance for unimpaired neurotransmission. Stimulated release of this highly concentrated ACh from vesicles of presynaptic neurons into the synaptic cleft results in local concentrations of up to 1 mM ACh which enable a prompt and efficient activation of postsynaptic receptors such as the nicotinic acetylcholine receptors (nAChRs) possessing  $EC_{50}$  values in the micromolar range (Zwart and Vijverberg 1998). Accordingly, VAcHT expressed in neurons of the central, peripheral, autonomic, and somatic nervous system is a significant component of the ACh release cycle. Due to the presence of VAcHT in particular brain regions (Erickson et al. 1994; Albin et al. 2018), it is obvious that dysfunction or attenuated expression affects cholinergic neurotransmission. Indeed, noninvasive imaging studies as well as postmortem investigations indicate a correlation between changes in the availability of intact VAcHT and neuropathological processes related to abnormal cholinergic function such as neurodegenerative or depressive disorders in both preclinical animal models and patients (Bohnen et al. 2019; Smith et al. 2006; Chen et al. 2011; Mazere et al. 2013; Ruberg et al. 1990; Gilmor et al. 1998) as well as age-related physiological changes (Albin et al. 2018; Parent et al. 2012). Accordingly, the potential of VAcHT as marker of the integrity of cholinergic pathways, in particular of presynaptic neurons, makes this protein an attractive target for the pharmacological research and the development of diagnostic techniques for a variety of brain diseases.

### 32.1.2 Vesamicol as Basis for the Development of Ligands Targeting the VACHT

The development of VACHT radioligands is mainly based on vesamicol (*trans*-2-(4-phenylpiperidino)cyclohexanol, Fig. 32.1) which was primarily developed already in the 1960s as an analgesic drug. Fundamental work was performed by Rogers et al. with the synthesis of a large number of structurally diverse vesamicol analogues and the determination of their potential to inhibit the [ $^3\text{H}$ ]ACh transport into synaptic vesicles isolated from the electric organ of *Torpedo californica* (Rogers et al. 1989). The group discovered some important features for obtaining a high binding affinity toward VACHT such as the *trans*-orientation of the hydroxy and the amino substituent at the cyclohexylring A of vesamicol (Fig. 32.1) and the enantioselectivity with *1R,2R*-(-)-vesamicol being 25-fold more potent than *1S,2S*-(+)-vesamicol. However, subsequent extensive research has shown that vesamicol additionally exhibits substantial affinity toward the  $\sigma_1$  and  $\sigma_2$  receptors, which are also expressed in cholinergic brain regions, thereby limiting the suitability of radio-labeled vesamicol itself as a specific VACHT imaging agent (Custers et al. 1997; Ogawa et al. 2013; Bouchard and Quirion 1997). Therefore, the VACHT ligand



**Fig. 32.1** Vesamicol, selected structurally diverse vesamicol analogs and spiroindolines as potential VACHT ligands

design has moved in the recent years away from focusing solely on VACHT binding toward additionally considering potential sigma receptor binding.

When summarizing the chemical efforts of the last three decades, it becomes obvious that structural modifications were performed at all three rings A, B, and C of the vesamicol skeleton (Fig. 32.1), which was also reviewed by Giboureau et al. (2010a). A tendency can be identified at which substitutions at ring A lead to considerably increased VACHT affinity while modifications in ring B and C are less favorable. This trend was supported by a 3D QSAR approach, giving more detailed insight into structural implications for high binding affinity (Szymoszek et al. 2008). By contrast, no general trend regarding sigma receptor affinity could be found based on the results of an extensive SAR study using data from a systematically modified compound library (Barthel et al. 2015).

Figure 32.1 shows some selected ligand classes which are characterized by quite interesting ring A modifications. One of the initial SAR findings was the favorable substitution in 4-position of the cyclohexylring A resulting in an increase of VACHT binding affinity. Known examples are fluoromethylvesamicol (FMV) (Rogers et al. 1993) and 4-*O*-fluorobenzylether vesamicol (Scheunemann et al. 2004) (class 1 in Fig. 32.1) as well as the 4-azavesamicol derivative fluorobenzyltrozamicol (FBT) (Efang et al. 1993). However, the obtained high binding affinity toward VACHT was accompanied by high affinity to at least one of the sigma receptors resulting in a low selectivity.

A milestone in the VACHT ligand development was the discovery of benzovesamicol (BV) which was achieved by the fusion of a benzene with the cyclohexylring A and resulting in one of the most potent VACHT ligands (Rogers et al. 1993). Despite the remaining non-negligible affinity toward the sigma receptors (Barthel et al. 2015), the selectivity of benzovesamicol significantly improved in comparison with vesamicol (Rogers et al. 1993) due to the 20-fold increase of VACHT affinity. Therefore this compound served later as basis for the development of several <sup>18</sup>F-labeled VACHT radioligands such as [<sup>18</sup>F]FEOBV. Ring fusions were also performed with saturated ring systems resulting in structurally similar vesamicol analogs with high VACHT binding affinity such as the *trans*-decalinvesamicol (Rogers et al. 1993) and the morpholinovesamicol (Sorger et al. 2008, 2009) derivatives. Using spiro-ring fusion as another approach for obtaining conformational restriction resulted in the azaspirovesamicol derivatives (Wenzel et al. 2011) showing slightly lower VACHT affinity in comparison with benzovesamicol but negligible affinity toward sigma receptors. Unexpectedly low binding affinities for the VACHT were found for the pyrrolovesamicols characterized by the fusion of ring A with a five membered heteroaromatic ring system (Wenzel et al. 2012).

To our knowledge along with vesamicol, the spiroindoline type (Fig. 32.1) represents the only other class of compounds being reported to potentially exhibit VACHT binding in the nanomolar range (Sluder et al. 2012). In order to examine the potential of these spiroindolines as a new lead compound candidate for the development of VACHT PET ligands, selected derivatives were recently synthesized

(Lindemann et al. 2017). However, the compounds demonstrated only moderate VACHT binding affinity not suitable for PET imaging purposes. Interestingly, also these non-vesamicol based derivatives have binding affinity toward the two sigma receptor subtypes.

### 32.1.3 Considerations on Binding Affinity Data for VACHT Obtained In Vitro

The evaluation and discussion of published in vitro binding affinity data for the VACHT should be carried out with some caution because different  $K_i$  values (inhibition constants) are reported for identical compounds due to the use of different experimental protocols as also mentioned by Ogawa et al. in the recently published review (Ogawa and Shiba 2018). In general, radioligand displacement assays are utilized with (–)-[<sup>3</sup>H]vesamicol as radioligand; however, the applied biological material differs regarding species and tissue. For example, the group of Rogers and Parsons initially used synaptic vesicles isolated from the electric organ of *Torpedo californica* and reported a  $K_i$  value of 1.0 nM for (±)-vesamicol (Rogers et al. 1993). By contrast, a  $K_i$  value of 33.9 nM was reported by the group of Shiba, using membrane preparations of the rat brain (Shiba et al. 2002; Kozaka et al. 2012). Furthermore, cell lines stably transfected with either rat (Barthel et al. 2015; Roghani and Carroll 2002) or human VACHT (Tu et al. 2009; Varoqui and Erickson 1996) are used by other research groups. As part of a comprehensive SAR study, our group recently devoted attention to this matter and investigated membranes obtained from the rat brain and PC-12 cells stably transfected with a cDNA of rat VACHT (rVACHT-PC12) and (–)-[<sup>3</sup>H]vesamicol as radioligand (Barthel et al. 2015). As the main finding of this study, considerable differences of  $K_i$  values became apparent. For example, for (±)-benzovesamicol,  $K_i$  values of 1.3 nM with rVACHT-PC12 cells and 23.4 nM with the rat brain were obtained. Also, for (±)-vesamicol a higher binding affinity to rVACHT-PC12 cells ( $K_i$  of 47.4 nM) was observed compared to rat brain membranes with a  $K_i$  value of 84.2 nM. Since the two VACHT membrane preparations applied are of rat origin, comparable affinity data of the vesamicol derivatives could be expected. However, in comparison with rVACHT-PC12 cell homogenates, the rat brain is characterized by a much lower concentration of VACHT and the presence of a complex protein mixture. The latter becomes evident by the higher nonspecific binding of (–)-[<sup>3</sup>H]vesamicol in brain membranes (>20%) compared to rVACHT-PC12 cell membranes (<5%). Furthermore, it has to be considered that (–)-[<sup>3</sup>H]vesamicol is not selective due to its high affinity to sigma receptors which are widely distributed in the brain but rarely present in rVACHT-PC12 cells.

Finally, for a reasonable interpretation of VACHT binding affinity data, both species differences and the source of biological material should be considered. If and how these differences of the in vitro data correlate with in vivo PET imaging results would be interesting to explore.

## 32.2 Fluorine-18- and Carbon-11-Labeled PET Radioligands for Imaging the VAcHT

Potential radiolabeled imaging agents for a quantitative visualization of the neuronal VAcHT in living systems have to fulfil several criteria such as (1) high VAcHT binding affinity in the low nanomolar range combined with high selectivity regarding sigma receptors, (2) low nonspecific binding, (3) no brain penetrant radiometabolites, (4) high blood-brain barrier permeability with a low potential for efflux transport, and (5) rapid clearance to reduce the radioactivity background in the target tissue. In order to fulfill the first criterion, the distribution of a VAcHT selective PET probe in the brain should correspond with the known protein expression. Immunohistochemistry studies of rat brain material have confirmed the exclusive location of VAcHT at cholinergic nerve terminals with high densities in the striatum, cortex, basal forebrain, hippocampus, thalamus, amygdala, and brain stem (Schäfer et al. 1998; Arvidsson et al. 1997; Weihe et al. 1996; Gilmor et al. 1996; Ichikawa et al. 1997).

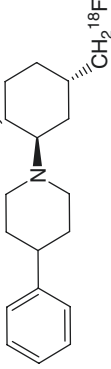
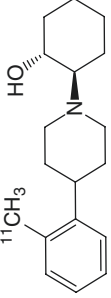
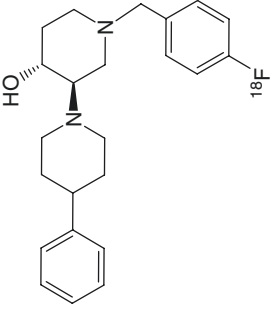
Despite the challenges in developing specific PET radioligands for imaging the VAcHT in the brain, a considerable number of fluorine-18- and carbon-11-labeled tracers were synthesized and evaluated mainly in preclinical studies. These efforts have been reviewed in 2010 by Giboureau et al. (2010b) and recently by Ogawa and Shiba (2018). A summary of the most important compounds is given in Table 32.1 and discussed below within selected structural groups.

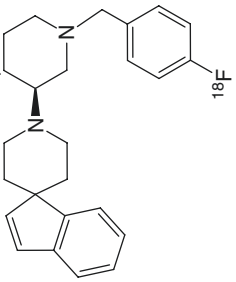
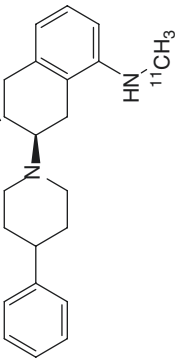
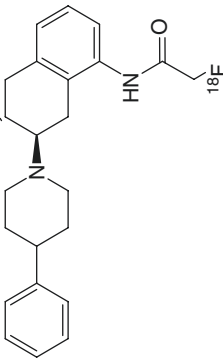
### 32.2.1 Vesamicol-Based PET Radioligands

(-)-[<sup>18</sup>F]FMV: The radiofluorinated fluoromethylvesamicol (-)-[<sup>18</sup>F]FMV was one of the first <sup>18</sup>F-labeled VAcHT PET radioligands and was developed by Widen and Rogers et al. in 1992 (Widen et al. 1992; Rogers et al. 1994). The (-)-enantiomer showed a distinctly higher affinity to the VAcHT ( $IC_{50} = 5.1$  nM) than the (+)-enantiomer ( $IC_{50} = 960$  nM), determined on vesicles of the electric organ of *T. californica* (Hicks et al. 1991). PET studies in rats and non-human primates showed a rapid and high uptake of (-)-[<sup>18</sup>F]FMV in the brain which could be blocked by pre-injection of vesamicol (Widen et al. 1992). Compared to the cerebellum, a substantially higher uptake of the radiotracer was observed in representative cholinergic brain regions such as the striatum and cortex indicating specific binding. However, the kinetic analysis of the (-)-[<sup>18</sup>F]FMV distribution in rats led the authors conclude that the pharmacokinetic profile of the radiotracer is not suitable for an accurate quantification of the VAcHT binding sites due to an irreversible binding pattern amongst others.

(-)-[<sup>11</sup>C]OMV: Carbon-11-labeled *o*-methylvesamicol (-)-[<sup>11</sup>C]OMV was reported in 2006 by Kawamura et al. (2006) and Shiba et al. (2006). In vitro binding studies revealed a high VAcHT binding affinity of (-)-OMV ( $K_i = 6.7$  nM) which was in the range of the binding affinity of (-)-vesamicol ( $K_i = 4.4$  nM) determined

**Table 32.1** In vitro binding affinity data and in vivo results of  $^{11}\text{C}$ - and  $^{18}\text{F}$ -labeled VAcHT PET ligands and the selected SPECT tracer (–) [ $^{23}\text{Na}/^{25}\text{Na}$ ]JBVM

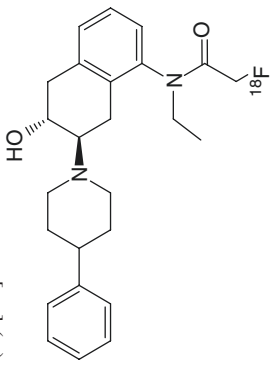
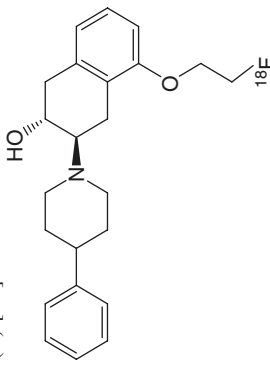
Compound name/structure	Binding affinity $K_i$ in nM		$\sigma_2$	Main findings
	VAcHT	$\sigma_1$		
(–) [ $^{18}\text{F}$ ]FMV 	$5.1 \pm 0.5^{\text{a}}$ ( $\text{IC}_{50}$ )	n.r.	n.r.	High cerebral uptake in rats and non-human primates. Unfavorable kinetics with irreversible binding (Widen et al. 1992)
(–) [ $^{11}\text{C}$ ]OMV 	$6.7 \pm 0.4^{\text{b}}$	$33.7 \pm 5.9^{\text{c}}$	$266 \pm 28^{\text{d}}$	Nonspecific accumulation in the brain of rats and non-human primates due to additional binding to sigma receptors (Kawamura et al. 2006; Shiba et al. 2006, 2009)
(+) [ $^{18}\text{F}$ ]FBT 	$0.22 \pm 0.05^{\text{a}}$	$21.6 \pm 3.9^{\text{e}}$	$35.9 \pm 6.3^{\text{d}}$	High accumulation in basal ganglia of non-human primates with suitable kinetic profile. No radiometabolites within the brain. No further studies are reported, e.g., on specificity (Efange et al. 1993, 1994a; Mach et al. 1997; Voytko et al. 2001; Gage et al. 2000)

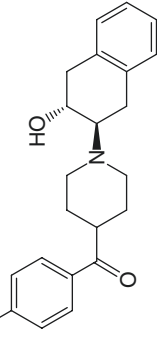
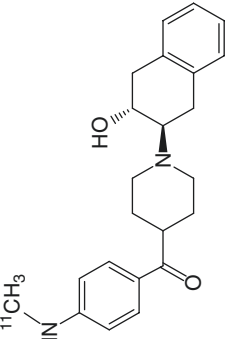
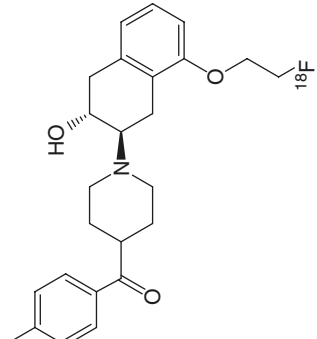
<p>(+)-[<sup>18</sup>F]Spiro-FBT</p> 	0.44 ± 0.04 <sup>a</sup>	48.7 ± 8.7 <sup>f</sup>	32.5 ± 1.3 <sup>f</sup>	Low basal ganglia-to-cerebellum ratio in non-human primates and rapid metabolism (Efange et al. 1994b, 1999)
<p>(-)-[<sup>11</sup>C]MABV</p> 	n.r.	n.r.	n.r.	High accumulation in the striatum, cortex, and hippocampus of the mouse brain with suitable kinetic profile. No radiometabolites in the brain. No further studies performed (Kilbourn et al. 1990)
<p>(-)-[<sup>18</sup>F]FAA</p> 	n.r.	n.r.	n.r.	Rapid metabolism in rats with the very likely formation of brain penetrant [ <sup>18</sup> F]fluoroacetate (Rogers et al. 1994)

(continued)



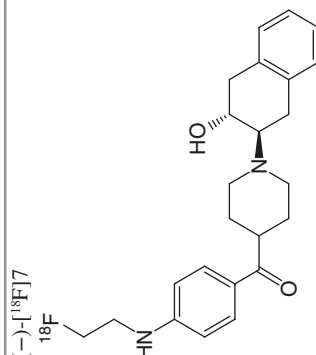
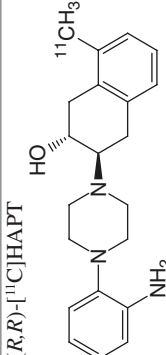
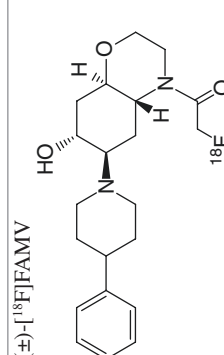
Table 32.1 (continued)

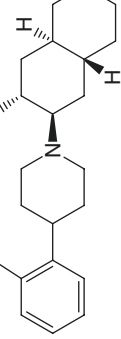
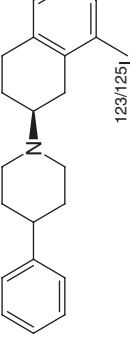
Compound name/structure	Binding affinity $K_i$ in nM		$\sigma_2$	Main findings
	VACHT	$\sigma_1$		
(-)-[ <sup>18</sup> F]NEFA 	0.32 ± 0.04 <sup>a</sup>	n.r.	n.r.	High and specific accumulation in the striatum of non-human primates with suitable tracer kinetics. High striatal but low cortical uptake in human healthy subjects. No significant difference of binding pattern between healthy controls and AD patients (Rogers et al. 1993; Ingvar et al. 1993; Widen et al. 1993)
(-)-[ <sup>18</sup> F]FEOBV 	6.75 ± 2.02 <sup>a</sup>	2275 ± 390 <sup>b</sup>	2118 ± 1058 <sup>c</sup>	High and specific binding in the striatum, thalamus, amygdala, hippocampus, neocortical regions, rostral brain stem, and cerebellar cortex in healthy humans. First successful studies in patients with AD, DLB, and PD (Albin et al. 2018; Bohnen et al. 2019; Schmitz et al. 2014; Aghourian et al. 2017; Schmitz et al. 2018; Nejad-Davarani et al. 2019; van der Zee et al. 2019; Mulholland et al. 1993, 1998; Helbert et al. 2019)

