# Review

# Amyloid- $\beta$ and Mitochondria in Aging and Alzheimer's Disease: Implications for Synaptic Damage and Cognitive Decline

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**Abstract**. This article reviews the role of amyloid- $\beta$  (A $\beta$ ) and mitochondria in synaptic damage and cognitive decline found in patients with Alzheimer's disease (AD). Recent molecular, cellular, animal model, and postmortem brain studies have revealed that A $\beta$  and mitochondrial abnormalities are key factors that cause synaptic damage and cognitive decline in AD. A $\beta$  is reported to accumulate in subcellular compartments and to impair the normal function of neurons in AD patients. Further, recent studies using biochemical methods and electron microscopy have revealed that the accumulation of A $\beta$  at nerve terminals affect synaptic activities, including the release of neurotransmitters and synaptic vesicles. Recent studies of the relationship between mitochondria and A $\beta$  in AD patients suggest that in mitochondria, structural changes caused by A $\beta$  result in increased mitochondrial fragmentation, decreased mitochondrial fusion, mitochondrial dysfunction, and synaptic damage. This paper discusses the latest research on A $\beta$ , mitochondria, age-dependent factors of AD in the brain, and synaptic damage in AD. This paper also briefly discusses potential mitochondrial therapeutics in the treatment of patients with AD.

Keywords: Amyloid- $\beta$ , Amyloid- $\beta$  precursor protein, mitochondrial therapeutics, synaptic pathology

# INTRODUCTION

Alzheimer's disease (AD) is a late-onset mental illness that is characterized by the loss of memory and an impairment of multiple cognitive functions [1–3]. The major pathological features in the brains of AD patients are the presence of intra-neurofibrillary tangles and extracellular protein amyloid- $\beta$  (A $\beta$ ) deposits, particularly in the regions related to memory and cogni-

tion [4]. Currently, 5 million Americans suffer from AD [5]. It is estimated that by the year 2050, 50% of people worldwide (approximately 370 million) who are 85 years of age or older will be afflicted with AD [4, 6]. With such a large, aged population poised to be afflicted, AD has become a major health concern that must be reckoned with. Despite tremendous progress that has been made in AD research, we still do not have early detectable markers, nor agents or drugs that can slow the progression of AD.

Causal factors for Alzheimer's disease

Genetic mutations in the amyloid- $\beta$  protein precursor (A $\beta$ PP), presenilin 1 (PS1), and PS2 genes cause

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Table 1
Genes and risk factors involved in Alzheimer's disease pathogenesis

Chromosome	Gene	Phenotype/pathology	Ref.
Genes in Famil	lial AD		
21	Amyloid- $\beta$ protein precursor	Mutations in A $\beta$ PP are causal factors of FAD and are involved in the increased production of all A $\beta$	[7]
14	Presenilin 1	Mutations in PS1 are causal factors of FAD and are involved in gamma secretase activity and in the increased production of $A\beta_{1-42}$	[8]
1	Presenilin 2	Mutations in PS2 are causal factors of FAD and are involved in gamma secretase activity and in the increased production of $A\beta_{-42}$	[9]
Genes Involved	d in AD as a Risk Factor	•	
19	Apolipoprotein E allele 4	E4 polymorphism is a risk factor for late-onset AD and is involved in the increased production of $A\beta$	[10]
11	Sortilin related receptor	Involved in increased production of $A\beta$	[11]
8	Clusterin	Clusterin is expressed abundantly in the brain and involved in clearing of $A\beta$ from brain to plasma. However, it is also reenter to the brain, and involved in decreased clearance of $A\beta$	[12,13]
1	Complement component receptor 1	Involved in clearance of $A\beta$ . However, variants in complement component receptor 1 interfere with $A\beta$ clearance	[12,13]

a small proportion (about 2%) of all AD cases (earlyonset or familial AD; Table 1); however, causal factors are still unknown for a vast majority of AD patients (late-onset AD). Recent genetic studies have identified several risk factors for late-onset AD, including genetic variants in the sortilin-related receptor 1 gene; clusterin (a protein associated with the clearance of cellular debris); the complement component receptor 1; and apolipoprotein E4 (ApoE  $\varepsilon$ 4) genotype [7–13] (Table 1). In the last decade, tremendous progress has been made in understanding the role of ApoE  $\varepsilon$ 4 involvement in AD progression and pathology [14–18, 18,20,21]. In addition, several other factors, including epigenetics, lifestyle, diet, and environmental exposure may contribute to the development of late-onset AD [3]. Recently, oxidative stress and mitochondrial abnormalities have been implicated in the development of late-onset AD [22–26], and aging has been identified as the 'number 1' risk factor in AD progression and pathology.

