REVIEW PAPER



Elucidating Critical Proteinopathic Mechanisms and Potential Drug Targets in Neurodegeneration

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Abstract

Neurodegeneration entails progressive loss of neuronal structure as well as function leading to cognitive failure, apathy, anxiety, irregular body movements, mood swing and ageing. Proteomic dysregulation is considered the key factor for neurodegeneration. Mechanisms involving deregulated processing of proteins such as amyloid beta (Aβ) oligomerization; tau hyperphosphorylation, prion misfolding; α-synuclein accumulation/lewy body formation, chaperone deregulation, acetylcholine depletion, adenosine 2A (A2A) receptor hyperactivation, secretase deregulation, leucine-rich repeat kinase 2 (LRRK2) mutation and mitochondrial proteinopathies have deeper implications in neurodegenerative disorders. Better understanding of such pathological mechanisms is pivotal for exploring crucial drug targets. Herein, we provide a comprehensive outlook about the diverse proteomic irregularities in Alzheimer's, Parkinson's and Creutzfeldt Jakob disease (CJD). We explicate the role of key neuroproteomic drug targets notably A\(\beta\), tau, alpha synuclein, prions, secretases, acetylcholinesterase (AchE), LRRK2, molecular chaperones, A2A receptors, muscarinic acetylcholine receptors (mAchR), N-methyl-D-aspartate receptor (NMDAR), glial cell line-derived neurotrophic factor (GDNF) family ligands (GFLs) and mitochondrial/oxidative stressrelated proteins for combating neurodegeneration and associated cognitive and motor impairment. Cross talk between amyloidopathy, synucleinopathy, tauopathy and several other proteinopathies pinpoints the need to develop safe therapeutics with ability to strike multiple targets in the aetiology of the neurodegenerative disorders. Therapeutics like microtubule stabilisers, chaperones, kinase inhibitors, anti-aggregation agents and antibodies could serve promising regimens for treating neurodegeneration. However, drugs should be target specific, safe and able to penetrate blood-brain barrier.

Keywords Chaperones · Drug targets · Neurodegeneration · Mitophagy · Free radicals

Introduction

Once degenerated, neurons of central nervous system (CNS) cannot replace themselves thus causing severe neurodegenerative disorders. The actual cause underlying regenerative failure of neurons is still not known. However, scientists report the presence of "road-blocks" in the injured or diseased CNS that prevent neurons from regenerating

themselves. These road-blocks are diverse in nature and include molecules found in extracellular matrix as well as the proteins that get expressed once the neurons are damaged. In addition, once the CNS development is completed, there occurs activation of certain genes in neurons that act as brakes and prevent the further development of CNS. Nowadays, scientists are trying to silence such genes or their products so as to turn on growth programs in neurons.

Neurodegenerative diseases (ND's) are growing at alarming pace due to rise in the elderly population (Heemels 2016). Some of these diseases damage person's memory and cognition while others affect the ability to move, speak and breathe (Gitler et al. 2017). In America alone, around 5 million people suffer from Alzheimer's disease (AD), 0.4 million from multiple sclerosis (MS), 1 million from Parkinson's disease (PD), 30,000 from amyotrophic lateral sclerosis (ALS) and 3000 from Huntington's disease (HD)

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(Agrawal and Biswas 2015). Globally, CJD affects about one person in every one million annually. In US alone, about 350 cases are detected annually. Cure for neurodegenerative diseases (NDs) is still under investigation and the current therapeutics only treat symptoms (Lang 2010). Necessity to develop effective drugs is the need of the hour; however, the task poses perplexing challenges owing to complex nature of NDs. Proteins act as important enzymes, biochemical regulators, electron transporters, signal transducers, receptors and scaffolding agents in diverse biological cascades. Their deregulated form can disrupt the normal cellular physiology and lead to NDs. Excessive accumulation of mutant or wild-type proteins with distorted conformations that aid aggregation is a characteristic feature of many neurodegenerative diseases including Parkinson's, Alzheimer's and CJD.

Strategy to strike proteomic drug targets represents an imperative approach for restricting neuronal damage. So far several drugs have been developed or being tested in various animal models (Fig. 1) for their neuroprotective action but the issues like (I) blood-brain barrier penetration, (II) toxicity and (III) limited drug efficacy often challenge the process. Conventional drug-discovery approaches based on one gene, one drug and one disease philosophy just modulate symptoms and do little to treat the root cause. Drugs that target only one protein are susceptible to drug resistance mainly due to the fact that even a single mutation in the target active site can substantially reduce the binding affinity of a drug and hence its efficacy. Drugs that strike multiple targets would require the unlikely event of concurrent mutations appearing in multiple protein targets. Thus, developing 'magic shotgun drugs" instead of 'magic bullet drugs" could play pivotal role in formulating effective regimens for neurodegenerative diseases and associated cognitive decline (Cornelis and Schyf 2011). This review provides mechanistic insight about diverse proteomic destabilisations that occur during neuronal damage and also elucidates potential proteomic drug targets (Fig. 2) to combat neurodegeneration.

Alzheimer's Disease

Alzheimer's disease (AD) is characterised by extreme dementia i.e. loss of overall cognitive functioning of brain like remembering, thinking or reasoning to such an extent that it interferes with person's daily activities. It is the leading cause of death in elderly. At present, about 24 million people suffer from AD across the globe (Chakrabarti et al. 2015). As per the Alzheimer's Association, about 5.4 million American people were affected by AD in 2016 (Rubenstein 2017). As per reports, by the year 2050, the population of AD patients in the United States will increase threefold over the number in 2000 (Hebert et al. 2003). Being an irreversible proteinopathy and a

progressive brain disorder, it causes slow destruction of brain's ability to memorise even simplest tasks. Unfortunately, there is no treatment to stop or reverse its progression, although some drugs may temporarily improve symptoms. AD is a complex neurological disorder primarily characterised by the accumulation of amyloid beta plaques $(A\beta)$ in brain, but amyloidopathy is not necessarily the ultimate cause of Alzheimer's disease. Infact, many other possible models have been proposed including N-APP-DR6-Caspase pathway, tauopathy, ApoE protein deregulation and environmental factors mediated proteinopathy, as illustrated below:

Amyloidopathy is considered the main cause of AD. Amyloid beta is not produced directly but indirectly from neuronal transmembrane amyloid precursor protein (APP). APP is encoded by app gene located on chromosome 21. APP protein is crucial for the growth, survival and postinjury repair of neuronal cells but its cleavage product amyloid beta $(A\beta)$ generated proteolytically by the mutual action of two aspartyl proteases i.e. beta and gamma secretases leading to neurotoxicity and neuronal death (Koo et al. 1990). Extracellular cleavage of transmembrane APP glycoprotein by beta secretase generates a soluble extracellular fragment known as sAPPB and a cell membrane-bound fragment called C99. Subsequent cleavage of C99 within its transmembrane domain by gamma secretase liberates an intracellular fragment called amino-terminal APP intracellular domain (AICD) and an extracellular fragment known as amyloid beta (Aβ) which is 40–42 amino acid long fibrils having molecular weight around 4kd. The later undergoes oligomerization to form abnormal clumps outside the neurons called Aβ plaques (Tiraboschi et al. 2004). These plaques trigger inflammatory response and oxidative stress in neurons thereby disrupting their normal communication and leading to their death (Bosco et al. 2011). Altogether these events constitute the amyloidogenic pathway of AD (Fig. 3). Formation of A oligomers can further activate microglia which in turn releases various neuroinflammatory cytokines that upregulate, and secretases increase APP levels and decrease A clearance in the brain thus causing further upsurge of A and subsequent formation of A oligomers and plagues in the brain (Alasmari et al. 2018). Amyloid beta plaques are of three types i.e. senile, diffuse and neuritic (D'Andrea and Nagele 2010). The diffuse plaques have a little effect on cognitive function, while the neuritic plaques are markedly involved in cognitive decline (Kokubo et al. 2009). It is hypothesised that diffused plaques appear during the early stages of Alzheimer's disease while the neuritic plaques appear in the advanced stages. Further, studies show that clathrin and clathrin adaptor proteins involved in the endocytosis of APP can lead to increased intracellular levels of amyloid beta peptide, contributing to the progression of Alzheimer's (Hallock et al. 2012).



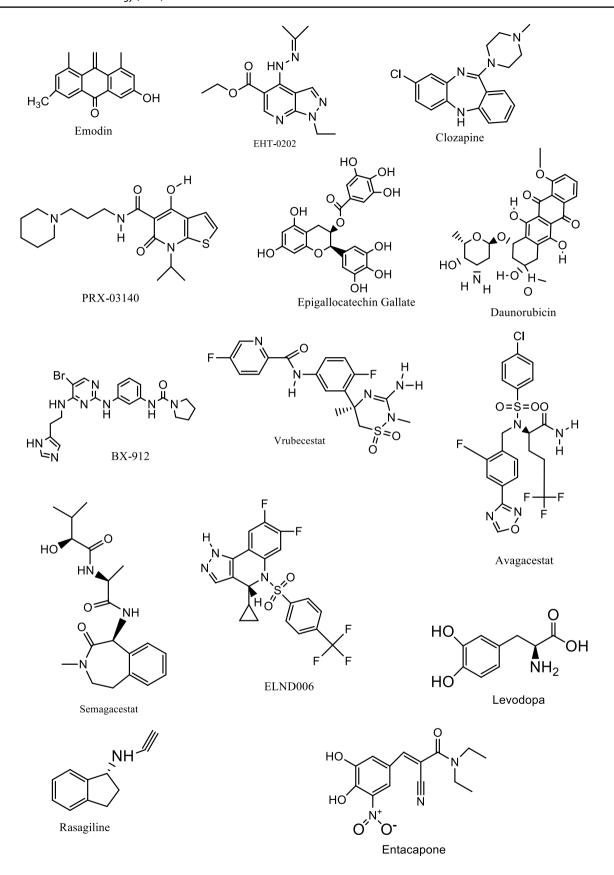


Fig. 1 Structures of drugs/molecules currently employed for the treatment of neurodegenerative diseases or being tested in clinical trials



Fig. 1 (continued)

Aβ is also involved in activation of apoptosis signal-regulating kinase-1 (ASK1) which is implicated in reactive oxygen species (ROS)-mediated activation of c-Jun N-terminal kinases (JNK) as well as apoptosis (Hashimoto et al. 2004;

Kadowaki et al. 2005a, b). Thus, $A\beta$ may indirectly lead to neuronal death via ASK1 activation. Amyloid beta also modulates redox factor-1 that plays crucial role in cell death signalling pathway and DNA repair by showing interaction



Fig. 2 Diverse proteins found deregulated during neurodegenerative diseases and the associated common pathological symptoms

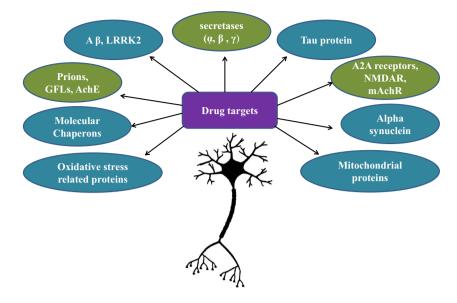
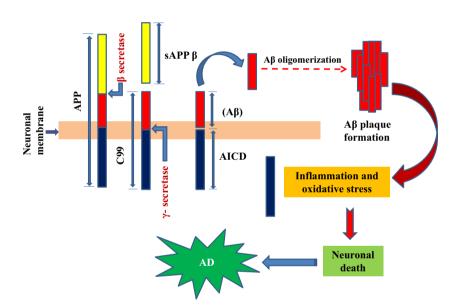


Fig. 3 Amyloidogenic pathway of Alzheimer's disease: Beta secretase cleaves Amyloid precursor protein into soluble amyloid precursor protein-beta (APP- β) and C99 fragment. The later in the presence of gamma secretase cleaves to amyloid beta and AICD fragment. Amyloid beta A β undergoes oligomerization and plaque formation leading to inflammation, oxidative stress and finally neuronal death



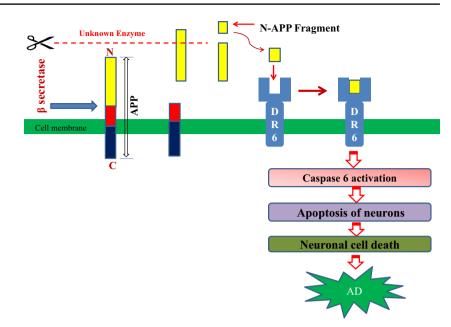
with transcription factors like nuclear factor kappa-light-chain-enhancer of activated B cells (NF-kappa B), Activator protein 1 (AP-1) and tumour protein p53 (Tan et al. 2009a, b). Amyloid beta may also induce oxidative damage, primarily through mitochondria also affects the cholesterol equilibrium, promoting neuronal toxicity due to generation of highly reactive free radical species (Harris et al. 1995; Colell et al. 2009). Overall, it seems $A\beta$ plays crucial role in inducing neurotoxicity through mechanisms like triggering plaque formation, neuroinflammation, apoptosis as well as oxidative stress.