<p>(-)-[<sup>18</sup>F]FBBV</p> 	4.1 ± 0.6 <sup>f</sup>	658 ± 118 <sup>e</sup>	319 ± 36 <sup>d</sup>	<p>Low brain uptake in rats. High striatal uptake in non-human primates with moderate striatum to cerebellum ratio. No further studies reported (Tu et al. 2009)</p>
<p>(-)-[<sup>11</sup>C]TZ659</p> 	0.78 ± 0.07 <sup>i</sup>	992 ± 21 <sup>e</sup>	11,443 ± 690 <sup>d</sup>	<p>High striatal and specific binding in non-human primates with moderate striatum-to-cerebellum ratio. Suitable pharmacokinetic profile. Further studies proposed reg. suitability in humans (Li et al. 2013; Jin et al. 2016)</p>
<p>(-)-[<sup>18</sup>F]VAT</p> 	0.59 ± 0.06 <sup>j</sup>	>10,000 <sup>e</sup>	>10,000 <sup>d</sup>	<p>High specific, striatal binding in non-human primates with suitable pharmacokinetics and dosimetry. Further studies are ongoing reg. suitability in humans (Jin et al. 2018; Tu et al. 2015)</p>

(continued)

Table 32.1 (continued)

Compound name/structure	Binding affinity $K_i$ in nM		Main findings	
	VACHT	$\sigma_1$		
<p>(-)-[<sup>18</sup>F]7</p> 	0.31 ± 0.03 <sup>j</sup>	1870 ± 250 <sup>e</sup>	5480 ± 140 <sup>d</sup>	High uptake in the striatum and thalamus in preliminary studies in non-human primates. Further characterization necessary (Yue et al. 2017)
<p>(R,R)-[<sup>11</sup>C]HAPT</p> 	27.5 <sup>k</sup>	>1000 <sup>e</sup>	n.r.	High, specific, and stereospecific uptake in the putamen and caudate of non-human primates. Suitable metabolism profile. Further studies needed (Nishiyama et al. 2014)
<p>(±)-[<sup>18</sup>F]FAMV</p> 	39.9 ± 5.9 <sup>j</sup>	–	(>1500) <sup>l</sup>	Low but specific accumulation in the rat brain. Fast metabolic degradation with formation of brain penetrable radiometabolites (Sorger et al. 2008)

<p>(±)-[<sup>18</sup>F]FBMV</p>  <p><sup>18</sup>F</p>	27.5 ± 8.8 <sup>i</sup>	–	(>3000) <sup>i</sup>	Moderate, specific uptake in the rat brain. Reduction of tracer binding in rat model with cholinergic immunolesion. Suitable metabolism profile Preliminary PET imaging data from pig (Sorger et al. 2009)
<p>(±)-[<sup>11</sup>C]OMDV</p>  <p><sup>11</sup>CH<sub>3</sub></p>	11.9 <sup>b</sup>	70.3 <sup>c</sup>	43.6 <sup>d</sup>	Relatively high brain uptake in rats. Moderate target to non-target ratios (Kitamura et al. 2016)
<p>(–)-[<sup>125</sup>I/123]IBVM</p>  <p>123/125<sub>I</sub></p>	n.r.	n.r.	n.r.	High and specific binding in the striatum, thalamus, hippocampus, pons, and cerebellar vermis in healthy human. Promising results from first studies in patients with early AD and PD (Kuhl et al. 1994, 1996; Mazere et al. 2008, 2012; Jung et al. 1990, 1996; Van Dort et al. 1993; Sorger et al. 2000)

n.r., not reported

<sup>a</sup> K<sub>i</sub> or IC<sub>50</sub> values are determined using vesicles from the electric organ of *Torpedo californica* and (–)-[<sup>3</sup>H]vesamicol

<sup>b</sup> K<sub>i</sub> values are determined using cerebral membranes of rat and (–)-[<sup>3</sup>H]vesamicol

<sup>c</sup> K<sub>i</sub> values are determined using cerebral membranes of rat and (+)-[<sup>3</sup>H]pentazocine

(continued)

**Table 32.1** (continued)

Compound name/structure	Binding affinity $K_i$ in nM		Main findings
	VACHT	$\sigma_1$	
<sup>d</sup> $K_i$ values are determined using liver membranes of rat and [ <sup>3</sup> H]DTG with blocking $\sigma_1$ receptor sites using (+)-pentazocine		$\sigma_2$	
<sup>e</sup> $K_i$ values are determined using brain membranes of guinea pig and (+)-[ <sup>3</sup> H]pentazocine			
<sup>f</sup> No exact assay information available			
<sup>g</sup> $K_i$ values are determined using PC12 cell membranes stably expressing rat VACHT and (-)-[ <sup>3</sup> H]vesamicol			
<sup>h</sup> $K_i$ values are determined using cortex homogenates of rat and (+)-[ <sup>3</sup> H]pentazocine			
<sup>i</sup> $K_i$ values are determined using liver membranes of rat and [ <sup>3</sup> H]DTG with blocking of $\sigma_1$ receptor sites using dextralorphan			
<sup>j</sup> $K_i$ values are determined using PC12 cell membranes stably expressing human VACHT and (-)-[ <sup>3</sup> H]vesamicol			
<sup>k</sup> $K_i$ values are determined using cerebral membranes of rat and [ <sup>3</sup> H]aminobenzovesamicol			
<sup>l</sup> $K_i$ values are determined using liver membranes of rat and [ <sup>3</sup> H]DTG without blocking the $\sigma_1$ receptor sites			

by using rat cerebral membranes (Shiba et al. 2006). However, this vesamicol analog possessed in vitro also considerable affinity toward the  $\sigma_1$  receptor ( $K_i = 33.7$  nM). The resulting low selectivity was reflected by further in vivo evaluation of (–)-[<sup>11</sup>C]OMV (Kawamura et al. 2006). Biodistribution studies revealed an uptake of (–)-[<sup>11</sup>C]OMV in the rat brain of 1.1%ID/g at 5 min p.i. which nearly remained on this level for 60 minutes. A significant inhibition of the uptake was not only observed by co-injection of (–)-vesamicol but also by the two sigma receptor ligands SA4503 and haloperidol (Kawamura et al. 2006). Furthermore, low striatum-to-cerebellum and cortex-to-cerebellum ratios of 1.5 and 1.2, respectively, have been obtained in subsequent PET studies in non-human primates (Shiba et al. 2009). The authors concluded that (–)-[<sup>11</sup>C]OMV showed a high brain uptake and appropriate brain kinetics for a carbon-11-labeled radiopharmaceutical and suggested to use the tracer in a simultaneous imaging setup together with the sigma receptor ligand [<sup>11</sup>C]SA4503 to differentiate between VAcHT and sigma receptor binding.

### 32.2.2 Trozamicol-Based PET Radioligands

Another approach to develop VAcHT radioligands was taken in the design and synthesis of vesamicol analogs by replacing the cyclohexyl ring A by a piperidine ring (see Fig. 32.1). Thus, the available amino nitrogen of the so-called trozamicols provides an easily accessible position for introduction of radiolabeled moieties.

(+)-[<sup>18</sup>F]FBT: Fluorobenzyltrozamicol (FBT) was synthesized by Efange et al. in 1993 as a first example of the trozamicols (Efange et al. 1993). Interestingly, the (+)-enantiomer showed a higher affinity ( $K_i = 0.22$  nM) than the (–)-enantiomer ( $K_i = 16.0$  nM) determined later by Mach et al. on synaptic vesicles isolated from the electric organ of *T. californica* (Mach et al. 1997). This phenomenon was only observed for the trozamicol compound class. For all other vesamicol analogs, the (–)-enantiomer was found to be the more potent stereoisomer. The binding affinity of (+)-FBT was slightly higher compared to the affinity of (–)-vesamicol ( $K_{i(\text{ves})} = 1.0$  nM). A lower, but not negligible binding affinity of (+)-FBT was observed for the  $\sigma_1$  ( $K_i = 21.6$  nM) and  $\sigma_2$  ( $K_i = 35.9$  nM) receptors resulting in 98- and 160-fold ratio of VAcHT/ $\sigma_1$  and VAcHT/ $\sigma_2$  affinities, respectively. In vivo evaluation of (+)-[<sup>18</sup>F]FBT in rats showed an initial brain uptake of 0.82%ID/g at 5 minutes decreasing to 0.27% ID/g at 3 h p.i. (Efange et al. 1994a). The highest accumulation of activity was observed in the striatum and hippocampus, followed by moderate uptake in cortex and low uptake in the cerebellum resulting in a maximum striatum-to-cerebellum ratio of 2.8 at 3 h p.i. Sigma receptor binding appeared to account for only a small fraction of (+)-[<sup>18</sup>F]FBT binding since pre-injected haloperidol had no significant effect on the binding in frontal cortex and hippocampus. PET imaging studies in non-human primates revealed a high accumulation and slow washout of (+)-[<sup>18</sup>F]FBT from the basal ganglia with a continuously increasing basal ganglia-to-cerebellum ratio reaching 1.75 at 3 h p.i. (Mach et al. 1997). In contrast, the time-activity curves of cortical regions are in the same range as for the cerebellum. In this study no blocking experiments were performed in order to verify

the specificity of the radiotracer. (+)-[<sup>18</sup>F]FBT was found to be similarly metabolized in both species, and the generated radiometabolites did not cross the blood-brain barrier (Mach et al. 1997). The reproducibility of the (+)-[<sup>18</sup>F]FBT PET measurements in the basal ganglia was demonstrated in a separate study (Gage et al. 2000). The authors concluded that (+)-[<sup>18</sup>F]FBT is a promising radiotracer for studying cholinergic terminal density with PET and indeed an age-dependent reduction of the binding of (+)-[<sup>18</sup>F]FBT in the basal ganglia of aged rhesus monkeys could be demonstrated (Voytko et al. 2001). However, obtained individual differences require further investigations. To our knowledge, there are no reports on the application of (+)-[<sup>18</sup>F]FBT in human studies despite the promising preclinical results. Tu and Mach et al. mentioned in a paper in 2009 (Tu et al. 2009) unpublished data which demonstrated unacceptably high nonspecific binding in human subjects.

**(+)-Spiro-FBT:** Fluorobenzyl spirotrozamicol (Spiro-FBT) was developed by Efange et al. in 1994 with the aim to design conformationally restricted vesamicol analogues maintaining an orthogonal orientation of the phenyl ring C and the piperidine ring B (Efange et al. 1994b). As also observed for FBT, (+)-Spiro-FBT showed a higher binding affinity ( $K_i = 0.44$  nM) toward the VACHT than (–)-Spiro-FBT ( $K_i = 30.5$  nM) (Efange et al. 1999). The affinity to sigma receptors was determined with  $K_i$  values of 48.7 and 32.5 nM for  $\sigma_1$  and  $\sigma_2$ , respectively. Biodistribution studies with (+)-[<sup>18</sup>F]Spiro-FBT in rats showed an initial brain uptake of 0.74% ID/g at 5 min p.i. The accumulation in the different brain regions striatum, hippocampus, cortex, and cerebellum was in a similar range of 0.66–0.84% ID/g at 5 min p.i. (Efange et al. 1999). However, due to different rates of washout, the striatum-to-cerebellum and hippocampus-to-cerebellum ratios increased from about 1.2 to 2.0 within 120 min p.i. Pretreatment with (+)-FBT reduced the uptake in most of the brain regions suggesting identical binding sites for these two ligands in vivo. PET studies in rhesus monkeys revealed a rapid accumulation of activity in the brain. However, the washout of (+)-[<sup>18</sup>F]Spiro-FBT from the basal ganglia was more rapid displaying a lower basal ganglia-to-cerebellum ratio in comparison with (+)-[<sup>18</sup>F]FBT. This unexpected result was explained by a fast metabolism of (+)-[<sup>18</sup>F]Spiro-FBT with only 10% of intact tracer in the plasma at 10 min p.i. This led the authors to conclude that this radiotracer might be not suitable for PET imaging of the VACHT.

### 32.2.3 Benzovesamicol-Based PET Radioligands

Most of the PET radioligands developed for imaging of the VACHT are derived from benzovesamicol as this vesamicol analog demonstrated a very high binding affinity ( $K_i = 0.055$  nM, on vesicles of *T. californica* Rogers et al. 1993).

**(–)-[<sup>11</sup>C]MABV:** One of the first representatives is the carbon-11-labeled 5-(N-methylamino)benzovesamicol (–)-[<sup>11</sup>C]MABV which was developed by Kilbourn et al. already in 1990 on the basis of 5-aminobenzovesamicol (ABV, Rogers et al. 1993; Kilbourn et al. 1990). Biodistribution studies in mouse showed

a very high radiotracer uptake in the striatum (8.1% ID/g at 45 min p.i.) followed by the cortex, hippocampus, and hypothalamus. A slower washout in these brain regions compared to the cerebellum resulted in striatum-to-cerebellum and cortex-to-cerebellum ratios of 13 and 5, respectively, at 75 min p.i. of (–)-[<sup>11</sup>C]MABV. The accumulation in these brain regions was lower after pretreatment of (±)-vesamicol and showed stereoselectivity with higher uptake of the (–)-enantiomer compared to the (+)-enantiomer. A radiometabolite analysis of homogenized mouse brain tissue at 45 minutes p.i. of (–)-[<sup>11</sup>C]MABV revealed the sole presence of the radiotracer. However, despite these promising results, the authors did not continue with this compound, and except an improved radiosynthesis protocol (Mulholland and Jung 1992), no further work on (–)-[<sup>11</sup>C]MABV has been published.

[<sup>18</sup>F]FAA and [<sup>18</sup>F]NEFA: The *N*-[<sup>18</sup>F]fluoroacetylated and *N*-ethyl-[<sup>18</sup>F]fluoroacetylated 5-aminobenzovesamicol derivatives [<sup>18</sup>F]FAA and [<sup>18</sup>F]NEFA ( $K_{i((\pm)\text{NEFA})} = 0.32$  nM Rogers et al. 1993) were developed by Rogers et al. (1994). PET studies of [<sup>18</sup>F]FAA in rats in combination with the monitoring of arterial blood revealed a rapid metabolism with the very likely generation of [<sup>18</sup>F]fluoroacetate, and therefore the authors excluded this radiotracer from further studies. By contrast, initial PET investigation of [<sup>18</sup>F]NEFA in rats indicated a specific uptake in cerebral tissues and a higher metabolic stability due to the *N*-ethylation. Subsequent PET imaging studies in non-human primates (Ingvar et al. 1993) showed a stereoselective binding of (–)-[<sup>18</sup>F]NEFA with a regional distribution pattern of striatum>cortex>cerebellum at late time points. The activity ratio of striatum to cerebellum increased continuously and reached a value in the range of 2.5–3 at 50 min p.i. The tracer binding could be selectively blocked but not displaced with vesamicol and remained unaffected by the pretreatment with the  $\sigma_1$  receptor ligand pentazocine. The twofold increase of the striatum-to-cerebellum ratio in response to the pretreatment with haloperidol or raclopride suggested that blocking of the dopamine-D2 receptors stimulated the cholinergic system of the striatum. Therefore, a human PET study was performed on two healthy subjects and one AD patient (Widen et al. 1993). However, despite a high accumulation in the striatum, (–)-[<sup>18</sup>F]NEFA showed no relevant binding in the cortex of the healthy subjects, and no difference was observed between healthy and diseased brain. Accordingly, the authors did not perform further studies with this radiotracer.

(–)-[<sup>18</sup>F]FEOBV: One of the most investigated VACHT PET radioligands is the 5-[<sup>18</sup>F]fluoroethoxy-substituted benzovesamicol [<sup>18</sup>F]FEOBV which was developed by Mulholland et al. (1993, 1998). In a very fundamental study in rodents, this group investigated the regional distribution in the brain, stereospecificity, dosimetry, toxicity, pharmacological blocking effects, and the *in vivo* metabolism. (–)-[<sup>18</sup>F]FEOBV demonstrated (1) a high brain uptake (2.68% ID/g at 5 min p.i.) with a strong cholinergic pattern of localization matching known distributions of other pre-synaptic cholinergic markers, (2) a favorable metabolism, and (3) no considerable interaction with dopamine or  $\sigma_1$  receptor binding sites. The clearance of the radiotracer from the cerebellum was fast and slow from the striatum where the highest tracer accumulation was reached between 45 and 180 min p.i. (8–9% ID/g) with a maximum striatum-to-cerebellum ratio of about 5. Interestingly, to our knowledge,



no in vitro binding affinity data were reported at this time. Recently, our group together with Elsinga et al. investigated (–)-FEOBV (Helbert et al. 2019) and determined a  $K_i$  value of 6.75 nM toward the VACht (on rVACht PC12 cells). For  $\sigma_1$  and  $\sigma_2$  receptors,  $K_i$  values of about 2000 nM were determined (Helbert et al. 2019). These data correspond to the specific binding observed in the initial in vivo studies. The early results obtained by Mulholland et al. were later confirmed and the studies expanded to non-human primates demonstrating the potential of this radiotracer for imaging the VACht in the brain (Kilbourn et al. 2009). Moreover, a correlation of reduced (–)-[ $^{18}\text{F}$ ]FEOBV binding in the rat brain with aging as well as with cholinergic deficits caused by lesions was reported (Parent et al. 2012; Cyr et al. 2014). Because of these very promising preclinical findings, in vivo PET studies in human followed. In the recently published reports (Albin et al. 2018; Petrou et al. 2014), the authors describe a tracer distribution and regional density pattern which is highly consistent with the known distribution of cholinergic nerve terminals. Accordingly, highest binding was found in the striatum, followed by particular regions of the thalamus, amygdala, hippocampus, some neocortical regions, rostral brain stem, and some regions of the cerebellar cortex. The data also demonstrated that (–)-[ $^{18}\text{F}$ ]FEOBV has an acceptable dosimetry and safety profile in human subjects (Petrou et al. 2014). Moreover, the results indicated an association of normal aging with regional reductions of (–)-[ $^{18}\text{F}$ ]FEOBV binding in the striatum and some cortical regions (Albin et al. 2018). Finally, the authors concluded that PET imaging using (–)-[ $^{18}\text{F}$ ]FEOBV provides more direct information about cholinergic terminal integrity than, for example, acetylcholinesterase-based PET methods as these are indirect markers because they reflect the activity of a hydrolytic enzyme which might be variable. With these satisfying results, a starting signal was given for using (–)-[ $^{18}\text{F}$ ]FEOBV in clinical studies related to neuropathologies. Thus, in the last 2 years, patients suffering from AD (Aghourian et al. 2017; Schmitz et al. 2018), dementia with Lewy bodies (DLB) (Nejad-Davarani et al. 2019), Parkinson's disease (Bohnen et al. 2019; van der Zee et al. 2019), or REM sleep behavior disorder (Bedard et al. 2019) were investigated with (–)-[ $^{18}\text{F}$ ]FEOBV PET. The therein described initial observations are very promising as it seems to be possible with this first selective VACht radiotracer to distinguish between the participation of certain cholinergic regions or projections in the different neuropathologies probably due to the fact that the targeted protein is solely present in cholinergic nerve terminals. Further studies are needed now to, e.g., (1) better characterize the (–)-[ $^{18}\text{F}$ ]FEOBV binding in diseases associated with cholinergic dysfunction, (2) to investigate the relationship between regional cholinergic alterations and clinical symptomatology, and (3) to compare (–)-[ $^{18}\text{F}$ ]FEOBV PET to other imaging approaches with regard to a reliable disease discrimination.