# Sites of pathology in Alzheimer's disease

Anatomical and immunohistochemical analyses of AD postmortem brains and the brains from AD transgenic mice revealed that a neurodegenerative process is initiated in layer 2 of the entorhinal cortex [4]. This process spreads to the hippocampus, temporal cortex, frontoparietal cortex, and, finally, to subcortical nuclei. Interestingly, in AD patients, these regions of the brain are involved in learning, memory, and cognitive functions [4].  $A\beta$  secretion also occurs mainly in these regions, as do  $A\beta$  deposits; however, the reasons for  $A\beta$  secretion and the formation of  $A\beta$  deposits in these areas are not fully understood.

# Cellular changes

More than two decades of intense research has revealed that AD is a complex, heterogeneous disease, with multiple cellular changes implicated in its pathogenesis [1,2,4,8,27,28]. Major cellular changes that have been implicated in AD are: 1)  $A\beta$  and amyloid cascade events; 2) hyperphosphorylation of tau and intracellular neurofibrillary tangles; 3) synaptic pathology and neuronal loss; 4) mitochondrial structural and functional abnormalities; and 5) inflammatory responses [22,29–46].

# Early events in Alzheimer's disease progression

Although multiple cellular changes have been reported to be involved in AD pathogenesis, synaptic pathology and mitochondrial oxidative damage have been identified as early events in AD progression and pathogenesis [3]. It is generally accepted that an accumulation of  $A\beta$  in synapses and in synaptic mitochondria, particularly in neurons affected by AD, cause synaptic degeneration and cognitive decline in AD patients [2, 3,47–50].

In this article, we first focus on  $A\beta$ , its generation, accumulation, and age-related factors of  $A\beta$ ; next we focus on mitochondrial structural and functional abnormalities, and synaptic damage in AD progression and pathology.

# AMYLOID- $\beta$

In AD,  $A\beta$ , the 39–43 amino acid residue protein, is a major component of neuritic plaques found in brain

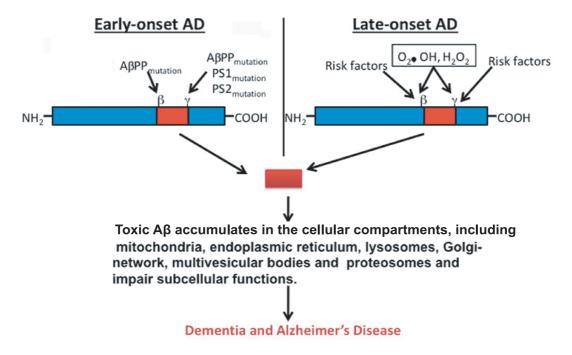


Fig. 1. Amyloid- $\beta$  secretion in AD neurons. In early-onset AD, mutations in A $\beta$ PP, PS1, and PS2 genes activate  $\beta$ - and  $\gamma$ -secretases, cleave A $\beta$ . In late-onset AD, oxidative stress related factors (O2 $_{\bullet}$ -, H<sub>2</sub>O<sub>2</sub>, OH), risk factors including ApoE4, sortilin related receptor 1, clusterin, and complement component either activate secretases or decrease A $\beta$  clearance in AD neurons. The cleaved A $\beta$  accumulate in subcellular compartments, including mitochondria, endoplasmic reticulum, Golgi-network, lysosomes, and multivesicular bodies, and disrupt the functions of these subcellular organelles and damage neuronal function.

regions known to be responsible for learning and memory [51]. A $\beta$  is generated by proteolysis of A $\beta$ PP by the sequential enzymatic actions of  $\beta$ -site A $\beta$ PP cleaving enzyme 1 (BACE 1),  $\beta$ -secretase, and  $\gamma$  secretase. In early-onset AD, genetic mutations in A $\beta$ PP, PS1, and PS2 genes activate  $\beta$ - and  $\gamma$ - secretases, and cleave  $A\beta$  [3].  $A\beta PP$  mutations that flank the  $A\beta$  domain increase the production of  $A\beta_{42}$  (Fig. 1). In late-onset AD, it has been hypothesized that factors related to oxidative stress may be involved in activating  $\beta$ - and  $\gamma$ - secretases, cleaving A $\beta$ PP, and releasing A $\beta$  [25] (Fig. 1). Recently, several in vitro and in vivo studies have provided experimental evidence to support to this hypothesis [51-56] (Fig. 1). In both early onset and late-onset AD, levels of A $\beta$  are steady-state and are controlled by the production of  $A\beta$ , the clearance of  $A\beta$ , and the degradation of  $A\beta$ . Decreased clearance of  $A\beta$  or the overproduction of  $A\beta$  may lead to an accumulation of  $A\beta$  in subcellular compartments and may initiate a cascade of events in the brain, a process referred to as "A $\beta$  cascade hypothesis" [57]. Interestingly,  $A\beta$  can self-aggregate into multiple forms, ranging from 4 kDa monomers to oligomers and to fibrils. These fibrils eventually form  $\beta$ -pleated sheets, insoluble fibers, and deposits (Fig. 2). Soluble oligomers are

