

An update on pathological implications of enzymatic dysregulation in Alzheimer's disease.

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Abstract

Alzheimer's Disease (AD) is the most prevalent non-reversible neurodegenerative disorder that affects the memory and cognitive centres of brain. It has been reported that, AD turns out to be prominent among the people aged ~65 or above and is regarded as the most common cause of dementia. Moreover, AD stands among the leading causes of death in the first world nations, accounting more than 60% incidence of progressive cognitive impairment in elderly people. Amyloid beta and neurofibrillary tangles are two putative cytotoxic entities that have been identified, aggregation of which has been associated with the pathological signature of AD. Beta secretases—an amyloid precursor protein cleavage enzyme, plays a pivotal role in such pathogenic process of AD. Several other enzymatic dysregulations have also been linked with AD. Involvement of enzymatic dysregulation is the most discussed pathological implication in AD and therapeutic approaches have been postulated targeting such anomalies. Together, global consequences of enzymatic dysregulation and related therapeutic possibilities in AD remain the prime focus of present time. Therefore, research and study for the eloquent insight into the AD pathology from enzymatic perspective is essential and the same endeavour has been carried out in the present study.

Keywords: Alzheimer's disease, Amyloid beta, Beta secretase, Glycogen synthase kinase 3 beta, Acetylcholinesterase, Rho kinase, Prolyl endopeptidase, Monoglycerol lipase, Catechol-O-methyl transferase.

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Introduction

Alzheimer's Disease (AD)—an age-related, insidious neurodegenerative disorder, characterised by cognitive and memory impairment. It has been reported that AD is frequent among people with an age group of 65 or more. Clinically, AD is the common cause of dementia and regarded as the most common cause of death in the first world nations [1]. Over 60%-70% incidents of cognitive impairment in elderly patients have been found to be associated with AD [2]. The overall prevalence of AD in USA alone is around 2.3 million as per the statistics made in the early 20th century [3]. The prevalence

of AD among men vs. women is observed in the ratio of 1:2 to 1:5 [4]. Two known forms of AD have been reported; namely, familial and sporadic variety. Familial AD is less prevalent and stands around 10% of the total AD patient population [5]. Regardless the variety of AD, medical care becomes excessively necessary in the final phases of AD i.e., the last three years before death due to the reason that, AD not only causes memory loss but also incorporates the symptoms like dramatic personality changes, lack of physical coordination, and disorientation [6]. Generally, the final stages of AD are even more dramatic where victims are bedridden, with loss of

control in urinary and bowel movement, and frequent epileptic attacks [7]. Clinically, loss of synaptic function, presence of amyloid plaque, neuro-fibrillary tangles (NFT) and brain atrophy in specific brain areas confirm the presence of AD [8]. In a living patient, dementia is hard to confirm accurately; however, near to accurate diagnosis is possible through cognitive tests such as delayed recall and exclusion of other conditions such as hypothyroidism, nutritional deficiency and stroke [9]. An effective therapy for dementia is yet to be discovered. Safety concerns over synthetic drugs are now being promoted and drugs based on natural origin, are trending globally.

Enzymatic dysregulation is the most common pathological scenario of AD and involvement of such dysregulation has been targeted in several therapeutic approaches for AD [10]. Beta-secretase (β -secretase) is considered as the most critical enzyme involved in the AD pathology. Amyloid precursor protein (APP) is the substrate for β -secretase and uncontrolled cleavage of APP generates amyloid beta ($A\beta$) [11]. $A\beta$ oligomers are aggregation prone, which in advanced stage forms toxic senile plaques both in cytosol and extracellular matrix. Another crucial enzyme is glycogen synthase kinase-3 β (GSK-3 β), which is having pathogenic role by assisting in the process of tau-protein hyperphosphorylation. Hyperphosphorylated tau is also having the tendency to aggregate and it has been identified as an ingredient of toxic senile plaques [12]. Moreover, hyperphosphorylated tau protein is known to modify other essential protein structures by phosphorylation and GSK-3 β plays determining role in such cases. Enzymes like acetylcholinesterase; Rho kinase; prolyl endopeptidase; monoglycerol lipase; catechol-O-methyl transferase, are also having a putative role in the pathogenesis of AD [13]. Pathogenic role of each enzyme has been discussed in the present study for the better understanding of AD pathology. It is notable that several synthetic and herbal therapeutic approaches have been postulated, where regulation of such enzymatic function is the centre of remedy [14]. Though, complete cure of AD is still unaccomplished.