**[ $^{18}\text{F}$ ]FBBV:** In 2009, Tu et al. developed with [ $^{18}\text{F}$ ]fluorobenzoylbenzovesamicol ([ $^{18}\text{F}$ ]FBBV) the first representative of a new class of benzovesamicol derivatives, containing a carbonyl group between the fragments B and C of the vesamicol scaffold (Tu et al. 2009). Until 2018, this group and others reported on numerous  $^{11}\text{C}$ - (Jin et al. 2016; Padakanti et al. 2014; Luo et al. 2018; Bergman et al. 2014) and  $^{18}\text{F}$ -labeled (Jin et al. 2018; Yue et al. 2017; Luo et al. 2018; Yue et al. 2018)

compounds belonging to this class of which only the most potent candidates are reviewed here. The (–)-enantiomer of FBBV exhibited with a  $K_i$  value of 4.1 nM (on hVACHT PC12 cells) a high binding affinity to the VACHT accompanied by a high selectivity with  $K_i$  values of 658 and 319 nM for the  $\sigma_1$  and  $\sigma_2$  receptors, respectively (Tu et al. 2009). In vivo studies in rats revealed a moderate brain uptake of (–)-[ $^{18}\text{F}$ ]FBBV with a maximum of 0.82% ID/g at 5 min p.i. reaching a striatum-to-cerebellum ratio of 1.8 at 30 min p.i. MicroPET imaging studies in non-human primates demonstrated a high uptake in the striatum ( $\text{SUV}_{\text{max}}(\text{caudate}) = 4.5$  and  $\text{SUV}_{\text{max}}(\text{putamen}) = 5$  at 30 min p.i.) with a maximum striatum-to-cerebellum ratio of 2.1 at 2 h. No attempts were made to perform blocking studies in order to prove the specificity of the radiotracer. The authors stated that (–)-[ $^{18}\text{F}$ ]FBBV has excellent potential to be a novel PET tracer for clinical imaging of the VACHT; however, no further studies have been reported so far.

(–)-[ $^{11}\text{C}$ ]TZ659: With the  $^{11}\text{C}$ -labeled methylaminobenzoylbenzovesamicol (–)-[ $^{11}\text{C}$ ]TZ659, Tu et al. developed another compound of the carbonyl group containing benzovesamicol class. This derivative is characterized by an excellent in vitro binding profile with  $K_i$  values of 0.78, 992, and 11,443 nM for the VACHT,  $\sigma_1$ , and  $\sigma_2$  receptors, respectively (Li et al. 2013). Biodistribution studies in rats showed highest radiotracer accumulation in the striatum (0.6% ID/g at 30 min p.i.) reaching a striatum-to-cerebellum ratio of 3.75. An about twofold increase of brain uptake was observed due to the injection of the P-glycoprotein (P-gp) inhibitor cyclosporine indicating that (–)-[ $^{11}\text{C}$ ]TZ659 might be a potential substrate of this drug efflux transporter (Li et al. 2013). Subsequent PET imaging studies in non-human primates showed a high retention of the radiotracer in the striatum with a maximum SUV of 4.5 at 20 min p.i. and a slow washout (Jin et al. 2016). This tracer binding could be blocked and displaced in a dose-dependent manner by (–)-vesamicol but not (+)-pentazocine. Fastest washout was observed for the cerebellum giving striatum-to-cerebellum ratios in the range of 1.8–2.4 from 60 to 110 min scanning time. From occupancy studies, an  $\text{ED}_{50}$  value of 0.0057 mg/kg of (–)-vesamicol for striatal VACHT was calculated (Jin et al. 2016). Based on these results, the authors intended to further investigate (–)-[ $^{11}\text{C}$ ]TZ659 regarding its suitability for quantifying VACHT levels in human subjects and characterizing expression changes during disease progression or therapeutic treatment.

(–)-[ $^{18}\text{F}$ ]VAT: Further exploration of the carbonyl-substituted benzovesamicols leads to (–)-[ $^{18}\text{F}$ ]VAT which can be regarded as a structural merge of FEOBV and FBBV. The new derivative has an excellent in vitro binding profile with a  $K_i$  value of 0.59 nM for the VACHT and no binding affinity toward the  $\sigma_1$  and  $\sigma_2$  receptors (Tu et al. 2015). Interestingly, the introduction of the fluoroethoxy group in 5-position of the benzene ring of FBBV strongly diminished the binding to the sigma receptors, a phenomenon which was already observed for FEOBV. Biodistribution studies in rats showed highest regional radiotracer uptake in the striatum (0.9% ID/g at 30 min) reaching a striatum-to-cerebellum ratio of about 3 at 30 min (Tu et al. 2015). PET imaging and dosimetry studies in non-human primates were performed demonstrating the great potential of this radiotracer (Jin et al. 2018; Tu et al. 2015). Highest (–)-[ $^{18}\text{F}$ ]VAT binding was observed in the striatum followed by the

thalamus, hippocampus, frontal cortex, and vermis corresponding to known distribution of cholinergic terminals. The peak of striatal uptake was already reached at 15 min p.i. of (–)-[<sup>18</sup>F]VAT (SUV of 5) indicating a faster equilibrium compared to (–)-[<sup>18</sup>F]FEOBV. Blocking with (–)-vesamicol significantly reduced the striatal uptake, while no change was observed in cerebellar regions. A metabolite analysis of arterial blood samples showed a decrease of parent radiotracer from 66% at 15 min to 16% at 90 minutes p.i. Moreover, different modeling methods were applied to estimate the VAcHT levels in non-human primates, and a good reproducibility and reliability in repeated measurements were found. According to the authors, the clinical investigation of (–)-[<sup>18</sup>F]VAT in patients with various neurodegenerative and neurological diseases is ongoing (Luo et al. 2018).

(–)-[<sup>18</sup>F]7: A recently published derivative belonging to the carbonyl-containing benzovesamicols was obtained by replacement of the methyl group of (–)-[<sup>11</sup>C]TZ659 by a fluoroethyl chain in order to obtain the corresponding <sup>18</sup>F-labeled radiotracer (–)-[<sup>18</sup>F]7 (Yue et al. 2017). Also this compound showed very high binding affinity to the VAcHT ( $K_i = 0.31$  nM) and negligible affinities to the  $\sigma_1$  and  $\sigma_2$  receptors. Surprisingly, the radiosynthesis of (–)-[<sup>18</sup>F]7 turned out to be rather challenging, and the deliverable activity was not sufficient for routine PET studies in non-human primates. Therefore, only two PET studies in a male macaque were performed with, however, quite promising results (Yue et al. 2017). The activity uptake peaked at about 10 min p.i. and was highest in the striatum ( $SUV_{max} \sim 8$ ) followed by the thalamus ( $SUV_{max} \sim 7.5$ ), cerebellum, hippocampus, and cortical regions. The washout was slowest from the striatum followed by the thalamus and very fast from the cerebellum resulting in striatum-to-cerebellum and thalamus-to-cerebellum ratios of about 3 and 2.3, respectively, at 60 minutes p.i. of (–)-[<sup>18</sup>F]7. The authors intend to further characterize this radiotracer after improving the radiosynthesis procedure.

(*R,R*)-[<sup>11</sup>C]HAPT: In 2014 Nishiyama et al. reported on [<sup>11</sup>C]HAPT another <sup>11</sup>C-labeled benzovesamicol derivative bearing as ring B a piperazine instead of the piperidine ring in the vesamicol core (Nishiyama et al. 2014). Interestingly, the <sup>11</sup>C-methyl group is substituted in 8-position of the tetralin ring system which was described to be unfavorable compared to the 5-position in case of amino substitution like in aminobenzovesamicol (Rogers et al. 1989). The enantiomer (*R,R*)-[<sup>11</sup>C]HAPT showed a considerably higher binding affinity for the VAcHT ( $K_i = 27.5$  nM on rat cerebral membranes) than the (*S,S*)-[<sup>11</sup>C]HAPT ( $K_i = 699$  nM) and a negligible affinity toward the  $\sigma_1$  receptor. This was well reflected by the PET imaging studies in non-human primates (Nishiyama et al. 2014). While (*R,R*)-[<sup>11</sup>C]HAPT was heterogeneously distributed in the brain with highest uptake in the putamen and caudate ( $SUV_{max}$  at 20 min: 4 and 3.4, respectively), intermediate uptake in the thalamus and amygdala, lower uptake in cortical regions, and lowest in the cerebellum, the (*S,S*)-[<sup>11</sup>C]HAPT showed a lower and homogeneous distribution in all brain regions. The preadministration of (–)-vesamicol induced a complete inhibition of (*R,R*)-[<sup>11</sup>C]HAPT binding in all regions except the cerebellum. By contrast, no significant changes in tracer uptake were observed after preadministration of SA4503, a specific  $\sigma_1$  receptor ligand. A metabolite analysis revealed a

fraction of 34% of parent radiotracer (*R,R*)-[<sup>11</sup>C]HAPT in plasma, at 45 min p.i. The authors concluded that this radiotracer is a potential PET probe for quantification of the VAcHT in the living brain.

### 32.2.4 Morpholinovesamicols

In an effort to develop selective VAcHT radioligands keeping the favorable binding profile of benzovesamicol but possessing a more moderate lipophilicity, new structurally similar analogs were designed by fusing a morpholino instead of a benzene ring on ring A of the vesamicol skeleton. Moreover, the nitrogen atom of the morpholine system offers the possibility of a straightforward introduction of different fluoro-carrying functionalities. Thus, different morpholinovesamicol derivatives were developed, and two of them are radiofluorinated and preclinically evaluated (Sorger et al. 2008, 2009).

(±)-[<sup>18</sup>F]FAMV: The [<sup>18</sup>F]fluoroacetyl-substituted derivative (±)-[<sup>18</sup>F]FAMV was published in 2008 by Sorger et al. and showed with a  $K_i$  value of 39.9 nM (on rVAcHT PC12 cells) and low binding affinities toward the sigma receptors a good in vitro binding profile (Sorger et al. 2008). Injected into rats, (±)-[<sup>18</sup>F]FAMV accumulated only to low amounts (0.12% ID/g at 5 min p.i.) but specifically in the brain as demonstrated by pretreatment with (–)-vesamicol (Sorger et al. 2008). Ex vivo autoradiography showed a heterogeneous uptake which was highest in the striatum followed by the cortical regions, hippocampus, and thalamus and lowest in the cerebellum. Region-to-cerebellum ratios at 1 h p.i. of (±)-[<sup>18</sup>F]FAMV were 4.5 and 2 for the striatum and cortical regions, respectively. Metabolism studies revealed a fast degradation with the formation of brain penetrable radiometabolites, limiting the utility of this compound for in vivo studies.

[<sup>18</sup>F]FBMV: With the [<sup>18</sup>F]fluorobenzoyl-substituted derivative (±)-[<sup>18</sup>F]FBMV, another morpholinovesamicol was developed exhibiting a slightly higher binding affinity toward the VAcHT ( $K_i = 27.5$  nM) and selectivity regarding the  $\sigma$  receptors compared to (±)-[<sup>18</sup>F]FAMV (Sorger et al. 2009). Biodistribution studies in rats showed highest brain uptake of 0.17% ID/g at 5 min p.i. of (±)-[<sup>18</sup>F]FBMV with a regional distribution at later time points which is typical for VAcHT-containing brain regions (Sorger et al. 2009). At 1 h p.i., striatum-to-cerebellum and cortex-to-cerebellum ratios of about 3.4 and 1.8, respectively, were determined in ex vivo autoradiography experiments. Pretreatment with (–)-vesamicol inhibited the observed binding quantitatively resulting in a homogeneous tracer distribution in the brain. Investigation of (±)-[<sup>18</sup>F]FBMV using a rat model with a specific cholinergic immunolesion revealed a reduced tracer binding by about 20% in particular in cortical brain regions. In metabolism studies in rats, parent radiotracer fractions of 79, 61, and 43% at 15, 60, and 120 minutes p.i. of (±)-[<sup>18</sup>F]FBMV, respectively, were determined in plasma, and mainly intact radiotracer was detectable in the brain demonstrating a considerably higher in vivo stability compared to (±)-[<sup>18</sup>F]FAMV. Preliminary PET studies were performed in pigs and revealed a desirable brain uptake of (±)-[<sup>18</sup>F]FBMV with a regional distribution corresponding to

cholinergic nerve terminals. Co-administration of ( $\pm$ )-vesamicol resulted in a 60% reduction of radiotracer binding in the striatum; however, further studies are needed to better characterize this radiotracer.

### 32.2.5 Other Vesamicol Analogs

[ $^{11}\text{C}$ ]OMDV: The ring-fused analog *trans*-decalinvesamicol (Fig. 32.1) was first synthesized by Rogers et al. (1993) and demonstrated an exceptional high binding affinity to the VAcHT ( $K_i = 0.009$  nM on vesicles of *T. californica*). However, the compound does not offer simple access for radiofluorination. Therefore, the group of K. Shiba (Kitamura et al. 2016) focused on  $^{11}\text{C}$ -labeling by substituting a methyl group in *ortho* position of the phenyl ring as this position was shown to be more suitable than the *para* position when the *o*-methylvesamicol [ $^{11}\text{C}$ ]OMV was developed (Shiba et al. 2006). Interestingly, the obtained OMDV showed still a high binding affinity toward the VAcHT ( $K_i = 11.9$  nM on rat cerebral membranes) but much lower than reported by Rogers et al. The binding affinities to the  $\sigma_1$  and  $\sigma_2$  receptors were determined to be 70.3 and 43.6 nM, respectively (Kitamura et al. 2016). Biodistribution studies in rats showed an initial uptake of 0.92% ID/g at 2 minutes p.i. of [ $^{11}\text{C}$ ]OMDV in the cerebrum which decreased within 60 minutes to 0.44% ID/g. The regional activity distribution at 30 minutes p.i. was highest in the striatum (0.69% ID/g) followed by the cortex (0.62% ID/g) and the cerebellum with 0.50% ID/g resulting in a striatum-to-cerebellum ratio of 1.4. The uptake was inhibited in all three brain regions by co-injection of (–)-vesamicol. When (+)-pentazocine or the  $\sigma_{1/2}$  receptor ligand (+)-3-PPP was administered, no significant changes were observed. These findings were confirmed by PET-CT studies in rats. However, the time-activity curves demonstrated rather low target-to-non-target ratios for the striatum and cortex during the scanning time of 35 minutes. In this study, the *in vivo* metabolism was not investigated. The authors concluded that [ $^{11}\text{C}$ ]OMDV may be suitable for PET imaging of dementia; however, further evaluation of the radiotracer is necessary.

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### 32.3 The Use of (–)-[ $^{123}/^{125}$ ]iodobenzovesamicol as SPECT Imaging Agent for VAcHT

Several radioiodinated vesamicol analogs were developed for the use as SPECT imaging agents. Since this review is focused on PET radiopharmaceuticals, they are not listed here. However, because of the important findings obtained with (–)-[ $^{123}$ ]iodobenzovesamicol in human studies, this section of our chapter concerns this SPECT tracer.

Jung et al. developed 5-[ $^{125}$ ]iodobenzovesamicol ([ $^{125}\text{I}$ ]IBVM) already in 1990 and investigated the biodistribution of this tracer in mice (Jung et al. 1990). The (–)-enantiomer showed a high accumulation in the striatum (6.26% ID/g) and the cortex (2.56% ID/g) and low uptake in the cerebellum (0.23% ID/g) at 4 h p.i. In

comparison, the accumulation of the (+)-enantiomer was much lower accompanied by a rather homogeneous distribution in the brain. Pretreatment with (–)-vesamicol reduced the striatal and cortical uptake of (–)-[<sup>125</sup>I]BVM considerably. A few years later, the same group established the radiosynthesis of the <sup>123</sup>I-labeled derivative (–)-[<sup>123</sup>I]BVM and demonstrated that the two radioiodinated compounds behave similarly in vivo (Van Dort et al. 1993). Sorger et al. evaluated (–)-[<sup>125</sup>I]BVM in a rat model with a cholinergic immunolesion showing a significantly reduced binding in the affected brain regions which corresponded to a reduced AChE binding as measured by AChE histochemical staining (Sorger et al. 2000). Moreover, (–)-[<sup>125</sup>I]BVM was found to specifically bind in cholinergic brain regions as demonstrated by blocking studies with (–)-vesamicol, haloperidol (dopamine D2 and sigma receptor ligand), and DTG ( $\sigma_{1/2}$  receptor ligand) (Sorger et al. 2000).

Kuhl et al. reported in 1994 and 1996 on human studies investigating (–)-[<sup>123</sup>I]BVM in healthy subjects (Kuhl et al. 1994) and in AD and PD patients (Kuhl et al. 1996). While in mildly demented AD patients with an early onset a reduced (–)-[<sup>123</sup>I]BVM binding in the neocortex and hippocampus was observed, reductions were significant in temporal cortex and hippocampus in mild AD patients with a late onset. By contrast, when the dementia was severe, terminal deficits were extensive in all AD patients independent from the time of onset. More than 10 years later, Mazère et al. also reported on (–)-[<sup>123</sup>I]BVM SPECT imaging in patients with early Alzheimer's disease and demonstrated a decrease of tracer binding of 47–62% relative to healthy subjects which was specifically localized in the cingulate cortex and parahippocampal-amygdaloid complex (Mazere et al. 2008). The authors concluded that the results suggest the occurrence of cholinergic degeneration in the early stage of AD which is likely related to the impairment of the cognitive function. Therefore, the data support the cholinergic hypothesis (Bartus et al. 1982) suggesting that human cholinergic dysfunction is an early pathological feature of cognitive decline in AD. Further SPECT imaging studies with (–)-[<sup>123</sup>I]BVM in patients with progressive supranuclear palsy (Mazere et al. 2012) and multiple system atrophy (Mazere et al. 2013) followed in later years to evaluate the integrity of central cholinergic pathways.

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## 32.4 Summary and Conclusion

The VAcHT is a very attractive target for imaging cholinergic innervation as this transporter is exclusively expressed in the presynaptic part of cholinergic neurons. This makes the VAcHT valuable for investigating, characterizing, and diagnosing brain diseases involving cholinergic dysfunction. As already concluded by Ogawa and Shiba (2018), this transporter additionally can be seen as an ideal imaging counterpart of other cholinergic elements such as acetylcholine receptors which are located post- and presynaptically and not only on cholinergic neurons.

However, the development of VAcHT-specific PET radioligands is very challenging as it is based on the only known lead compound vesamicol. Because of the structural similarity of vesamicol to the pharmacophore of sigma receptor ligands,

the design of suitable derivatives has to be directed not only to the VAcHT binding affinity but also to a high selectivity versus sigma receptors. This becomes evident by the numerous vesamicol-based ligands which often show binding affinities even in the nanomolar range in particular to  $\sigma_1$  receptors. Sigma receptors are also distributed in cholinergic brain regions such as the thalamus, hippocampus, or striatum (Bouchard and Quirion 1997) and would therefore impair specific in vivo imaging. However, although structure-affinity relationship studies considering the three competing targets have been performed (Barthel et al. 2015), no clear indicators for constructive structural modifications could be revealed so far as also concluded by Giboureau et al. (2010a). A tendency for high affinity to VAcHT was disclosed, but results for  $\sigma_1/\sigma_2$  receptor binding in dependence on the molecular structure were inconsistent.