the most toxic form of  $A\beta$  for neurons. Anatomical analyses of AD postmortem brains and AD transgenic mice revealed that  $A\beta$  secretion occurs mainly in the entorhinal cortex, hippocampus, temporal cortex, and frontoparietal cortex of the brain. These areas are important for learning, memory, and cognitive functions. The reasons for  $A\beta$  secretion and formation of amyloid deposits in these areas are not fully understood. However, it is possible that the capability of  $A\beta$  clearance is low in these areas or that oxidative stress is high, in which case oxidatively damaged neurons may produce more  $A\beta$ . This loop – increased production of  $A\beta$  and decreased clearance – may lead to an excessive accumulation of  $A\beta$  in brains.

Recent studies have revealed that  $A\beta$  is secreted wherever  $A\beta PP$  and the  $\beta$ - and  $\gamma$ -secretases are present.  $A\beta PP$ , and the  $\beta$ - and  $\gamma$ -secretases, are localized to several cellular compartments, including the endoplasmic reticulum, plasma membrane, trans-Golgi network, and multivesicular bodies [2,3]. Because of the presence of  $A\beta PP$  and the  $\beta$ - and  $\gamma$ -secretases in these subcellular compartments,  $A\beta$  is known to be present in these regions. In addition, several studies reported that  $A\beta$  is localized to mitochondrial membranes [37,38,46].

# Multiple Forms of Aβ Aβ monomers 1-40 1-42 Fibrils Oligomers Protofibrils Deposits

Fig. 2. Formation of amyloid- $\beta$  species and deposits in AD brain.

Age-dependent increase in the production of  $A\beta$ 

Multiple lines of evidence suggest that aging is a key factor for the increased production of  $A\beta$  and the decrease in  $A\beta$ -degrading enzymes in the AD brain. A time-course analysis of  $A\beta$  in AD transgenic mouse lines revealed that  $A\beta$  levels and deposits increase in AD-affected regions of the brain in an age-dependent manner [2,38]. In addition, studies of postmortem brains from aged humans with mild cognitive impairment (MCI) and patients with AD found an age-dependent increase of  $A\beta$  levels in the aged persons with MCI and in AD patients [58]. These studies suggest that aging plays a key role in the production and accumulation of  $A\beta$  in the brains of AD patients and in AD transgenic mice.

In the last decade, a large body of research has been devoted to understanding  $A\beta$  toxicity, particularly intracellular  $A\beta$ . It is now generally accepted that extracellular  $A\beta$  deposits are the by-products of AD pathology. Recent studies of AD patients and AD transgenic mice found intracellular  $A\beta$  present in AD-affected brain regions [38,58–67].  $A\beta_{1-42}$  participates mainly in fibrillogenesis and the formation of  $A\beta$  deposits. It is generally accepted that intracellular  $A\beta$  has been found to precede extracellular  $A\beta$  deposits in AD brains [68]. In addition, several studies of AD transgenic mice reported that intracellular  $A\beta$  accumulates early in AD progression [2,3].

Aging and deceased levels of  $A\beta$ -degrading enzymes

Several molecular and cellular studies have revealed that  $A\beta$ -degrading enzymes, including nephrilysin (NEP) and the insulin-degrading enzyme (IDE), decrease as disease progresses in AD patients [69–72].

Hellström-Lindahl and colleagues [69] investigated whether decreased NEP levels contribute to the accumulation of  $A\beta$  in AD patients and in aged persons without AD. Protein levels of NEP were reduced in the temporal and frontal cortex of brains from AD patients and aged patients without AD. They found an inverse correlation between NEP and insoluble  $A\beta$  levels in both groups, suggesting that NEP is involved in the clearance of  $A\beta$ . The observed, age-dependent decline in NEP may be related to the increased levels of  $A\beta$  found in aged patients without AD, during normal aging.

Mohajeri et al. [70] measured NEP levels in AD mice. Neuronal upregulation of NEP in young AD transgenic mice expressing the mutant  $A\beta PP$  led to the reduction of  $A\beta$  levels and the delayed formation of  $A\beta$  deposits. In contrast, a comparable increase of NEP levels in the brains from aged  $A\beta PP$  mice (swe) with pre-existing  $A\beta$  deposits did not result in a significant reduction of plaque pathology. They suggested that the use of NEP for AD therapy might be most effective early in the course of AD pathophysiology since NEP is age-dependent.

Apelt and collaborators [71] measured the levels of mRNA and proteins of NEP in  $A\beta PP$  transgenic mice during postnatal maturation and aging. NEP levels were decreased in the cerebral cortex of mice 2–22 months old, independent of their transgene status. Immunocytochemistry revealed few NEP-positive dystrophic neurites around  $A\beta$  plaques and an upregulation of NEP in plaque-surrounding reactive astrocytes, which suggests a role for  $A\beta$  deposit-mediated astrogliosis in  $A\beta$  degradation.

