

Interactions between β -amyloid and central cholinergic neurons: implications for Alzheimer's disease

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Alzheimer's disease is an age-related neurodegenerative disorder that is characterized by a progressive loss of memory and deterioration of higher cognitive functions. The brain of an individual with Alzheimer's disease exhibits extracellular plaques of aggregated β -amyloid protein (A β), intracellular neurofibrillary tangles that contain hyperphosphorylated tau protein and a profound loss of basal forebrain cholinergic neurons that innervate the hippocampus and the neocortex. A β accumulation may trigger or contribute to the process of neurodegeneration. However, the mechanisms whereby A β induces basal forebrain cholinergic cell loss and cognitive impairment remain obscure. Physiologically relevant concentrations of A β -related peptides have acute, negative effects on multiple aspects of acetylcholine (ACh) synthesis and release, without inducing toxicity. These data suggest a neuromodulatory influence of the peptides on central cholinergic functions. Long-term exposure to micromolar A β induces cholinergic cell toxicity, possibly via hyperphosphorylation of tau protein. Conversely, activation of selected cholinergic receptors has been shown to alter the processing of the amyloid precursor protein as well as phosphorylation of tau protein. A direct interaction between A β and nicotinic ACh receptors has also been demonstrated. This review addresses the role of A β -related peptides in regulating the function and survival of central cholinergic neurons and the relevance of these effects to cholinergic deficits in Alzheimer's disease. Understanding the functional interrelations between A β peptides, cholinergic neurons and tau phosphorylation will unravel the biologic events that precede neurodegeneration and may lead to the development of more effective pharmacotherapies for Alzheimer's disease.

La maladie d'Alzheimer est une maladie neurodégénérative reliée à l'âge caractérisée par une perte progressive de la mémoire et la détérioration des fonctions cognitives supérieures. Le cerveau d'une personne atteinte de la maladie d'Alzheimer contient des plaques extracellulaires de protéine β -amyloïde (A β) agrégée et des enchevêtrements neurofibrillaires intracellulaires qui contiennent de la protéine tau hyperphosphorylée, et présente une perte profonde de neurones cholinergiques du cerveau antérieur basal qui innervent l'hippocampe et le néocortex. L'accumulation de protéine A β peut déclencher le processus de neurodégénération ou y contribuer. Les mécanismes par lesquels la protéine A β provoque la perte de cellules cholinergiques du cerveau antérieur basal et la déficience de la cognition demeurent

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toutefois obscurs. Des concentrations physiologiquement pertinentes de peptides reliés à la protéine A β ont des effets négatifs aigus sur de multiples aspects de la synthèse et de la libération de l'acétylcholine (ACh) sans provoquer de toxicité. Ces données indiquent que les peptides ont peut-être une influence neuromodulatrice sur les fonctions cholinergiques centrales. L'exposition chronique à des protéines A β micromolaires provoque la toxicité des cellules cholinergiques, peut-être par l'hyperphosphorylation de la protéine tau. On a par ailleurs démontré que l'activation de certains récepteurs cholinergiques altère la transformation de la protéine précurseur amyloïde, ainsi que la phosphorylation de la protéine tau. On a aussi démontré l'existence d'un lien direct entre la protéine A β et les récepteurs de l'ACh nicotiques. Cette étude porte sur le rôle des peptides reliés à la protéine A β dans la régulation de la fonction et de la survie des neurones cholinergiques centraux et la pertinence de ces effets pour les déficits cholinergiques dans les cas de maladie d'Alzheimer. La compréhension des liens fonctionnels entre les peptides A β , les neurones cholinergiques et la phosphorylation de la protéine tau précisera les éléments biologiques qui précèdent la neurodégénération et pourra peut-être déboucher sur la mise au point de pharmacothérapies plus efficaces contre la maladie d'Alzheimer.