Global research in search of therapeutic values of various plant species is going on and few plants have been identified with potential efficacy to improve symptomatic AD pathology [14]. Countless mentions from ayurvedic medicinal recipe also have offered better therapeutic promises [15]. Another advantage of herbal therapy is that, it contains no or less side effects and is available at a very low cost [16]. Since immortal time of mankind, traditional medicines served as potential therapeutic means against mental disorders and success of such approach has gained immense interest and popularity across the globe [17]. Together, enzymatic dysregulation is the crucial point of interest for therapeutic intervention against AD.

Overview of AD

AD is the most common neurodegenerative disorder and also the main cause for dementia in humans. Initial symptom of AD includes short term memory loss that develops into profound memory failures at subsequent stages [18]. AD is having two

distinct pathological signatures namely senile plaques (containing deposits of $A\beta$ protein) and NFT; consisting of hyperphosphorylated tau protein. In the early stages of disease, huge number of neuronal cells dies causing neuronal losses in the brain's memory regions. Studies in this regard, showed that aberrant proteolytic processing of the APP leads to $A\beta$ 1-40 and $A\beta$ 1-42 fragments capable of initiating cytotoxicity in neuronal environment in AD brain [19]. Nonetheless, the knowledge concerning mechanism behind plaque deposition, neuronal cell death, and NFT by $A\beta$ proteins still remains unclear.

Incidence and Prevalence of AD

More than four million people in the United States are presently affected by AD. It is now causing more number of deaths compared to those caused due to stroke. Together, stroke and AD stand as the third most common cause of deaths in USA [20]. Statistically, the prevalence of AD is found to be more common among women. As per the data published in Diagnostic and Statistical Manual of Mental Disorders (4th Edn.), prevalence of dementia is observed in the range of 1% in developing countries like India, Peru, etc. to almost 6.4% in countries like Cuba. As compared to developed countries, developing countries like India, Nepal, Brazil, Nigeria, and Taiwan show lower annual incidence estimates in the range 1%-2% of total reported cases [21-24]. Mean survival time for AD patients in developed countries is perceived to be 5 y to 9.3 y while that in developing countries, it is 3.3 y from the time dementia sets in. For AD patients, the mean survival time further reduces to meagre 2.7 y [23].

AD Pathology

Amyloid beta peptides ($A\beta$) production and clearance determines molecular pathogenesis of AD. Primarily, generation of $A\beta$ occurs through cleavage of APP by beta- and gamma- processing enzymes. These are then cleared from the brain through diffusion, exported to vascular system, and followed by degradation or phagocytosis [25]. Experimental studies in this regard have provided some insights into the molecular mechanism of AD; identifying APP as the causative gene for Familial AD (FAD). In a mouse-based animal model, endogenous apolipoprotein E4 (APOE-4) was observed to enhance $A\beta$ deposition [26,27]. Presence of aggregates of hyperphosphorylated tau proteins commonly referred to as NFT serves as the primary markers for diagnosis of AD. These insoluble NFT have an abnormal conformation and are deposited in the neuronal cell bodies. These are capable of forming specific insoluble structures called paired helical filaments (PHF) [28,29]. Alternate splicing of *tau* gene gives rise to six tau protein isoforms; primarily used for stabilization and binding of microtubules thereby promoting microtubule polymerization. The mechanism behind aggregation or PHF formation is attributed to hyperphosphorylation of tau which induces disassociation of tau from microtubules [30,31].

Role of Oxidative Stress in AD

On account of high lipid content, brain is prone to oxidative damage. In addition to this, high rate of metabolic function and supply of the required transition metals makes it an easy target for free radical attacks [32]. Direct evidences in this regard are, increased composition of Fe, Al, and Hg in the brain, reduction in levels of polyunsaturated fatty acids in the brain, increased lipid peroxidation as well as rise in levels of ventricular fluids such as 4-hydroxynonenal which is produced in the process, increased DNA and protein oxidation in the brain, decreased cytochrome c oxidase in the brain along with diminished energy metabolism [33], advanced glycation end products (AGE), malondialdehyde, carbonyls, peroxyxynitrite, heme oxygenase-1 and superoxide dismutase-1 (SOD-1) in NFT and AGE, heme oxygenase-1, SOD-1 in senile plaques, and generation of free radicals by A β .