Iwata et al. [72] sought to determine whether spatial changes in NEP correlate with  $A\beta$  in AD-affected regions of brains from AD transgenic mice. When

NEP levels in various brain regions of 10-, 80-, and 132-week-old AD transgenic mice were evaluated by an NEP-dependent, endopeptidase-activity assay and Western blot quantitative analysis, a clear change in NEP levels was observed in the hippocampal formation, levels reduced by 20% at 132 weeks, compared to the 10-week group. In addition, quantitative immunohistochemical analysis confirmed the reduction of NEP levels in the outer molecular layer and in the polymorphic layer of the dentate gyrus, and in the stratum lucidum of the hippocampus, by 56%, 82%, and 83% respectively in the mice at 132 weeks, compared to the 10-week group. NEP levels were decreased at the terminal zones in axons of the lateral perforant path and in mossy fibers. These are also the brain sites that exhibit disease pathology in mutant A $\beta$ PP transgenic mice and synaptic loss in AD patients.

Similar to NEP, IDE is also an important enzyme that is involved in  $A\beta$  clearance and that has been found to be decreased in the brains of aged persons without AD [73,74]. The concentration of IDE and its activity were significantly decreased in the hippocampus in the brains from aged humans without MCI, compared to humans who were mildly cognitively impaired and aged persons who were considered at high risk to develop AD [73]. Membrane-bound IDE concentrations and IDE activity in the hippocampus continued to decrease as the patients progressed from MCI to mildsevere AD. Most interestingly, IDE activity in the hippocampal membrane negatively correlated with  $A\beta_{42}$ in the brains from aged persons with MCI and with AD. Findings from the Zhao et al. [73] study suggest that interventions aimed at promoting membrane-bound IDE activities in the brain of aged persons with MCI may help to prevent the onset and possibly the progression of AD through mechanisms involving the clearance of monomeric  $A\beta$  from the brain.

Farris and colleagues [74] recently studied the connection between IDE gene and  $A\beta$  using a rat model for IDE. In a well-characterized rat model of type 2 diabetes mellitus, they found naturally occurring IDE missense mutations, which decreased catalytic efficiency, and they found a significant deficit (about 15 to 30%) in the degradation of both insulin and  $A\beta$ . Endogenously secreted  $A\beta_{40}$  and  $A\beta_{42}$  were significantly elevated in primary neuronal cultures from animals with the IDE mutations. These researchers concluded that naturally occurring, partial loss-of-function mutations in IDE were sufficient to cause diabetes mellitus 2 and impaired neuronal regulation of  $A\beta$  levels. However, they noted that the brain apparently compensates for the partial deficit during the life span of the rat [74].

# MITOCHONDRIA AND ALZHEIMER'S DISEASE

Mitochondrial dysfunction in AD pathogenesis was described two decades ago, but its underlying mechanisms were not clear until recently. Mitochondrial dysfunction has been found and described in postmortem brains from patients with AD [39,75–77], in their platelets [78], in AD transgenic mice [34,37,38, 79–81], and in cell lines that express mutant  $A\beta$ PP and/or cells treated with  $A\beta$  [82–84].

Increasing evidence suggests that mitochondrial abnormalities play a large role in AD pathogenesis. Decreased mitochondrial enzymes, including cytochrome oxidase activity, pyruvate dehydrogenase, and  $\alpha$ ketodehydrogenase were found in fibroblasts, lymphoblasts, and postmortem brains from AD patients relative to age-matched control subjects (reviewed in [3]). Further, a recent study described abnormal mitochondrial dynamics in fibroblasts from AD patients, indicating that impaired mitochondrial dynamics are involved in AD pathogenesis [85]. Several other studies found increased free radical production, lipid peroxidation, oxidative DNA damage, oxidative protein damage, decreased ATP production, and decreased cell viability in brains from AD patients compared to those from age-matched control subjects [39,75–77,86].

In the 1990s, Swerdlow and colleagues [88] studied mitochondrial function using a cytoplasmic hybrid (cybrid) approach to determine the role of mitochondrial DNA (mtDNA) in AD pathogenesis [87,88]. They isolated platelets from AD and age-matched control subjects, and fused those platelets with both human neuroblastoma (SH-SY5Y) cells and human teratocarcinoma (NT2) cells depleted of their endogenous mtD-NA [87,89]. The cells lacking mtDNA did not exhibit mitochondrial functional activities; that is, only those cells with mtDNA exhibited intact mitochondrial functional activities. However, cells containing cybrids of AD and control subjects exhibited mitochondrial respiratory activities, with a difference between the cybrid cell lines containing AD subject mitochondria and cybrid cell lines with control subject mitochondria. AD cybrid cell lines exhibited increased A $\beta_{42}$  production and mitochondrial dysfunction: the cytochrome oxidase activity was lower, free radical production and oxidative stress markers were elevated, calcium homeostasis was altered, the mitochondrial membrane potential was reduced, and apoptosis pathways were altered [88]. Findings from these cybrids studies further support that mitochondria are involved in AD pathogenesis.