Introduction

First described in 1906 by Alois Alzheimer,¹ Alzheimer's disease (AD) is the most common cause of dementia today, accounting for about 50%–60% of all age-related dementia that affects individuals over 65 years of age. It is characterized clinically by progressive memory loss that begins early in the disease process. Other cognitive (disorientation, confusion and problems with reasoning) and behavioural (agitation, anxiety, delusions, depression and insomnia) disturbances appear as the disease progresses and impair function in activities of daily living.² The average course of AD is a decade, but the rate of progression is variable. Epidemiological data have shown that AD afflicts about 8%–10% of the population over 65 years of age, and its prevalence doubles every 5 years thereafter.³ With the proportion of elderly people in the population increasing steadily, AD is expected to pose an increasing economic challenge to Western economies and be a burden to health care delivery systems over the coming decades.^{2,3}

Both genetic and environmental factors can contribute to the development of AD. A minority of cases have an obvious genetic origin and demonstrate an autosomal dominant pattern of inheritance. Linkage studies indicate that point mutations in the gene for the amyloid precursor protein (APP), on chromosome 21, are associated with a subset of early onset (< 65 yr) familial AD cases. However, most early onset cases have been linked to alterations in 2 other genes: presenilin 1 (PS1) on chromosome 14 and presenilin 2 (PS2) on chromosome 1.^{4–6} Although these findings are of immense importance in elucidating the biologic pathogenesis of AD, it is important to recognize that mutations in these 3 genes may only account for 30%–50%

of all autosomal dominant early onset cases. Familial clustering is also found in individuals with late-onset AD. Various factors including concomitant pathology and limited sample sizes make it difficult to identify genetic causes of late-onset disease by conventional linkage analysis. However, association studies have identified candidate genes that significantly increase the risk for late-onset disease. The ϵ 4 allele of the apolipoprotein E gene, on chromosome 19, is one such risk factor. Possessing a single copy of the allele may increase the chance of developing AD 2–5 times, whereas having two ϵ 4 alleles raises this probability more than 5 times. Conversely, expression of the ϵ 2 allele appears to protect against development of AD.^{6–8} A polymorphism in an intronic region of the α -2 macroglobulin was found to segregate with the AD phenotype in some subjects with late-onset disease.⁹ However, follow-up study found no association in 2 independent familial AD data sets, both of which had shown earlier evidence for linkage with a locus on chromosome 12. Thus, it is now thought that the locus of interest is situated elsewhere in the chromosome.^{10,11} Given that the vast majority of cases of AD have not been associated with any of the genes implicated to date, it is highly likely that additional causative mutations and genetic risk factors remain to be identified.^{6,11} Other factors that may play an important role in the pathogenesis of AD include age, head injury and oxidative stress.¹²

Neuropathologic features of AD

Neurofibrillary tangles

Neuropathologic hallmarks of both familial and sporadic AD include intracellular neurofibrillary tangles,

extracellular parenchymal and cerebrovascular amyloid deposits, and loss of neurons and synaptic integrity in specific brain areas. These features are also seen in the brains of individuals with Down's syndrome (age < 40 yr) and, to a limited extent, in the normal aging brain.^{11,13-15} Neurofibrillary tangles are composed of paired helical filaments (PHF) and occasional single straight filaments, mainly containing an abnormal hyperphosphorylated form of the microtubule-associated protein, tau. In healthy neurons, tau binds and stabilizes microtubules, which make up the cytoskeleton of the cell, by reversible enzymatically mediated phosphorylation and dephosphorylation processes. If the phosphorylated tau is not dephosphorylated, it is unable to bind other microtubules. This results in polymerization of the phosphorylated tau into straight filaments, which are then cross-linked by glycosylation to form PHF-tau.^{13,15-17} Intraneuronal PHF-tau aggregates are often found in conjugation with ubiquitin.¹⁶ Neurofibrillary tangles in the brain of an individual with AD are particularly abundant in the entorhinal cortex, hippocampus, amygdala, association cortices of the frontal, temporal and parietal lobes, and certain subcortical nuclei that project to these regions. Formation of PHF-tau reduces the ability of tau to stabilize microtubules, leading to disruption of neuronal transport and eventually to the death of affected neurons.¹⁶⁻¹⁸ The extent of neurofibrillary pathology, and particularly the number of cortical neurofibrillary tangles, correlates positively with the severity of dementia. However, tangles are also found in a variety of other neurodegenerative diseases that exhibit neither amyloid deposits nor neuritic plaques.^{11,15,16,19,20}