Oxidative stress induces serious damage to formation of biological macromolecules like malondialdehyde and lipofuscin [34]. As per experimental studies, oxidative damage primarily involves lesion on account of low protein outputs [35]. In lesions of AD, adduction production of hydroxynomenal [36] and acrolein [37] is also found due to lipid peroxidation. Nevertheless, lesions are not the dominant sites of damage, rather neuronal cytoplasm of neurons are susceptible damage sites of death and damage in AD. Inside the cell, cross linked products of lipid peroxidation and glycation undergo oxidative modification thus, becoming more and more resistant to breakdown. Crosslinking not only hinders proteasome activity [38] but also renders proteins resistant to removal by proteasomes. Neurons that face oxidative damage begin to have protein-based reactive carbonyl and nitro tyrosine in the cytoplasm. Evidences of its formation in amyloid β or τ deposits suggest cytoplasm rather than lesions as the source of reactive oxygen species (ROS) [39]. Among vulnerable neuronal populations, 8-Hydroxyguanosine (8-OHG) a nucleic acid modification obtained through hydroxyl free radical attack on guanidine is greatly increased in the cytoplasmic RNA [40]. 8-OHG decoration was observed more in the endoplasmic reticulum with majority of mitochondria showing scarce 8-OHG.

Role of Inflammation in AD

Pathophysiology of AD involves extensive neuronal death accompanied by deposition of amyloid in various regions of the brain. These amyloid deposits lead to accumulation of several proteins along with underlying inflammatory reactions, thereby resulting in extracellular senile plaques [41]. Intracellular deposition of hyperphosphorylated degenerate filaments results in NFT that form due to the aggregation of micro tubular protein tau. Such cellular progresses results in heavy amounts of neuronal losses in the hippocampus, entorhinal cortex, and associated regions of brain neocortex. Although, the reason behind neuronal cell death still remains unknown yet postulates in this context suggest 'apoptosis' as a possible reason [42]. Inflammation is also seen to be actively involved in the progression and pathology of AD. Association

of microglia with the senile plaque leads to amyloid plaque deposition which results in the phenotypic activation and subsequent elaboration of neurotoxic as well as pro-inflammatory factors [43].

Neurons which are located nearby chronically activated astrocytes and microglial cells die out due to toxins such as ROS intermediates, proteolytic enzymes, excitatory amino acids, nitric oxide, complimentary factors etc. Proinflammatory cytokines not only enhances A β 40 and A β 42 peptide production but also decreases APP solubility alongside providing neuroprotective effects [44]. In the onset of a neurological disease, APP modification begins long before disease symptoms begin to appear. During disease period, astrocyte and microglial cells activate following binding of A β to the CD14 receptor and its co-receptor—toll-like receptor 4 (TLR4). After which morphological changes in microglial cells is observed, these now turn into tissue macrophages producing inflammatory molecules [45]. This has generated enormous interest towards the *in vitro* study of the anti-inflammatory effect of selected plant extracts.

Enzyme Involved in AD Pathobiology

Beta secretase

A large type-I membrane protein called APP, when gets endoproteolysed, it generates the A β peptide [46,47]. A β is a normal catabolic product of APP metabolism in cells showing ubiquitous expression of APP. APP cleavage by β -secretase form the amino terminus Asp+1 residue of the A β sequence. This process generates two cleavage products, *viz.*, a secreted ectodomain of APP, named APPs β and second being the C99 fragment, the membrane-bound C-terminal 99 amino acids residue (Figure 1) [48]. After the first cleavage, the C99 get acted upon by a second protease called γ -secretase. This enzyme cleaves C99 to generate the carboxyl terminus of A β along with a mature peptide which is then secreted from the cell. The non-precise γ -secretase cleavage results in a spectrum of A β peptides that vary in length by a few amino acids at the carboxyl terminus, however, majority of A β terminates at 40th amino acid residue. Another type of protease called α -secretase, cleaves APP in the mid-A β domain (at Leu+17) and hence, precludes the formation of A β [49]. The α -secretase produces two cleavage products: the secreted APPs α ectodomain and the membrane-bound C-terminal fragment C83. The C83 is then cleaved by γ -secretase to form a 3 kDa fragment, p3 [50]. Intriguingly, the APP mutations leading to FAD occur at APP near the site of cleavage and as a result, the mutations directly affect cleavage efficacy. For example, the Swedish mutation is the amino acid substitution of lysine and methionine to asparagine and leucine at the P2-P1 positions at N-terminal of the β -secretase cleavage site in APP [51]. This double mutation makes APP a much better substrate for β -secretase thus dramatically increasing the cleavage efficacy. Many more examples of FAD mutations have been identified that are positioned near the γ -secretase cleavage site and these shift the balance of γ -secretase cleavage towards generation of

the more toxic A β 42 (similar to the action of FAD mutations in the presenilins) [52]. Moreover, efficiency of α -secretase cleavage has been reduced by mutations in FAD near the enzyme cleavage site. This alteration results in furnishing more APP as substrate for β -secretase cleavage and consequent generation of A β .