In other studies, increased mitochondrial DNA changes were found in postmortem brain tissue from AD patients and age-matched control subjects, compared to DNA changes in brain tissue from young control subjects without AD [90,91]. These findings suggest that the accumulation of mitochondrial DNA in AD pathogenesis is age-related.

The Reddy laboratory [34,92] and others [93–95] found that mitochondrial encoded genes were abnormally expressed in AD postmortem brains and in those from AD transgenic mice. Recently, we [34] investigated gene expression profiles in brain slices from A $\beta$ PP transgenic mice at 3 stages of AD progression: long before (2 months of age), immediately before (5 months), and after (18 months) the appearance of A $\beta$  plagues in the cerebral cortex [34]. We compared those profiles to those of age-matched wild-type mice. Our analysis revealed that the genes related to mitochondrial energy metabolism and apoptosis were upregulated in the 2-month-old A $\beta$ PP transgenic mice and that the same genes were upregulated in these mice at 5 and 18 months of age. In another study, we found decreased cytochrome oxidase, increased free radicals, and increased carbonyl proteins in the 2-month-old A $\beta$ PP transgenic mice compared to the age-matched wild-type mice [38]. Taken together, these results suggest that mitochondrial energy metabolism is impaired by mutant A $\beta$ PP and/or A $\beta$ , and that the upregulation of mitochondrial genes may be a compensatory response to this impairment. Further, we found abnormal mitochondrial gene expression in the 2-month-old A $\beta$ PP transgenic mice, suggesting that mitochondrial dysfunction is an early event in AD progression.

Using quantitative RT-PCR techniques, the Reddy laboratory also analyzed mRNA expression in 11 mitochondrial-encoded genes from the frontal cortex of 3 subject groups: patients with early AD, patients with definite AD, and age-matched control subjects [92]. This analysis revealed a down-regulation of mitochondrial genes in complex I of electron transport chain genes in both early and definite AD brain specimens, but not in the control subjects. In the brain specimens from both the early and definite AD patients, complex I showed a down-regulation of mitochondrial genes, but complexes III and IV showed increased mRNA expressions, suggesting a great demand for energy production in the brains from AD patients. These results suggest that mitochondrial dysfunction is an early event in AD progression and continues into later-stage AD progression, and that abnormal mitochondrial gene expression may be a compensatory response to mutant  $A\beta$ -PP- and  $A\beta$ -initiated mitochondrial toxicity.

Further, the Reddy laboratory [38] and others [37,39, 41,46,96] found that  $A\beta PP$  and  $A\beta$  are localized to mitochondrial membranes and is responsible for generating free radicals and initiating mitochondrial dysfunction. Other groups found presequence peptidase, a peptidase that is known to degrade  $A\beta$  species, in the mitochondria of AD neurons [97], further supporting the association of  $A\beta$  with mitochondria and mitochondrial dysfunction in AD. In addition, recent studies of mitochondrial structure and of neuronal cells expressing mutant  $A\beta PP$  in brain tissues from AD patients found that  $A\beta$  fragments mitochondria and causes structural changes in neurons [55,98,99].

Overall, findings from these studies suggest that mitochondrial abnormalities occur early in AD progression.

# $A\beta PP$ , $A\beta$ , AND ABNORMAL MITOCHONDRIAL DYNAMICS IN ALZHEIMER'S DISEASE

Increasing evidence suggests that mutant  $A\beta PP$  and/or  $A\beta$  overexpression cause mitochondrial fragmentation in neurons affected AD [98–100]. In a recent gene expression study of AD transgenic mice, the Reddy laboratory [34] found increased expression of mitochondrial – encoded genes in AD affected regions of the brain that may be due to the excessive production of mitochondria. This overproduction of mitochondria may be due to mutant  $A\beta PP$  and  $A\beta$  toxicity in neurons affected by AD [34,38,99].

The relationship between the overexpression of mutant  $A\beta PP$  and  $A\beta$ , and the increased production of mitochondria is supported by several studies.