In conclusion, numerous PET radioligands for the VAcHT have been developed, but with [ $^{18}\text{F}$ ]NEFA and [ $^{18}\text{F}$ ]FEOBV, only two of them were investigated in humans so far. Just recently, [ $^{18}\text{F}$ ]FEOBV was translated into clinical application for investigation of patients with dementia. Additionally, human studies with the SPECT tracer (–)-[ $^{123}\text{I}$ ]BVM demonstrated the relevance of the VAcHT for imaging cholinergic dysfunction already in the early stage of neurodegenerative diseases. Finally, the first published human data are very promising and give hope for a breakthrough of successful VAcHT PET imaging in the brain and the implementation of this method in diagnosing and understanding cholinergic pathologies as appearing, e.g., in Alzheimer's disease.

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# PET Imaging of Vesicular Monoamine Transporters

# 33

Michael R. Kilbourn

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## Abstract

The vesicular monoamine transporter type 2 (VMAT2) is a protein specifically located in the membrane of neurotransmitter storage vesicles of monoaminergic neurons (dopamine, serotonin, norepinephrine, and histamine) and was considered as a potential marker for noninvasive *in vivo* imaging of changes in monoaminergic innervation in human neurological diseases. Development efforts towards suitably radiolabeled compounds for *in vivo* imaging using positron emission tomography (PET) yielded optimized carbon-11 ((+)- $\alpha$ -[<sup>11</sup>C]dihydro-tetrabenazine, DTBZ) and fluorine-18 (9-O-(3-[<sup>18</sup>F]fluoropropyl)dihydro-tetrabenazine, AV-133) radioligands. Applications of VMAT2 radioligand imaging in animal models (mouse, rat, pig and nonhuman primate) and human studies of numerous neurodegenerative diseases (e.g., Parkinson's and related movement disorders, dementia with Lewy bodies), psychiatric diseases (e.g., schizophrenia), and drug abuse (cocaine, methamphetamine) have demonstrated that *in vivo* imaging of the VMAT2 provides a powerful tool to investigate changes of monoaminergic innervation in human disease conditions.

## Abbreviations

AD	Alzheimer's disease
AMPH	Amphetamine
AMPT	Alpha-methyl-para-tyrosine
AV-133	9-(3-Fluoropropyl)-10-methoxy-3-(2-methylpropyl)-1,3,4,6,7,11b-hexahydrobenzo[a]quinolizin-2-ol
COC	Cocaine
DASB	3-Amino-4-(2-dimethylaminomethylphenylsulfanyl)-benzonitrile
DLB	Dementia with Lewy bodies
DOPA	3,4-Dihydroxyphenylalanine
DTBZ	Dihydro-tetrabenazine: 9,10-dimethoxy-3-(2-methylpropyl)-1,3,4,6,7,11b-hexahydrobenzo[a]quinolizin-2-ol
FDG	2-Fluoro-2-deoxy-D-glucose
FluoroDOPA	6-Fluoro-3,4-dihydroxyphenylalanine
FPBM	2-(2'-((Dimethylamino)methyl)-4'-(3-fluoropropoxy)phenylthio)benzenamine
FTD	Frontotemporal dementia

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HD	Huntington's disease
HED	Hydroxyephedrine
IBVM	Iodobenzovesamicol
MDMA	3,4-Methylenedioxy- <i>N</i> -methylamphetamine
METH	Methamphetamine
MPH	Methylphenidate
MPP+	1-Methyl-4-phenylpyridinium
MPTP	1-Methyl-4-phenyl-2,3,5,6-tetrahydropyridine
MSA	Multiple system atrophy
MTBZ	Methoxytetraabenazine
OPCA	Olivopontocerebellar atrophy
PD	Parkinson's disease
PDD	Parkinson's disease with dementia
PET	Positron emission tomography
PIB	Pittsburgh compound B
PMP	<i>N</i> -methylpiperidiny propionate
PSP	Progressive supranuclear palsy
RAC	Raclopride
SCZ	Schizophrenia
SPECT	Single photon emission computed tomography
TBZ	Tetraabenazine: 3-isobutyl-9,10-dimethoxy-1,3,4,6,7,11b-hexahydro-pyrido[2,1- <i>a</i> ]isoquinolin-2-one
TS	Tourette's syndrome
VACHT	Vesicular acetylcholine transporter
VGAT	Vesicular GABA transporter
VGLUT	Vesicular glutamate transporter
VMAT	Vesicular monoamine transporter

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### 33.1 Introduction

The simplified view of neurotransmission in the brain involves the biosynthesis of neurotransmitters, transporter-mediated uptake, and concentration of the neurotransmitter molecules within vesicles located in presynaptic terminals, stimulated release of the vesicular contents through calcium-induced fusion of synaptic vesicles with the neuronal membrane and exocytotic release of neurotransmitters into the synapse and binding of the released neurotransmitters to post-synaptic receptors. The removal of the neurotransmitter from the synapse can result from transporter-mediated reuptake of neurotransmitter back across the neuronal membrane into the cytosol, intra-synaptic catabolism, or simply diffusion out of the synaptic space. Those neurotransmitter molecules recovered from the synapse and transported back into the presynaptic terminal can then be metabolically degraded by cytosolic enzymes or repackaged into vesicles for future release. Radiopharmaceutical chemists have targeted every one of these steps for



development of in vivo radioligands (Zimmer and Luxen 2012): in the dopaminergic system, for example, this has produced useful radiotracers for neurotransmitter synthesis (e.g., [ $^{18}\text{F}$ ]fluoroDOPA), post-synaptic receptor binding (e.g., [ $^{11}\text{C}$ ]raclopride), presynaptic reuptake (e.g., [ $^{11}\text{C}$ ]methylphenidate), and catabolism (e.g., [ $^{11}\text{C}$ ]deprenyl) (Brooks and Pavese 2011). The last target for PET radiotracer development has been the vesicular neurotransmitter transporters.

As a critical part of this process of neurotransmission, the vesicular neurotransmitter transporters are synaptic vesicle proteins that actively transport newly synthesized or recovered cytosolic neurotransmitter molecules across the vesicle lipid bilayer and into the lumen. Separate transporters are found for each of the major classical neurotransmitters: VMAT1 (*SLC18A1*) and VMAT2 (*SLC18A2*) for the monoamines, VAcHT (*SLC18A3*) for acetylcholine, VGAT (*SLC32A1*) for inhibitory amino acids, and VGLUT (*SLC17A6*, *SLC17A7*, *SLC17A8*) for the vesicular glutamate transporters (Eiden et al. 2004; Gasnier 2004; Omote et al. 2011). These vesicular transporters are located in the membranes of vesicles located in presynaptic terminals and have been targeted as potential biomarkers of specific neuronal populations (although the possibility of neurons expressing more than one vesicular transporter cannot be discounted (Munster-Wandowski et al. 2016; Silm et al. 2019)). As such, vesicular transporter targets complement the more extensive efforts expended in radioligand development for the neuronal membrane neurotransmitter transporters (Brooks and Pavese 2011), a structurally unrelated family of proteins responsible for the reuptake of intra-synaptic neurotransmitters across the synaptic membrane and into the cytosol.

Only two vesicular neurotransmitter transporters, the VMAT2 and VAcHT, have been successfully targeted for development of in vivo radioligands for human studies. The development and applications of VMAT2 radioligands are described in the following sections. In contrast to VMAT2 radiotracer development, a wider variety of compounds have been radiolabeled as prospective in vivo imaging agents for the VAcHT, but only two ([ $^{123}\text{I}$ ]IBVM for single photon emission computed tomography, [ $^{18}\text{F}$ ]FEOBV for positron emission tomography) have reached the stage of achieving approvals for human studies (Kuhl et al. 1994; Petrou et al. 2012a). At present the development of in vivo radioligands for the VGAT and VGLUT transporters is severely limited by a lack of structures known to be high affinity inhibitors that might serve as templates for radiotracer design (Thompson et al. 2005).

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## 33.2 Biology and Pharmacology of the Vesicular Monoamine Transporter Type 2

### 33.2.1 Molecular Biology of the VMAT2

The VMAT2 is part of the family of amine transporters (Eiden et al. 2004) and transports single positively charged amine substrates into vesicles with a concomitant

extrusion of two protons. The accumulation of monoamines is thus dependent on a pH gradient maintained by the action of a separate membrane-bound ATPase. A crystal structure of the VMAT2 has not yet been achieved, but homology modeling has suggested a 12-transmembrane domain configuration with both the amino and carboxyl terminal ends within the cytoplasm and a large intracellular loop between transmembrane domains I and II (Wimalasena 2010; Yaffe et al. 2018). The structure of VMAT2 is highly conserved across mammalian species (Schafer et al. 2013). VMAT2 is the form found in the neurons of the brain, although it is also expressed in nerves and exocrine cells in a variety of peripheral organs. The structurally related VMAT1 (with 40% sequence homology) is predominantly located in organs outside the brain with highest concentrations in the adrenals, although recent studies have suggested the presence of VMAT1 in the mammalian brain (Lohoff et al. 1712). A functional monoaminergic neuron consists of the VMAT2 co-expressed with the necessary neurotransmitter-specific biosynthetic enzymes and corresponding neuronal membrane reuptake transporter (e.g., dopaminergic neurons have tyrosine hydroxylase, DOPA decarboxylase, and the dopamine transporter (DAT)). Homozygous knockout of the VMAT2 is lethal in mice, but heterozygous knockouts survive with reduced filling of neuronal vesicles (Miller et al. 2001). Genetic variants in the human *SLC18A2* gene are rare and produce a severe infantile parkinsonian disorder termed brain dopamine-serotonin vesicular transport disease (Jacobsen et al. 2016).

### 33.2.2 Localization of the VMAT2 in Mammalian Brain

In vitro studies of rodent and human brain tissues using the VMAT2 ligand [<sup>3</sup>H]dihydro-tetrabenazine ([<sup>3</sup>H]DTBZ, originally termed [<sup>3</sup>H]TBZOH) have demonstrated the concentration of the transporter in brain regions rich in monoaminergic neuronal synapses, with the highest concentrations in the basal ganglia, intermediate in such cell bodies as the substantia nigra, and low in the cortex (Scherman and Raisman 1988) (Table 33.1). Radioligand binding in the striatum is predominantly (>95%) to dopaminergic terminals; in other brain regions binding is likely due to a mixture of dopaminergic, serotonergic, and adrenergic innervation. The high concentration of VMAT2 in the basal ganglia has also been demonstrated using immunochemical techniques and a specific hVMAT2 antibody (Tong et al. 2011). Postmortem studies in human brains using radioligand binding or immunochemical staining techniques have demonstrated the significant loss of VMAT2 in the basal ganglia in neurodegenerative diseases such as Parkinson's disease (Scherman et al. 1989; Miller et al. 1998). The extremely low levels of VMAT2 in regions such as the occipital cortex or cerebellar cortex have proven advantageous for in vivo imaging, as these regions can then be used to estimate nonspecific distribution of radioligands for pharmacokinetic modeling purposes.

**Table 33.1** Regional in vitro and in vivo binding data for VMAT2 radioligands in mouse, rat, and human brains ( $DV_{total}$  = total distribution volume)

<i>Mouse</i>	<i>fmol/mg protein (Scherman and Raisman 1988)</i>	<i>[<sup>11</sup>C]TBZ % inj dose/g (DaSilva and Kilbourn 1992)</i>
Striatum	1345	4.96
Thalamus	111	2.06
Hippocampus	141	2.01
Hypothalamus	411	2.96
Pons-medulla	167	1.96
Cortex (frontal)	115	1.91
Cerebellum	56	1.76
<i>Rat</i>	<i>fmol/mg protein (Vander Borgh et al. 1995a)</i>	<i>[<sup>11</sup>C]DTBZ <math>DV_{total}</math> (Kilbourn et al. 2002)</i>
Striatum	1199	7.35
Thalamus	111–669	2.80
Hippocampus	124	2.51
Hypothalamic nuclei	838	5.13
Pons-medulla	–	–
Cortex (frontal)	103	2.18
Cerebellum	60	2.07
<i>Human</i>	<i>fmol/mg protein (Scherman and Raisman 1988)</i>	<i>[<sup>11</sup>C]DTBZ <math>DV_{total}</math> (Vander Borgh et al. 1995b)</i>
Caudate	742	12.5
Putamen	766	11.7
Hypothalamus	245	–
Hippocampus	–3	–
Thalamus	–	4.75
Cortex (frontal)	–7	4.16
Cerebellum	–	4.06

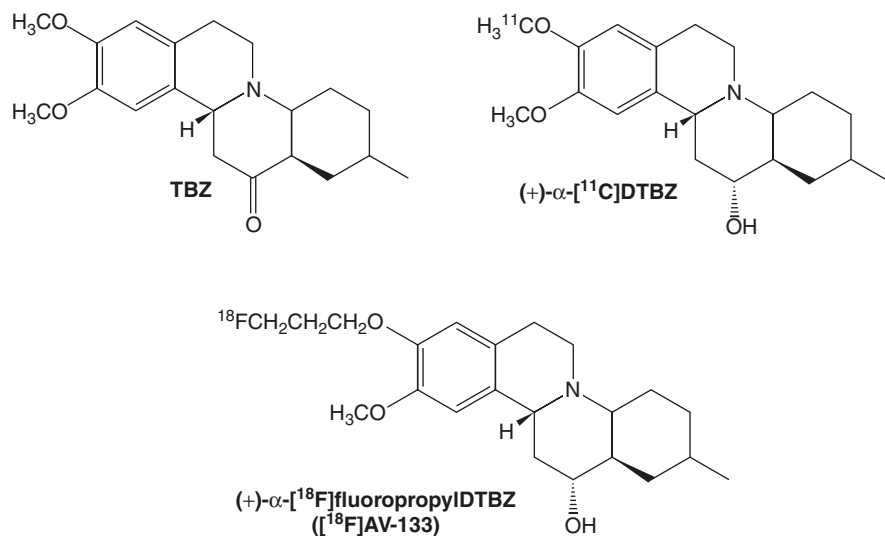
### 33.2.3 VMAT2 Substrates and Inhibitors

The VMAT2 transports a wide variety of molecules, including not only the classical neurotransmitters DA, NE, and 5-HT but also many biogenic amines, drugs of abuse (e.g., amphetamines), and neurotoxins (e.g., 1-methyl-4-phenylpyridinium, MPP+). Substrates for the VMAT2 have typical low binding affinities (>100  $\mu$ M) but high rates of transport. The wide variety of compounds known to be antagonists for the receptors and neuronal membrane transporters for the monoamines are ineffective inhibitors of either VMAT2 transport function or high affinity radioligand binding to the VMAT2 (Wimalasena 2010; Gonzalez et al. 1994).

When development of in vivo VMAT2 radioligands was begun, the list of known medium to high affinity ligands (agonist or antagonist) for the VMAT2 was limited to reserpine, tetrabenazine, and ketanserin, and decades later this situation has not appreciably changed. Reserpine is a large (MW = 608) complex alkaloid with an extremely high affinity for VMAT2 ( $K_D$  = 0.3 nM) and is essentially an irreversible inhibitor with a protracted effect on brain VMAT2 (Naudon et al. 1995): those

properties made it a poor candidate for in vivo radioligand development. Ketanserin has a moderate affinity for the VMAT2 ( $K_D = 45$  nM) but also high affinity for serotonin receptors, making it also a less than optimal target for development of an in vivo imaging agent. Attempts at modification of the ketanserin structure have failed to yield compounds with potential as higher affinity and more specific radioligands (Darchen et al. 1989; Provencher et al. 2018). A series of compounds derived from lobelane (1,4-diphenylalkylpiperidines) were reported to be potent inhibitors of vesicular dopamine uptake, but they have only moderate affinities ( $>200$  nM) for inhibition of [ $^3\text{H}$ ]DTBZ binding (Zheng et al. 2005; Nickell et al. 2016) and did not inhibit in vivo VMAT2 binding in a rat model (Kilbourn et al. 2010). Most recently, a new lead compound GZ-11610 (*R*-3-(4-methoxyphenyl)-*N*-(1-phenylpropan-2-yl)propane-1-amine) was reported as a high affinity ( $K_i = 8.7$  nM) inhibitor of DA uptake by the VMAT2 (Lee et al. 2018), but the potential use as an in vivo radioligand has not been investigated.

Tetrabenazine (TBZ: 3-isobutyl-9,10-dimethoxy-1,3,4,6,7,11b-hexahydro-pyrido[2,1-a]isoquinolin-2-one, Fig. 33.1) is one of a series of benzoisoquinolines initially developed by Hoffman-LaRoche (Quinn et al. 1959). For decades after its clinical introduction, there had been little further examination of the structure-affinity relationships of such benzoisoquinolizines for the VMAT2, but prompted by the interest in developing optimal radioligands for imaging, a larger sets of derivatives of tetrabenazine have been recently synthesized and evaluated in vitro and in vivo. As described in the following sections, these efforts have led to stereospecific syntheses of TBZ and its primary metabolite dihydrotetrabenazine (DTBZ: 9,10-dimethoxy-3-(2-methylpropyl)-1,3,4,6,7,11b-hexahydrobenzo[*a*]quinolizin-2-ol, Fig. 33.1) and



**Fig. 33.1** Chemical structures for tetrabenazine (TBZ) and the two VMAT2 radioligands which have been most used for animal and human PET imaging studies, (+)- $\alpha$ -[ $^{11}\text{C}$ ]DTBZ, and (+)- $\alpha$ -[ $^{18}\text{F}$ ]fluoropropylDTBZ ([ $^{18}\text{F}$ ]AV-133)

the modification of DTBZ with 2-alkyl substituents (Lee et al. 1996), altered 3-alkyl groups (Zheng et al. 2011), and substitutions of longer alkyl or fluoroalkyl chains for the 9-O-methyl substituent (Goswami et al. 2006; Hostetler et al. 2007).

### 33.3 VMAT2 Radioligands for Autoradiography

#### 33.3.1 [<sup>3</sup>H]Tetrabenazine ([<sup>3</sup>H]TBZ)

Tritiated tetrabenazine was only recently reported (Rhee et al. 2011) using the catalytic reduction of a suitably constructed alkene precursor (1,3,4,6,7,11b-hexahydro-3-(2-methyl-2-propenyl)-9,10-dimethoxy-2-oxo-2H-benzo[a]quinolizine) with tritium gas to yield the [<sup>3</sup>H<sub>2</sub>]tetrabenazine with a specific activity of 1850 GBq/mmol.

#### 33.3.2 α-[<sup>3</sup>H]Dihyrotetrabenazine ([<sup>3</sup>H]DTBZ)

The tritiated form of α-dihyrotetrabenazine (originally termed TBZOH) was first reported in 1981 and prepared by the sodium borotritide reduction of the 2-ketone group of TBZ (Scherman et al. 1981). The radioligand was obtained as a mixture of the α- and β-isomers resulting from a non-stereospecific reduction of the ketone, and chromatographic purification likely gave the α-[<sup>3</sup>H]DTBZ (specific activity of 185–555 GBq/mmol) as a racemic mixture of enantiomers. The synthesis of the single isomer (+)-α-[<sup>3</sup>H]DTBZ (2*R*,3*R*,11*bR* configuration) was accomplished by O-methylation of (+)-α-9-O-desmethylDTBZ using [<sup>3</sup>H]methyl iodide, with a specific activity of 2923 GBq/mmol (Kilbourn and Sherman 1997). [<sup>3</sup>H]DTBZ remains the preferred in vitro radioligand for the VMAT2.