Establishment of the fact that APP endoproteolysis produces A β [53,54], resulted in information hunt on the properties of β -secretase activity in cells and tissues. Concomitant studies utilized the data in various ways to validate certain β -secretase 1 (BACE-1) attributes. In this context, β -secretase activity has been briefly reviewed. This enzyme has been detected in majority of the body cells and tissues [55], but its maximum activity has been found in neural tissue and neuronal cell lines [56]. One interesting fact is expression of β -secretase activity in astrocytes is less compared to neurons [57]. The research reports strongly indicated that β -secretase might be widely expressed in many tissues and cell lines but its maximum expression is found in neurons of the brain. β -secretase activity in cells results in efficient cleavage of membrane-bound APP only, whereas APP constructs lacking the transmembrane domain did not get cleaved when transfected into cells [58]. The fact implied that β -secretase might be a membrane-bound protease or, alternatively, it is strongly associated with a membrane protein for its function. At an acidic pH, β -secretase shows maximum activity and hence, the agents that disrupt intracellular pH also inhibit β -secretase activity [59,60]. Additionally, cellular level studies found maximum β -secretase activity in the acidic subcellular compartments of the secretory pathway, including the Golgi apparatus and endosomes [61,62]. From these results, it has been postulated that the active site of β -secretase might be situated within the lumen of acidic intracellular compartments. Amino acids surrounding the cleavage site in APP when changed by site directed mutagenesis leads to identification of sequence preferences of β -secretase [63]. Substitution of larger hydrophobic amino acids (such as Leu found in the Swedish FAD mutation) for the Met residue at P1 site improved β -secretase cleavage efficiency. Conversely, substitution of the smaller hydrophobic amino acid Val at the same position showed cleavage inhibitory effect. Some more substitution studies at this site and at some surrounding positions showed a decrease in cleavage efficacy and hence establish the fact that β -secretase is highly sequence specific. Studies from radiosequencing demonstrates that A β isolated from amyloid plaques and those produced in cell lines, predominantly begins at the Asp+1 residue of A β [64]. But, some A β species were found to begin at Val-3, Ile-6, and Glu+11 residues also [65]. Studies utilizing inhibitors suggest that the Val-3 and Ile-6 species are generated by a protease other than β -secretase [66]. However, the Glu+11 species is found to be produced in parallel with Asp+1 A β which suggests that β -secretase can cleave at both positions. Interestingly, the Glu+11 species is found to be the predominant form of A β made in rat primary neuron cultures [67]. Finally, β -secretase shows no sensitivity towards pepstatin which is an inhibitor of many (but not all) aspartic proteases.

Glycogen synthase kinase 3 β

Glycogen synthase kinase 3 β (GSK3 β) has been linked as a central player in AD by many studies. It has been found that deregulation of GSK3 β shows numerous pathological hallmarks of the disease in both sporadic and FAD cases [68]. These crucial findings led to ultimate formulation of the 'GSK3 β hypothesis of AD'. Evidences from studies showed that GSK3 β is intimately involved in the hyperphosphorylation of tau, memory impairment, increased production of A β and also in inflammatory responses (Figure 1). GSK3 β has reductive effect in acetylcholine synthesis, which is in accordance with the cholinergic deficit character of AD [69]. Moreover, GSK3 β is a key mediator of apoptosis and hence involves the possibility to contribute towards neuronal cell death in AD [70].

Naturally if GSK3 β is central to AD pathogenesis, its increased activity in AD patients should be a common occurrence. However, there is little such evidence, as it is technically difficult, if not impossible, to measure enzymatic activity in post-mortem brain tissue. Certain indirect evidences from many studies do support the role of GSK3 β in disease and it shows co-localization with dystrophic neurites and NFTs [71-73]. Active GSK3 β appears in neurons with pre-tangle changes [74] and in AD pathology, there is increased GSK3 β activity in the frontal cortex which was evidenced by immunoblotting for GSK3 β phosphorylated at Tyr216 [75]. Furthermore, the hippocampus region of AD patients shows up-regulated GSK3 β expression [76]. In case of both AD and mild cognitive impairment, GSK3 β expression is up-regulated in circulating peripheral lymphocytes [77]. It has recently been reported that a polymorphism in the GSK3 β promoter is a risk factor for late onset AD [78] and the same might account for alterations in GSK3 β expression in disease. Collectively, these findings suggest that GSK3 β activity might be increased in AD and the process changes in phosphorylation state as well as in expression levels. But still direct evidence for this mechanism is equivocal and some studies have found no change in GSK3 β activity [79] or even reduced GSK3 β activity in AD [80].