1)  $A\beta PP$ , and monomeric and oligomeric forms of  $A\beta$  have been found in mitochondrial membranes [37–39,46,79,96,101]. In the Lustbader et al. study [79], they found  $A\beta$  normally interacting with the mitochondrial matrix protein ABAD, leading to mitochondrial dysfunction. Caspersen et al. [37] found an accumulation of  $A\beta$  in the mitochondria from postmortem brain specimens of AD patients and  $A\beta PP$  transgenic mice. Recently, the Reddy laboratory found  $A\beta$  monomers and oligomers in mitochondria isolated from the cerebral cortex of  $A\beta PP$  transgenic mice [38] and from N2a cells expressing  $A\beta PP$ . Our digitonin fractionation analysis of isolated mitochondria from  $A\beta PP$  trans-

genic mice revealed  $A\beta$  in the outer and inner membranes and matrix of mitochondria. Our study also found that mitochondrial  $A\beta$  decreases cytochrome oxidase activity and increases free radicals and carbonyl proteins. Yao et al. [46] found  $A\beta$  mitochondrial membranes in cortical tissues from triple transgenic mice.

- 2) Using confocal and electron microscopy, and human neuroblastoma (M17) cells transfected with wild-type or mutant A $\beta$ PP, Wang and coworkers [98] investigated the effects of A $\beta$ PP and A $\beta$  on mitochondrial structural changes. Confocal and electron microscopic analysis revealed that about 40% of M17 cells overexpressing wild-type A $\beta$ PP and more than 80% of M17 cells overexpressing mutant A $\beta$ PP displayed alterations in mitochondrial morphology, particularly fragmented mitochondria. They also found that increased levels of Fis1 are critical for mitochondrial fission in A $\beta$ PPwt and A $\beta$ PPswe M17 cells. The overexpression of A $\beta$ PP and/or A $\beta$ -derived diffusible ligand treatment also led to mitochondrial fragmentation and morphological changes.
- 3) Using electron and confocal microscopy, gene expression analysis, and biochemical methods, the Reddy laboratory studied mitochondrial structure and function, and neurite outgrowth in neurons treated with  $A\beta$  [99]. In neurons treated with only  $A\beta$ , we found increased expressions of mitochondrial fission genes (Drp1 and Fis1) and decreased expressions of fusion genes (Mfn1, Mfn2, and Opa1), indicating abnormal mitochondrial dynamics in AD neurons. mRNA expression of antioxidant enzyme-encoded genes (peroxiredoxins 1-6) was significantly decreased in neurons treated with  $A\beta$  relative to untreated neurons. Our electron microscopy of neurons treated with A $\beta$  revealed a significant increase in mitochondrial fragmentation, further supporting abnormal mitochondrial dynamics. We also found significantly decreased neurite outgrowth and decreased mitochondrial function in cells treated with A $\beta$  [99]. These findings suggest that  $A\beta$  fragments mitochondria and causes abnormal mitochondrial dynamics, leading to mitochondrial dysfunction.
- 4) Zhao and colleagues [102] studied the effects of wild-type and an arctic form of  $A\beta_{42}$  using neurons from adult flies. They performed extensive time-course analyses to determine the function and structure of both axon and presynaptic terminals of individual neurons. They found  $A\beta$  accumulated intracellularly, and they found a wide range of age-dependent changes, including the depletion of presynaptic mitochondria, a slow-down of bi-directional transports of axonal mitochon-

dria, decreased synaptic vesicles, increased large vacuoles, and elevated synaptic fatigue. These structural and functional synaptic changes correlated with agedependent deficits in the motor behavior of the flies. Such changes were accelerated in flies expressing the arctic form of  $A\beta$ . The depletion of presynaptic mitochondria was the earliest phenotype that they were able to detect in the fly. Zhao et al. [102] determined this depletion was not caused by the change in axonal transport of mitochondria. They also found a dramatic reduction in the number of axonal mitochondria and also a significant increase in their size, in aged A $\beta$ -expressing flies, suggesting a global depletion of mitochondria in the neuron and an impairment of mitochondria fission. These results suggest that A $\beta$  accumulation depletes presynaptic and axonal mitochondria, leading to other presynaptic deficits.

Taken together, these findings suggest  $A\beta$  enters mitochondria and causes abnormal mitochondrial dynamics in neurons that are known to be affected in AD, and that such abnormal mitochondrial dynamics cause mitochondrial dysfunction and abnormal mitochondrial trafficking in AD neurons.

# Abnormal mitochondrial trafficking in AD

In a process called mitochondrial trafficking, mitochondria travel along the axons and dendrites to supply energy to nerve terminals for normal neural communication; then they travel back to the cell body [103]. Mitochondria are transported from the cell body back to nerve terminals via an anterograde mechanism and from nerve terminals to the cell body via a retrograde mechanism. In healthy neurons, anterograde and retrograde transport of mitochondria are equal and active. In AD neurons, both anterograde and retrograde transport of mitochondria are slow because of the presence of large number of defective and functionally inactive mitochondria [104,105]. As discussed earlier, AD mitochondria with an accumulation of A $\beta$  disrupt mitochondrial function and inhibit ATP production. These  $A\beta$ -laden mitochondria are not able to supply sufficient levels of energy to the nerve terminals, which may impair neurotransmission and may ultimately result in synaptic damage, neurodegeneration, and cognitive decline in AD patients [103].