An updated version of amyloid hypothesis came in 2009 which suggests that amyloid beta protein may not be the real culprit of disease but a close relative of it namely N-amyloid precursor protein (N-APP). Accordingly, amyloid precursor

protein (APP) is enzymatically cleaved by an unknown mechanism at its N terminal to generate N-terminal amyloid precursor protein (N-APP) fragment. This fragment triggers the self destructive pathway by binding to a neuronal receptor known as death receptor 6 (DR6). This receptor is expressed majorly in developing neurons of brain and is required for normal cell body death and axonal pruning both in vivo and after trophic factor deprivation in vitro. N-APP fragment acts as a ligand and interacts with DR6 causing neurodegeneration via activation of caspase-6 enzyme contributing to the pathology of Alzheimer's disease (Nikolaev 2009). Therefore, N-APP/DR6/caspase 6 signalling pathway (Fig. 4) represents an important cascade implicated in AD pathophysiology that might be targeted and hijacked in the ageing AD brain. The strategy to either prevent the



Fig. 4 Demonstration of N-APP-DR6-Caspase-6 pathway of Alzheimer's disease AD. Beta secretase cleaves amyloid precursor protein (APP), generating two fragments. Then some unknown enzyme cleaves one of the fragments towards N terminal of APP to generate N- terminal fragment amyloid precursor protein known as N-APP fragment. The later binds to death receptor-6 (DR6) located in the neuronal membrane. N-APP/DR6 interaction triggers the apoptotic pathway by activating caspase-6 that causes neuronal death leading to Alzheimer's disease



formation of N-APP fragment or block its interaction with death receptor 6 (DR6) may have better therapeutic outcome for AD.

Tauopathy has been long debated in the pathogenesis of AD. Studies based on biomarker analysis show tau protein is more closely linked to AD progression than Aβ. Tau protein is an axonal protein that regulates microtubule stability (Johnson and Hartigan 1999). Normally, tau protein when phosphorylated interacts with and stabilises the microtubules, therefore also known as microtubule-associated protein (MAP). During AD, Tau protein undergoes certain biochemical changes becoming hyperphosphorylated. Hyperphosphorylated form of tau protein dissociates from the microtubules and then aggregates to form neurofibrillary tangles (NFTs) inside nerve cell bodies (Mudher and Lovestone 2002). NFTs disintegrate microtubules within neuronal cells thus destroying their cytoskeleton and ultimately damaging neuronal transport system (Iqbal et al. 2005). Tau protein contains at least forty-five phosphorylation sites, majority of them lie in the C-terminal tail region (residues 368-441) and proline-rich region (residues 172-251) (Hanger et al. 2009). Phosphorylation at C-terminal of tau protein plays more important role so far as AD is concerned. Phosphorylation at Ser262 selectively disrupts binding of tau to microtubules (Fischer et al. 2009), while phosphorylation at Ser202 promotes tau aggregation to form NFTs (Rankin et al. 2005). It seems that phosphorylation at multiple sites of tau protein rather than at single site plays a critical role in the pathology of Alzheimer's disease (Alonso et al. 2001). Abnormal phosphorylation of tau protein is also implicated in increasing the activity of certain kinases notably glycogen synthase kinase 3-beta (GSK-3β), mitogen-activated protein kinase (MAPK) p38, stress-activated protein kinases, cyclin-dependent kinase-5, p42/44 MAP kinase and mitotic protein kinases; while decreasing the activity of some other enzymes like protein phosphatases (PPs) such as PP1, PP2a, PP2b and PP5 (Mondragón-Rodríguez et al. 2012). Protein phosphatases PP1, PP2A, PP2B and PP5 have been found to de-phosphorylate tau protein at Ser199, Ser202, Thr205, Thr212, Ser214, Ser235, Ser262, Ser396, Ser404 and Ser409. Among all the mentioned PPs, PP2A is considered to be the strongest protein phosphatase (Liu et al. 2005). These entire phenomenons initially hamper the biochemical communication between neurons and later cause neuronal death. Therefore, research should focus on developing drugs with ability to block the abnormal phosphorylation of tau protein which in turn could prevent microtubular destruction as well as neuronal death.

A recent study by Koren and co-workers showed tau protein interacts with ribosomes and aids translation between global transcripts and 5' Terminal oligopyrimidine (5'TOP) mRNA sequences. But in diseased condition, abnormal tau proteins diminish the effectiveness of translation and foil phosphorylation of ribosomal protein S6 (rpS6) at S240/S244. This reduced activity of rpS6 in turn impairs global translation since the translation of 5'TOP mRNAs coding for ribosomal and translational machinery is affected/reduced (Koren et al. 2019). Thus, there may be some definite link between tau pathology and ribosomal dysfunction in AD which needs to be investigated in future studies.

In addition, studies targeting tau-ribosomes interaction are warranted in exploring effective treatment regimens. Besides phosphorylation, tau protein may also undergo other modifications such as lysine ubiquitylation, lysine acetylation, O-linked N-acetylglucosamine (O-GlcNAc) modification, lysine dimethylation and arginine monomethylation



(Du et al. 2018). Nevertheless under pathological conditions, hyperphosphorylation renders tau protein more prone to aggregation thus reducing its affinity to neuronal microtubules and affecting the plasticity of neurons.

Recent research also highlights the role of Apolipoprotein E (apoE) in the pathophysiology of Alzheimer's disease (Liraz et al. 2013). The ApoE protein encoded by apoE gene is produced by astrocytes in the brain. It is an important protein that transports lipids. ApoE has three alleles namely apoE2, apoE3 and apoE4. ApoE protein is composed of 299 amino acids and its isoforms differ only at amino acid location 112 and 158, which may be arginine or cysteine (Liu et al. 2013). This minor difference is enough to alter the tertiary structure of ApoE protein which disrupts its ability to interact with lipids, $A\beta$ and receptors. Among all the known isoforms of ApoE protein, ApoE4, a hypolipidated version of the protein has been linked to an increased risk of developing Alzheimer's. ApoE4 allele shows higher percentage in families with sporadic and late-onset Alzheimer's. Patients with elevated levels of ApoE4 reportedly have an elevated level of accumulated Aβ, hyperphosphorylated tau, reduced neural plasticity and neuropathology (Liraz et al. 2013). Since there occurs accumulation of Aβ in apoE4positive patients, this led researchers to believe that the two cascades interact with each other and the cross talk between them causes the pathological effects. Occurrence of apoE4 allele negatively hampers the interaction of ApoE4 to lipids thus leading to the proposal that the disease caused due to lipid-related mechanisms (Liraz et al. 2013). ApoE4 not only causes the formation of senile plaques, but also obstructs the body's means to eliminate it. Normally, the Amyloid plaques and NFTs are removed from the brain through Autophagy (Simonovitch et al. 2016). Under normal conditions, astrocytes of central nervous system (CNS) produce apoE, which protect the brain from harmful protein buildup. Recent research shows that the presence of the apoE4 allele disrupts body's mechanisms of naturally clearing protein build up through the process of autophagy (Rubenstein 2017). Researchers need to develop drugs with ability to reduce the levels of apoE4 that may allow normal autophagy and prevent abnormal accumulation of proteins in neurons, hence reduce neurotoxicity.

Environmental factors are also implicated in the pathogenesis of various neurodegenerative diseases including Alzheimer's disease. Many natives of the Island of Guam known as Chamorro suffer from a debilitating neurodegenerative disease with symptoms similar to multitude of neurodegenerative diseases including Alzheimer's disease (Cox et al. 2016). Persons with this mystery disease showed the presence of Amyloid- β plaques and neurofibrillary tangles in their brains. The non-native people who adopted Chamorro lifestyle also got affected with this disease. The cause of this disease is believed to be

an environmental toxin. Initially, there was difficulty in identifying the nature of this neurotoxin because it took years for symptoms to appear after the exposure. Finally, researchers achieved success in isolating this toxin from the seeds of cycad and it was named β-N-methylamino-L-alanine (BMAA). Chamorro people used cycad seeds for making flour, a staple food for them. During 1980s, this toxin was given to macaques which developed acute neurological symptoms. Later in 1990s, Paul Alan Cox, an ethnobotanist proposed that BMAA causes ALS/PDC seen in the natives (Holtcamp 2012). According to him, BMAA is synthesised by cyanobacteria that live in symbiotic relation with the roots of cycad. BMAA gets accumulated in the gametophytes of the cycad seeds, that when ground to flour is consumed by Chamorro people. Later cox showed that BMAA also reaches Chamorro people via animals such as feral deer, flying foxes and pigs who feed on cycad seeds. BMAA undergoes ten thousand times biomagnifications in these animals and thus gets accumulated in the food chain that results in neurological symptoms in Chamorro people. Cyanobacteria that produce BMAA live in many marine ecosystems thus its biomagnifications take place in many marine dwellers like sharks, shellfish and bottom-dwelling fish. The people who consume these animals get BMAA accumulation in their brain and hence develop neurological problems. BMAA can easily cross the blood-brain barrier and reach brain. Cellular mechanism mistakenly takes BMAA for L-serine and incorporates it into proteins thus leading to misfolding, aggregation and apoptotic cell death (Cox et al. 2016). Normally, proteins have an interior hydrophobic part and an exterior hydrophobic part but incorporation of BMAA leads to conformation changes making the exterior part hydrophobic and the interior structure hydrophilic. These external hydrophobic regions of proteins are highly sticky and therefore clump together to form small neurodegenerative protein aggregates. These small aggregates oligomerise to form larger ones that impair cellular functions (Holtcamp 2012). BMAA also promotes phosphorylation of tau protein by causing activation of mGluR5, a metabotropic glutamate receptor, which reduces the activity of protein phosphatase 2A (PP2A). There is a significant decrease of PP2A in Chamorro ALS/PDC brains that increase phosphorylated tau (Cox et al. 2016). Research shows that dietary exposure to BMAA causes deposition of A β and formation of neurofibrillary tangles thus acting as an initiator for sporadic Alzheimer's disease. There is possibility to use L-serine as a therapeutic tool for treatment of AD since it may reduce formation of plagues and NFTs in brain tissue.

Overall, AD seems to be a syndrome rather than a disease since its pathophysiology involves multiple possible models and thus there is requirement of a multifaceted therapeutic



approach striking multiple targets in AD for its better clinical and therapeutic outcome. This can be achieved by focusing on 'magic shotgun drugs' rather than 'magic bullet drugs'.

Targets for Alzheimer's Disease

Amyloid Beta

Aggregation of soluble monomeric form of Aβ to form different states of soluble aggregates like dimers, oligomers and polymers is looked as an important factor that causes synaptic loss and cognitive impairment (Lue et al. 1999). Clearing off these toxic aggregates is an imperative therapeutic approach; however, there are certain challenges for instance developing drugs to disintegrate Aβ aggregates can lead to the accumulation of Aß monomers in the brain thereby causing neurotoxicity; also the strategy to block the formation of Aβ aggregates can lead to the accumulation of monomeric structures that will again trigger neurotoxicity. Nowadays, the best alternative to deal this concern is the development of immunity against Aß plaques that will completely clear them from brain. Working on same lines, Aβ immunisation performed in mouse model efficiently cleared Aß plaques from brain but could not improve cognition (Rinne et al. 2010; Wang et al. 2011). In another study, patients suffering from mild-to-moderate AD when vaccinated with Aß during phase II of clinical trial underwent inflammatory response in CNS (Kuzuhara 2010). However, this inflammatory response has not been fully proven to be responsible for the failure of the anti-Aβ therapy. Janssen and Pfizer's bapineuzumab were the first attempts to obtain humanised monoclonal antibodies (mAB) with capability to clear-off the amyloid beta from the brain. This drug was tried for thousands of patients with mild-to-moderate AD (Gravitz 2011) but failed to prove its therapeutic potential in phase III of clinical trial. It could not improve the cognition, functional ability and clinical outcome in patients with AD. Research is still going on to find effective antibody against alzheimer's and in this regard a total of five different anti-Aβ antibodies namely crenezumab, gantenerumab and aducanumab, solanezumab and BAN2401 are being currently tried in preclinical stage of familial AD and asymptomatic individuals at high risk of AD (Salloway et al. 2014; Hung and Fu 2017; Francesco et al. 2019). Gantenerumab, an entirely human recombinant IgG1 monoclonal antibody interacts both with the middle regions as well as the amino-terminal of A β . As compared to A β monomers, Gantenerumab exhibits higher affinities for $A\beta$ oligomers and $A\beta$ fibrils (Bohrmann et al. 2012). This point is supported by the fact that gantenerumab does not change the plasma Aβ levels in vivo. Intravenous administration of gantenerumab was found to decrease brain Aβ plaques (Jacobsen et al. 2014). However, the cognitive effects of gantenerumab still need to be investigated. Currently, the drug is being investigated in phase II/III trial. Aducanumab, a recombinant human IgG1 monoclonal antibody targets the amino- terminal residues 3-7 of Aβ (Arndt et al. 2018). Aducanumab shows 10,000-fold greater selectivity for soluble Aβ aggregates and insoluble fibrils than for Aβ monomers. Administration of the drug decreased Aß plaque size in a dose-dependent fashion in young (9-month old) Tg2576 mice but not in aged (22-month old) (Kastanenka et al. 2016). Presently, there are no reports on the cognitive or behavioural effects of Aducanumab. Crenezumab, a humanised anti-Aβ monoclonal IgG4 antibody interacts with multiple species of Aβ, especially fibrillary 16mer assemblies of aggregated Aβ and pentameric oligomeric Aβ (Adolfsson et al. 2012; Zhao et al. 2017). This drug promotes disaggregation and blocks aggregation of Aβ (Ultsch et al. 2016). When injected in Tg2576 mouse model of AD, the drug exhibited no notable inflammatory changes (Fuller et al. 2015). There is deficit of studies pertaining to the cognitive effects of crenezumab in animal models of AD. Currently, the drug is being tested in a double-blind, placebo-controlled clinical trial which may be completed in March 2022 (Tariot et al. 2018). Solanezumab, a humanised IgG1 monoclonal antibody recognises the central region of A β (A β 13–28). Studies carried out in mice and human models indirectly indicate that solanezumab targets the soluble monomeric $A\beta$ because it caused a considerable increase in the levels of total plasma A\u03c3. Furthermore, studies carried in human brain tissue suggest that solanezumab recognises to Aβ plaques (Bouter et al. 2015). The mouse version of solanezumab known as m266 when injected into the PDAPP transgenic mouse model of AD reversed the memory deficits without affecting brain amyloid plaques (Dodart et al. 2002). There is a possibility that the antibody could target the soluble pool of A β (DeMattos et al. 2001). The drug might come out as a potent anti-Aβ regimen in future. BAN2401, another monoclonal antibody specifically binds to soluble Aβ protofibrils (Logovinsky et al. 2016). The antibody showed promising results in an 18-month adaptive phase IIb study carried out in 856 persons with mild or prodromal form of AD. When administered at a higher dose of 10 mg twice a week, the drug caused a significant reduction of brain Aβ accumulation compared to placebo (Osswald 2018) thus indicating the drug could prove effective therapeutic regimen in future.