Genetic and epidemiological studies indicate that GSK3 β deregulation in AD is due to alterations in upstream Wnt and insulin signalling pathway intermediates. The low-density lipoprotein receptor related protein 6 (LRP6), a co-receptor for Wnt signaling, has recently been identified as a gene of risk for late onset AD in apolipoprotein E4-e4 negative individuals [81], implicating aberrant Wnt signalling in AD pathology. In addition, an association of AD with diabetes and insulin resistance has been reported [82] and genetic studies find insulin signalling genes to be potential target loci for AD [83,84]. As studies suggested, the positive association of Wnt pathway with AD may help in studying neurological alterations during carcinogenesis or conversely pre-AD stage neuronal alterations, which may be served as a marker for ongoing and future onco-events.

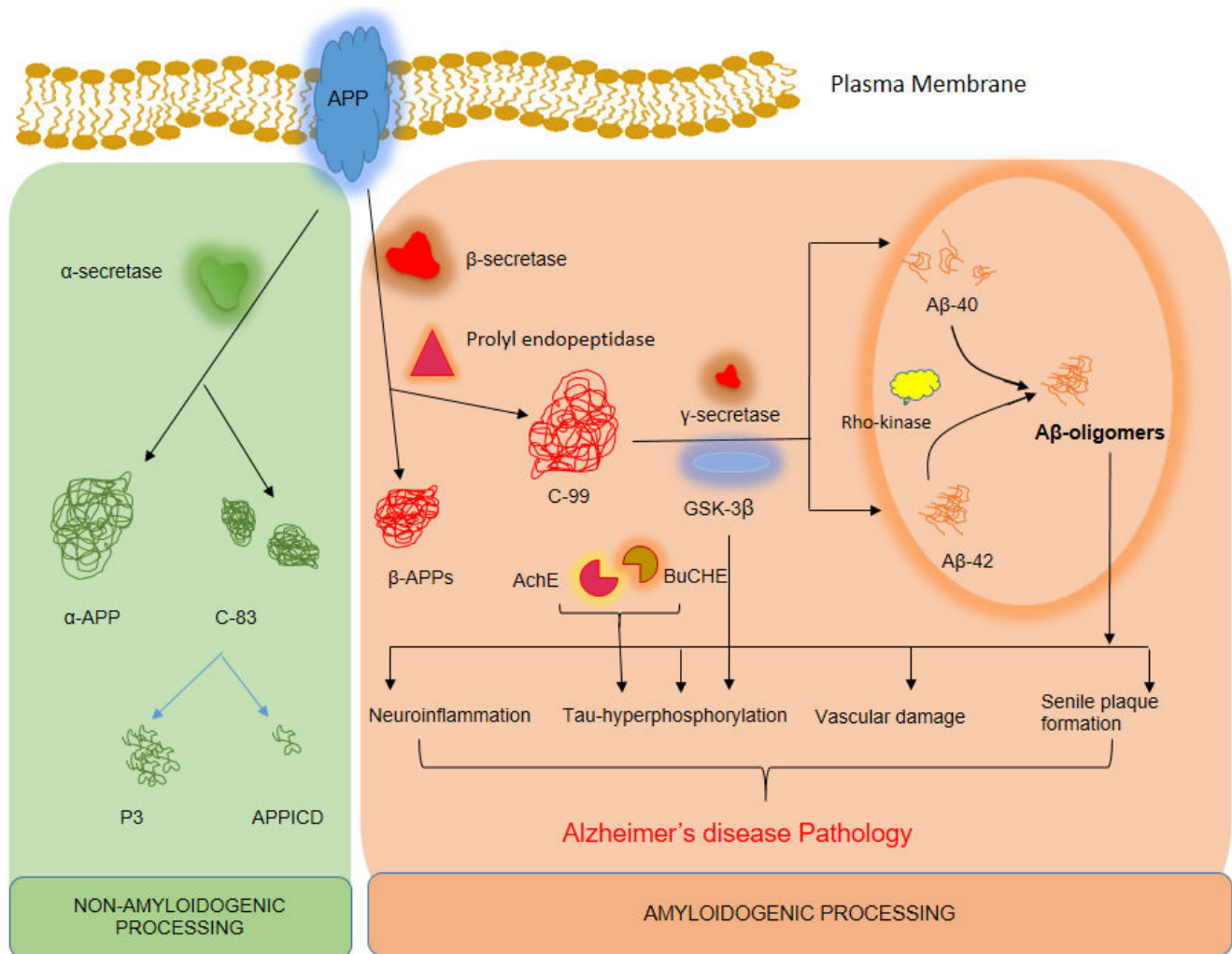


Figure 1. Schematic presentation of enzymes involved in amyloidogenic and non-amyloidogenic processing of APP.

Acetylcholinesterase (AChE) and butrylcholinesterase (BuChE)

Cholinergic transmission in the central and peripheral nervous system is controlled by a neurotransmitter—Acetylcholine (ACh). ACh hydrolysis is catalysed by Acetylcholinesterase (AChE). Butyrylcholinesterase (BuChE) is another similar group of neurotransmitter which is present in selected areas of central and peripheral nervous system. The cognitive impairment faced by AD patients is associated with accumulation of NFT, loss of cholinergic functions along with constitution of hyperphosphorylated Tau protein. Both of these enzymes participate in collaboration with plaques, and tangles in AD [85]. In AD subjects, levels of AChE and BuChE are found to be elevated in various regions of the brain. Severity of AD is expressed in terms of higher activity of AChE and BuChE enzymes along with increased number of cortical and neocortical amyloid rich tangles and plaques [86]. Neuronal losses in AD patients' especially in terms of cholinergic neurons; are associated with apoptotic cell death thereby inducing cortical shrinkage in the brain. The cholinergic neurons are responsible for maximum expression of AChE as

compared to remaining neuronal cells [87]. Prolonged availability of AChE in the neuronal cells serves in amelioration of AD symptoms [88]. Contemporary works suggest that inhibition of AChE and BuChE enhances cholinergic transmission in AD. Therefore, current therapeutics involves inhibition of these enzymes with the help of phytochemicals *viz.* galanthamine, rivastigmine, and donepezil.

Rho kinase