Neuritic plaques

Neuritic plaques are multicellular lesions that contain a compact deposit of amyloid peptides surrounded by dystrophic neurites, activated microglia and reactive astrocytes. The major amyloid peptides that are found in the plaques are β -amyloid₁₋₄₂ ($A\beta_{1-42}$) and $A\beta_{1-40}$, peptides that are generated by proteolytic cleavage of APP. In the brain with AD, $A\beta_{1-42}$ is deposited first and is the predominant form in senile plaques, whereas $A\beta_{1-40}$ is deposited later in the disease process. Other proteins, such as components of the complement cascade, apolipoprotein E, α -1-antichymotrypsin, lysosomal proteases and antioxidant enzymes, constitute minor components of the neuritic plaques. These plaques are

most prominent in areas affected by neurodegeneration, such as the entorhinal cortex, hippocampus and association cortices.^{11,21-23} Neuritic plaque number does not itself correlate with the severity of dementia, although a clinical correlation between elevated levels of $A\beta$ peptide in the brain and cognitive decline has been reported.²⁴ Recent investigations in animal models and human brain samples have placed a special emphasis on soluble $A\beta$.^{25,26}

Several lines of evidence suggest that accumulation of $A\beta$ peptide in the brain may, over time, initiate and/or contribute to the pathogenesis of AD. These include the association of some cases of AD with inherited APP mutations,^{11,13,15} the elevation of $A\beta$ peptides and the appearance of amyloid plaques in advance of other pathology in the brains of individuals with AD or Down's syndrome,^{27,28} the increased production of $A\beta_{1-42}$ in vivo and in vitro by pathogenic mutations in *PS1* and *PS2*,¹¹ and the in-vitro neurotoxic potential of fibrillar $A\beta$ peptides.^{11,29,30} Overproduction or reduced clearance, or both, of $A\beta$ peptides are likely key to amyloid aggregation, which in turn contributes to the development of neurofibrillary tangles and subsequent neuronal degeneration.^{11,31-33} Recent studies of APP transgenic mice³⁴⁻³⁷ and of intrathecally administered $A\beta$ in nontransgenic adult animals³⁸⁻⁴¹ reinforce the notion that overexpression of $A\beta$ peptide, or injection of aggregated $A\beta$, induces subcellular alterations or neuronal loss in selected brain regions. It has been suggested that overexpression or injection of $A\beta$ peptide may potentiate the formation of neurofibrillary tangles in tau transgenic mice,^{42,43} a relation first inferred from consideration of kindreds with familial AD. Although these results suggest a role for $A\beta$ peptides in the neurodegenerative process, both the role of $A\beta$ in the normal brain and the mechanisms by which it causes neuronal loss and tau abnormalities in AD remain poorly understood.

Loss of basal forebrain cholinergic neurons

Degenerating neurons and synapses in the brain of individuals with AD are located predominantly within regions that project to or from areas that display high densities of plaques and tangles. Severely affected regions include the hippocampus, entorhinal cortex, amygdala, neocortex, and some subcortical areas such as basal forebrain cholinergic neurons, serotonergic neurons of the dorsal raphe and noradrenergic neurons

of the locus coeruleus.⁴⁴⁻⁴⁷ Biochemical investigations of tissues from biopsy and autopsy indicate that various neurotransmitters and modulators including acetylcholine (ACh), serotonin, noradrenaline and somatostatin are differentially altered in the brains of individuals with AD.^{14,45,48} The early and most consistently reproduced finding is a profound reduction in the activity of the ACh-synthesizing enzyme, choline acetyltransferase (ChAT), in the neocortex, which correlates positively with the severity of dementia.^{45,47,49} Reduced choline uptake, ACh release and loss of cholinergic neurons from the basal forebrain region further indicate a selective presynaptic cholinergic deficit in the hippocampus and neocortex of brains of individuals with AD.^{48,50} Cholinergic neurons in the brain stem and striatum are either spared or affected only in late stages of the disease.^{45,47,48} Together with pharmacologic evidence of cholinergic involvement in the affected cognitive processes, these findings led to the development of a "cholinergic hypothesis" of AD. This hypothesis posits the degeneration of the cholinergic neurons in the basal forebrain and the loss of cholinergic transmission in the cerebral cortex and other areas as the principal cause of cognitive dysfunction in patients with AD.^{47,48,50-52} The hypothesis is supported by evidence that drugs that potentiate central cholinergic function (such as donepezil, rivastigmine and galantamine) have some value in symptomatic treatment during early stages of the disease.^{47,53}