In addition to the above-mentioned antibody regimens, active anti-Aβ vaccine (antigen) namely CAD106 has also been developed. The vaccine consists of many copies of Aβ1–6 fragment attached to a virus-like particle (an adjuvant carrier). Administration of CAD106 decreased brain amyloid accumulation in two APP transgenic mouse lines without any evidence of inflammatory reactions or increased microhemorrhages (Osswald 2018). In cell cultures, antibodies generated in response to CAD106 administration



interacted with A β monomers and oligomers and blocked amyloid toxicity (Winblad et al. 2012). Still there is no information available on cognitive effects of CAD106 in animal models of AD. At present, CAD106 is in phase III clinical trials and may come with promising outcome in the near future.

Tau Protein

There are various approaches to target tau protein toxicity notably (I) manipulating phosphatases and kinases, (II) tau vaccination, (III) stabilising microtubules and (IV) blocking tau aggregation. Earlier tau aggregation blocker namely TRx0237 was tested but it failed in phase III clinical trials (Gauthier et al. 2016). Currently, inhibitors like methylene blue and curcumin are being tested. Methylene blue is one of the promising Tau polymerization inhibitors that block the in vitro tau aggregation by trapping monomeric tau in an aggregation-incompetent conformation (Wischik et al. 1996; Panza et al. 2016). In transgenic mouse models of tauopathy, this drug effectively reduced tau pathology and also enhanced cognition (Panza et al. 2016). Methylene blue passes through blood-brain barrier and has been used long in humans; hence, it could serve a promising therapeutic candidate in future. Curcumin, a plant-derived compound isolated from *Curcuma longa* is another tau polymerization inhibitor that interacts with proteins in β -sheet conformation and block aggregation. The compound has been found to decrease tau as well as AB pathology and improve cognitive impairment in animal models (Hamaguchi et al. 2010; Hu et al. 2015). Currently, curcumin is being tested in an 18-month-long phase II study (https://clinicaltrials.gov/ show/NCT01811381). In addition, another clinical trial based on combined study of curcumin and exercise is also being recruited (Fitzgerald et al. 2012). Both these trials will assess patients for biomarkers for AD biomarkers as well as for changes on MRI and PET scans. Many other drugs targeting tau aggregation without interfering the stabilisation of microtubules have also been introduced notably emodin, daunorubicin, adriamycin, PHF016 and PHF005 (Pickhardt et al. 2005). Targeting kinases and phosphatases is also looked as a potential candidate for treating neurodegenerative disorders especially AD (Gong and Iqbal 2008). Recent studies have shown sodium-selenate as a possible drug that could treat neurodegenerative disorders by stabilising PP2A in transgenic AD models (Van-Eersel et al. 2010). Moreover, neuroleptics such as trifluoperazine, clozapine and chlorpromazine have also been developed to treat AD (Gong et al. 1996). Among microtubule stabilising drugs, Abeotaxane (TPI 287) is currently being studied in two clinical trials (https://clinicaltrials.gov/show/NCT01966666; https://clini caltrials.gov/show/NCT02133846). Abeotaxane has shown the ability to pass through blood-brain barrier in mouse model (Fitzgerald et al. 2012). So far as vaccine therapy is concerned, IVIG, the only passive vaccine (Intravenous immunoglobulin) was tried in patients with mild-to-moderate AD but failed in phase III trials (Du et al. 2018). At present, two passive vaccines namely RG6100 and ABBv-8E12; and two active vaccines namely ACI35 and AADvac-1 are being tried in phase I and II clinical trials.

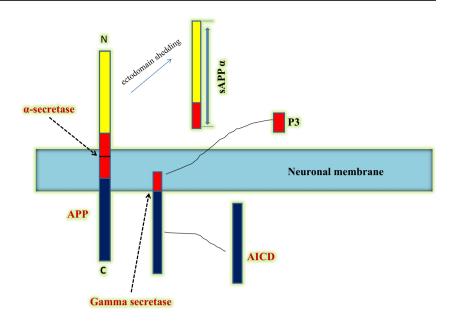
Currently, the preponderance of tau-targeting therapies being tested in trials includes immunotherapies, which have shown promising results in several preclinical trials and may prove effective therapeutics in the near future (https://clini caltrials.gov/show/NCT01811381). C2 N-8E12 is an antibody that targets amino acids of the tau protein. The drug effectively blocks the pathological tau seeding triggered by exogenous tau aggregates when infused into the brain of transgenic mouse model of tauopathy. C2 N-8E12 diminished the hyperphosphorylated tau and aggregated and also improved cognition (Kfoury et al. 2012; Erin and Congdon 2018). Analogous results were observed when the drug was incorporated systemically. Studies showed no evidence of any adverse immune reaction and furthermore drug administration resulted in the reduction of microglial activation (Yanamandra et al. 2013, 2015). The drug showed promising results in the first phase of clinical trial and is currently being tested in second phase (Congdon et al. 2014; West 2016; Barthélemy et al. 2016). RO 7105705, an antibody most likely recognises pSer409 on the tau protein (Giacobini and Gold 2013). The antibody has not been disclosed so far and is currently being tested in phase I clinical trial. LY3303560 is an anti-tau antibody that possibly recognises a conformational epitope, although this information has not been disclosed so far. Eli lilly has started two phase I clinical trials to test the pharmacokinetics and safety of LY3303560. One trial will be carried in healthy persons and patients with AD, and other trial in patients with MCI or AD (Erin and Congdon 2018). These two trials are expected to reach completion in few years. In the recent times, two new antibodies namely JNJ-63733657 and UCB0107 are also entering the clinical trials. JNJ-63733657 possibly targets the middle region of tau, while UCB0107 this seemingly targets amino acids 235-246 in the proline-rich region of tau (Rogers 2018; https://clinicaltrials.gov/ct2/show/NCT03375697). The field of tau immunotherapy is progressing rapidly and many new trials may come into lime light in the near future.

Alpha Secretases

As far as the role of alpha secretase is concerned it is not involved in the generation of amyloid beta. In fact, α -secretases participate in non-amyloidogenic pathway (Fig. 5) to cleave APP within the amyloid sequences generating a soluble APP ectodomain (sAPP α) through the process termed as "ectodomain shedding". Function of P3



Fig. 5 Non-amyloidogenic pathway of Alzheimer's disease: Alpha secretase cleaves APP to generate the neuroprotective fragment $_{\rm S}{\rm APP}_{\alpha}$ towards N terminal via ectodomain shedding. The other remaining fragment of APP fragment is cleaved by gamma secretase into two subfragments P3 and amino-terminal APP intracellular domain (AICD)



fragment generated due to the action of gamma secretase in non-amyloidogenic pathway is not clear. Soluble amyloid precursor protein-alpha (sAPPα) is believed to be neuroprotective in the perspective of AD since it hinders amyloid beta peptides formation (Furukawa et al. 1996). sAPPα prevents cultured neurons against the oxygen-glucose deprivation as well as excitotoxicity by blocking calcium currents, enhancing potassium currents and thus stabilising the resting membrane potential. α -secretase activity is mediated by one or more enzymes from the family of disintegrin and metalloproteinase domain proteins (ADAM) with ADAM 9, ADAM 10, ADAM 17 and ADAM 19 being the most probable candidates (Chow et al. 2010). Moreover, alpha secretase causes proliferation of embryonic neural stem cells and promotes synaptic density and improves memory retention (Vella and Cappai 2012). Drugs that enhance alpha secretase activity are considered as promising therapeutics to tame AD. Various drugs acting as indirect α -secretase activators are being tried in different phases of clinical trials to find a treatment for AD. For instance etazolate (EHT 0202), the GABA receptor modulator previously proven to stimulate sAPPα production and protect against Aβ induced toxicity in rat cortical neurons has reached the Phase II of human clinical trials (Marcade et al. 2008; Vellas et al. 2011). PRX-03140, the 5-HT4 agonist proven to stimulate the activity of α -secretase although showed positive results in phase II of clinical trials in 2008 but was not examined further (Sabbagh 2009). Epigallocatechin-gallate (EGCG), a polyphenolic compound obtained from green tea has also proven to act as prominent stimulator of α -secretase through the protein kinase-C (PKC) pathway besides reducing cerebral amyloid deposition in the brain of AD mice. Currently, the efficacy of this drug is being evaluated in phase II/III clinical trial investigations (MacLeod et al. 2015). Recent reports from "Nature Medicine" show that the activity of necrosis factor-converting enzyme (TACE), the stimulator of α-secretase is decreased both in AD patients as well as on the surface of the neurons isolated from mouse model of AD. In addition they also observed an increased activity of 3-phosphoinositide-dependent protein kinase 1 (PDK1) enzyme in the brains of AD patients. It is believed that the later prompts the internalisation of TACE and thus disrupts TACE-mediated α-secretase activity. By blocking the enzyme PDK1 in mouse models using BX912 (PDK1 inhibitor), researchers observed a considerable decrease in the formation of amyloid plagues with an elevation in the levels of sAPPα (Pietri et al. 2013). These results indicate that inhibition of PDK1 enzyme could serve a promising therapeutic approach in future and therefore efforts to develop PDK1 inhibitors may serve a promising therapeutic regimen.

Beta Secretase

Beta secretase enzyme playing an important role in $A\beta$ formation is also looked as a possible target to tame neurological disorders but is associated with serious side effects. A study analysing the effect of BACE1 knockout in mice demonstrated significantly higher mortality rate in mice in their 1 weeks of life (Dominguez et al. 2005). Thus, strategy to block this enzyme could prove fatal. Moreover, this enzyme has many important/alternative substrates including the proteins involved in sodium homeostasis and myelination (Wong et al. 2005; Hu et al. 2006). These studies confirm a higher risk of morbidity with the use of BACE1 inhibitors. Now, there arises a question whether the positive modulatory effects of BACE1 inhibitors outweigh its ill effects in



the elderly aged people or not. A deep discussion among scientists has lead to an interesting assumption that the partial inhibition rather than complete inhibition of BACE1 could prove beneficial in reducing the levels of A_B. Various BACE1 inhibitors have been tried in clinical trials but had to be abandoned due to their toxicity. For instance, Eli Lily's LY2886721 exhibited an initial decrease in the levels of Aβ in plasma but showed liver toxicity due to which its clinical trial was stopped immediately after phase II. Similarly, Eisai's E2609 demonstrated a considerable reduction in plasma Aß levels during Phase I of clinical trial but its Phase II trials have not been tried yet (Wolfe 2012; Folch et al. 2018). In an another attempt, the development of HPP854 by TransTech Pharma and RG7129 by Roche had to be stopped immediately after phase I clinical trial due to some undisclosed reasons. The ongoing clinical trial sponsored by Merck Sharp and Dohme Corp evaluates the efficacy of Verubecestat in patients with prodromal Alzheimer's disease. This clinical trial (NCT01953601) is expected to generate its outcome in February 2019. In addition, Astra-Zeneca and Eli Lilly and Company have joined their hands together for co-developing lanabecestat (AZD3293). The crucial Phase II/III clinical trial of lanabecestat started in late 2014 and is intended to recruit 1500 patients and will probably end in May 2019 (https://www.reuters.com/artic le/2014/12/01/health-alzheimers-astrazeneca-eli-lilly-idUSL 6N0TL0ST20141201).

Gamma Secretase

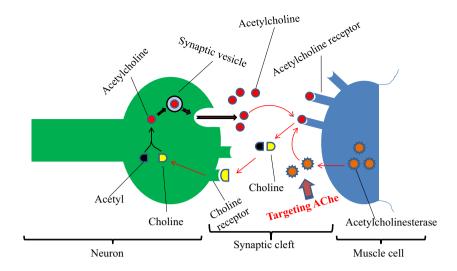
Gamma secretase, although a much anticipated target for treating AD, has been shown to cleave a wide range of substrates. The most prominent substrates for this crucial enzyme include Notch, an important cell surface receptor pivotal for cellular development and differentiation, whose role is also implicated in tumour suppression (Sorensen and Conner 2010). Inhibition of γ -secretase has been found associated with potential toxic effects, for example, Eli Lily's semagacestat (LY450139) was abandoned after phase III due to its failure to stop disease progression and worsening the cognition (Doody et al. 2013). Studies also warn about the side effects like skin cancer due to γ -secretase inhibition, most probably due to concomitant inhibition of Notch signalling. Bristol-Myers-Squibb's Avagacestat (BMS708163) was abandoned after Phase II trials due to side effects like worsening of cognition and the risk of skin cancer (Coric et al. 2012; Golde et al. 2013). Here it is worth mentioning that even the Notch-sparing inhibitor ELND006 developed by Elan Corporation when subjected to clinical trial was found to cause liver toxicity and was abandoned (Hopkins 2011). Another drug, Tarenflurbil used as γ-secretase modulator although showed promising outcome during phase II clinical trial, was discontinued due to poor response in third phase of clinical trial (Green et al. 2009). Similarly, CHF-5074 (γ -secretase modulator) developed by Chiesi successfully reached phase II of clinical trial (NCT01303744) but was abandoned due to some hidden reasons. The futuristic strategy of γ -secretase inhibition will largely depend on the development of APP-specific γ -secretase inhibitors with property to spare Notch signalling.