Rho kinase is a serine or threonine kinase which is activated by binding to active GTP-bound from Rho. Two known isoforms, ROCK II (ROK α /Rho kinase α) and ROCK I (Rho kinase β /ROK β) are existent which phosphorylates various substrates such as myosin-binding subunit (MSB) of myosin phosphatase and myosin light chain (MCL) [89]. ROCK I is expressed abundantly in non-neuronal tissues such as the stomach, liver, and kidney. Whilst ROCK II is preferentially expressed in the brain and muscle tissues. Several functions of the central nervous system (CNS), for instance, regulation of axonal growth [89], formation of branched dendrites [90], long-term spatial memory [91], regulation of the level of amyloidogenic A

β 42 [92], neurotransmitter release [93], etc. is attributed to Rho/Rho-kinase signalling. In both vertebrates and invertebrates, morphological changes in neurons such as change in dendritic length, size, shape and number of synapses are perceived as hallmarks of long-term memory [89]. Neuronal morphology is controlled by actin filaments and its regulation influences memory. Subsequently, Rho-ROCK pathway is an attractive target of interest for drug discovery works in CNS related disorders. It is recently found that ROCK pathway is closely related to the pathogenesis of several CNS disorders like stroke, AD, spinal cord issues, etc. Poor regeneration of injured axons in adults is observed which is attributed to the presence of myelin-associated growth inhibitors for example Nogo, oligodendrocyte-myelin glycoprotein (OMgp), repulsive guidance molecule (RGM), and myelin-associated glycoprotein (MAG). It is believed that the blockade of Rho-ROCK pathway promotes axonal regeneration and functional recovery of injured CNS which might further help in reversing the effects of these inhibitors. The Rho-ROCK pathway is also an important regulator of cell growth, apoptosis, and migration *via* regulation of actin cytoskeleton assembly [89,92]. Reduced cholesterol dependent and independent mechanisms mediate inhibition of antibody production by statins. Even though precise molecular mechanism behind reduction of $A\beta$ by statins is yet undetermined, it is believed to be associated with enhancement of α -secretase activity. Irrespective of depletion in cellular cholesterol levels, statins inhibit small GTPases such as Rho by lowering protein isoprenylation *via* reduction of mevalonate synthesis [94]. Statin-mediated inhibition of Rho-ROCK results in either the activation of α -secretase cleavage [95] or enhancement in APP lysosomal degradation [96], both of which ultimately result in inhibition of $A\beta$ production. Contemporary works report inhibition of neurite outgrowth by $A\beta$ which is caused through the activation of the Rho-ROCK pathway in H-SY5Y neuroblastoma cells [97]. Inhibitory effects of $A\beta$ is mediated partly through induction of an alternatively spliced form of collapsin response mediator protein-2 (CRMP-2) i.e. CRMP-2A and partly by the upregulated phosphorylation of CRMP-2 by ROCK. Such findings suggest that Rho-ROCK pathway is not only involved in $A\beta$ production but also in $A\beta$ induced neurite outgrowth inhibition. This advocates the beneficiary possibilities of Rho-ROCK blockers in AD treatment.