# $A\beta$ , SYNAPTIC ALTERATIONS, AND MITOCHONDRIAL DAMAGE IN ALZHEIMER'S DISEASE

Synapses are the sites of high-energy demand [3].

Synaptic damage is considered the earliest cellular event in AD pathogenesis, and synaptic loss is the best correlate of cognitive impairment in AD [29–31, 106]. Damaged synapses due to insufficient mitochondrial ATP lead to synaptic degeneration [3]. Synapses and neurites are mostly damaged in the vicinity of  $A\beta$  plaques [107,108].

In healthy subjects, synaptic terminals transmit signals between cells in order to process information. During aging, the number of synapses and their transmissions of signals dramatically decrease [109,110]. The decrease in synapses has been documented in different brain regions of aged persons, supporting the hypothesis that synaptic changes are ubiquitous features of normal brain aging [3].

# Synaptic loss and Alzheimer's disease

Several studies using electron microscopy and AD postmortem brains revealed a loss of synapses in the hippocampus of AD patients compared the number of synapses in control subjects [39,111,112]. This loss correlates with cognitive decline in AD patients.

Bretoni-Freddari et al. [111] studied the number of synapses per neuron in cerebellar and hippocampal brain tissues from adult and aged control subjects and from AD-affected and unaffected brain tissues in patients with AD. The synapse-to-neuron ratio varied according to the brain regions from which the samples were taken and the individual's health. No significant differences in the synapse-to-neuron ratio were found in samples taken from the cerebellum of adult and aged persons without AD and of aged AD patients. However, in the hippocampal samples, the synapse-to-neuron ratio decreased more than 50% in the adult and aged persons without AD, compared to the ratio in the aged AD patients.

In several studies investigating the extent that synaptic loss correlates with cognitive decline in AD patients [31,112] researchers found a 25–30% decrease in synapses in the cortex and a 15–35% decrease in synapses per cortical neuron, suggesting that synaptic loss in AD patients may correlate more with cognitive decline than with the number of A $\beta$  plaques, neurofibrillary tangles, neuronal loss, or the extent of cortical gliosis.

# Loss of synaptic proteins and Alzheimer's disease

Using immunoblotting and immunohistochemical analyses to determine synaptic proteins, several stud-

ies revealed decreased levels of presynaptic (synaptophysin) and postsynaptic proteins (synaptopodin and PSD95) in AD patients compared to age-matched control subjects [113–118], suggesting that presynaptic and postsynaptic proteins are critically involved in AD progression. Further, immunoblotting analysis of postmortem brain tissues from the cerebral cortex of AD transgenic mice also revealed decreased levels of synaptic proteins in AD transgenic mice [119], suggesting that the loss of synaptic proteins are confined to brain regions known to be affected in AD.

Oligomeric  $A\beta$ , synaptic damage, and impairments in long-term potentiation

Several recent in vitro and in vivo studies revealed that oligomeric  $A\beta$  is responsible for a decrease in the long-term potentiation (LTP) and disruption of synaptic plasticity [63,120–125], in addition to A $\beta$  pathology. Intracellular A $\beta$  was detected in 5XFAD mice 1.5 months of age, and cognitive impairments and LTP abnormalities were found in 5XFAD mice 5-6 months of age [63]. In a triple transgenic mouse model of AD, intracellular  $A\beta$  was found in mice 3 months of age and LTP impairments in mice 6 months of age, indicating that intracellular  $A\beta$  may be critical for synaptic damage and cognitive impairments [66]. In a wellcharacterized AD transgenic mouse model (Tg2576), significantly decreased synapse density was observed in the outer molecular layer of the dentate gyrus in mice 6-9 months of age and in layers II and III of the cortex in mice 15–18 months of age [106]. These results suggest that synaptic changes caused by soluble  $A\beta$ may contribute to the loss of synapses and of synaptic proteins and may be responsible for cognitive decline in AD patients.

In addition to synaptic pathology [120–125], mitochondrial alterations were also observed in several lines of transgenic mice [37,38,46], indicating that both synaptic damage and mitochondrial abnormalities, particularly the accumulation of  $A\beta$  in mitochondria, may have a role in triggering cognitive deficits in AD transgenic mice. Recent mitochondrial and synaptic studies revealed that mitochondria are distributed abnormally, suggesting that abnormal mitochondrial distribution may contribute to synaptic damage in AD [102,126]

Further, using electron microscopy and a rapid Golgi method, Baloyannis and collaborators [127] investigated synaptic alterations, including synapses and dendritic spines in subjects with early AD and in control subjects. They found substantial loss of synapses and

synaptic alterations in the medial geniculate bodies in neurons from the AD subjects. In particular, mitochondrial alterations and fragmentation of Golgi apparatus were seen in the neurons of the medial geniculate bodies and of the inferior colliculi. These findings suggest that mitochondrial and synaptic alterations in the medial geniculate bodies and inferior colliculi are involved in the impairment of neuronal communication and symbolic sound perception in the early stages of AD progression.