Acetylcholinesterase

Acetylcholinesterase (AChE) belongs to carboxylesterase family of enzymes. It is the primary cholinesterase in the body that catalyses the breakdown of neurotransmitters such as acetylcholine and other choline esters. Acetylcholinesterase is a key enzyme in the cholinergic nervous system synthesised in the muscle cell and then released into the neuromuscular junction/synaptic cleft where it functions as the terminator of synaptic transmission. AChE cleaves acetylcholine molecules into acetyle and choline. Later, the choline re-enter the neuronal cell where it is re-used to synthesise acetylcholine (Fig. 6). During the pathology of AD, there is deterioration of many cholinergic neurons in the brain which results in the profound decline of acetylcholine (Du et al. 2018). Thus, AD patients produce less acetylcholine (compared to normal person), which hampers the normal neurotransmission and thus producing clinical manifestations of AD like memory loss. The strategy to inhibit the function of acetylcholinesterase using acetylcholinesterase inhibitors is looked as a promising approach to raise the level of acetylcholine and promote continued stimulation of the muscles and glands that potentiate the parasympathetic activities such as constriction of pupils of the eyes, vasodilatation, increased production of saliva, sweat, and tears, slow heart slow heart rate, constriction of bronchioles and mucus secretion in the respiratory tract (Samii et al. 2004). Drugs like donepezil, galantamine and rivastigmine are commonly used as AChE inhibitors for providing symptomatic relief (Jacobsen et al. 2005; Mehta et al. 2012). There has been a continued search for obtaining efficient inhibitors of acetylcholinesterase and thus researchers have explored many new/modified synthetic and natural compounds/drugs with potential anti-acetylcholinesterase activity. The presence of peripheral anionic site (PAS), in addition to the catalytic site (CS) on acetylcholinesterase, has been implicated in promoting formation of amyloid fibril and its co-localization. Novel flavonoid derivates capable of binding to both the PAS and CS site of acetylcholinesterase have been designed and can inhibit acetylcholinesterase much better than conventional rivastigmine and donepezil (Sheng et al. 2009). Khan and co-workers used chemical as well as computation-based approaches to evaluate the derivatives of four flavonoids namely quercetin, rutin, kaempferol and macluraxanthone. They found macluraxanthone and quercetin derivatives as



Fig. 6 Prolonged activation of acetylcholine esterase caused decline of acetylcholine (neurotransmitter). The strategy to inhibit acetylcholinesterase using acetylcholinesterase inhibitors is looked as a promising approach to raise the level of acetylcholine and promote continued stimulation of the muscles and glands



very good inhibitors of acetylcholine esterase (Khan et al. 2009). Various modified novel carbamates tested in silico and in vitro have been found to possess prominent AChE inhibitory activity (Roy et al. 2012). In addition, pyridopyrimidine, a novel natural compound has shown greater AChE inhibitory action than the conventional drug galantamine (Kumar et al. 2016). The hybrids of donepezil and aminopyridine namely pyridonepezil and 6-chloro-pyridonepezil, respectively, have been found to be more potent inhibitors of AchE than the single donepezil molecule (Varadaraju et al. 2013). These molecules are considered as the dual inhibitors of AchE that bind to both the PAS and CS site of acetylcholinesterase. Developing such therapeutics with better efficacy and safety could serve as a promising strategy in ameliorating AD.

N-methyl-D-aspartate Receptor

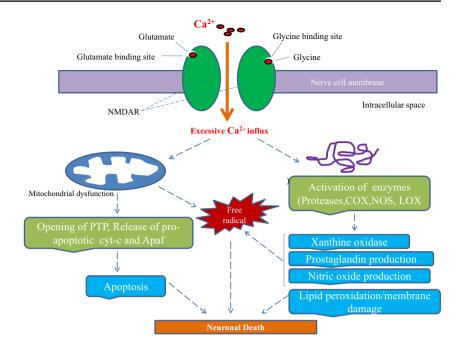
N-methyl-D-aspartate receptor also called as the NMDA receptor or NMDAR is an ion channel protein and a glutamate receptor present in the membrane of nerve cells. NMDAR is activated when glutamate and/or glycine (or D-serine) binds to it; and in its activated form NMDAR permits the flow of positively charged ions through the cell membrane of nerve cells. This receptor is very important for controlling synaptic plasticity as well as the memory function. During Alzheimer's disease, there is hyperactivation of NMDAR due to glutamate binding which leads to excessive influx of Ca²⁺ ions through it (Lipton 2005). Here it is worth mentioning that glutamate acts as the most important excitatory neurotransmitter in the brain and glutamate-mediated synaptic transmission is vital for the normal functioning of the nervous system (Dingledine et al. 1999). Calcium influx triggers hyperactivation of enzymes including the phospholipases, cyclooxygenase-2 (COX-2), lipoxygenases (LOX), proteases and nitric oxide synthase (NOS) (Tan et al. 2009a, b; Ezza and Khadrawy 2014). Glutamate-mediated overactivation of NMDAR and subsequent Ca²⁺ influx is associated with mitochondrial dysfunction, free radical generation and neurodegeneration (Fig. 7).

Activated phospholipases cause hydrolysis of essential phospholipids from neuronal membrane and subsequent accumulation of free fatty acids (FFAs), diacylglycerol (DAG), platelet activating factor (PAF's) and lysophospholipids. FFAs, DAG and lysophospholipids have detergentlike effect on normal membranes and can uncouple oxidative phosphorylation thereby producing changes in membrane permeability. PAF is pro-aggregatory and may cause adhesion and activation of leukocytes and hence produce an inflammatory reaction at endothelial cell surface. FFAs will trigger arachidonate pathway and generate inflammatory mediators including prostaglandins, leukotrienes, leukotrienes and thromboxanes. Prolonged activation of arachidonate pathway triggers increased production of free radicals, lipid peroxidation and hence oxidative damage to membrane-bound proteins. Phospholipase mediated phospholipid degradation also sets stage for calcium influx and PKC activation, which has deeper implications in neurodegeneration (Asaoka et al. 1992).

Activated lipoxygenases (LOXs) catalyse oxidation of arachidonic (AA) to various bioactive lipids implicated in neurodegeneration. Briefly, Lipoxygenases (LOXs) oxidise AA to hydroperoxyl derivatives including 5-hydroperoxyeicosatetraenoic acids (HPETEs). These derivatives upon reduction form corresponding hydroxyeicosatetraenoic acids (HETEs) and leukotriene (LT) via 5-lipoxygenase, lipoxins and hepoxilins. LOXs peroxidize membrane lipids and lead to structural changes in the cell. 15-LOX-1 is the major enzyme which is responsible for membrane lipid peroxidation (Brash 1999). 5-HPETE inhibits synaptosomal membrane Na⁺, K?-ATPase activity (Foley 1997).



Fig. 7 Glutamate induced activation of NMDAR, calcium influx, free radical generation and subsequent neuronal death



Activated COX-2 oxidises dopamine to a highly reactive form dopamine-quinone which can react with cysteinyl residues of target proteins, and transform them. Transformation could in turn alter the function of proteins leading to cell death. COX2 also increases production of prostaglandin E2, which in turn leads to increased production of reactive oxygen species (Bazan 1999).

Activated protease converts xanthine dehydrogenase to xanthine oxidase. The latter is a key enzyme in the production of free radicals such as hydroxyl and superoxide. The brain is highly sensitive to free radicals since it lacks normal free radical scavengers. Free radical acts on the phospholipid of neuronal membrane and destroys them. Ca²⁺ also activates NOS which subsequently increases nitric oxide production. The latter is a gaseous free radical and mediates the excitotoxicity and neurodegeneration (Gagliardi 2000).

Excess of Ca²⁺ influx due to NMDAR overactivation also triggers opening of mitochondrial permeability transition pore (PTP) through which cytochrome c (Cyt c) and apoptosis-inducing factor (AIF) comes out into cytoplasm and stimulates apoptotic pathway leading to programmed cell death (Hibiki and Giovanni 2017). Prolonged calcium influx through continuous opening of NMDA receptors disrupts mitochondrial membrane potential, disrupts the oxidative phosphorylation thus reducing ATP production and making cell more susceptible to death insults (Fiskum et al. 2003). Due to disruption of energy metabolism, the energetically compromised neurons become depolarized and cannot maintain ionic homeostasis (Zhou and Danbolt 2014).

Drugs/molecules which act as antagonists of NMDA receptor are therapeutically considered beneficial in many neurological disorders like dementia, stroke and neuropathic

pain syndromes (Gitto et al. 2014). At present, memantine is the only drug marketed as NMDA receptor antagonist. A wide range of molecular docking studies are being carried out to obtain novel/active ligands against this receptor in AD. Examples of some prominent ligands confirmed by docking experiments include triazolylamidine derivatives (Espinoza-Moraga et al. 2012); 3-substituted-1H-indoles (Colotta et al. 2012); phenyl-amidine; 1-benzyl-1,2,3,4tetrahydro-b-carboline (Parson et al. 2013); and 3-hydroxy-1H-quinazoline-2, 4-dione derivatives (Abreu et al. 2013). The amino acid glycine binds to NR1 subunit of NMDA receptor and has been identified as a co-agonist of NMDA. Studies are being conducted to find out the molecules that could block the glycine binding NR1 subunit of NMDA receptor (Krueger et al. 2009). In addition, studies based on molecular docking have revealed that ifenprodil and similar compounds posses strong activity in blocking NR2B unit of NMDA (Gitto et al. 2008).

Muscarinic ACh Receptors

Muscarinic acetylcholine receptors (mAChR) form one of the G-protein-receptor complexes in the cell membranes of certain neurons and other cells and act as ACh receptors at various locations including the central nervous system (CNS). They are stimulated by neurotransmitter Ach, released from postganglionic fibres in the parasympathetic nervous system (PNS). mAChR are of five types represented as M1-M5 and are implicated in motor control and learning process. The M1-type mAChR present in the cerebral cortex and hippocampus play important role in memory, cognitive processing and learning which become impaired



in Alzheimer's disease (Caulfield and Bridsall 1998). These cholinergic deficits observed in AD can be restored via cholinergic activation, which can be done by the use of muscarinic agonists. Over many years, M1 subtype agonists/drugs such as AF102B, AF150, AF267B and AF292 have been tried in AD patients. Among these compounds AF267B has been found to have excellent pharmacokinetic profile and can also cross the blood-brain barrier, whereas the compounds namely AF150(S), AF102B and AF267B are associated with neurotrophic effects, elevated non-amyloidogenic APP and decreased Aβ (Davie et al. 2013). In Alzheimer's disease, formation of amyloid beta reduces the potential of mAChR receptors to transmit signals, thereby leading to diminished cholinergic activity. Reports suggest that the strategy to activate M1 mAChRs could ameliorate AD pathology besides restoring cognitive functions and decreasing hyperphosphorylated form of tau protein (Foster et al. 2014). Here, it is worth mentioning that even though a few M1 agonists improved cognition in the initial phase, but failed after reaching clinical trial due to their non-specific nature (Melancon et al. 2013). 77-LH-28-1, an M1 allosteric candidate from GlaxoSmithKline (Harlow, UK) has shown great CNS penetration and promising pharmacological profile. In addition, two other M1 selective agonists namely VU0364572 and VU0357017 developed by "Vanderbilt Centre for Neuroscience Drug Discovery", (Nashville, TN, USA) have been tested on various animal models and cell lines and found effective on many parameters (Kumar et al. 2016). Although still in infancy, the strategy to develop mAChRs activators may serve a good approach in taming AD.

Parkinson's Disease

Parkinson's disease (PD) is the second most common neurodegenerative disorder damaging mainly the motor system and affects about 1.5% of the world population over the age of sixty (Lozano et al. 1998). This disease is characterised by the death of 70–80% of dopaminergic neurons in the substantia nigra—the important part of mid-brain, thus resulting in the decreased secretion of a neurotransmitter i.e. "dopamine" in these areas of brain. Since neurons of this region of brain control the voluntary movements therefore there occurs degeneration in four cardinal symptoms namely postural imbalance, bradykinesia, muscular rigidity and resting tremor (Bhat et al. 2015). All together these motor symptoms represent "Parkinsonism" or "Parkinsonian syndrome" (Fig. 8).

Sometimes also called as syndrome, Parkinson's disease has a very complex pathophysiology and involves not only the dopaminergic loss of neuronal cells but also the loss of noradrenergic, cholinergic and serotonergic neurons.

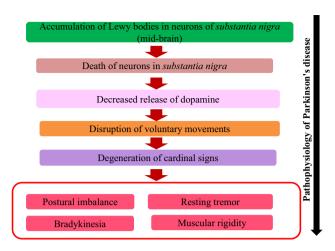


Fig. 8 Pathophysiology of Parkinsonism and its cardinal signs

Research shows that the additional loss of noradrenaline (NA) neurons of the locus coeruleus, the chief source of NA in the brain could cause motor and non-motor dysfunctions. Furthermore, the use of selective agonists and antagonists of noradrenaline alpha receptors, scientists have revealed that $\alpha 2$ receptors have an important role in controlling motor activity and that targeting $\alpha 2$ receptor with antagonists could prove pivotal in improving the motor symptoms as well as L-Dopa-induced dyskinesia. Damage or loss of NA neurons in PD influences all PD symptoms and the strategy to add NAergic drugs to dopaminergic medication seems a promising approach in the treatment of the disease.