Prolyl endopeptidase

Prolyl endopeptidase (PEP) is a serine protease enzyme known to cleave peptide substrates at the C-terminal end of proline residues. PEP is actively involved in the metabolism of neuropeptides containing proline such as arginine vasopressin, thyrotropin-releasing hormone, substance P which controls learning and memory process [98-100]. It is found widely distributed among various organs, particularly in the brain of patients with amnesic disorders. Post-mortem analysis of an AD patient's brain reveals significantly increased PEP activity which suggests PEP inhibitor might serve useful as a therapeutic target for an anti-amnesic drug. As per recent

findings, involvement of prolyl endopeptidase-a cytosolic enzyme belonging to a distinct class of serine peptidase in processing of C-terminal region of APP has been found in AD subjects. PEP activity in AD subjects is found to be higher as compared to that of a healthy person [101]. Reports also suggest that prevention of memory loss and enhanced attention span in subjects suffering from senile dementia can be prohibited through specific PEP inhibition. This was also reported in scopolamine-treated and dorsal hippocampal-lesioned rats where memory and learning capability was found to improve [102]. The memory enhancing effects of PEP inhibitor (JTP-4819) is attributed to inhibition of metabolic degradation of brain neuro-peptides by PEP other than enhancement of ACh release. In addition, release of ACh from specific brain regions such as frontal cortex, hippocampus, and regions closely associated with memory is observed as well. Study of PEP inhibitory activity in phytochemicals has therefore turned into a necessity.

Monoglycerol lipase

Monoglycerol lipase (MGL) is a serine hydrolase which converts monoglycerides into fatty acid and glycerol thereby participating in 2-Arachidonoylglycerol (2-AG) inactivation [103]. Likewise, in case of pathologies like neuroinflammation, pain modulation, and neuro-protection; endocannabinoids system has been postulated as the most suitable target [104]. CB1 and CB2 receptors are endogenous endocannabinoid (EC) which along with ligands anandamide (AEA), 2-arachidonylethanolamide, degradation causing enzyme like fatty acid amide hydrolase (FAAH), and monoglyceride lipase (MGL) serve as key elements of the EC system [104]. These are useful in a number of physiological functions such as immune response, cognitive activity, and motor function. In the event of a neurological disorder such as AD, Huntington's disease (HD), multiple sclerosis (MS), etc., study of the neuroprotective roles of EC systems as well as the modulations observed in neurotransmission are studied with equal emphasis [87]. It has been observed in various experimental AD models that EC system faces imbalance in terms of decreased neuronal cannabinoid CB1 receptors, increased glial cannabinoid CB2 receptors, and over-expression of FAAH in astrocytes. Post-mortem report of AD brain and AD animal models have clearly suggested a protective efficacy of EC system [105].

Catechol-O-methyl transferase

Catechol-O-methyl transferase (COMT) affects levels of catecholamines *viz.* dopamine, epinephrine, and norepinephrine by degradation. It is mediated through dopamine signalling in the frontal lobes which is the cause of cognitive impairment [106]. COMT is also a suggested candidate for Alzheimer-related psychosis (AD-P) susceptibility as well as a functional association between valine and methionine polymorphism [106]. Neurotropic factors, like nerve growth factor (NGF), neurotrophin 3 (NT-3) or brain derived neurotrophic factor (BDNF) promote neuron functioning in the peripheral and central nervous system. These are synthesized in the glial and

neuronal cells induced by dopamine and other biogenic amines. Clinical studies have helped to understand the role of monoamine oxidase B (MAO-B) inhibitors which slows down the progression of neurological and cognitive deficits in PD and AD [107]. An increase in biosynthesis of neurotropic factors mentioned herein is noted upon inhibition of activity for the extra-neuronal and neuronal located monoamine oxidase (MAO) enzyme and/or the predominantly glial situated COMT.