Overall, findings from studies of postmortem brains from AD patients and AD transgenic mice revealed synaptic damage in neurons affected by AD, and suggest that an accumulation of oligomeric  $A\beta$  at synapses and synaptic mitochondria cause synaptic damage and cognitive impairments. Therefore, therapeutics targeting  $A\beta$  and mitochondria may be helpful in reducing the progression of AD in AD patients.

### MITOCHONDRIAL THERAPEUTICS

Currently, several laboratories are involved in developing therapeutic strategies to delay or prevent progression and development of AD. Several groups are focused on developing and testing antioxidants that target mitochondria [128–132] and several others are involved in developing and testing drugs that target A $\beta$ , such as drugs that involve immunotherapy, and  $\beta$ - and  $\gamma$ -secretase inhibitors. Both approaches have shown promise at preclinical levels, meaning they have been successful in AD transgenic mice but essentially unsuccessful in clinical trials.

Since synaptic damage and mitochondrial dysfunction have been reported as early pathogenic events associated with aging and AD, it may be possible to treat these pathogenic events by: 1) developing molecules that treat mitochondria (by targeting ROS); these molecules would decrease free radical production and oxidative damage, and boost overall mitochondrial function, which would ultimately increase synaptic branching of neurons; such increased branching would increase neural communication; and 2) therapeutically boosting ATP levels in mitochondria, which would ultimately increase synaptic outgrowth and neuronal connectivity.

Given the significant involvement of mitochondrial dysfunction in aging and AD, it seems reasonable to delay their progression in patients with neurodegenerative diseases such as AD, through antioxidant treatment or a diet supplemented diet with antioxidants. However, recent studies of AD patients' intake of natural antioxidants gave mixed results. Findings from some epidemiologic studies point to the increased intake of antioxidant vitamins, including vitamin E, vitamin C, and beta carotene, to possibly reduce the risk of developing AD [3]. However, findings from other studies do not. They found that their antioxidant approaches did not decrease the risk of developing AD in elderly people, which suggests that antioxidant approaches will not be effective in treating neurodegenerative diseases because naturally occurring antioxidants, such as vitamins E and C, do not cross the blood-brain barrier and so cannot reach the relevant sites of free radical generation [3]. To overcome these problems and to better assess whether antioxidant approaches may be valuable therapeutic treatments, improved delivery of antioxidants to the brains of AD patients is needed.

In the last decade, tremendous progress has been made in developing mitochondrially-targeted antioxidants that are capable of crossing the blood-brain. To increase the delivery of antioxidants into mitochondria, several antioxidants have been developed: triphenylphosphonium-based antioxidants (MitoQ, MitoVitE, and MitoPBN); the cell-permeable, small peptide-based antioxidant SS31; and mitochondrial-permeability transition pore inhibitors such as Dimebon [128–133]. Several laboratories across the world are investigating neuroprotective molecules including those that can target antioxidants to mitochondria, such as SS31, MitoQ, and Dimebon, but the research is at its infant stages and is currently focused on animal models of AD.

# CONCLUSIONS AND FUTURE DIRECTIONS

Increasing evidence suggests that  $A\beta$ , mitochondrial dysfunction, and synaptic damage are critically involved in AD progression and development. The latest research into AD revealed that  $A\beta$  and mitochondrial abnormalities are key factors that cause synaptic damage in AD neurons.  $A\beta$  is reported to accumulate in subcellular compartments and to impair the normal function of neurons. Further, recent *in vitro* and *in vivo* studies of  $A\beta$  using biochemical methods and electron microscopy revealed that the accumulation of  $A\beta$  at nerve terminals damages synaptic activities, including the release of neurotransmitters and synaptic vesicles. Further, recent discoveries of mitochondria in AD suggest that structural changes in mitochondria, including increased mitochondrial fragmentation and

decreased mitochondrial fusion, are critical factors associated with mitochondrial dysfunction and synaptic damage in AD. Despite tremendous progress that has been made in AD research, we still do not have drugs or other agents to prevent or slow down disease progression. Further, we still do not know the precise toxic effects that are caused by  $A\beta$  and mitochondrial abnormalities in neurons, particularly at synapses, that are involved in cognitive decline. Further research is needed to develop drugs capable of crossing the bloodbrain barrier and targeting mitochondria, and to develop the agents to boost mitochondrial function and decrease  $A\beta$  toxicity and improve synaptic branching and cognitive functions in elderly people and patients with AD.

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