Serotonin (also known as 5-Hydroxytryptamine), an another neurotransmitter plays important role in developing PD by affecting several motor and non-motor functions thereby causing symptoms like tremor, cognition impairment, depression, psychosis, as well as L-DOPA-induced dyskinesia (Huot et al. 2017). Declined levels of 5-HT were observed in the prefrontal cortex (PFC) up to 18 weeks following an acute injection of 1-methyl-4-phenyl-1,2,3,6tetrahydropyridine (MPTP) in mice (Ansah et al. 2011). In another study, reduction in the levels of 5-HT transporter (SERT) in the cortex and anterior cingulate was reported post unilateral striatal lesions in the macaque monkey (Sanchez et al. 2013). Furthermore, decline of serotonin transporter (SERT) immunoreactive axons in the (prefrontal cortex) PFC declined 5-HT-immunoreactivity in the median raphe neurons, or reduced PFC SERT binding capacity has also been observed in PD brains (Guttman et al. 2007; Haapaniemi et al. 2001). PD patients suffer a loss of about 25% serotonergic receptors (HT1A) at median raphe nucleus, and this is concurrent with the severity of resting tremor (Doder et al. 2003), which indicates that 5-HT projections in mid-brain have more implication in triggering PD tremor than loss of nigrostriatal DA-projections.



Experiments in MPTP model of PD suggests that 5-HT turnover in the PFC may have an important role in executive dysfunction (Maiti et al. 2016). Several investigators in PD show a strong relation between decline of 5-HT and depression (Tan et al. 2011); however, further research is needed in elucidating the role of 5-HT with the progression of PD.

A broad band of cell clusters called nucleus basalis of Meynert (nbM) present within the basal forebrain subventricular region is principally cholinergic in nature. Acetylcholine is found downregulated in many neurological diseases, including PD (Tagliavini and Pilleri, 1983, Tan et al. 2011)). In fact, evidences revealing neuronal loss in the nbM region of PD patients indicate a definite involvement of cholinergic system in PD pathophysiology (Tagliavini and Pilleri 1983; Liu et al. 2015). In addition, PD patients with cognitive decline show the presence of lewy bodies as well as neuronal loss in nbM region of brain of suggesting the involvement of cholinergic system in the cognitive loss.

PD involves accumulation of abnormal protein aggregates called lewy bodies inside the neuronal cells of substantia nigra (mid-brain) (Kalia and Lang 2015). Lewy bodies appear as spherical masses that displace other cell components. They are composed of eosinophilic cytoplasmic inclusions forming a dense "core" surrounded by a "halo" of radiating fibrils composed of alpha-synuclein. Parkinson's disease (PD) is sometimes known as synucleinopathy due to an abnormal accumulation of alpha-synuclein protein in the brain in the form of lewy bodies (Galpern and Lang 2006). In addition to alpha-synuclein, Lewy bodies also contain proteins like ubiquitin, neurofilament protein and alpha-B-crystalline (Engelender 2008). Sometimes tau proteins may also be present in lewy bodies (Arima et al. 1999; Ishizawa et al. 2003). Alpha-synuclein, the major culprit in the pathogenesis of PD seems to get transferred from one neuronal cell to another cell through many pathways such as direct membrane penetration, endocytosis, exosome-mediated transfer, trans-synaptic dissemination and receptor-dependent uptake (Lashuel et al. 2013). An elevated level of alpha-synuclein in the brain neurons of PD patients is the indicative of defective alpha-synuclein clearance which contributes to its pathogenesis. Normally, alpha-synuclein can undergo degradation in neurons by two ways i.e. through ubiquitin-proteasome system (UPS) and the autophagy/lysosomal pathway (ALP) (Dehay et al. 2015). There is presence of a chaperone-mediated autophagy (CMA) recognition motif 95VKKDQ99 (KFERQ like) in the alpha-synuclein protein that allows its interaction with cytosolic chaperone "HSC-70" and then translocation into the lysosome via lysosome-associated membrane receptor protein, LAMP2a (Cuervo et al. 2004). In vivo studies confirm that the normal soluble form of α -synuclein is mainly degraded by UPS while the complex aggregated form is degraded by ALP (Ebrahimi-Fakhari et al. 2011).

Research has shown depletion in proteasome components as well as reduced lysosome number in sporadic PD brains and in both toxic and genetic rodent models (Dehay et al. 2010). There occurs a vicious cycle leading to accumulation of alpha-synuclein in brain cells and the formation of lewy bodies which in turn hinders protein clearance thereby leading to neuronal damage. Currently, there is no cure for Parkinson's disease. However, some medications like dopamine agonists notably L-DOPA (levodopa) are being used, which becomes less effective after long-term use and produces side effects marked by involuntary writhing movements. With the passage of time there occurs massive neuronal loss. These medications become less effective while at the same time they produce complications marked by involuntary writhing movements (Sveinbjornsdottir 2016). Scientists mostly consider PD a non-genetic disorder; however, around 15% of individuals with PD show first degree relative that has the disease (Samii et al. 2004). Classical treatment strategies for Parkinson's disease include the use of three types of therapeutic approaches namely dopamine agonists, dopamine antagonist (e.g. inhibitors of monoamine oxidase) and inhibitors of catechol-O-methyltransferase. In 2005, Rascol and co-workers reported potential therapeutic role of MAO-B inhibitor "rasagiline" in PD patients with motor fluctuations. Rasagiline drug was compared against an agent from another class-the catechol-O-methyltransferase inhibitor "entacapone". Both drugs exhibited comparable benefit (Rascol et al. 2005). In addition, techniques like gene therapy and cell transplantation are being tested in animal models to find an effective treatment for PD. However, such treatment strategies are still in infancy and may take several years to become successful.

Targets for Parkinson's Disease

Alpha-Synuclein

The first predominantly hereditary mutation described in PD was that of A53T encoding alpha-synuclein (Polymeropoulos et al. 1997). Mutated form of alpha-synuclein protein aggregates inside the dopamine neurons of the substantia nigra and other brain-stem neurons leading to lewy body formation thereby making neurons prone to oxidative stress/cell death. Evidences indicate that toxicity of alpha-synuclein is mediated by Sirtuin-2 protein (an important member of HDAC family) under both the in vitro as well as in vivo transgenic Drosophila model of PD (Outeiro et al. 2007). Being a crucial protein in the pathogenesis of PD, alphasynuclein is looked as a novel drug target for the possible treatment of this dreadful disease. Currently, one of the most trusted strategies to tame PD is the use of antibodies that target alpha-synuclein protein. Positive results from in vivo mouse model studies have led various pharmaceutical



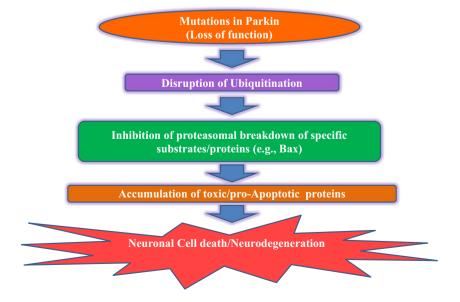
companies like Roche to start clinical trial for PRX002, the monoclonal antibody directed against α -synuclein. The antibody was initially developed by Elan Pharmaceuticals (patent#US7910333). In another example, active immunisation with Affitope PD01 (Affiris, patent O2009103105) in 32 PD patients was found safe in a first pilot study (Dehay et al. 2015). Moreover, passive Immunisation involving the use of a novel monoclonal α -syn antibody (9E4) against the carboxy-terminus of alpha-synuclein was shown to traffic into the central nervous system (CNS) and improves the deficits caused by alpha-synuclein aggregation. Various in vivo experiments showed that 9E4 cross into the CNS and bind to cells that display alpha-synuclein accumulation and promotes alpha-synuclein clearance via the lysosomal pathway (Masliah et al. 2011). In future, drugs preventing alpha-synuclein aggregation may become an effective strategy to combat PD.

Parkin

Ubiquitination of substrates is a highly regulated biochemical mechanism involving three important enzymes namely E1 ubiquitin-activating enzyme, E2 ubiquitin-conjugating enzyme and E3 ubiquitin ligase 9 (Hershko and Ciechanover 1998). E3 ubiquitin ligases play an essential role in the recognition of substrate and thus contribute to the specificity of ubiquitin reaction. Defective parkin-mediated ubiquitination may disrupt the targeting of specific substrates for degradation (Tan et al. 2009a, b), thereby causing accumulation of toxic proteins in the cell followed by cell death (Fig. 9). Parkin is composed of 465 amino acids and acts as E3 ligase in the ubiquitin–proteasome system (Shimura et al. 2000; Kahle and Haass 2004) and is widely neuroprotective in action. Parkin deficiency may affect cell survival through

complex mechanisms. Parkin mediates autophagic degradation of mitochondria during mitochondrial depolarization (Narendra et al. 2008). Autosomal recessive loss-of-function mutations in PARK2 gene cause functional inactivation of parkin thus leading to degeneration of catecholaminergic neurons and a familial form of Parkinson's disease. There are evidences which indicate that there may be some correlation between the mitochondrial function of parkin and its neuroprotective role. The role of parkin in blocking apoptosis has been widely reported and may involve the basic alterations in the threshold for the release of apoptotic cytochrome c. Parkin blocks the basal and apoptotic stress induced translocation of Bax to the mitochondria. Furthermore, its apoptotic function was retained by an engineered ubiquitinationresistant form of Bax. In addition, bax knocked out cells complemented with lysine-mutant bax did not manifest the antiapoptotic effects of parkin that were observed in cells expressing wild-type bax. Also the parkin-deficient neurons are more sensitive to apoptosis, exhibiting 74% increase in the levels of caspase 3/7 (Johnson et al. 2012). In addition to its role in ubiquitin-proteasome system (UPS), parkin has many other functions for instance it leads to monoubiquitylation of epidermal growth factor receptor pathway substrate 15 (Eps15). This monoubiquitylation event in turn blocks the interaction between Eps15 and ubiquitinated EGF receptor (EGFR) and promotes EGFR internalisation. Impaired function of parkin may disrupt this process leading to enhanced endocytosis and degradation of the EGF receptor thereby decreasing neuronal survival (Fallon et al. 2006). Blocking parkin expression promotes oxidative damage while increasing parkin expression diminishes oxidative damage (Shinichiro et al. 2013). Based on these lines, strategy to enhance the non-proteasomal ubiquitinating functions of parkin will contribute significantly to neuronal survival. Gene therapy

Fig. 9 Figure depicting the altered/mutated parkin affects normal ubiquitination process thereby inhibiting the proteasome-mediated breakdown of selected proteins. Accumulation of the latter causes neurotoxicity and hence neurodegeneration





has been proposed as a promising approach to modulate the effects of aberrant protein aggregation and misfolding for instance virus-mediated delivery of parkin effectively reduces the neuronal toxicity caused due to overexpression of alpha-synuclein and it is supposed that lentiviral delivery of beta-synuclein may also confer similar neuroprotective benefits.

Molecular Chaperones

Molecular chaperones are an important class of proteins that interact with other proteins and help them in acquiring their native/stable conformation (Hart et al. 2011). In neurons, there is an extensive network of chaperones and co-chaperones that mediate protein folding and re-folding. They also interact with protein degradation cascades like ubiquitin-proteasome pathway or autophagy to eliminate the wrongly folded and potential pathogenic proteins. Thus, deregulation of chaperone expression is critical for neurodegeneration. Studies demonstrated the role of chaperones like Hsp70, Hsp90, Hsp40, Hsp60 and Hsp27 as part of lewy bodies in PD (Ebrahimi-Fakhari et al. 2013). Hsp70 is believed to have critical role in maintaining α -synuclein's folding (Auluck et al. 2002) and also mitigates α -synuclein toxicity. Exogenous overexpression of Hsp70 and other chaperones has critically proven neuroprotective in PD models. GRP78/BiP (an HSP70 located in the endoplasmic reticulum) diminishes α-synuclein-mediated neurotoxicity by downregulating Endoplasmic Reticulum (ER) stress mediators and the level of apoptosis. GRP78/BiP also promotes the survival of nigral tyrosine hydroxylase (TH)-positive cells and elevates striatal DA levels. There is formation of a complex between GRP78/BiP and α -syn that may possibly contribute to prevent neurotoxicity caused due to α-synuclein. Molecular chaperones GRP78/BiP have a definite neuroprotective role in α-synuclein-triggered neurodegeneration (Marina et al. 2012). These data, together with the results from age-related studies, highlight the importance for developing drugs to induce elevation of endogenous GRP78 in order to increase cellular survival and extend functional longevity. BIX (a selective inducer of BiP, BiP Inducer X (BIX) has been shown to selectively induce the GRP78 mRNA and to modulate the ER stress response in cells, thus promoting the survival of neuronal cells undergoing degeneration associated with activation of the UPR. This has further implied that augmentation of GRP78 is a feasible therapeutic approach for the treatment of neurodegeneration (Marina and Oleg 2013). Co-expression of TorsinA (a protein with homology to Hsp104) (McLean et al. 2002a, b), Hsp27 (Zourlidou et al. 2004), Hsp40 (McLean et al. 2002a, b) or Hsp70 (Klucken et al. 2004) remarkably diminishes α-synuclein levels and neurotoxicity. Studies prove that there occurs a transient and reversible interaction between the substrate-binding domain of HSP-70 and the core hydrophobic region of soluble α-synuclein intermediates (Luk et al. 2008). While elucidating the role of small Hsps (Hsp20, Hsp27, HspB8, HspB2B3 and αB-crystallin) with both mutant and wild-type α -synuclein, it was found that all HSPs interacts transiently with different forms of α -synuclein and blocks the formation of mature α -synuclein (Bruinsma et al. 2011). Molecular chaperones might thwart neurotoxicity via different ways like e.g. by facilitating disease protein degradation or sequestration, blocking inappropriate protein interactions or by blocking downstream signalling events responsible for neurodegeneration. Thus, cellular depletion and the subsequent loss of chaperone function may promote neurodegeneration. Targeting chaperones with drugs that increase their expression is regarded as a promising strategy for treating PD (Muchowski and Wacker 2005a, b). Chemical chaperones or the compounds with direct chaperone activity like geldanamycin, trehalose, 4-phenylbutyrate, celastrol and mannosylglycerate are being tested as potential anti-PD therapeutics. For instance, trehalose (a disaccharide) demonstrates direct interaction with client proteins and can also promote autophagy-mediated protein clearance in different models of neurodegenerative diseases. The compounds mannosylglycerate, 4-phenylbutyrate and mannitol can markedly ameliorate α-synuclein aggregation and toxicity in different PD models (Ebrahimi-Fakhari et al. 2013). Chemical chaperones with low toxicity could be the ideal candidates for future drug development to treat PD.