#### 33.3.3 (±)-[<sup>3</sup>H]Methoxytetrabenazine ([<sup>3</sup>H]MTBZ)

In an attempt to prepare a radioligand with better long-term radiochemical stability than [<sup>3</sup>H]DTBZ, the derivative 2-(O-[<sup>3</sup>H]methyl)dihyrotetrabenazine ([<sup>3</sup>H]MTBZ, *K*<sub>i</sub> = 3.9 nM) was prepared by alkylation (±)-α-DTBZ with tritiated methyl iodide (Vander Borgh et al. 1995a). This tritiated molecule was considerably more stable and showed less radiolytic decomposition upon storage as compared to tritiated DTBZ.

#### 33.3.4 [<sup>3</sup>H]/[<sup>125</sup>I]Reserpine

Tritiated reserpine (292 Gbq/mmol) has been synthesized from [2,6-<sup>3</sup>H]3,4,5-trimethoxybenzoic acid and used for ex vivo autoradiography in rat brain. A radioiodinated reserpine derivative, 18-O-[3-(3-[<sup>125</sup>I]iodo-4-azidophenyl-propionyl)methyl reserpate, has been used for photoaffinity labeling experiments with the VMAT2 (Sievert et al. 2007).

### 33.3.5 Radioiodinated Tetrabenazines

A radioiodinated derivative of tetrabenazine with an iodovinyl substituent at the 2-position was prepared in very high-specific activities (81,400 GBq/mmol) by an iododestannylation reaction (Kung et al. 1994) and obtained as a mixture of two isomers of which only one showed high affinity binding to the VMAT2 ( $K_D = 0.22$  nM). The stereochemistries of the radioiodinated ligands have not been determined.

More recently, a 9- $^{[131]I}$ iodovinyl-DTBZ derivative was reported to show specific binding to the VMAT2 in rat brain using in vitro autoradiography (Cao et al. 2018). No in vivo studies have been reported.

### 33.3.6 Ketanserin Derivatives

Ketanserin is a moderate affinity ( $K_i = 45$  nM) but nonspecific ligand for the VMAT2 (Darchen et al. 1988), and with appropriate pharmacological blocking of its other binding sites (e.g., serotonin receptors), two radioiodinated derivatives, 7-amino-8- $^{[125]I}$ iodoketanserin and 7-azido-8- $^{[125]I}$ iodoketanserin, have been used for in vitro autoradiography and photoaffinity labeling experiments (Darchen et al. 1989).

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## 33.4 VMAT2 Radioligands for PET Imaging Studies

The development of VMAT2 radioligands for positron emission tomography (PET) began with the practical choice of TBZ as the initial target compound. The TBZ molecule has two phenolic methoxy groups suitable for radiolabeling using carbon-11 methyl iodide, and the resultant radioligand would be unchanged from the drug that had been in human use for decades; this avoided any worries about toxicology as the pharmacology and metabolism of the drug were very well-known.

### 33.4.1 $^{[11C]}$ Tetrabenazine ( $^{[11C]}$ TBZ)

The labeling of tetrabenazine simply required removal of one of the O-methyl groups and replacement with the carbon-11 ( $t_{1/2} = 20$  min) labeled methyl group (DaSilva and Kilbourn 1992; DaSilva et al. 1993a). Although simple in concept, the cleavage of the phenolic methoxy group using reagents such as boron tribromide proved to be a difficult reaction and led to complex mixtures of the mono- and bis-demethylated products. Careful chromatography yielded a single pure product assigned as the 9-O-desmethyltetrabenazine, with the identification of the demethylation site (9 vs. 10) based on proton and carbon-13 NMR spectroscopy; that was later proved correct through a synthesis of 9-O-desmethyltetrabenazine by an independent and unambiguous route. The reaction of the free phenolic compound with  $^{[11C]}$ methyl iodide under basic conditions gave the desired  $^{[11C]}$ tetrabenazine in

modest yields and good specific activities (74000–92,500 GBq/mmol) (DaSilva et al. 1993a). At the time of its preparation and use in humans, [ $^{11}\text{C}$ ]TBZ was obtained as a mixture of stereoisomers; based on later studies of DTBZ isomers (see below), the starting material and the final radiochemical product were both likely racemic mixtures of 3*R*,11*bR* and 3*S*,11*bS* enantiomers.

### 33.4.2 ( $\pm$ )- $\alpha$ -(9-O-[ $^{11}\text{C}$ ]Methyl)Dihydrotetrabenazine ([ $^{11}\text{C}$ ]DTBZ)

The consideration of the metabolic fate of TBZ in the human body led to the rapid realization that what was really desired for an in vivo imaging agent was the carbon-11 labeled DTBZ, which is formed by the action of carbonyl reductase in the liver and is the predominant chemical species in the bloodstream after administration of TBZ to rats or humans (Mehvar et al. 1987). That metabolite (previously termed TBZOH) had been used for years in tritiated form for in vitro studies of the VMAT2 (Scherman and Raisman 1988). The reduction of the ketone of TBZ by in vivo metabolism, or by chemical means in the laboratory, produces two isomers termed  $\alpha$ - and  $\beta$ -DTBZ to denote the configuration of the newly created hydroxyl group. The  $\alpha$ -isomer was identified as having the higher affinity in vitro for the VMAT2. The required precursor for radiolabeling was ( $\pm$ )-9-O-desmethylDTBZ, prepared in two steps from TBZ by reduction of the ketone followed by demethylation (Jewett et al. 1997). Alkylation with [ $^{11}\text{C}$ ]methyl iodide produced ( $\pm$ )- $\alpha$ -(9-O-[ $^{11}\text{C}$ ]methyl)DTBZ in moderate yields and useful specific activities (>59,200 GBq/mmol) but still as a racemic mixture of enantiomers.

### 33.4.3 2-(O-[ $^{11}\text{C}$ ]Methyl)Dihydrotetrabenazine ([ $^{11}\text{C}$ ]MTBZ)

In an attempt to alter and perhaps improve the metabolic disposition of VMAT2 radioligands, a new derivative with a [ $^{11}\text{C}$ ]methoxy substituent at the 2-position was synthesized (DaSilva et al. 1993b). Prepared by reaction of ( $\pm$ )-DTBZ with [ $^{11}\text{C}$ ]methyl iodide under strongly basic conditions, the radioligand 2-(O-[ $^{11}\text{C}$ ]methyl)dihydrotetrabenazine (MTBZ;  $K_i = 3.9$  nM) was successfully introduced into human studies (Vander Borgh et al. 1995b). However, the radiochemical synthesis proved more difficult than the alkylation of the phenol (as done for [ $^{11}\text{C}$ ]DTBZ), and with the development of (+)- $\alpha$  [ $^{11}\text{C}$ ]DTBZ, MTBZ was not utilized further.

### 33.4.4 (+)- $\alpha$ -(9-O-[ $^{11}\text{C}$ ]Methyl)Dihydrotetrabenazine (+)- $\alpha$ -(9-O-[ $^{11}\text{C}$ ]Methyl)Dihydrotetrabenazine ([ $^{11}\text{C}$ ]DTBZ)

The last step in improvement of carbon-11 VMAT2 radiotracers was the development of (+)- $\alpha$ -(9-O-[ $^{11}\text{C}$ ]methyl)DTBZ ((+)- $\alpha$ -[ $^{11}\text{C}$ ]DTBZ, Fig. 33.1). That TBZ and its metabolite DTBZ were mixtures of stereoisomers which had been known for many years (Schwartz et al. 1966), but the isomers had not been isolated, and their

absolute configurations had not been determined. The enantiomers of racemic  $\alpha$ -DTBZ were finally isolated by preparative chiral HPLC (Kilbourn et al. 1995a; Kilbourn et al. 1997) providing a high affinity (+)-isomer ( $K_i = 1$  nM) and low affinity (–)-isomer ( $K_i = 2200$  nM). Crystals of (–)- $\alpha$ -DTBZ were successfully grown for X-ray structural analysis, and the absolute stereochemistry of the isomers finally assigned: (2*R*,3*R*,11*bR*) for the (+)-isomer and (2*S*,3*S*,11*bS*) for the (–)-isomer (Kilbourn et al. 1997; Liu et al. 2012; Liu et al. 2013). The interest in the stereoisomers of DTBZ as imaging agents and potential drugs stimulated numerous new and improved synthetic routes to TBZ and DTBZ (Boldt et al. 2008; Paek et al. 2010; Rishel et al. 2009a) including the preparation and characterization of all eight possible DTBZ stereoisomers (Tridgett et al. 2008; Yu et al. 2010; Yao et al. 2011). Determination of the in vitro binding affinities have consistently shown that the isomer with the 2*R*,3*R*,11*bR* absolute stereochemistry is the highest affinity DTBZ ligand for the VMAT2 (Fig. 33.1).

The synthesis of (+)- $\alpha$ -[<sup>11</sup>C]DTBZ required the preparation of the resolved (+)-isomer of the 9-O-desmethylDTBZ, which was first accomplished by a low-yield sequence using reduction of TBZ, isolation of the racemic  $\alpha$ -isomer, chiral column HPLC resolution, and a modified demethylation reaction condition using sodium hydride/*N*-methylaniline/hexamethylphosphoramide (Kilbourn et al. 1997). The more recent improved synthetic routes to TBZ, (+)- $\alpha$  DTBZ, and (+)- $\alpha$ -9-O-desmethylDTBZ have made synthesis of the precursor much more practical. Alkylation of the desmethyl precursor with [<sup>11</sup>C]methyl iodide (Jewett et al. 1997) or [<sup>11</sup>C]methyl triflate (Zhang et al. 2012) provided a reliable route to preparation of (+)- $\alpha$ -[<sup>11</sup>C]DTBZ in high average specific activities (370,000 GBq/mmol) and a defined 2*R*,3*R*,11*bR* stereochemistry.

### 33.4.5 (+)- $\alpha$ -(10-O-[<sup>11</sup>C]Methyl)Dihydrotrabenzazine

The synthesis and in vivo evaluation of DTBZ labeled in the 10-methoxy position was recently accomplished (Huang et al. 2016). Racemic 10-O-desmethylDTBZ was resolved using fractional crystallization and then reduced to yield the desired (+)- $\alpha$ -(10-desmethyl)DTBZ. Alkylation with [<sup>11</sup>C]methyl iodide formed (+)- $\alpha$ -(10-[<sup>11</sup>C]methyl)DTBZ. MicroPET imaging in rat brains showed specific uptake in the striatum with striatum-to-cerebellum ratios reportedly higher than for the corresponding 9-[<sup>11</sup>C]methoxyDTBZ. No human studies have been reported for this radiotracer.

### 33.4.6 Copper-64 Labeled Dihydrotrabenzazine

In an attempt to develop a VMAT2 radioligand labeled with a longer-lived PET radionuclide, a 2-amino derivative of DTBZ was conjugated with the chelating group 2,2'-(1,4,8,11-tetraazabicyclo[6.6.2]hexadecane-4,11-diyl)diacetic acid



(Kumar et al. 2014). Complexation of copper-64 yielded a radioligand with an in vitro  $IC_{50}$  of 16.8 nM. No in vivo studies of this radioligand have been reported.

### 33.4.7 Fluorine-18 Labeled Dihydrotrabenazines

The application of VMAT2 imaging to widespread clinical care would require the availability of radioligands labeled with longer half-life radionuclides. Fortunately, development of fluorine-18 ( $t_{1/2} = 109.6$  min) labeled VMAT2 radioligands was able to build on the experience developed in the carbon-11 series.

The substitution of the 9-methoxy group of DTBZ with [ $^{18}F$ ]fluoroalkoxy groups resulted in the syntheses of 9-O- [ $^{18}F$ ]fluoromethyl-, 9-O-(2- [ $^{18}F$ ]fluoroethyl)- and 9-O-(3- [ $^{18}F$ ]fluoropropyl)DTBZ. The [ $^{18}F$ ]fluoromethyl derivative ( $K_i = 0.87$ ) was prepared by O-alkylation of (+)- $\alpha$ -9-O-desmethylDTBZ with [ $^{18}F$ ]FCH<sub>2</sub>Br under basic conditions (Hostetler et al. 2007). The [ $^{18}F$ ]fluoroethyl ( $K_i = 0.76$  nM) and [ $^{18}F$ ]fluoropropyl ( $K_i = 0.56$  nM) derivatives were first prepared in racemic form by [ $^{18}F$ ]fluoride ion displacement of the corresponding mesylates in specific activities >74,000 GBq/mmol (Goswami et al. 2006; Kung et al. 2007). Subsequently, stereospecific syntheses of (+)- $\alpha$ -9-O-(3- [ $^{18}F$ ]fluoropropyl)DTBZ (also termed [ $^{18}F$ ]AV-133: (2*R*,3*R*,11*bR*)-9-(3- [ $^{18}F$ ]fluoropropoxy)-10-methoxy-3-(2-methylpropyl)-1,3,4,6,7,11*b*-hexahydro-2*H*-benzo[*a*]quinolizin-2-ol)) from the corresponding chiral mesylate or tosylate were developed (Li et al. 2017), yielding the very high affinity ( $K_i = 0.1$  nM) single isomer with specific activities of 55,000–74,000 GBq/mmol. As was expected, the (–)- $\alpha$ -(3-fluoropropyl)DTBZ isomer showed negligible affinity for VMAT2 (>3000 nM). Recently, attempts were made to reduce in vivo metabolism of the [ $^{18}F$ ]fluoroalkyl DTBZ radioligands using deuterium substitutions on the fluoroalkyl groups, but significant improvements of the radiotracer imaging properties were not achieved (Varone et al. 2012; Liu et al. 2018).

As of the start of 2019, the only fluorine-18 radioligand that had reached human studies was the [ $^{18}F$ ]fluoropropyl derivative [ $^{18}F$ ]AV-133. A variety of other compounds had, however, been synthesized and radiolabeled with fluorine-18. Replacement of the 2-hydroxyl group of DTBZ with a 2-fluoropropylamino group yielded a derivative with an in vitro  $K_i$  of 7 nM (as racemate) (Zhu et al. 2009). Substitution of the 2-hydroxyl function of 2- [ $^{18}F$ ]fluoropropyl-(+)-DTBZ with an epoxide ring provided a derivative with an excellent in vitro affinity ( $K_i = 0.08 \pm 0.01$  nM) (Kung et al. 2008). Attempts to change the 3-alkyl group resulted in reduced in vitro binding affinities (Zheng et al. 2011). A completely different approach to fluorine-18 labeled DTBZ molecules was reported in a series of patents (Kankanamalage et al. 2009; Rishel et al. 2009; Amarasinghe et al. 2009) that claimed a large number of fluorine-substituted (+)- $\alpha$ -DTBZ derivatives, but in vitro biological information is only available for a few of the compounds claimed: the 2-fluoroethyl substituted DTBZ derivative showed reasonable affinity

( $K_i = 19$  nM), consistent with the earlier demonstration of bulk tolerance of reasonably small 2-alkyl substituents (Lee et al. 1996), and the 2-fluoroethoxy ether derivative had a good in vitro binding affinity of 3.2 nM, which was not significantly different than that previously obtained for the slightly smaller methyl ether derivative (Vander Borgh et al. 1995a).

### 33.5 Evaluation of VMAT2 Imaging Radioligands in Animals

The evaluation of VMAT2 radioligands for pharmacological specificity, kinetics, sensitivity to endogenous neurotransmitters, propensity for up- or downregulation, and changes in models of human disease has been extensively pursued in both small and large animal models (Table 33.2).

#### 33.5.1 Mouse Brain VMAT2 Studies

The mouse was utilized for the initial evaluation of [ $^{11}\text{C}$ ]TBZ, where it was demonstrated that (a) the radioligand localized in regions of the brain containing high concentrations of the VMAT2, (b) localization was likely not specific only to dopaminergic neurons but, as expected, also due to noradrenergic and serotonergic innervation (Table 33.1), and (c) binding could be blocked by pharmacological doses of TBZ but not drugs that bound to the neuronal membrane dopamine transporter (DaSilva and Kilbourn 1992).

Subsequent studies (Table 33.2) of VMAT2 radioligands ([ $^{11}\text{C}$ ]DTBZ, [ $^{11}\text{C}$ ]MTBZ, and [ $^{18}\text{F}$ ]AV-133) binding in the mouse brain demonstrated that simple

**Table 33.2** Animal studies using VMAT2 radioligands

Species	Radioligands	Topic	Ref.
Mouse	[ $^{11}\text{C}$ ]TBZ	First in vivo studies	DaSilva and Kilbourn (1992)
	[ $^{11}\text{C}$ ]TBZ	Brain distribution in vivo, metabolism, dosimetry	DaSilva et al. (1994)
	[ $^{11}\text{C}$ ]MTBZ	Study in <i>tottering</i> mice	Kilbourn et al. (1995b)
	(+)- and (-)-[ $^{11}\text{C}$ ]DTBZ	Stereospecificity of in vivo binding	Kilbourn et al. (1995a)
	( $\pm$ )-[ $^{11}\text{C}$ ]MTBZ	Binding in different mouse strains	Kilbourn and Frey (1996)
	( $\pm$ )-[ $^{11}\text{C}$ ]MTBZ	Evaluation of DTBZ derivatives	Lee et al. (1996)
	( $\pm$ )-[ $^{11}\text{C}$ ]MTBZ	Binding site recovery after TBZ dose	Kilbourn (1997)
	(+)-[ $^{11}\text{C}$ ]DTBZ, [ $^{11}\text{C}$ ]MPH	Differential VMAT2, DAT binding in MPTP mice	Kilbourn et al. (2000)
	[ $^{18}\text{F}$ ]FE-DTBZ, [ $^{18}\text{F}$ ]FP-DTBZ	Evaluation of new fluoroalkyl DTBZ derivatives	Goswami et al. (2006)
	[ $^{18}\text{F}$ ]AV-133	PET and in vitro autoradiography in MPTP mice	Chao et al. (2012)
	[ $^{18}\text{F}$ ]AV-133	PET imaging of MPTP mice	Toomey et al. (2012)
	[ $^{18}\text{F}$ ]AV-133	MPTP mice treated with magnolol	Weng et al. (2017a)
	(+)-[ $^{11}\text{C}$ ]DTBZ	Effect of methcathinone	Asser et al. (1652)

(continued)