The loss of basal forebrain cholinergic neurons has prompted extensive study of ACh receptors in the brains of individuals with AD.^{45,47,48,50,54} ACh exerts effects on the central nervous system by interacting with G-protein-coupled muscarinic and ligand-gated cation channel nicotinic receptors. Five distinct muscarinic receptor subtypes, m_1 - m_5 , have been cloned and shown to correspond to 5 pharmacologically defined M1-M5 muscarinic receptors. It is generally believed that M2 receptors, most of which are located on presynaptic cholinergic terminals, are reduced in the brains of individuals with AD.^{47,54} The density of postsynaptic M1 receptors remains unaltered, but there is some evidence for disruption of the coupling between the receptors, their G-proteins and second messengers.^{54,55} The profiles of M3 and M4 receptors in the brains of individuals with AD remain equivocal.^{56,57}

For the nicotinic receptor family, 11 genes that encode 8 alpha (α_2 - α_9) and 3 beta (β_2 - β_4) receptor subunits have been identified.^{47,58,59} High-affinity central nervous

system binding sites of the agonist nicotine are mostly composed of $\alpha_4\beta_2$ subunits, whereas homomers of the α_7 receptor subunit contribute to the high-affinity binding of the antagonist α -bungarotoxin (α -BgTx).^{59,60} Epi-batidine, a potent nicotine agonist, binds with high affinity to a subtype of nicotinic receptor containing the α_3 subunit.⁵⁹ Nicotinic receptors are predominantly located on cholinergic terminals. High-affinity nicotinic binding sites are markedly reduced in the hippocampus and cortex of postmortem brains of individuals with AD, and these observations have been confirmed by in-vivo positron emission tomography.^{48,61} There is also evidence of a significant decrease in α_7 protein expression and α -BgTx binding sites in the hippocampus of brains of individuals with AD.⁶² However, a recent immunocytochemical study demonstrated an increase in the proportion of astrocytes expressing α_7 immunoreactivity in the hippocampus and entorhinal cortex of the brain of individuals with AD relative to age-matched controls.⁶³ Notwithstanding these data, cholinomimetics only delay the cognitive decline in a subset of patients at early stages of the disease. Whether the early changes in the cholinergic system might play a pathogenic role in further dysregulating APP processing or promoting tau phosphorylation are important issues that remain to be addressed.

Cholinergic system and APP processing

APP processing

A β peptides, the principal component of amyloid deposits, are a group of hydrophobic peptides of 39-43 amino acid residues. These peptides are derived by proteolytic cleavage of APP, a single transmembrane glycoprotein with a long N-terminal extracellular region and a short C-terminal cytoplasmic tail.^{11,14,23,64} Nine isoforms are produced from a single APP gene by alternative mRNA splicing and encode proteins ranging from 365 to 770 amino acids. Two of these isoforms (APP₃₆₅ and APP₅₆₃) do not contain A β peptides. Some variants of APP (e.g., APP₇₇₀) contain a 56-amino acid insert that is homologous to the Kunitz family of serine protease inhibitors (KPI) along with a 19-residue segment homologous to the thymocyte OX2 antigen. Others contain only the KPI domain (e.g., APP₇₅₁) or neither segment (e.g., APP₆₉₅).^{11,22,23} APP expression occurs ubiquitously, and the primary isoform varies according to cell and tissue type. In the nervous system, APP₆₉₅ is

expressed predominantly in neurons, whereas APP₇₇₀ and APP₇₅₁ are found in neuronal as well as nonneuronal cells.^{11,22,23,65} Two additional genes encoding for the APP-like proteins (i.e., APLP1 and APLP2) have been identified. The N-terminal regions of these proteins are homologous with APP. However, they lack the A β C-terminal domain. Recent data from mice with combined ablations of APP and APLP genes revealed that these proteins serve both overlapping and distinct functions in vivo.⁶⁶