A2A Receptors

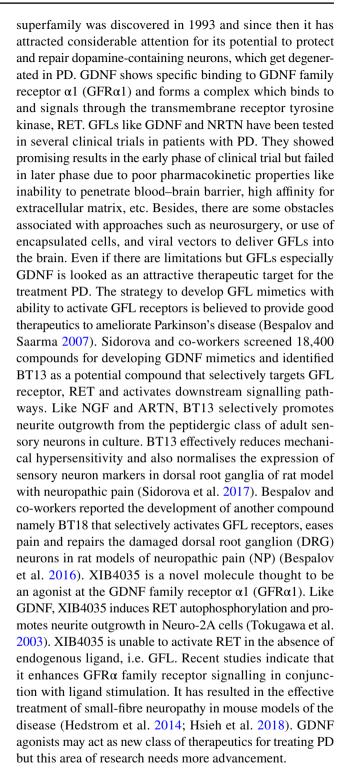
In addition to dopamine, many other neurotransmitters such as 5-hydroxytryptamine, norepinephrine, adenosine, glutamate and acetylcholine are implicated in PD. Researchers try to explore the role of non-dopaminergic therapies for PD. Modulation of neurotransmitters that act downstream of dopamine in the striatal outflow pathways is a promising target. A good example of such a strategy includes the inhibition of adenosine 2A (A2A) receptors. Briefly, the A2A receptor belongs to the G-protein-coupled adenosine receptor (GPCR) family and is expressed majorly in striatum (Cristalli et al. 2009; De-Lera-Ruiz et al. 2013). A2A receptors are also expressed at lower levels in other parts of brain including cerebral cortex, hippocampus, motor nerve terminals, nucleus tractus solitarius and glial cells. Activation of A2ARs enhances the release of several neurotransmitters, for instance dopamine, acetylcholine and glutamate but inhibits the release of gamma aminobutyric acid (GABA). A2A receptor activation also modulates neuronal excitability and synaptic plasticity, and affects various behaviours including sleep-wake cycle, locomotor activity, depression, anxiety, learning and memory. At the cellular level, A2ARs



are localised predominantly in the soma of GABAergic (enkephalin-containing, dopamine D2 receptor-expressing) striato-pallidal projection neurons and to a lesser extent in asymmetrical excitatory synapses at the dendrites of corticostriatal terminals. At the molecular level, the A2AR has been shown to interact with other neurotransmitters and neuromodulator receptors (possibly through molecular dimerization) including metabotropic glutamate receptor subtype 5 (mGluR5), adenosine A1 receptor (A1R), dopamine D2 receptor (D2R), cannabinoid receptor type 1 (CB1) and facilitatory nicotinic acetylcholine (Ach) receptor. These interactions expand the range of possibilities used by adenosine to interfere with neuronal function and communication (De-Lera-Ruiz et al. 2013). The strategy of blocking A2A receptors itself has shown a moderate effect on the symptoms of PD but when used in combination with levodopa and dopamine has improved the therapeutic outcome significantly (Bara-Jimenez et al. 2003). Results obtained from animal models also indicate that the use of adenosine receptor blockers could protect neurons in PD (Chen et al. 2001). Till now, clinical research testing A2A antagonists as therapeutic candidates for PD continues to evolve from drugs formerly discontinued (i.e. preladenant and vipadenant) to new derivatives in development (like tozadenant, ST1535, ST4206, PBF-509 and V81444) and the relatively older drug istradefylline, which has been already licensed in Japan as an anti-parkinsonian drug (Pinna 2014).

GFLs

GFLs are the proteins that regulate the development and maintenance of the nervous system. GFLs support and restore multiple neuronal populations such as dopaminergic, hippocampal, sensory, motor, enteric, basal forebrain, sympathetic and parasympathetic neurons. Thus, GFLs have a great potential for curing diseases caused due to neurodegeneration. Till now four GFLs namely GDNF, neurturin (NRTN), artemin (ARTN) and persephin (PSPN) have been characterised. GFLs mediate their effects by a receptor complex which consists of the signal-transducing module receptor tyrosine kinase known as RET and the ligand-binding co-receptor i.e. GFRα. Like in the case of other RTKs, RET has an intracellular tyrosine kinase domain (Gill et al. 2003) and a unique extracellular domain (ECD) composed of a cysteine-rich region and four cadherin-like regions (He et al. 2005; Slevin et al. 2005). RET does not interact with GFLs directly, but GFLs bind with GFRa co-receptors. GDNF preferably binds to GFRα1; ARTN binds to GFRα3; NRTN binds to GFRα2 and PSPN binds to GFRα4. Additionally, GDNF can activate RET through GFRα2, and NRTN through GFRa1 while as ARTN shows weak binding to GFRα1 in vitro (Sidorova and Saarma 2016). GDNF, a distant member of the transforming growth factor-beta (TGFβ)



LRRK2

Leucine-rich repeat kinase 2 (LRRK2) is a multidomain and multifunctional protein expressed widely. It is also named as dardarin and is encoded by PARK8 gene in humans. Besides having kinase activity, LRRK2 can also take part in intracellular signalling as a scaffold protein by



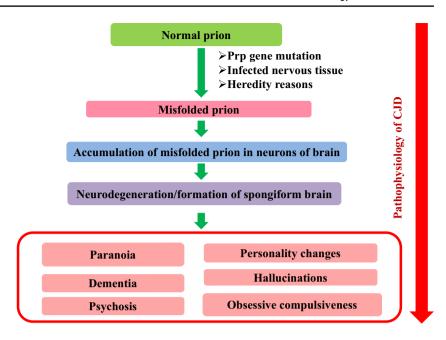
bringing together heterologous complexes via its various protein-protein interacting (PPI) domains. Mutations in LRRK2 gene are the major cause of inherited and sporadic form of Parkinson's disease. There is substantial overlapping between clinical familial LRRK2-linked PD and idiopathic PD. Exploring LRRK2 function may provide deep insights into both familial as well as idiopathic form of PD (Haugarvoll et al. 2008; Healy et al. 2008). The Gly2019Ser mutation is one of the prominent LRRK2 mutations that lead to Parkinson's disease. Recently, genome-wide gene expression study was conducted to compare G2019S-associated PD and idiopathic PD. This study identified deregulations in mechanisms associated with complement and coagulation cascades, cell adhesion molecules, extracellular matrix organisation and hematopoietic cell lineage. Although this study generated deep insights into the probable LRRK2-associated mechanisms in Parkinson's disease, the large difference between the blood transcriptome profiles and G2019S-associated PD and idiopathic PD supports the idea that PD is a pathogenically heterogeneous state (Infante et al. 2016). Nowadays, there is pressing demand for the development of effective drugs for PD and the option of using LRRK2 kinase inhibitors is looked as a promising approach. However, there are some critical issues that need to be taken care off while employing LRRK2 kinase inhibitors in clinical practice, such as (I) there must be evaluation of the effect of LRRK2 kinase inhibition on its expression, (II) researchers must know whether the inhibition of LRRK2 will affect its non-kinase function (e.g. the interaction of LRRK2 with microtubule) and (III) does LRRK2 has a self-regulatory function associated with its autophosphorylation event. Various compounds with LRRK2 inhibitory activity have been developed (Taymansa and Greggio 2016). First-generation LRRK2 inhibitors such as LRRK2-IN-1, CZC-25146 and CZC-54252 had good pharmacokinetic profile but they could not penetrate blood-brain barrier. Additionally, LRRK2-IN-1 also had some off-target effects relating to inflammatory pathways. Second-generation LRRK2 inhibitors can cross the blood-brain barrier and include drugs like HG-10-102-1, GSK2578215A, TAE684, JH-II-127, GNE-7915, GNE-0877, GNE-9605, PF-06447475 and MLi-2. Various second-generation LRRK2 inhibitors notably GNE-0877, GNE-7915 and GNE-9605 have good pharmacokinetic properties and reasonable half-life, but they show side effect like lung abnormalities. Among numerous LRRK2 kinase inhibitors developed so far, PF-06447475 has promising features since as it has been tested in various mammalian species and shows least side effects. The major disadvantage of this compound is its variable half-life in different species (Chan and Tan 2017). We expect better LRRK2 inhibitors with better pharmacokinetic/clinical profile will be developed in the near future.

Creutzfeldt Jakob Disease

Creutzfeldt-Jakob disease (CJD) is a fatal neurodegenerative disease sometimes known as human form of mad cow disease. This disease is caused due to prion misfolding. Prion, a membrane anchored glycoprotein normally exists in the neurons of central nervous system (CNS) and is believed to affect several signalling pathways causing neurodegeneration that leads to spongiform appearance in the affected brain (Sattar 2013). Mutations in the prion protein encoding gene "PrP" located on chromosome 20p13 is considered one of the main reasons for altered prion folding, leading to CJD (Peoc'h et al. 2000). CJD prion is dangerous because it initiates a cyclic conversion of native prion protein into the diseased state. Once misfolded, the prion proteins interact with normally folded prions, also changing them to misfolded structures thereby complicating the disease furthermore (Clarke et al. 2001). Misfolded prion proteins are highly resistant to proteolytic cleavage and get aggregated in the brain causing damage to neurons (Prusiner et al. 1998). There occurs an exponential increase in the number of misfolded proteins in neurons and their aggregation forms insoluble CJD protein clumps leading to severe damage to neurons (Fig. 10). Accumulated CJD prions destroy brain by developing tiny holes in it thereby changing its shape to "spongy form". A CJD patient often goes into coma and dies within a short span of 1–2 years. CJD is the most common human prion disease and is initially characterised by symptoms like dementia, personality changes, hallucinations, psychosis, paranoia and obsessive compulsive symptoms (Murray et al. 2012). Currently, there is no cure for this disease. Actual cause of the disease is still hidden from the eye of scientific community. However, researchers believe that mutations in the gene encoding prion protein can lead to misfolding of dominant alpha helical regions into beta pleated sheets thus introducing conformational changes in the prion protein. The defective prion proteins invade the brain and induce other prion protein molecules to misfold in a self-sustaining feedback loop. CJD has three forms, first one "acquired" i.e. transmissible from an affected person to a healthy person via exposure to brain or nervous-system tissue, mainly through certain medical procedures. In fact iatrogenic CJD can be transmitted by grafting infected tissue from a person who is either ill with the disease or in incubation. For example, some cases of dura matter grafts, a few exploratory neurosurgical interventions (stereo-EEG electrodes previously done in patients with CJD) and a case of corneal graft, are already on record (Fantini and Yahi 2015). CJD is transmitted through contact with tissues from organs like brain, eye (cornea), lung, kidney, liver, lymph,



Fig. 10 Pathophysiology of Creutzfeldt Jakob disease (CJD): Factors like mutations in prion gene, infections of nervous tissue and heredity transmission of infected prions can change the normal prion to infected/misfolded prion. Accumulation of infected/misfolded prion in the brain causes neurodegeneration characterised by spongiform brain with clinical manifestations like paranoia, dementia and obsessive compulsiveness



spleen or CSF and not through contact with blood (Scott 2013). Second form of CJD is the "sporadic form" that occurs without any known cause. By far sporadic form is the most common type of CJD and accounts for at least 85 percent of total cases. Sporadic CJD (sCJD) is rarely found in juvenile patients, and it usually affects adults in their 60s (De-Villemeur 2013). sCJD patients have a very short survival i.e. just 6 months. In fact, more than 90% of patients suffering from sCJD die within a year of symptom onset. The peak incidence of sCJD is in the 7th decade, while the disease shows less incidence in younger (20-40 s) or older (>80) cases. As per the hypothesis, sCJD results either from the random structural change in the PrP protein causing the formation of PrPSc, or it results from a somatic prion protein (PRNP) gene mutation. Clinically, sCJD is characterised by features like ataxia, rapid cognitive decline, and myoclonus terminating in an akinetic mute state. The incidence rate of sCJD is reported as one per million populations per annum. However, mortality rates have gradually increased in the UK and in many other countries (Mackenzie and Will 2017). The third form of CJD is "hereditary form" where a person has a familial history of disease. If there is alteration in prion protein gene in a person's sperm or egg cells, the mutation can be transmitted to the person's offspring and these mutations in the prion gene are inherited as dominant traits. Variant CJD (vCJD), the novel human prion disease is a spontaneous neurodegenerative disease which occurs predominantly in the United Kingdom (UK). It has been associated to the consumption of beef products contaminated with the agent of the cattle disease, bovine spongiform encephalopathy (BSE) (Will et al. 2000; Will 2003; Ward et al. 2006). Moreover, it is now confirmed that that the secondary transmission of vCJD occurs through blood

transfusion as seen in three clinical cases of vCJD linked to the transfusion of non-leucodepleted erythrocytes derived from individuals who themselves went on to develop _vCJD (Urwin et al. 2016). Also, a recipient of an implicated transfusion who died of a non-neurological disorder was found to have PrP^{Sc} positivity in the spleen, indicating preclinical infection (Peden et al. 2004, Mackenzie and Will 2017). The field of CJD diagnosis has improved to a great extent with the advent of improved brain imaging and the potential for diagnostic tests in plasma and urine in vCJD and also the development of specific CSF tests in sCJD.