**Table 33.2** (continued)

Species	Radioligands	Topic	Ref.
Rat	[ <sup>125</sup> I]Iodoviny]TBZ	In vitro autoradiography	Kung et al. (1994)
	[ <sup>3</sup> H]MTBZ	Autoradiography: Effects of drug treatments	Vander Borgh et al. (1995c)
	[ <sup>3</sup> H]MTBZ	Autoradiography in 6-OHDA lesioning	Vander Borgh et al. (1995a)
	(±)-[ <sup>11</sup> C]DTBZ	Ex vivo studies: DA-ergic drug treatments	Kilbourn et al. (1996)
	[ <sup>3</sup> H]DTBZ	Bolus vs. equilibrium rat studies	Kilbourn and Sherman (1997)
	[ <sup>3</sup> H]DTBZ	Autoradiography in METH-treated rats	Frey et al. (1997)
	[ <sup>3</sup> H]MTBZ	In vitro studies: Effects of DA-ergic drug treatments	Kemmerer et al. (2003)
	(+)-[ <sup>3</sup> H]DTBZ, [ <sup>3</sup> H]MPH	Equilibrium infusion technique	Kilbourn et al. (2002)
	(+)-[ <sup>3</sup> H]DTBZ, (+)-[ <sup>11</sup> C]DTBZ	Long-term reproducibility ex vivo	Kilbourn (2004)
	(±)-[ <sup>11</sup> C]DTBZ	6-OHDA treated rats	Strome et al. (2000)
	[ <sup>18</sup> F]FP-DTBZ, (+)-[ <sup>11</sup> C]DTBZ	microPET imaging: Pharmacokinetics	Kilbourn et al. (2007)
	[ <sup>18</sup> F]FP-DTBZ (AV-133)	First brain uptake studies	Kung et al. (2007)
	(+)-[ <sup>11</sup> C]DTBZ	Unilateral 6-OHDA reproducibility	Sossi et al. (2007)
	(±)-[ <sup>11</sup> C]DTBZ	Effects of AMPT, AMPH	Tong et al. (2008)
	(+)-[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]MPH	Sensitivity to DA levels	Kilbourn et al. (2010a)
	(+)-[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]MPH, [ <sup>11</sup> C]RAC	Unilateral 6-OHDA: Effects of pramipexole and levodopa	Sossi et al. (2010a)
	[ <sup>18</sup> F]AV-133, [ <sup>18</sup> F]FPBM	Study of VMAT2 and SERT in 6-OHDA lesioned rat	Wang et al. (2010)
	(+)-[ <sup>11</sup> C]DTBZ	Competition by lobelane derivative	Kilbourn et al. (2010)
	(±)-[ <sup>11</sup> C]DTBZ	Intranigral lactacystin	Mackey et al. (2013)
	(±)-[ <sup>11</sup> C]DTBZ, [ <sup>19</sup> F]FDOPA	Longitudinal studies in unilateral 6-OHDA rats	Walker et al. (2013a)
	(±)-[ <sup>11</sup> C]DTBZ, [ <sup>19</sup> F]FDOPA	PET and DA microdialysis in 6-OHDA rats	Walker et al. (2013b)
	[ <sup>18</sup> F]AV-133, [ <sup>19</sup> F]AV-133	Compare LC-MS/MS for cold ligand vs. in vivo [ <sup>18</sup> F]ligand	Deng et al. (2014)
	(±)-[ <sup>11</sup> C]DTBZ, [ <sup>18</sup> F]FDOPA, [ <sup>11</sup> C]RAC	PET imaging normal rats	Lara-Camacho et al. (2014)
	10-[ <sup>11</sup> C]methoxy-DTBZ	PET in 6-OHDA rats	Li et al. (2014)
	(±)-[ <sup>11</sup> C]DTBZ	PET in over-expression of α-synuclein	Phan et al. (2017)
	[ <sup>18</sup> F]AV-133	Effect of impurity on brain binding	Wu et al. (2015)
	(±)-[ <sup>11</sup> C]DTBZ	Uni- and bilateral 6-OHDA lesions PET	Molinet-Dronda et al. (2015)
	[ <sup>18</sup> F]FP-DTBZ	Effects of anesthesia on imaging	Chen et al. (2016)
	10-( [ <sup>11</sup> C]methoxy)-DTBZ	In vivo comparison with 9-( [ <sup>11</sup> C]methoxy)DTBZ	Huang et al. (2016)
	[ <sup>18</sup> F]FP-DTBZ	PET after unilateral lactacystin injection	Weng et al. (2017b)
	(±)-[ <sup>11</sup> C]DTBZ	6-OHDA and lactacystin lesions	Mejias et al. (2016)
	(±)-[ <sup>11</sup> C]DTBZ	Test-retest reproducibility	Avendaño-Estrada and Ávila-Rodríguez (2018)
d <sub>6</sub> -[ <sup>18</sup> F]FP-DTBZ	Effects of deuteration on in vivo binding and metabolism	Liu et al. (2018)	
[ <sup>18</sup> F]FP-DTBZ	Longitudinal PET study	Jiang et al. (2018)	

**Table 33.2** (continued)

Species	Radioligands	Topic	Ref.
Minipigs	(+)-[ <sup>11</sup> C]DTBZ	Before and after lactacystin treatment	Lillethorup et al. (2018a)
	(+)-[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]PK11195	Over-expression of $\alpha$ -synuclein	Lillethorup et al. (2018b)
	(+)-[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]yohimbine, [ <sup>11</sup> C]DASB, [ <sup>11</sup> C]PK11195	Chronic lactacystin	Lillethorup et al. (2018c)
Guinea pigs	[ <sup>18</sup> F]FP-DTBZ	Retina imaging and experimental myopia	Sun et al. (2018)
Primate	( $\pm$ )-[ <sup>11</sup> C]TBZ	Normal and unilateral MPTP lesion	DaSilva et al. (1993c)
	[ <sup>3</sup> H]DTBZ	In vitro after METH in baboons	Villemagne et al. (1998)
	( $\pm$ )-[ <sup>11</sup> C]DTBZ	PET in METH treated monkeys	Fantegrossi et al. (2004)
	(+)-[ <sup>11</sup> C]DTBZ, [ <sup>18</sup> F]FDOPA	Normal and MPTP lesioned	Doudet et al. (2006)
	(+)-[ <sup>11</sup> C]DTBZ, ( $\pm$ )-FP-DTBZ, (+)-FP-DTBZ	Comparison of pharmacokinetics via PET	Kilbourn et al. (2007)
	(+)-[ <sup>11</sup> C]DTBZ	Dosimetry study in baboon	Murthy et al. (2008)
	(+)-[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]RAC, [ <sup>11</sup> C]PK11195, [ <sup>11</sup> C]WIN35428	Baboon chronic MPTP study	Chen et al. (2008)
	(+)-[ <sup>11</sup> C]DTBZ, [ <sup>18</sup> F]FDOPA	Normal and MPTP lesioned	Collantes et al. (2008)
	(+)-[ <sup>11</sup> C]DTBZ, [ <sup>18</sup> F]FDOPA	MPTP treated	Blesa et al. (2010)
	(+)-[ <sup>11</sup> C]DTBZ, [ <sup>18</sup> F]FDOPA	Slow MPTP intoxication	Blesa et al. (2012)
	(+)-[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]CFT	In vitro studies—Unilateral MPTP	Tian et al. (2012)
	(+)-[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]CFT, [ <sup>18</sup> F]FDOPA	MPTP monkey and study of apathy	Brown et al. (2012)
	(+)-[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]CFT, [ <sup>18</sup> F]FDOPA	MPTP monkey and in vitro nigral cell counts	Karimi et al. (2013b)
	(+)-[ <sup>11</sup> C]DTBZ, [ <sup>18</sup> F]FDOPA	MPTP monkey the carboxyfullerene treatment	Dugan et al. (2014)
	[ <sup>18</sup> F]AV-133	Progressive losses in MPTP monkey	Liu et al. (2014)
	(+)-[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]CFT, [ <sup>18</sup> F]FDOPA	MPTP monkey and in vitro biochemistry	Tian et al. (2014)
	(+)-[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]CFT, [ <sup>18</sup> F]FDOPA	MPTP monkey, various doses, apathy study	Tian et al. (2015)
	(+)-[ <sup>11</sup> C]DTBZ	Changes in chronic cocaine self-administration	Narendran et al. (2015)
	(+)-[ <sup>11</sup> C]DTBZ	Long-term reproducibility of PET measures	Kilbourn and Koeppe (2017)

measurement of tissue ratios between high (striatum) and low (cerebellum) regions of the mouse brain could be used to demonstrate the stereospecificity (Kilbourn et al. 1995a) and reversibility of [<sup>11</sup>C]DTBZ binding to the VMAT2 [37]; measure decreases of VMAT2 binding sites after MPTP (1-methyl-4-phenyl-1,2,3,4-tetrahydropyridine) lesioning (Kilbourn et al. 2000; Chao et al. 2012; Toomey et al. 2012; Weng et al. 2017a) and shows increased in vivo binding in *tottering* mutant mice known to over-express the VMAT2 (Kilbourn et al. 1995b). In vivo measure of VMAT2 binding in MPTP-treated mice can also be used to evaluate potential therapeutic drug approaches (Weng et al. 2017a).

### 33.5.2 Rat Brain VMAT2 Studies

A larger number of studies of VMAT2 radioligands have been conducted in rats (for refs. see Table 33.2). These have utilized a variety of techniques, including (a) *ex vivo* measurements of tissue distributions following bolus injections, (b) *ex vivo* autoradiography of rat brain slices, (c) *ex vivo* determination of regional brain distributions after equilibrium infusions (Kilbourn et al. 2002), and (d) *in vivo* PET imaging studies including dynamic multi-tracer approaches (Sossi et al. 2009). Such studies consistently demonstrated the expected regional distribution of VMAT2 binding sites in the rat brain (Table 33.1), and studies of both the active and inactive stereoisomers of [<sup>11</sup>C]DTBZ have shown that uptake and retention of VMAT2 radioligands in regions of the rat brain such as the cerebellum and the occipital cortex represents essentially negligible concentrations of the VMAT2 (Kilbourn et al. 2002), and thus those brain regions are suitable for estimations of nonspecific distributions of these radiotracers.

The sensitivity of measures of VMAT2 radioligand binding to regulation by chronic drug use has been examined both *in vitro* and *in vivo*, with decidedly mixed results. Several studies have demonstrated that chronic administration of such drugs as L-DOPA/benserazide, apomorphine, tetrabenazine, pramipexole, paroxetine, amphetamine, or cocaine did not alter *in vitro* levels of VMAT2 radioligand binding, nor was *in vivo* DTBZ binding altered by repeated administration of L-DOPA, deprenyl, or tetrabenazine (Kemmerer et al. 2003; Kilbourn et al. 1996; Schwartz et al. 2006; Vander Borght et al. 1995c; Vilpoux et al. 2000; Wilson and Kish 1996). These results supported the hypothesis that VMAT2 was a less regulated site than, for example, the neuronal membrane dopamine transporter for which changes could be seen following such drug treatments (Vander Borght et al. 1995c). In contrast, other studies have reported increases or decreases of VMAT2 radioligand binding after administration of some of the same or related drug substances (Schwartz et al. 2006; Frey et al. 1997; Sossi et al. 2010a). Drug effects can be dose-dependent, as seen for methamphetamine where reductions of VMAT2 binding are observed after neurotoxic but not lower behaviorally effective doses (Frey et al. 1997). For methcathinone (an analog of methamphetamine), an increase in VMAT2 radioligand binding was reported; that was attributed to a decrease of tissue dopamine levels and less competition of the neurotransmitter for the TBZ binding site on the VMAT2 (Asser et al. 1652). That was consistent with a few earlier studies in rats that supported a sensitivity of *in vivo* VMAT2 radioligand binding to drugs which directly decrease endogenous dopamine levels (e.g.,  $\alpha$ -methyl-p-tyrosine) (Kilbourn et al. 2008; Kilbourn et al. 2010a; Tong et al. 2008) or indirectly raise dopamine levels (pramipexole) (Sossi et al. 2010a). Those studies have indicated that alterations of vesicular dopamine concentrations may affect *in vivo* VMAT2 radioligand binding to the TBZ site located on the lumen surface of the vesicle, although it is noted that modest changes of *in vivo* radioligand binding (36% increase) required significant depletion of endogenous dopamine concentrations (>75%) in the rat brain.

To demonstrate losses of VMAT2 radioligand ([<sup>11</sup>C]DTBZ or [<sup>18</sup>F]AV-133) binding in rat models of neurodegeneration, investigators have widely used the

unilateral injection of 6-hydroxydopamine directly into the medial forebrain bundle (MFB: unilateral lesion) or the third ventricle (bilateral lesion) (for refs. see Table 33.2). This technique can produce a wide range of striatal lesions, from mild to severe, with a corresponding range of losses of VMAT2 radioligand binding. The decreased VMAT2 radioligand binding correlates with estimates of denervation from behavioral studies or in vitro autoradiographic studies of VMAT2 or DAT ligand binding. Using dynamic microPET imaging, the losses of radioligand binding were shown to be due to changes in numbers of VMAT2 binding sites ( $B_{\max}$ ) and not the apparent binding affinity ( $K_d^{\text{app}}$ ) (Sossi et al. 2007). Used in combination with microPET studies with 6- $^{18}\text{F}$ fluoroDOPA, the longitudinal changes in DA synthesis, and turnover and estimates of the compensatory changes in dopamine turnover (synthesis + vesicular trapping) after 6-OHDA lesions have been made (Walker et al. 2013a; Walker et al. 2013b).

Alternative models of dopaminergic degeneration in the rat brain have also been examined using VMAT2 radioligands. The stereotactic unilateral injection of lactacystin (a ubiquitin proteasome system inhibitor) into the substantia nigra produced dose-dependent decreases in VMAT2 radioligand binding (PET using  $^{11}\text{C}$ ]DTBZ or  $^{18}\text{F}$ ]AV-133) in the ipsilateral striatum, with losses of tyrosine hydroxylase immunoreactive cells (Mackey et al. 2013; Weng et al. 2017b). As a model of striatal dopaminergic terminal loss without nigral cell loss, the local unilateral injection of recombinant adeno-associated viral vectors coding for human  $\alpha$ -synuclein produces  $\alpha$ -synuclein deposits, motor impairments, and significant reductions in VMAT2 radioligand binding demonstrated both in vitro ( $^3\text{H}$ ]DTBZ autoradiography) and in vivo ( $^{11}\text{C}$ ]DTBZ/PET) (Phan et al. 2017).

### 33.5.3 VMAT2 Radioligand Studies in Non-human Primates

A comprehensive autoradiographic study of the in vitro distribution of VMAT2 radioligand binding sites in the non-human primate brain is still missing from the literature, but specific binding in the caudate and putamen has been demonstrated (Chen et al. 2008; Tian et al. 2012; Blesa et al. 2012). The in vivo distribution of radiotracer binding would be expected to be consistent with the general distribution in mammalian brain as no significant species differences in brain VMAT2 distributions have been noted (Schafer et al. 2013). In contrast to the use of VMAT2 radioligands in rodent brain, all primate studies using PET radioligands have utilized in vivo imaging techniques (for refs. see Table 33.2). Such PET studies in normal primate brain have shown high binding in the striatal regions, with intermediate levels of radioactivity retention in the thalamus, hypothalamus, and midbrain cell bodies (raphe nucleus, locus ceruleus, substantia nigra), and negligible binding in cortex and cerebellum. The binding of  $^{11}\text{C}$ ]DTBZ and  $^{18}\text{F}$ ]AV-133 is stereospecific for the (+)-isomer (2*R*,3*R*,11*bR*-configuration) with low and uniform brain distribution of the corresponding inactive (–)-isomers (Kilbourn et al., unpublished data). A direct comparison of (+)- $^{11}\text{C}$ ]DTBZ and  $^{18}\text{F}$ ]AV-133 demonstrated a higher striatal value for the distribution volume ratio (DVR, a measure of the specific

binding) for the fluorinated radioligand. A longitudinal study of (+)-[<sup>11</sup>C]DTBZ binding in monkey brain has shown good reproducibility with little age effect over a 6-year span (8- to 14-year-old rhesus monkey) (Kilbourn and Koeppe 2017), although an age-dependent reduction of [<sup>11</sup>C]DTBZ was noted by others when the comparison is made between young (8-year old) and older (22-year old) monkeys (Doudet et al. 2006).

To study dopaminergic degeneration in the monkey brain, investigators have uniformly utilized the MPTP lesion model. The initial demonstration of complete loss of VMAT2 radioligand binding in a monkey with a severe unilateral MPTP-induced lesion (DaSilva et al. 1993c) has been followed by numerous investigations in monkeys (Table 33.2) with either unilateral or bilateral lesions. Several studies have employed multi-tracer PET studies combining a VMAT2 radioligand ([<sup>11</sup>C]DTBZ or [<sup>18</sup>F]AV-133) together with [<sup>18</sup>F]fluoroDOPA (a measure of dopamine synthesis and storage) and a DAT radioligand (e.g., [<sup>11</sup>C]CFT), with some studies including postmortem tissue analyses (VMAT2 and DAT autoradiography, TH histochemistry, stereological cell counts in nigra: for references see Table 33.2). All studies have shown decreases of VMAT2 radioligand binding in MPTP-treated animals. The extent of the decreased VMAT2 binding correlates with the severity of the lesions, as indicated by losses of striatal dopamine (Karimi et al. 2013a) as well as changes in DAT radioligand binding or [<sup>18</sup>F]fluoroDOPA uptake and retention. In several studies, there is evidence to support that the *in vivo* measures with VMAT2 radioligands provided an earlier indication of loss of striatal dopaminergic terminals, as compared to DAT radioligands or [<sup>18</sup>F]fluoroDOPA uptake (Chen et al. 2008; Blesa et al. 2012; Doudet et al. 2006; Blesa et al. 2010).

The MPTP monkey model replicates most of the biochemical and behavioral changes of human PD, and the dosing of nonhuman primates with MPTP (Masilamoni and Smith 2018) provides a readily accessible and useful model to evaluate *in vivo* imaging radiotracers. The MPTP model provides a primate option for evaluation of potential pharmacological interventions to prevent or ameliorate dopaminergic degeneration, where PET imaging of the VMAT2 binding can serve as a biomarker (Dugan et al. 2014).

The MPTP intoxication model is thus widely used, but the potential of drugs of abuse to produce dopaminergic terminal losses has also been of interest. A single study of the potential changes of *in vivo* [<sup>11</sup>C]DTBZ binding in a primate model of self-administration of methylenedioxymethamphetamine (MDMA) showed no evidence of dopaminergic degeneration in the striatum (Fantegrossi et al. 2004). In contrast, a postmortem study of amphetamine-treated baboons showed evidence for reduced binding of [<sup>3</sup>H]DTBZ to the VMAT2 (and reduced radioligand binding to the DAT) (Riquarte et al. 2005). A PET imaging study of [<sup>11</sup>C]DTBZ binding in monkeys after 16 months of cocaine self-administration demonstrated a significant (−25%) loss of striatal VMAT2 binding sites (Narendran et al. 2015): it was not determined if this was due to a downregulation of the VMAT2 or loss of dopaminergic terminals.

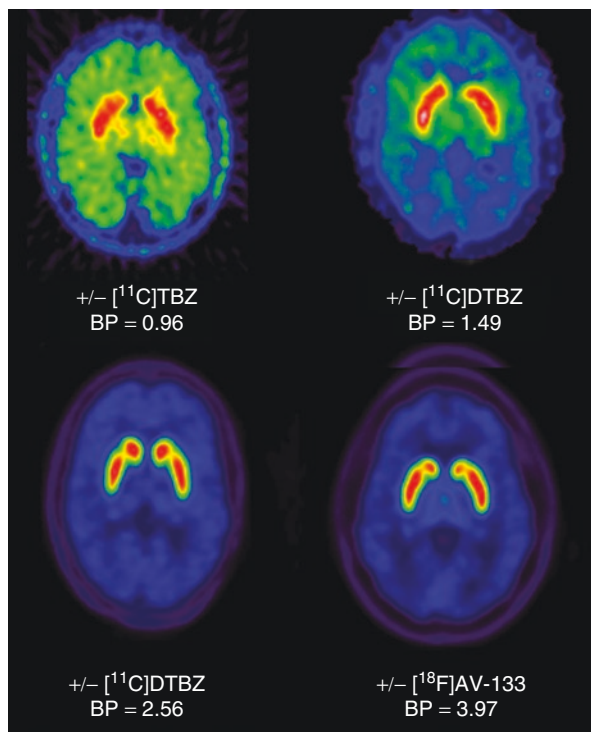
### 33.6 Applications of VMAT2 Radioligands in Human Imaging Studies

The first human studies of the VMAT2 using in vivo imaging with PET were reported in 1993 using racemic [ $^{11}\text{C}$ ]tetrabenazine (Kilbourn et al. 1993), prompting additional years of radiotracer development culminating in the 2008 introduction into human use of a fluorine-18 labeled derivative, 9-(3- $^{18}\text{F}$ fluoropropyl)- $\alpha$ -(+)-dihydro-tetrabenazine ( $^{18}\text{F}$ ]AV-133) (Frey et al. 2008). The combination of improved VMAT2 radioligands with advancements in PET imaging instrumentation and image analysis resulted in significantly improved imaging of the transporter sites in the human brain (Fig. 33.2). As of the beginning of 2019, there were nearly 100 peer-reviewed publications using PET radioligands for VMAT2 imaging in humans, with thousands of subjects (Table 33.3). The majority of such studies have utilized [ $^{11}\text{C}$ ]DTBZ, but there has been an increasing use of the fluorine-18 radioligand [ $^{18}\text{F}$ ]AV-133.