AD and Therapeutic Approaches

Over the years, traditional medicines especially those of herbal origin have gained tremendous popularity in AD treatment. At present, extensive research on different plants species is underway globally [108]. Economic viability, higher therapeutic window, along with no or less side effects has helped herbal medicines gain acceptance and fame worldwide [109]. Plants like St John's wort, Kava-kava, Valerian, *Bacopa monnieri*, and *Convolvulus pluricaulis* are among the ones most studied for their effectiveness in the treatment of neurological disorders [110]. Plant studies for AChE activity include *Withania somnifera*, *Semecarpus anacardium*, *Embelia ribes*, *Tinospora cordifolia*, *Ficus religiosa*, and *Nardostachys jatamansi* [111]. Experimental studies confirm the usefulness of plant extract of *Ginkgo biloba* in early stages of AD. In the early stages of AD, it helps patients live a reasonably sound life. In the same direction, works conducted by Selkoe et al. confirm that *Ginkgo biloba* extract has normalising effects on the ACh receptors found at the severely affected hippocampus region of the brain, particularly for older subjects [112]. Selective and competitive inhibition of ACh is also possible with the help of galantamine enzyme. Trails conducted by researchers have emphasised on the significant improvement in disease symptoms of AD and dementia as observed upon treatment with huperzine A [113,114]. Similarly; strong evidences concerning cognitive function improvement, reduced agitation, and related therapeutic activity is shown by *Melissa officinalis* (also called as 'lemon balm') which shows both nicotinic and muscarinic binding properties in the CNS [115,116].

As of now, treatment of AD involves drugs which slow down its progression thereby improving the patient's cognitive functions. The pharmaceutical industry at present offers 'memantine' as the only available drug approved for the treatment of mild to severe AD symptoms. It interferes with the functioning of hippocampal neurons by controlling glutamatergic excitotoxicity [117]. Other drugs available are rivastigmine, galantamine, tacrine, and donepezil. Mechanism of action comprises modulation of brain ACh levels through anti-cholinesterase inhibition. Treatment with anti-inflammatory drugs like prednisone [118], diclofenac [119], rofecoxib [120], and naproxen [121,122] show side effects which is presently the major drawback in AD treatment. Side effects include nausea and vomiting. Hepatotoxicity was detected especially in case of treatment with tacrine, therefore it is rarely used. Donepezil, when given once a day is easily tolerated along

with improving mental capabilities in AD patients. Similarly, rivastigmine works well when given twice a day.

Conclusion

The lack of adequate knowledge on the molecular mechanisms of AD pathogenesis remains a leading cause underlying inadequate treatment and mortality. Progressive research has significantly identified some of the crucial developments of AD pathology; however, the molecular mechanisms, enzymatic actions, signaling pathways involved in AD pathogenesis are not eloquently established. Hence, finding potential biomarkers, drug targets, development of drugs and novel delivery systems is a major challenge. Many studies showed positive and satisfactory outcomes with the usage of herbal medicine and antioxidants but the clinical effectiveness and scope of progression in research are limited due to various factors associated with traditional and herbal medicine. This indicates the need of early diagnostic modalities, potential biomarker identification and designing novel inhibitors or drugs.

The on-going researches in the fields of molecular biology and biomedical engineering are attempting to unfold the underlying molecular events and to explore the mechanistic link between enzymes and AD pathogenesis. It is believed that, in AD pathogenesis amyloid-degrading enzymes play a significant role in enzymatic pathway and serve in A β clearance. Among the number of enzymes involved, it is very crucial and important to know which of these enzyme(s) are critical and can be served as potential drug targets and easy to regulate. Considering the fact that AD pathology develops over many years, therefore, the early manipulation of these enzymes is essential as it could reduce the disease progression. Identifying the link between AD and other pathological events is important in finding targets. In addition, development of modifiers of AD may also serve as therapeutics for AD. Pharmacological approaches are also important to provide economical treatment options for AD; in addition, development of effective drug delivery systems other than gene delivery systems such as exosomes help in avoiding safety and ethical concerns. Within last two decades, the advancements in biomedical sciences have provided promising insights into targets, designing of drugs and other potent therapeutic molecules associated with AD; therefore, search should be continued to produce 'anti-amyloid' to combat the progression of AD.

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