Mature APP is processed proteolytically by distinct α -secretase or β -secretase pathways (Fig. 1). The α -secretase activity cleaves the A β domain within Lys¹⁶ and Leu¹⁷ residues to prevent formation of full-length A β peptide. This pathway yields a soluble N-terminal APP α and a 10-kd C-terminal APP fragment that can be further processed by γ -secretase to generate A β ₁₇₋₄₀ or A β ₁₇₋₄₂, which are also known as the P3 peptides. The α -secretase cleavage occurs mostly at the cell surface, although it can be mediated to some extent during the secretory intracellular trafficking of APP.^{11,22} The β -secretase pathway, which results in the formation of intact A β peptide, is mediated by the sequential actions of β -secretase (β -APP cleaving enzyme [BACE]) and γ -secretase enzymes (Fig. 1). The β -secretase cleavage

generates a truncated soluble APP β and a membrane-bound A β -containing C-terminal fragment. Further proteolysis of the C-terminal fragment by γ -secretase yields the full-length A β _{1-40/42} peptide.^{11,22,23,67,68} γ -Secretase activity resides in a multimeric protein complex that contains PS, which is considered to be a putative aspartyl protease,⁶⁹ along with at least 3 components (nicastrin, PEN-2 and APH-1) that are required for substrate recognition, complex assembly and targeting the complex to its site of action.⁷⁰ Cell culture studies have suggested that most A β _{1-40/1-42} is generated in the endosomal recycling pathway. A minority of A β _{1-40/1-42} is produced in the secretory pathway, within the endoplasmic reticulum and Golgi apparatus.^{11,22,23} Under normal conditions, about 90% of secreted A β peptides are A β ₁₋₄₀, which is a soluble form of the peptide that only slowly converts to an insoluble β -sheet configuration and thus can be eliminated from the brain. In contrast, about 10% of secreted A β peptides are A β _{1-42/43/} species that are highly fibrillogenic and deposited early in individuals with AD and Down's syndrome.^{11,14,27,28} Relative activities of the discrete APP processing pathways can be influenced by a variety of factors, including the stimulation of receptors for ACh, serotonin, glutamate, estrogen, neuropeptides and growth factors.^{71,72} The influence of cholinergic stimulation on amyloid formation is of particular interest in view of the early targeting of the cholinergic basal forebrain in AD and the possibility that maintenance of this cholinergic tone might slow amyloid deposition in cholinergic terminal fields.

Cholinergic regulation of APP processing

A clear connection has been established between the cholinergic system and APP metabolism. Nitsch et al⁷³ first demonstrated cholinergic regulation of APP processing in human embryonic kidney (HEK) 293 cell lines that were stably transfected with human muscarinic m₁, m₂, m₃ and m₄ receptors. Carbachol, a nonselective muscarinic receptor agonist, significantly increased the release of soluble APP α in cells expressing m₁ and m₃, but not in cells expressing m₂ or m₄ receptor subtypes. This response was both sensitive to atropine and blocked by staurosporine, indicating the mediation of intracellular protein kinases in receptor-controlled APP α secretion.⁷³ Activation of muscarinic m₁-receptor-transfected cells not only enhanced soluble APP α secretion but also reduced the secretion of A β . Similarly,

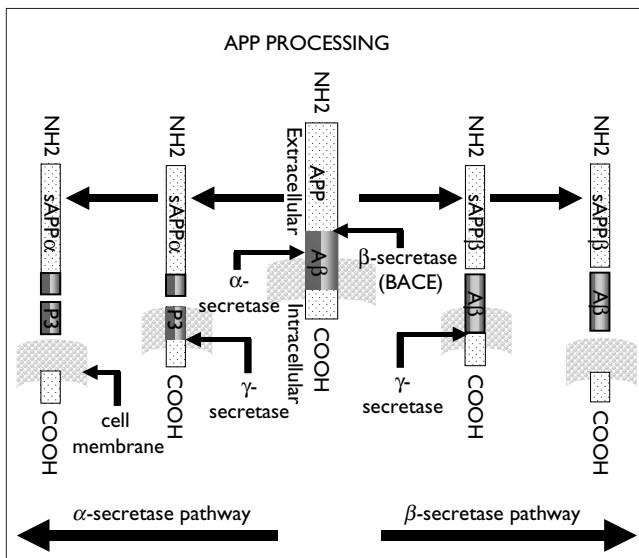


Fig. 1: Amyloid precursor protein (APP)-processing pathways. APP is a type I transmembrane protein that is processed by 2 distinct pathways. α -Secretase and β -secretase generate soluble APP α (sAPP α) and sAPP β , respectively. The remaining C-terminal fragment in the membrane is cleaved by γ -secretase to yield P3 following α -secretase cleavage, or β -amyloid (A β) subsequent to β -secretase cleavage.