#### 33.6.1 Pharmacokinetic Studies of VMAT2 Radioligands in Normal Brain

The in vivo regional distribution of VMAT2 binding (Table 33.1) and pharmacokinetics of (+)- $\alpha$ - $^{11}\text{C}$ ]DTBZ in the normal human brain have now been thoroughly studied. Initial evaluation of radioligand binding was done using bolus radiotracer

**Fig. 33.2** Representative human PET images demonstrating how the development of better radioligands resulted in improvements of image quality and the estimation of specific binding (BP, binding potential) in normal subjects through the years





**Table 33.3** Human studies using VMAT2 radioligands

Subjects	Radioligands	Topic	Year	Ref
3 NC	(±)[ <sup>11</sup> C]TBZ	First VMAT2 PET study	1993	Kilbourn et al. (1993)
6 NC	(±)[ <sup>11</sup> C]MTBZ	Evaluation of new VMAT2 ligand	1995	Vander Borghet et al. (1995b)
15 NC, 7 PD	(±)[ <sup>11</sup> C]DTBZ	First PET study of VMAT2 in PD	1996	Frey et al. (1996)
9 NC, 8 OPCA, 4 MSA	(±)[ <sup>11</sup> C]DTBZ	Non-PD movement disorders	1996	Gilman et al. (1996)
7 NC	(+)[ <sup>11</sup> C]DTBZ	Eval of pharmacokinetic models	1996	Koeppel et al. (1996)
9 NC	(+)[ <sup>11</sup> C]DTBZ	Eval of equilibrium infusion PET	1997	Koeppel et al. (1997)
6 NC	(+)[ <sup>11</sup> C]DTBZ, (-)[ <sup>11</sup> C]DTBZ	Extrastriatal binding	1999	Koeppel et al. (1999)
22 NC, 8 TS	(+)[ <sup>11</sup> C]DTBZ	VMAT2 in Tourette's	1999	Meyer et al. (1999)
10 NC	(±)[ <sup>11</sup> C]DTBZ	Reproducibility studies	1999	Chan et al. (1999)
4 PDC	(+)[ <sup>11</sup> C]DTBZ	Paroxysmal dystonic choreoathetosis	1999	Bohnen et al. (1999)
64 NC, 19 HD	(+)[ <sup>11</sup> C]DTBZ	Huntington's disease	2000	Bohnen et al. (2000)
10 NC	(±)[ <sup>11</sup> C]DTBZ	Eval of pharmacokinetic models	2000	Sossi et al. (2000)
16 NC, 35 PD	(±)[ <sup>11</sup> C]DTBZ, [ <sup>18</sup> F]FDOPA, [ <sup>11</sup> C]MPH	Multi-tracer PET study in PD	2000	Lee et al. (2000)
16 NC, 16 BPD	(+)[ <sup>11</sup> C]DTBZ	Bipolar disorder	2000	Zubieta et al. (2000)
12 NC, 26 SCZ	(+)[ <sup>11</sup> C]DTBZ	Schizophrenia	2000	Taylor et al. (2000)
15 NC, 15 BPD, 12 SCZ	(+)- <sup>11</sup> C]DTBZ	Comparison of VMAT2 in bipolar vs. schizophrenia	2001	Zubieta et al. (2001)
27 NC, 19 TS	(+)[ <sup>11</sup> C]DTBZ	VMAT2 in Tourette's	2003	Albin et al. (2003)
21 NC, 5 DRD	(±)[ <sup>11</sup> C]DTBZ, [ <sup>18</sup> F]FDOPA, [ <sup>11</sup> C]MPH	Multi-tracer study in Dopa-responsive dystonia	2003	De la Fuente-Fernandez et al. (2003a)
20 PD	(±)[ <sup>11</sup> C]DTBZ, [ <sup>18</sup> F]FDOPA, [ <sup>11</sup> C]MPH	Multi-tracer study in early vs. late onset PD	2003	De la Fuente-Fernandez et al. (2003b)
11 PD	(±)[ <sup>11</sup> C]DTBZ	Asymmetric PD and levodopa	2003	Kumar et al. (2003)
13 MSA	(+)[ <sup>11</sup> C]DTBZ, [ <sup>125</sup> I]IBVM	VMAT2 and VAcHT in MSA	2003	Gilman et al. (2003)
19 NC, 20 DLB, 25 AD	(+)[ <sup>11</sup> C]DTBZ	VMAT2 in AD vs. DLB	2004	Gilman et al. (2004)

**Table 33.3** (continued)

Subjects	Radioligands	Topic	Year	Ref
20 NC, 67 PD	(±)[ <sup>11</sup> C]DTBZ	Disease duration and striatal anteroposterior gradient	2004	Lee et al. (2004)
33 NC, 67 PD, 15 <i>LRRK2</i>	(+)[ <sup>11</sup> C]DTBZ, [ <sup>18</sup> F]FDOPA, [ <sup>11</sup> C]MPH	Multi-tracer study in <i>LRRK2</i> subjects	2005	Adams et al. (2005)
19 NC, 25 AD, 20 DLB, 7FTD	(+)[ <sup>11</sup> C]DTBZ	Use of BP and K <sub>1</sub> to differentiate dementias	2005	Koeppel et al. (2008)
10 NC, 9 PD, 8 PSP, 10 MSA	(+)[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]HED	Losses of brain VMAT2 and heart adrenergic innervation	2006	Raffel et al. (2006)
18 NC, 16 METH	(+)[ <sup>11</sup> C]DTBZ	Abstinent METH users	2006	Johanson et al. (2006)
75 NC, 31 PD	(+)[ <sup>11</sup> C]DTBZ	Age-related decline in NC: Losses in PD	2006	Bohnen et al. (2006)
2 <i>SPG3A</i>	(+)[ <sup>11</sup> C]DTBZ	<i>SPG3A</i> genetic mutations	2008	Albin et al. (2008)
14 NC, 16 METH	(+)[ <sup>11</sup> C]DTBZ	Binding in methamphetamine users	2008	Boileau et al. (2008)
33 NC, 27 PD	(+, -)[ <sup>11</sup> C]DTBZ	VMAT2 loss in early PD	2008	Martin et al. (2008)
15 NC, 44 PD	(+)[ <sup>11</sup> C]DTBZ	Study of VMAT2 and AChE in PD	2009	Bohnen et al. (2009)
28 NC, 33 TS	(+)[ <sup>11</sup> C]DTBZ	VMAT2 in Tourette's	2009	Meyer et al. (1999)
6 PD	(+)[ <sup>11</sup> C]DTBZ	Binding after acute levodopa	2009	De la Fuente-Fernandez et al. (2009)
36 PD	(+)[ <sup>11</sup> C]DTBZ	Increased loss in PD w/ dyskinesia	2009	Troiano et al. (2009)
78 PD	(±)[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]MPH, [ <sup>18</sup> F]FDOPA	Longitudinal multi-tracer study	2009	Nandhagopal et al. (2011)
58 PD	(+)[ <sup>11</sup> C]DTBZ	Olfactory dysfunction in PD	2010	Bohnen et al. (2010)
33 NC	(±)[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]MPH	Aging effects VMAT2 and DAT	2010	Troiano et al. (2010)
80 NC, 19 AD, 9 FTD, 5 DLB, 16 MCI	(+)[ <sup>11</sup> C]DTBZ, [ <sup>18</sup> F]FDG	Use of K <sub>1</sub> from DTBZ as surrogate for blood flow	2010	Albin et al. (2010)
9 NC	(+)[ <sup>11</sup> C]DTBZ	Effects of low-dose amphetamine	2010	Boileau et al. (2010)
9 NC	[ <sup>18</sup> F]AV-133	Human dosimetry study	2010	Lin et al. (2010)
8 <i>LRRK2</i>	(±)[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]MPH [ <sup>18</sup> F]FDOPA	Multi-tracer study in <i>LRRK2</i> subjects	2010	Sossi et al. (2010b)

(continued)

**Table 33.3** (continued)

Subjects	Radioligands	Topic	Year	Ref
6 NC, 17 PD	[ <sup>18</sup> F]AV-133	Losses in PD	2010	Okamura et al. (2010)
73 PD	(+)[ <sup>11</sup> C]DTBZ	Leucoaraiosis in PD	2011	Bohnen et al. (2011)
75 dementia	(+)[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]PIB	VMAT2 and amyloid to classify dementias	2011	Burke et al. (2011)
15 NC, 34 PD	(+)[ <sup>11</sup> C]DTBZ	Rhinorrhea in PD	2011	Chou et al. (2011)
35 NC, 78 PD	(±)[ <sup>11</sup> C]DTBZ	Progression of changes in young vs. older PD	2011	De la Fuente-Fernandez et al. (2011)
4 NC, 9 PD	[ <sup>18</sup> F]AV-133	Eval of imaging times for single-scan protocol	2011	Lin and Lin (2011)
10 NC, 10 AD, 20 PD, 9 DLB	[ <sup>18</sup> F]AV-133	Using VMAT2 to differentiate AD form DLB	2011	Villemagne et al. (2011)
10 NC, 9 DLB, 11 AD, 20 PD	[ <sup>18</sup> F]AV-133	Using VMAT2 to differentiate AD form DLB	2012	Villemagne et al. (2012)
35 NC, 78 PD	(±)[ <sup>11</sup> C]DTBZ, [ <sup>18</sup> F]FDOPA, [ <sup>11</sup> C]MPH	Longitudinal study of compensatory changes	2011	Nandhagopal et al. (2009)
4 NC, 9 PD	(+)[ <sup>18</sup> F]AV-133	Optimization of scan protocol	2011	Lin et al. (2011)
26 <i>LRRK2</i>	(±)[ <sup>11</sup> C]DTBZ	Losses in symptomatic <i>LRRK2</i>	2012	Aasly et al. (2012)
12 NC, 12 COC	(+)[ <sup>11</sup> C]DTBZ	Small losses in cocaine users	2012	Narendran et al. (2012a)
13 NC	(+)[ <sup>11</sup> C]DTBZ	Omega-3 supplementation	2012	Narendran et al. (2012b)
80 PD	(+)[ <sup>11</sup> C]DTBZ	Rapid eye movement disorder	2012	Kotagal et al. (2012)
51 PD	(+)[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]DASB	Sleep disordered breathing	2012	Lelieveld et al. (2012)
40 PD-MCI	(+)[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]PIB	VMAT2 and amyloid in PD w/MCI	2012	Petrou et al. (2012b)
32 NC, 125 PD	(+)[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]PMP	VMAT2 and AChE –Gait speed correlation	2013	Bohnen et al. (2013)
22 NC	[ <sup>18</sup> F]AV-133	Normal brain distributions	2013	Lin et al. (2013)
148 PD	(+)[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]PMP	PD and diabetes	2013	Kotagal et al. (2013)
18 PD	(+)[ <sup>11</sup> C]DTBZ	VMAT2 losses and levodopa	2013	Kwak et al. (2013)
44 PD	(+)[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]PIB	Postural instability and gait	2013	Muller et al. (2013a)

**Table 33.3** (continued)

Subjects	Radioligands	Topic	Year	Ref
25 NC, 124 PD	(+)[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]PIB	VMAT2 and AChE in thalamus	2013	Muller et al. (2013b)
5 NC, 5 PD, 11 DLB, 10 AD	[ <sup>18</sup> F]AV-133 Florbetapir-F18	VMAT2 and amyloid in cognition	2014	Siderowf et al. (2014)
17 NC, 30 PD	[ <sup>18</sup> F]AV-133	Losses in early PD	2014	Lin et al. (2014)
8 NC, 10 PD	[ <sup>18</sup> F]AV-133 [ <sup>99m</sup> Tc] TRODAT-1	Comparison of VMAT2 and DAT losses in PD	2014	Hsiao et al. (2014a)
17 NC, 53 PD	[ <sup>18</sup> F]AV-133	Losses in advanced PD	2014	Hsiao et al. (2014b)
148 PD	(+)[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]PMP	VMAT2 and AChE in PD w/diabetes	2014	Bohnen et al. (2014)
3 <i>DCTN1</i>	(+)[ <sup>11</sup> C]DTBZ, [ <sup>18</sup> F]FDOPA, [ <sup>11</sup> C]RAC, [ <sup>11</sup> C]DASB	Multi-tracer study in <i>DCTN1</i> genetic mutation	2014	Felicio et al. (2014)
49 NC, 83 PD	(+)[ <sup>11</sup> C]DTBZ	Cardiovascular risk	2014	Kotagal et al. (2014)
29 PD, 18 PD-MCI	(+)[ <sup>11</sup> C]DTBZ	Cognitive tests in MCI	2014	Chou et al. (2014)
8 NC, 8 PD, 4 Mn-intoxication	[ <sup>18</sup> F]AV-133	VMAT2 in Mn-intoxication	2015	Huang et al. (2015)
143 PD	(+)[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]PMP	VMAT2 and AChE in cognition	2015	Bohnen et al. (2015)
9 NC, 18 PD	[ <sup>18</sup> F]AV-133	Decrease in CO-induced parkinsonism	2015	Chang et al. (2015)
48 PD	(+)[ <sup>11</sup> C]DTBZ	Exercise and symptom severity	2015	Snider et al. (2015)
133 PD	(+)[ <sup>11</sup> C]DTBZ	VMAT2 loss and fatigue in PD	2016	Chou et al. (2016)
22 NC, 28 METH, 9 COC	(+)[ <sup>11</sup> C]DTBZ	VMAT2 losses in cocaine and methamphetamine use	2016	Boileau et al. (2016)
4 NC, 22 PD	[ <sup>18</sup> F]AV-133	Correlations with CSF biomarkers	2016	Gao et al. (2016)
62 PD	(+)[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]PIB, [ <sup>11</sup> C]PMP	Correlations with amyloidopathy	2016	Shah et al. (2016)
12 NC, 14 <i>22q11.2</i>	(+)[ <sup>11</sup> C]DTBZ	<i>22q11.2</i> genetic mutation effects on VMAT2	2017	Butcher et al. (2017)
49 PD	(+)[ <sup>11</sup> C]DTBZ	Color discrimination and PD	2017	Bohnen et al. (2017)
36 PD	(+)[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]PIB	Hoeh and Yahr stage and amyloid burden	2017	Kotagal et al. (2017)

(continued)

**Table 33.3** (continued)

Subjects	Radioligands	Topic	Year	Ref
47 PD	[ <sup>18</sup> F]AV-133	Clinical impact in uncertain PD	2017	Alexander et al. (2017)
5 NC, 12 PD	[ <sup>18</sup> F]AV-133	Reproducibility study	2017	Wu et al. (2017)
13 PD, 16 <i>LRRK2</i>	(+)[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]DASB	Dual VMAT2 and SERT study	2017	Wile et al. (2017)
9 NC, 15 PD, 17 <i>LRRK2</i>	(+)[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]DASB	Pattern analysis of dual study	2018	Fu et al. (2018)
85 PD	[ <sup>18</sup> F]AV-133	Clinical impact in uncertain PD	2018	Xu et al. (2018)
26 NC, 39 PD	[ <sup>18</sup> F]AV-133	Losses in PD	2018	Lung et al. (2018)
14 NC, 14 ND	(+)[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]PMP	Study of peripheral neuropathy	2018	Beaulieu et al. (2018)
39 PD	(+)[ <sup>11</sup> C]DTBZ	Diagnostic utility in uncertain PD	2018	Perez-Lohman et al. (2018)
12 NC, 30 PD	(+)[ <sup>11</sup> C]DTBZ	VMAT2 loss in globus pallidus	2019	Cho et al. (2019)
75 NC, 129 PD	(+)[ <sup>11</sup> C]DTBZ	Combined with MRI diffusion imaging	2019	Yang et al. (2019)
8 PD	(+)[ <sup>11</sup> C]DTBZ	Implants of neural progenitor cells	2019	Madrazo et al. (2019)
7 PD	(+)[ <sup>11</sup> C]DTBZ	Deep brain stimulation study	2019	Smith et al. (2019)
15 PD	(+)[ <sup>11</sup> C]DTBZ, [ <sup>11</sup> C]MPH	Joint pattern analysis VMAT2 + DAT	2019	Fu et al. (2019)
21 NC, 37 PD	[ <sup>18</sup> F]AV-133	Correlations with nonmotor symptoms	2019	Shi et al. (2019)
83 NC, 84 PD	[ <sup>18</sup> F]AV-133	Constipation and inflammation	2019	Zhou et al. (2019)

injections, arterial blood sampling and determination of metabolite-corrected blood curves, and compartmental modeling to obtain estimates of binding potentials (BP) for [<sup>11</sup>C]DTBZ in multiple regions of the human brain (Koeppel et al. 1996). (+)- $\alpha$ -[<sup>11</sup>C]DTBZ is a reversibly binding radioligand also well suited for use in equilibrium infusion applications (Koeppel et al. 1997). Subsequent studies using injections of both the active and inactive stereoisomers, each in carbon-11 labeled form, provided direct evidence of very low binding of the radioligand in non-target regions such as the cerebellum and the occipital cortex (Koeppel et al. 1999), consistent with studies of immunoreactivity for VMAT2 (Tong et al. 2011) and in vitro autoradiographic studies (Scherman and Raisman 1988). The reproducibility of binding measures using [<sup>11</sup>C]DTBZ in humans has been examined and found reliable, provided care is exercised in selecting appropriate reference regions (Chan et al. 1999).

The in vivo pharmacokinetics of the fluorine-18 labeled radiotracer ([<sup>18</sup>F]AV-133) have similarly been evaluated using a variety of analysis methods (Frey et al. 2008;

Lin and Lin 2011; Lin et al. 2013; Okamura et al. 2010). Compartmental modeling of [ $^{18}\text{F}$ ]AV-133 kinetics can be used to estimate regional binding potentials (BP) (Okamura et al. 2010), but the longer half-life of this radiotracer also permits a proposed use of delayed single-frame imaging protocols to estimate standard uptake value ratios (SUVr) as reasonable estimates of the specific binding of the radioligand (Lin and Lin 2011). Good reproducibility of such SUVr values has been demonstrated (Wu et al. 2017).

### 33.6.2 Aging

As many of the potential applications of VMAT2 imaging in neurodegenerative disease involve elderly populations (e.g., PD, AD), the possible effect of age-related declines in radioligand binding must be considered. Studies of aging effects in normal controls have however provided inconsistent results, showing both no decline of VMAT2 radioligand binding with age (Troiano et al. 2010) as well as a small but steady decrease of 0.5–0.77%/year (Frey et al. 1996; Bohnen et al. 2006). In a single study, a longitudinal study of the age-dependent losses of VMAT2 radioligand binding in early versus late onset PD subjects demonstrated a slower progression in the younger onset patients (De la Fuente-Fernandez et al. 2011). Age-related effects on radioligand binding remain a potential confound in imaging studies, and for that reason well-designed studies of VMAT2 changes in disease require the inclusion of appropriate age-matched control subjects.