Targets for Creutzfeldt Jakob Disease

Theoretically, there are many possible targets for blocking CJD which include blocking conversion of PrPⁿ to PrPres, stopping prion progression in secondary lymphoid organs, accelerating the clearance of amyloid PrP deposits in peripheral tissues and brain, promoting neuronal healing and reducing brain inflammation. Current therapeutic approaches rely mostly on blocking the conversion of PrP proteins to PrPres form for preventing the subsequent accumulation of PrPres in the central nervous system and peripheral nervous system. Based on these lines, recent targets include direct inhibition of this conversion, altering PrPⁿ expression, blocking its cell surface localization, degradation of PrPres and interference with vital accessory molecules like glycosaminoglycans and fragment antigen binding. Factors like exposure to biomaterials contaminated with PrPres and mutations in PrP gene are believed as important causal agents of prion disease (Tamgüney et al. 2008). CJD can be dealt with the drugs inhibiting the aggregation of prion proteins in brain. Many such drugs have been developed, for



example, pentosan sulphate, vidarabine, quinacrine, amantadine, acyclovir, curcumin, doxycycline, astemizole and flupirtine. These drugs have the ability to reduce the symptoms of familial CJD. Aggregation of prion proteins in the body does not trigger any immune response since prion proteins exist normally in the body and it is the point mutation that generates abnormal prion resistant to protease. Strategies like targeting with miRNA, gene silencing and antibodies can serve better to tame this disease (Linden et al. 2008; Manuelidis 2013). Targeting laminin, a high affinity prion receptor (LPR/LR) is regarded as a potential candidate for CJD therapy. Studies indicate that LRP/LR acts as a receptor for both the normal (PrPn) as well as the diseased form of prion (PrPres) (Gauczynski et al. 2006). This receptor is believed to play crucial role in Prpres binding and its intracellular internalisation. Gauczynski and co-workers reported that polysulfated glycans can block the synthesis of abnormal PrP by inhibiting of LPR/LR-dependent binding of prions to target cells (Gauczynski et al. 2006; Mbazima et al. 2010). In an another experiment, Leucht and co-workers (Leucht et al. 2004) employed antisense RNA to knockdown the LPR/LR in mouse brain so that such animals could serve as experimental tools for elucidating the role of LRP/LR in pathogenesis of prion diseases. In another study using lentiviral vectors to generate small interfering RNA (siRNAs) against the laminin receptor precursor mRNA, there was a prolonged extension in the preclinical phase of scrapie (a prion disease)-infected mice. Moreover, stereotactic intracerebral microinjection of recombinant lentiviral vectors expressing siRNA to LRP7 and LRP9 into the hippocampus was found to effectively extend the preclinical phase in scrapie-infected mice (Pflanz et al. 2009). More research is needed to identify the ligands/compounds targeting this receptor so as to tame CJD. Various other studies have tried to block prp, for example, RNAi-mediated knockdown of PrP and the resultant inhibition of PrPSc (Tilly et al. 2003). Virally expressed RNAi has been employed for reducing the levels of PrP^C in goats, cattle and mice (Golding et al. 2006; White and Mallucci 2009). Transgenic mice produced by lentiviral transduction of embryos stably express anti-PrP shRNAs and have increased resistance to prion infection due to resultant RNAi of endogenous PrP (Pfeifer et al. 2006). The major hindrance to the use of therapeutic RNAi in taming neurodegenerative disease is its delivery to the brain. Blood-brain barrier (BBB) restricts the passive entry of molecules from the peripheral circulation thus forcing researchers to use alternative ways for delivering RNAi into brain like the transient disruption of the BBB's impermeability, delivery through active transport (across this barrier), or direct injection into the brain. Various new technologies have recently been recently developed which include utilising viruses capable of transversing the BBB (unaided) or targeting receptors in the BBB to mediate the transport of RNAi (Pfeifer et al. 2006). Transcription factors namely SP1 and MTF-1 have been found to significantly increase prion protein levels and upregulate prion gene expression in copper-replete Menkes protein (MNK) deletion cells. Also, siRNA-mediated "knockdown" of SP1 or MTF-1 in MNK deletion cells reduces prion protein levels and down-regulates prion gene expression. These results support a novel mechanism through which SP1 and MTF-1 act as copper-sensing transcriptional activators to regulate human prion gene expression (Bellingham et al. 2009; Panegyres and Armari 2013). Therefore, SP1 and MTF-1 represent the new targets for developing key therapeutics to modulate the expression of the cellular prion protein and subsequently the prevention of prion disease like CJD.

Anti-prion antibodies directed against PrP^C represent another strategy to treat CJD. However, this strategy is still in its infancy and needs more advancement. Earlier, monoclonal antibody like D13 was reported to exhibit strong target-related toxicity. Yet another antibody ICSM18, with an epitope that overlaps with antibody POM1, was reported to be safe when injected into brains of mouse. However, the subsequent studies contradicted these findings. In fact several antibodies targeted against certain epitopes of PrP^C, including antibody POM1, were found extremely neurotoxic. When assessed with dose-escalation studies using diffusionweighted magnetic resonance imaging and various histological techniques, Reimann and co-workers found that both D13 and ICSM18 induce rapid, dose-dependent, on-target neurotoxicity. They concluded that antibodies directed to this region may not be suitable as therapeutics. However, they also reported that no such toxicity was observed when antibodies were administered against the flexible tail present at amino-terminus of the prion protein (Reimann et al. 2016), thus suggesting that any attempt aimed at immunotherapy of prion diseases should take into account such potential untoward effects/threats.

Targeting Mitochondrial Proteins in Neurodegeneration

Human brain requires abundant energy supply and utilises about 20% of total bodily consumption of oxygen for its normal functioning. Mitochondria being the power house generate this huge energy through the process called oxidative phosphorylation (OXPHOS) (Moreira et al. 2009). Major portion of this energy is used to maintain ion gradients across the plasma membranes of neurons for generating action potentials. Making neurons deficient of oxygen or glucose even for a brief interval of time may result in neuronal death. OXPHOS although a necessary process for energy generation, also produces various toxic by-products in the form of free radicals like superoxide (O_2^-) , hydrogen peroxide (H_2O_2) and hydroxyl (OH^-) . These free radicals

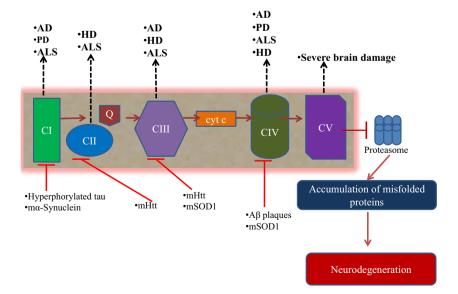


could damage various cellular components if not cleared immediately. Various mitochondrial enzymes/proteins detoxify such free radicals, for instance O₂⁻ is detoxified by the mitochondrial manganese superoxide dismutase (MnSOD) producing H₂O₂ which in turn can be converted to H₂O by another enzyme–glutathione peroxidase (GPx) (Moreira et al. 2010). Deregulation in the normal expression of such crucial enzymes can drastically elevate the levels of reactive oxygen species thereby damaging neurons and predisposing them to neurodegeneration Mitochondrial electron transport chain (ETC) is composed of various enzyme complexes namely complex-I (NADH-dehydrogenase), complex-II (succinate-dehydrogenase), complex-III (ubiquinolcytochrome-C-oxidoreductase), complex-IV (cytochrome-C-oxidase) and complex-V (ATP-synthase) (Coussee et al. 2011). Deficiency of these mitochondrial complexes/enzyme systems has clear association with the pathogenesis of neurodegenerative diseases like PD, AD, HD and ALS.

Reports suggest that increase in the expression of hyperphosphorylated tau and Aß plaques, mitochondrial α-synuclein, mitochondrial Htt and mitochondrial SOD1 lead to combined deficiencies of mitochondrial complexes (Mojtaba et al. 2017) thus promoting the accumulation of aggregated/misfolded form of such proteins through the inhibition of the proteasome activity during AD, PD, HD and ALS, respectively (Fig. 11). The mechanism of mitochondrial dysfunction in Alzheimer's disease shows the involvement of various proteins/enzymes. Initially, Aβ precursor protein (APP) blocks the translocase enzyme of mitochondrial outer membrane (TOM40), and inner membrane (TIM23) thus blocking the entry of nuclear-encoded subunits of COX/CIV into mitochondria and declining cytochrome-oxidase (COX/CIV) activity. Alternatively, Aβ can enter the mitochondria through TOM complex, interact with the heme groups of COX/CIV, decrease its catalytic activity and thus elevate ROS production which in turn elevates Aß production as well as cyclophilin D-mediated (CypD) opening of permeability transition pore (PTP). This is followed by the release of pro-apoptotic factors such as cytochrome c (Cyt c) and apoptosis-inducing factor (AIF) through PTP into the cytosol where it activates the apoptotic pathway thus leading to death of cells/neurons (Hibiki and Giovanni 2017). In case of PD, mitochondrial dysfunction has been associated with inhibition of mitochondrial complex-I (CI) activity. Mutations in parkin and tensin homologue (PTEN)-induced kinase 1 (PINK1) involved in mitochondrial function are considered as the genetic causes of PD. PINK1 causes phosphorylation of parkin on threonine 175 thus eliciting its translocation to mitochondria. Overexpression of PINK1 seems to protect mitochondria by inhibiting release of cytochrome c (Cyt c) from mitochondria, increasing expression of mitochondrial CI subunits, and reducing reactive oxygen species production. All such beneficial effects are found lost in parkin-deficient experimental models which show decline in CI and complex-IV (CIV) subunits and an increase in ROS production. ROS generation potentiates the cyclophilin D (CypD)-mediated permeability transition pore (PTP) opening and consequent release of Cyt c from mitochondria thus promoting apoptotic pathways and cell death (Moreira et al. 2010). Therefore, strategies to increase ETC capacity or to reduce mitochondrial ROS production by modulating expression of proteins like parkin, PINKY, Cyt c and mitochondrial CI subunits may prove useful for the treatment of brain ageing and neurodegenerative diseases.

Mitophagy, an important phenomenon associated with neurodegenerative diseases refers to the selective degradation of mitochondria due to autophagy. Compromised

Fig. 11 Proteinopathy triggered mitochondrial complex deficiencies during neurodegenerative disorders like amyotrophic lateral sclerosis (ALS), Huntington's disease (HD), Alzheimer's disease (AD) and Parkinson's disease (PD)





mitophagy lead to the accumulation of protein aggregates which ultimately conclude with neurodegeneration. In mammals, mitophagy is activated by ROS generation and mitochondrial permeability transition pore (PTP) induction. One of the most studied pathways for mitophagy in mammalian cells is the PINK1/Parkin pathway (Youle and Narendra 2011; Pickrell and Youle 2015). In this pathway, mitochondrial damage decreases the mitochondrial membrane potential which leads to the stabilisation of the ubiquitin kinase (PTEN)-induced kinase 1 (PINK1) on the outer mitochondrial membrane. Here it phosphorylates ubiquitin, thus leading to the recruitment of the E3 ubiquitin ligase parkin. PINK1 phosphorylation activates parkin which polyubiquitinates mitochondrial proteins leading to their association with the ubiquitin-binding domains of autophagy receptors and the formation of the autophagosome. The autophagosome then fuses with the lysosome, leading to degradation of the mitochondria (Lazarou et al. 2015). Alternatively, PINK1 can recruit autophagy receptors directly in a parkinindependent manner, leading to low levels of mitophagy. Parkin-mediated mitophagy can be suppressed by deubiquitination of its substrates. USP8 (Ubiquitin-Specific Peptidase 8) was found to regulate mitophagy in human cell lines (Durcan et al. 2014), while the reduction of USP30 (Bingol et al. 2014) and USP15 (Cornelissen et al. 2014) can increase mitophagy and rescue mitochondrial phenotypes in fly models. Mutations in Parkin (Kitada et al. 1998) and PINK1 (Valente et al. 2004) lead to the autosomal recessive form of Parkinson's disease. Growing number of studies indicate the protective role of mitophagy in several deleterious situations such as CoQ10 deficiency (Gegg et al. 2009), hypoxia (Zhang et al. 2008) and rotenone exposure (Pan et al. 2009) thereby making mitophagy and the associated proteins a good candidate(s) for therapeutic intervention.

Parkin being a possible link between mitochondrial dysfunction, mitochondrial fission and mitophagy during PD plays crucial role in the turnover of damaged mitochondria by mitophagy (McBride 2008) rendering this protein a potential target for therapeutic intervention to improve mitophagy during PD. Studies also show that Parkin is selectively recruited to dysfunctional mitochondria promoting their fragmentation and destruction in the autophagosomes (Narendra et al. 2008). These results suggest that mutations of parkin can impair mitophagy which in turn can enhance the accumulation of dysfunctional mitochondria. Thus, targeting mitophagy-related proteins in neurodegenerative disorders may open the avenues for potential targets for discovering effective therapies.