### 33.6.3 Parkinson's Disease

By far, the single largest application of VMAT2 radioligand imaging has been in Parkinson's disease, with more than 50 published studies (for references see Table 33.3). The PET studies done with [ $^{11}\text{C}$ ]DTBZ, and more recently [ $^{18}\text{F}$ ]AV-133, have consistently shown decreases in radioligand binding in the striatal regions (caudate, putamen, nucleus accumbens, globus pallidus) (Table 33.4). Significant losses of radioligand binding can be demonstrated in the presymptomatic phase, often with a unilateral presentation and observations of an anteroposterior gradient (losses in posterior putamen before the caudate nucleus and ventral striatum/nucleus accumbens) and progression to more severe and complete bilateral losses throughout the striatum in advanced disease (Fig. 33.3). The losses of VMAT2 radioligand binding can also be correlated with a variety of clinical observations (dyskinesias, olfaction, sleep disorders, gait, fatigue).

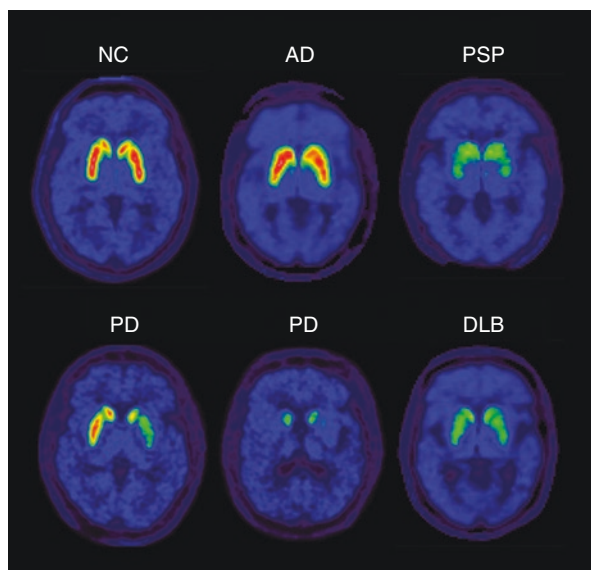
In order to more fully understand the biochemical changes in the human brain in the course of Parkinson's disease, numerous studies have utilized a multi-radiotracer approach by combining VMAT2 measures ([ $^{11}\text{C}$ ]DTBZ or [ $^{18}\text{F}$ ]AV-133) with imaging studies using radiotracers for the neuronal membrane dopamine transporter (DAT: [ $^{11}\text{C}$ ]methylphenidate (MPH) or [ $^{99\text{m}}\text{Tc}$ ]Trodar-1), dopamine synthesis

**Table 33.4** In vivo binding estimates for (+)- $\alpha$ -[ $^{11}\text{C}$ ]DTBZ and [ $^{18}\text{F}$ ]AV-133 ((+)- $\alpha$ -(3-[ $^{18}\text{F}$ ]fluoropropyl)DTBZ) in healthy controls (HC), Parkinson's disease (PD), dementia with Lewy bodies (DLB), and Alzheimer's disease (AD)

Ligand	Disease	Region	BP (binding potential) Mean (SD)
(+)-[ $^{11}\text{C}$ ]DTBZ (Koeppel et al. 2008)	HC	Caudate nucleus	1.69 (0.23)
		Anterior putamen	2.00 (0.25)
		Posterior putamen	1.91 (0.25)
	PD	Caudate nucleus	0.93 (0.26)
		Anterior putamen	0.65 (0.21)
		Posterior putamen	0.43 (0.15)
	DLB	Caudate nucleus	0.87 (0.30)
		Anterior putamen	0.71 (0.21)
		Posterior putamen	0.57 (0.22)
	AD	Caudate nucleus	1.74 (0.35)
		Anterior putamen	2.06 (0.41)
		Posterior putamen	1.92 (0.42)
[ $^{18}\text{F}$ ]AV-133 (Okamura et al. 2010)	HC	Caudate nucleus	2.53 (0.3)
		Anterior putamen	2.56 (0.3)
		Posterior putamen	2.53 (0.3)
	PD	Caudate nucleus	1.20 (0.5)
		Anterior putamen	0.66 (0.3)
		Posterior putamen	0.30 (0.2)
	DLB	Caudate nucleus	1.24 (0.6)
		Anterior putamen	0.90 (0.5)
		Posterior putamen	0.62 (0.5)
	AD	Caudate nucleus	2.83 (0.9)
		Anterior putamen	3.01 (0.9)
		Posterior putamen	2.87 (0.9)

([ $^{18}\text{F}$ ]fluoroDOPA), dopamine receptors ([ $^{11}\text{C}$ ]raclopride), serotonin reuptake (SERT: [ $^{11}\text{C}$ ]DASB), acetylcholinesterase ([ $^{11}\text{C}$ ]PMP), and amyloid burden ([ $^{11}\text{C}$ ]PIB). Combination of [ $^{11}\text{C}$ ]DTBZ with [ $^{18}\text{F}$ ]fluoroDOPA and [ $^{11}\text{C}$ ]MPH, all three of which measure aspects of presynaptic dopaminergic neurons, has demonstrated differential losses when comparing PD subjects at various stages of disease: investigators have proposed that VMAT2 measures appear the least subject to regulatory effects and may represent the most accurate measure of DA-ergic terminals, whereas these studies support that dopamine synthesis is upregulated (result: increased [ $^{18}\text{F}$ ]fluoroDOPA uptake and retention) and the DAT is downregulated (result: decreased [ $^{11}\text{C}$ ]MPH binding) (Nandhagopal et al. 2011; Lee et al. 2000). Such compensatory changes would work to maintain synaptic dopamine levels as terminals are lost to the disease process, but that conclusion is dependent on VMAT2 as a reliable marker of terminal density. A part of the rationale for development of VMAT2 radioligands was the hypothesis that the vesicular transporter, in comparison to measures of dopamine synthesis or the neuronal membrane dopamine transporter, would be more resistant to regulation of total numbers ( $B_{\text{max}}$ ) in response to

**Fig. 33.3** Representative human PET images of (+)-alpha-[ $^{11}\text{C}$ ]DTBZ binding in normal controls (NC), Alzheimer's disease (AD), progressive supranuclear palsy (PSP), early and late Parkinson's disease (PD), and dementia with Lewy bodies disease (DLB)



pharmacological treatments or compensatory changes in a disease process. Animal studies to evaluate that hypothesis have been inconsistent (see Sect. 33.5.2), and studies intended to perturb brain biochemistry are of course much harder to perform in living human subjects. The administration of amphetamine could lead to depletion of vesicular dopamine levels due to stimulant-induced release, but a low acute dose of amphetamine in human subjects was found to produce only a small (5%) decrease in radioligand binding to the VMAT2 (Boileau et al. 2010); whether the dose of amphetamine administered was adequate to reduce vesicular dopamine levels to an extent sufficient to subsequently reduce competition for radioligand binding remains an open question, as the effects of amphetamine on the VMAT2 are complex and dose-dependent (Reith and Gnegy 2020). It should be noted that in animals, demonstration of an effect of amphetamine required extensive depletion of tissue dopamine concentrations. The clinical equivalent of depletion of synaptic vesicle dopamine is the condition known as DOPA-responsive dystonia: the observation that *in vivo* binding of [ $^{11}\text{C}$ ]DTBZ is increased in those subjects (De la Fuente-Fernandez et al. 2003a) was an indication that vesicular dopamine may compete for binding of radioligands to the TBZ site located on the inner surface of the vesicular membrane. However, in both these patients and in studies of dopamine-depleted rat brains, alterations of VMAT2 radioligand binding *in vivo* were shown after extensive or complete neurotransmitter depletion was accomplished, and the sensitivity of VMAT2 radioligand binding to smaller changes in dopamine levels remains to be determined. The potential effects of increased endogenous dopamine levels affecting human [ $^{11}\text{C}$ ]DTBZ studies have been examined through acute administration of L-DOPA to subjects with advanced Parkinson's disease (De la Fuente-Fernandez et al. 2009), where significant decreases in radioligand binding in caudate and putamen were proposed to be due to the increases in vesicular



dopamine. Taken together, the animal and human studies involving manipulations of endogenous dopamine levels suggest that care should be taken to account for use of L-DOPA, or other dopamine-modifying drugs, when performing studies with VMAT2 radioligands.

Several studies have examined particular subsets of patients to probe different aspects of the disease. Examination of VMAT2 binding in a cohort of untreated early, untreated PD patients demonstrated that even in these subjects, there was a significant loss (>50%) of VMAT2 binding sites particularly in the putamen (Martin et al. 2008); a loss of DTBZ binding in the clinically unaffected side of PD patients has also been reported (Bohnen et al. 2006). In studies of unilateral Parkinsonism, the greater decrease of DTBZ binding on the more clinically affected side correlated with responsiveness to L-DOPA treatment (Kumar et al. 2003). A study of both the VMAT2 and the DAT in controls versus Parkinson's disease with dyskinesias showed no significant differences of DTBZ binding between groups but a difference in the ratio of dopamine transporter (DAT) to VMAT2 radioligand binding in the putamen, with ratios being significantly lower in the group having dyskinesias. The comparison of VMAT2 radioligand binding measurements with matched studies of the dopamine neuronal membrane transporter (DAT) and 6-[<sup>18</sup>F]fluoroDOPA uptake in a large cohort of Parkinson's disease patients (De la Fuente-Fernandez et al. 2011) provided evidence that younger-onset subjects had better compensatory mechanisms operating to maintain the biochemical balance of the dopaminergic system as compared to late-onset PD patients. A longitudinal multi-tracer study of PD patients provided evidence for compensatory upregulation of the DAT radioligand binding and 6-[<sup>18</sup>F]fluoroDOPA uptake in earlier stages of the disease, but that compensatory mechanisms decline with advanced neuronal losses as shown by extensive losses of [<sup>11</sup>C]DTBZ binding. The numerous studies that have now compared multiple presynaptic radiotracers have clearly demonstrated differential rates and extents of losses between the measures of the VMAT2, the DAT, and dopamine synthesis.

The VMAT2 radioligands [<sup>11</sup>C]DTBZ and [<sup>18</sup>F]AV-133 have not reached routine use in the clinical management of Parkinson's disease, but studies have begun to evaluate the value of such imaging studies in situations where the diagnosis of PD is uncertain (Alexander et al. 2017; Xu et al. 2018).

### 33.6.4 Non-Parkinson Movement Disorders

Losses of *in vivo* VMAT2 radioligand binding have been observed in studies of less common neurodegenerative movement disorders (progressive supranuclear palsy (PSP), multiple system atrophy (MSA), and olivopontocerebellar atrophy (OPCA)) that share traits with idiopathic parkinsonism but express additional clinical symptoms. Involvement of the basal ganglia is generally observed, but the pattern of VMAT2 radioligand binding changes are different: for example, whereas in

idiopathic Parkinson's disease losses are earlier, asymmetric, and more predominant in the posterior putamen, in PSP there was observed a more uniform involvement of both the caudate and the putamen (Fig. 33.3) (Raffel et al. 2006). Similarly, VMAT2 radioligand binding is altered in MSA, OPCA, and other degenerative diseases involving monoaminergic nerve terminals of the basal ganglia (Gilman et al. 1996; Gilman et al. 2003).

### 33.6.5 Huntington's Disease

Huntington's disease (HD) is characterized by significant neuronal loss and atrophy in the caudate nucleus and putamen. A single study of [ $^{11}\text{C}$ ]DTBZ binding in HD subjects showed reductions in total radioligand binding in caudate nucleus, anterior and posterior putamen, with losses most prominent in the posterior putamen (Bohnen et al. 2000). Within the study group, individuals with the akinetic-rigid phenotype (ADr) showed more significant losses of total radioligand binding than the subgroup with the chorea phenotype (HDc).

### 33.6.6 Dementia

The underlying biochemical changes in different clinical manifestations of dementia (Alzheimer's disease (AD), Parkinson's disease with dementia (PDD), and dementia with Lewy body disease (DLB)) do not uniformly involve the same degenerative pathways, and losses of VMAT2 are not uniform in the different dementias (Table 33.4). In Alzheimer's disease the striatal monoaminergic innervation is generally intact (Fig. 33.3), and no losses of in vivo VMAT2 radioligand binding have been reported (Gilman et al. 2004; Koeppe et al. 2008). In contrast, the basal ganglia are significantly affected in dementia with Lewy bodies, with losses of radioligand binding clearly evident throughout the caudate nucleus and putamen (Fig. 33.3); use of imaging techniques to clearly identify DLB vs. AD patients (Koeppe et al. 2008; Villemagne et al. 2011; Villemagne et al. 2012) has clinical impact, as DLB patients are subject to adverse reactions from dopaminergic treatments that might otherwise be useful for treatment of hallucinations or behavioral disturbances in AD patients. Finally, Parkinson's disease with dementia patients show the expected significant losses of striatal VMAT2 radioligand binding (Koeppe et al. 2008); the dementia is proposed to be associated with concurrent changes in the cholinergic system, which can sometimes be demonstrated by separate imaging studies with radiotracers for that neurotransmitter system (Bohnen and Frey 2007; Bohnen and Albin 2011). The combination of [ $^{11}\text{C}$ ]DTBZ imaging studies with PET imaging of amyloid deposits using such radioligands as [ $^{11}\text{C}$ ]PIB provides a method for classifications of dementia subjects which may not always agree with consensus clinical evaluations (Burke et al. 2011).

### 33.6.7 Genetic Diseases

There are several genetic mutations that have been examined using PET and VMAT2 radioligands. Mutations of the *LRK2* protein (leucine-rich repeat kinase 2) produce L-DOPA responsive parkinsonism, and PET imaging was used to show decreases of [ $^{11}\text{C}$ ]DTBZ, [ $^{18}\text{F}$ ]fluoroDOPA, and [ $^{11}\text{C}$ ]MPH in symptomatic subjects (Adams et al. 2005; Aasly et al. 2012). The Parkinson's disease phenotype of *22q11.2DS* produces clinical symptoms and response to treatment similar to PD, and preliminary imaging studies with [ $^{11}\text{C}$ ]DTBZ showed the interesting results of increased radioligand binding in non-symptomatic gene carriers and the expected severe losses of VMAT2 binding in a subject with parkinsonism (Butcher et al. 2017). In a preliminary study (Felicio et al. 2014), three individuals with a mutation in the dynactin gene (*DCTN1*) that produces an autosomal dominantly inherited parkinsonism (and other clinical signs, including depression, forming Perry syndrome) were examined by PET with [ $^{11}\text{C}$ ]DTBZ, [ $^{18}\text{F}$ ]fluoroDOPA, [ $^{11}\text{C}$ ]raclopride, and [ $^{11}\text{C}$ ]DASB. In two clinically affected subjects, there was a large decrease in [ $^{11}\text{C}$ ]DTBZ binding, a modest loss of [ $^{18}\text{F}$ ]fluoroDOPA uptake, and normal to increased [ $^{11}\text{C}$ ]RAC.

### 33.6.8 Environmental Toxins

PET imaging of [ $^{11}\text{C}$ ]DTBZ binding has been utilized to evaluate the potential losses of dopaminergic innervation after environmental insults. Chronic excessive exposure to manganese produces some clinical signs similar to idiopathic Parkinson's disease, but the involvement of dopaminergic degeneration has not been firmly established. PET studies with [ $^{18}\text{F}$ ]AV-133 showed no decreased VMAT2 binding, supporting the distinction between manganese intoxication and PD (Huang et al. 2015). In contrast, studies of [ $^{18}\text{F}$ ]AV-133 binding in subjects with carbon monoxide-induced parkinsonism showed decreases in caudate and putamen, although the sparing of radioligand binding in the substantia nigra suggested a differential pathophysiology as compared to idiopathic PD (Chang et al. 2015).

### 33.6.9 Psychiatric Disease

A few studies have been completed looking at VMAT2 radioligand binding in schizophrenia and bipolar disorder type I. In schizophrenia, no differences in striatal or thalamic binding of VMAT2 radioligands was observed, but small increases in binding in the ventral brainstem of schizophrenics were observed relative to controls (Taylor et al. 2000; Zubieta et al. 2001). The importance of that observation is difficult to ascribe to any single neurotransmitter system, as that portion of the brain has cell bodies and projections for both serotonergic and dopaminergic neurons. For the bipolar disorder type I group, increases of VMAT2 binding in the thalamus relative to controls can at least be proposed not to be due to changes in dopaminergic

innervation, which is very low in that brain region, but to changes in the serotonergic and noradrenergic innervation (Zubieta et al. 2000).

The *in vivo* binding of VMAT2 radioligands has also been examined in Tourette's syndrome, and inconsistent results have been obtained, with the discrepant observations of both no changes (Meyer et al. 1999) and increased [<sup>11</sup>C]DTBZ binding (Albin et al. 2003) attributed to the heterogeneity of Tourette's syndrome patient populations.

### 33.6.10 Drug Abuse

As many drugs which are abused are known to interact with the monoaminergic systems of the brain (e.g., amphetamine, methamphetamine, cocaine), the effects of such drug use on the *in vivo* binding of VMAT2 radioligands to monoaminergic terminals have been examined, with particular interest in the potential neurotoxic effects on monoaminergic terminals. Postmortem studies of chronic cocaine users have demonstrated both no changes in VMAT2 ligand binding or immunoreactivity (Staley et al. 1997) as well as modest decreases in VMAT2 radioligand binding (Wilson et al. 1996a; Little et al. 2003). In a PET study of cocaine users, reduced *in vivo* VMAT2 binding was shown in several striatal regions, suggesting either a compensatory downregulation of VMAT2 storage systems after drug use or a loss of dopaminergic terminals (Narendran et al. 2012a); which of these is occurring, and the clinical impact needs further study.

The effects of methamphetamine on striatal VMAT2 radioligand binding have been examined both *in vitro* and *in vivo* with conflicting results (Boileau et al. 2016, 2008; Wilson et al. 1996b; Johanson et al. 2006). Postmortem studies of methamphetamine users have not shown decreases in VMAT2 binding, but PET studies have shown both decreased (suggesting neurodegeneration) and increased [<sup>11</sup>C]DTBZ binding (suggesting higher VMAT2 site availability due to lower DA levels). The effects of methamphetamine on the dopaminergic system and the VMAT2, and the potential neurotoxicity, are thus poorly understood (Kish et al. 2017).

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## 33.7 Conclusions

In contrast to the development of radiopharmaceuticals for *in vivo* PET imaging of most other biochemical targets, where multiple competing radioligands have been synthesized and advanced through to human studies (e.g., dopamine receptors, amyloid imaging), the development of an *in vivo* imaging radiotracer for the VMAT2 has been focused on derivatives of a single basic structure that of tetrabenzazine. The currently available radiotracers, (+)- $\alpha$ -[<sup>11</sup>C]DTBZ and [<sup>18</sup>F]AV-133, present very favorable characteristics: good brain permeability, reversible kinetics, lack of interfering metabolites, well-known pharmacology, acceptable radiation dosimetry, and simple radiochemical preparation. *In vivo* human imaging of the vesicular

monoamine transporter 2 (VMAT2) will continue to be useful in studies of diseases involving changes of monoaminergic neuronal densities in the brain.

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