Mitochondrial biogenesis is also seen as an important factor linked to neurodegeneration. It is associated with expression of genes including peroxisome proliferator-activated receptor-c coactivator 1 alpha (PGC- 1α), mitochondrial transcription factor A (TFAM), nuclear respiratory factor 1 and

2 (NRF1 and NRF2), and mitochondrial transcription factor B1 (TFB1 M) (Golpich et al. 2015). It is highlighted that impaired mitochondrial biogenesis potentially contributes to the mitochondrial dysfunction and neurodegenerative diseases. Evidences suggest the possible role of various signal transduction proteins, transcription factors and transcription co-activators in the regulation of mitochondrial mass and number inside neurons (Onyango et al. 2010; Dominy and Puigserver 2013). Mitochondrial biogenesis is dependent on several signalling cascades and transcriptional complexes that promote the formation and assembly of mitochondria. It is a process that is heavily dependent on timely and coordinated transcriptional control of genes encoding for mitochondrial proteins.

PGC-1 family members (e.g. PGC-1α and PGC-1β) coactivate genes encoding proteins for transcription and replication of mtDNA as well as importation of mitochondrial protein (Wu et al. 2006; Mojtaba et al. 2017). They also have contribution in the physiological integration of mitochondrial biogenesis with oxidative metabolism and provide an overlapping and amplifying regulation of several nuclear-encoded mitochondrial genes (Scarpulla 2011). As a co-transcriptional regulation factor, the PGC-1α provokes mitochondrial biogenesis by activating several transcription factors, including NRF1 and NRF2. Furthermore, it is also called GA-binding protein A (GABPA) that regulates the expression of multiple nuclear genes to encode mitochondrial proteins such as TFAM. The TFAM in turn contributes to the mtDNA maintenance as well as motivates the replication and transcription of both mtDNA and PPARs (Scarpulla 2008; Yin et al. 2008; Sharma et al. 2014). As a transcriptional coactivator, PGC-1 functions together with combination of other transcription factors such as peroxisome proliferator-activated receptors (PPARs) in the regulation of mitochondrial biogenesis. PPARs, in particular PPAR-c, may be a major signalling pathway involved in neuroinflammation. PPAR-c is also an important regulatory factor in the modulation of genes having PPAR response element (PPRE) in their promoters, such as those encoding for oxidative stress, inflammation (COX-2), inducible nitric oxide synthase (iNOS), nuclear factor-kappaB (NF-kB), and apoptosis (Kiaei 2008). In addition, mitochondrial biogenesis involves transcription factors such as sirtuins. Sirtuins (SIRTs) are a family of the nicotinamide adenine dinucleotide (NAD⁺)-dependent protein deacetylases which contribute in cellular processes such as cell cycle, transcription, energy metabolism, mitochondrial functions, ageing, apoptosis and cell survival. Mammals possess seven sirtuins with different activities, localised to the nucleus (SIRT1, SIRT6 and SIRT7), cytosol (SIRT2) and mitochondria (SIRT3, SIRT4 and SIRT5). Sirtuins can influence the progression of neurodegenerative disorders by modulating transcription factor activity (Kanfi et al. 2012; Johri and Beal 2012;



Min et al. 2013; Herskovits and Guarente 2013). Sirtuins as the transcriptional regulators may have potential therapeutic effects on a several chronic age-related and aggregate-forming neurodegenerative diseases including AD, PD, HD and ALS.

Antioxidant peptides like SS31 and SS20 represent a novel therapeutic approach that can scavenge mitochondrial free radicals and also inhibit mitochondrial permeability transition and cytochrome c release which prevents oxidant-induced cell death (Cho et al. 2007; Johri and Beal 2012; Hou et al. 2016;). Recently, preclinical studies support potential use of the mitochondria-targeted antioxidants as an effective treatment for neurodegenerative disorders (Szeto 2008). Manczak and her colleagues stated that SS31 prevents Aβ toxicity as well as decreases learning and memory deficits (Manczak et al. 2010). Moreover, SS31 and SS20 demonstrated significant neuroprotective effects on dopaminergic neurons of MPTP-treated mice (Yang et al. 2009). SS31 may also be used to treat HD due to its property to promote mitochondrial function and neuronal viability (Chandra et al. 2014). SS31 is considered a novel therapeutic approach to treat neuronal damage induced by oxidative stress. It targets the ROS production at the inner mitochondrial membrane and prevents further mitochondrial damage (Petri et al. 2006). Taken together, proteins involved in mitochondrial ETC functioning, mitophagy and biogenesis represent a novel therapeutic target and confirm a modern neuroprotective approach for most of diseases such as AD, PD. HD and ALS in the near future.

Oxidative Damage, Proteinopathy and Neurodegeneration

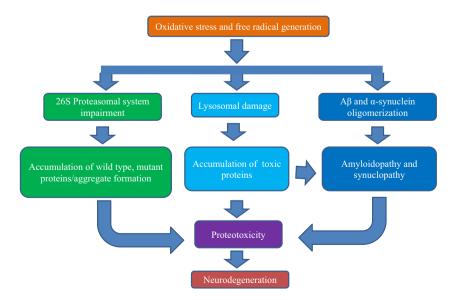
Studies with transgenic animal models reveal that oxidative modifications of brain proteins result in their functional

loss and hence neurodegeneration. Reactive oxygen species cause proteotoxicity and neurotoxicity mostly via three major mechanisms i.e. (1) impairing proteasomal system, (2) damaging lysosomes and (3) promoting $A\beta$ and α -synuclein oligomerization (Fig. 12).

Proteasomal system especially the 26S proteasome represents one of the pivotal cellular mechanisms damaged by oxidative modifications. Impaired proteasomal system prevents the degradation of toxic protein aggregates (Aiken et al. 2011; Pajares et al. 2015). Excessive accumulation of mutant or wild-type proteins inside cells triggers conformational changes in proteins (e.g. helix to β -strand conformation) which promote more self-aggregation and loss of function. Heat shock proteins like Hsp40, Hsp70, Hsp90 and others normally avert misfolding of intracellular proteins but their excess accumulation or redox modifications may disrupt the normal expression of proteins and inflict toxic effects on various cellular organelles including lysosomes and mitochondria (Morimoto 2008; Niforou et al. 2014).

Lysosomes play critical role in the removal of excessive/ toxic proteins from a cell. Their destruction due to ROS promotes intracellular accumulation of several proteins (Pajares et al. 2015). For instance, oligomerisation of Aβ and α-synuclein leads to proteotoxicity (Snead and Eliezer 2014; Xiang et al. 2015). The proteotoxicity in turn causes impairment of mitochondrial function. Studies based on analysis of postmortem brain samples, isolated mitochondria and cultured cells suggest that oligomerised proteins or even the monomers of α-synuclein or Aβ can impair mitochondria by altering mitochondrial fusion/fission processes, mitophagy and bioenergetic (Devi et al. 2006; Wang et al. 2009; Banerjee et al. 2010; Protter et al. 2012). The fact that Aβ production triggers ROS generation has been proved by various invivo studies, cell-free chemical systems and cell cultured experiments. Both A\u00e440 and A\u00e442 can interact

Fig. 12 Oxidative stress, free radical generation and neuronal damage





with transition elements via different amino acid residues including His14, His13 and His6. These bound elements can participate in various redox-cycling reactions thus producing ROS (Smith et al. 2007). Postmortem evidences show the presence of elevated levels of metals like Zn, Fe and Cu in the brain of AD patients. These metals are located mainly near the plaque which suggests the putative role of A β -mediated ROS production in AD pathogenesis.

Aβ may also increase intracellular ROS production via the involvement of apoptosis signal-regulating kinase 1 (ASK1) (Kadowaki et al. 2005a, b). Neuronal cell culture experiments show that Aβ promotes ROS production seemingly through the activation of reduced NADPH oxidase (Shelat et al. 2008; Hu and Li 2016). The other mechanism through which Aβ induces ROS production includes microglial activation via fibrillar Aβ as observed in microglial cell cultures and coculture of microglia and neurons (Qin et al. 2002; Qin et al. 2006). Microglial cells possess a B-type of scavenger receptor known as CD36. Fibrillar Aβ acts as a ligand and binds CD36 to stimulate microglial ROS production, cytokine expression (such as IL-6, IL-1β, TNF)- α) and phagocytosis in AD brain (Coraci et al. 2002; Doens and Fernández 2014).

PD brain contains elevated levels of oxidative damage markers and transition metals such as Fe (Jenner and Olanow 1996; Sofic et al. 1988). Increased expression of α -synuclein during PD could presumably enhance the production of ROS as observed in several experimental models (Bir et al. 2014; Di-Maio et al. 2016). Overexpression of mutant or wild α-synuclein has been found to be associated with elevation in the levels of ROS in SH-SY5Y cells (Anandhan et al. 2015). Studies based on biophysical analysis indicate the binding of elements like Cu and Fe to α -synuclein that may increase the cytotoxicity of α-synuclein (Carboni and Lingor 2015). Iron which promotes ROS production and catalyses peroxidative damage to biomolecules also causes translational upregulation of α -synuclein (Febbraro et al. 2012) in PD brains. Fe can prompt the formation of large oligomers of α-synuclein that can form membrane-spanning channels and thus lead to intracellular toxicity. Interaction of Fe with α-synuclein may even trigger the formation of H₂O₂ through various redox reactions (Carboni and Lingor 2015).

Since PD involves loss of DAergic neurons, it is believed that the oxidation products of DA notably the toxic quinines and ROS could act as contributing factors in the pathology of PD (Hastings 2009; Jana et al. 2011). DA oxidation products seem to have a modulatory effect on the oligomerisation and cytotoxicity of α -synuclein (Leong et al. 2009). Rat DAergic cell lines and human DAergic neurons when exposed to paraquat lead to increased aggregation of α -synuclein and this aggregation is critically dependent on NADPH oxidase, thus giving a clear indication that ROS has an important role in PD pathology. There also occurs gathering of oxidative

damage markers in the substantia nigra, which can be eliminated by using adeno-associated virus-mediated overexpression of a specific shRNA to knockdown NADPH oxidase (Nox1) gene (Cristóvão et al. 2012).

Superoxide dismutase (SOD) is an important free radical scavenger/antioxidant enzyme. Earlier reports suggest an increased expression of SOD1 immunoreactivity in Bergmann glia and astrocytes in cerebellum of CJD cases. Also there was an elevated expression of SOD1 in the cerebral cortex of CJD cases (Freixes et al. 2006). Recently, Brazier and his colleagues used an agent with actions similar to SOD in a rodent model which extended the life of scrapieinfected animals thus raising the possibility of using free radical scavengers and antioxidants as the possible therapeutics in the treatment of prion diseases. To provide further insights into the relative pathogenic importance of oxidative stress, Brazier and his colleagues used a potent manganese-superoxide dismutase/catalase mimetic, EUK-189, as a therapeutic agent in the mouse model of human prion disease. They observed a significant but relatively modest prolongation of survival due to EUK-189-treatment, which correlated with reductions in ROS and reactive nitrogen species (RNS) mediated damage to proteins when compared to untreated disease controls. In addition, EUK-189 also caused reductions in spongiform change in specific brain regions of terminally sick mice as confirmed by lesion profiling (Brazier et al. 2008). These results suggest a prominent role of antioxidants in ameliorating/modulating prion diseases. (nbr2). There also occurs decreased expression of nNOS along with its misfolding and abnormal localization in neuroblastoma cells as well as the brain during scrapie infection (Ovadia et al. 1996; Keshet et al. 1999; Petersen et al. 2005). Furthermore, reports indicate increased levels of inducible nitric oxide synthase and endothelial and in the cerebral cortex in sporadic CJD (Freixes et al. 2006). Thus, oxidative stress-mediated derangements of brain proteins/ enzymes have deeper implications in the pathology of neurodegenerative diseases and could serve as premier source for targeting of drugs.

Conclusion

Neurodegenerative diseases have complex pathophysiology and result due to multiple proteinopathies as discussed in this review. Treating damaged neurons poses perplexing challenges notably the blood-brain barrier, non-specificity/off-targets, side effects and weak pharmacokinetic profile of proposed drugs. Research should focus on developing therapeutics that are target specific, have ability to cross blood-brain barrier, possess least possible side effects and show cost effectiveness. Since many neurodegenerative diseases involve cross talk or co-existence of several



proteinopathies therefore designing drugs with ability to strike multiple targets in the pathophysiology of ND's seems to be an effective strategy. Interactions between synucleinopathy and tauopathy can promote neurodegeneration via distinct mechanisms such as (I) α-Synuclein can bind directly to tau and tubulin protein and thus block the normal interaction between the two proteins thus interfering with their physiological function, (II) α -Synuclein could promote hyperphosphorylation of tau protein by recruiting various kinases, (III) α-Synuclein can promote tau aggregation and the formation of tau aggregates, (IV) α -synuclein oligomers may in turn seed tau fibrillation and formation of insoluble NFTs thus triggering tauopathy. Therapeutics including synthetic chaperones, kinase inhibitors, anti-aggregation agents, antibodies, LRRK2 inhibitors, mAChRs activators, ApoE4 inhibitors, dopaminergic and NAergic drugs could prove effective medicines in taming various ND's; however, there is need for shift from 'magic bullet' to 'magic shotgun' strategy. While combing several drug entities that act independently on different etiological targets of a disease researches must take utmost care of difference in (1) bioavailability of each individual entity of a single drug, (2) pharmacokinetic parameters, (3) metabolic profile and overall (4) the unpredicted drug-drug interactions that will result in the development of ideally effective drugs. Prospective strategies need to focus on combinatorial progress and advancement in neuro-proteome-based therapeutic regimen, clinical data, bioinformatics and neuro-imaging technology that can verily improve the diagnosis, prognosis, as well as the therapeutic outcome of neurodegenerative diseases.

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Compliance with Ethical Standards

Conflict of interest Authors declare that there are no conflict of interest.

Ethical Approval This article does not contain any studies with human participants or animals performed by any of the authors.

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