muscarinic m_1 - and m_3 -receptor agonists stimulated soluble APP α release from rat cortical slices.⁷⁴ Both m_1 and m_3 receptors activate signalling cascades involving phosphatidylinositol hydrolysis and protein kinase C (PKC). Treating cells with phorbol esters mimicked the effect of agonist administration on soluble APP α secretion, and this effect was blocked by PKC inhibitors.^{71,75} Moreover, other G-protein-coupled receptors that activate PKC-dependent signalling pathways, including the vasopressin, bradykinin, serotonin and metabotropic glutamate receptors, share this capacity to stimulate soluble APP secretion and inhibit A β formation.^{71,75} The mechanism whereby PKC activity increases soluble APP α secretion is still unknown, but it may involve additional kinase steps and the eventual activation of the proteases that mediate APP cleavage.^{71,72,75}

In contrast to the muscarinic influence on APP processing, few studies have examined the contribution of nicotinic mechanisms. Treatment of PC12 cells with nicotine increases the release of soluble APP α in a concentration-dependent ($> 50 \mu\text{mol/L}$) and time-dependent ($> 2 \text{ h}$) manner, without affecting the expression of APP mRNA or A β secretion.⁷⁶ The relative increase in soluble APP α was attenuated by the α_7 -nicotinic receptor antagonist methyllycaconitine and also by EGTA, a Ca^{2+} chelator. The nicotine antagonist chlorisondamine blocked in-vivo elevation of total soluble APP induced by exposure to a high dose (8 mg/kg per day) of nicotine.⁷⁷ A nicotine-induced increase in Ca^{2+} influx was found to correspond with the increase in soluble APP secretion, suggesting that Ca^{2+} influx through nicotinic receptors may be involved in enhanced secretion. This result is in agreement with the findings from several studies that show that increased cytoplasmic Ca^{2+} levels can stimulate soluble APP secretion.^{72,75,78}

The effects of acetylcholinesterase (AChE) inhibitors on the level of soluble APP α differ between cell types and depend upon the specific drug, duration of treatment and the dose tested. For example, metrifonate did not alter soluble APP or A β levels in human SK-N-SH neuroblastoma cells,⁷⁹ whereas short-term treatment with the inhibitor was able to increase the secretion of soluble APP α in SH-SY5Y neuroblastoma cells, presumably by increasing the availability of ACh and thereby stimulating muscarinic receptors.^{80,81} Physostigmine has been shown to elevate soluble APP α secretion in rat cortical slices⁸² but decreased soluble APP

secretion without altering A β levels in SK-N-SH neuroblastoma cells.⁷⁹ Tacrine, a potent cholinesterase inhibitor, was found to attenuate secretion of soluble APP α in glial, fibroblast, PC12 and neuroblastoma cells. Other AChE inhibitors such as phenserine, cymserine and tolserine decreased soluble APP α levels, whereas 3,4-diaminopyridine failed to affect soluble APP α levels in SK-N-SH neuroblastoma cells.⁷⁹ The differential effects of the AChE inhibitors on APP processing appear to be unrelated to their selectivity for the cholinesterase enzymes but may depend upon other mechanisms, such as their influence on APP synthesis, expression, turnover or trafficking, or the regulation of APP-processing enzymes.^{75,79,83}

Modulation of cholinergic functions by A β peptides

Short-term effects of A β on cholinergic neurons

Concentrations of A β peptides in the picomolar–nanomolar range can negatively regulate various steps of ACh synthesis and release, without apparent neurotoxicity (Table 1,^{84–99} Fig. 2). The high potency and reversible nature of this effect, together with the fact that picomolar–nanomolar concentrations of A β peptides are found constitutively in normal brain cells, suggest that A β -related peptides may act as a modulator of cholinergic function under normal conditions.^{50,75,92,100–102} A 1-hour exposure to picomolar–nanomolar concentrations of A β can inhibit K^+ -evoked or veratridine-evoked endogenous ACh release from rat hippocampal and cortical slices. This effect is insensitive to tetrodotoxin, suggesting that the A β peptide may act at the level of cholinergic terminals.^{84,86} Structure-activity studies reveal that several A β fragments, including A β_{1-42} , A β_{1-40} , A β_{1-28} and A β_{25-35} , similarly inhibit ACh release from rat hippocampal slices, indicating that the activity resides within the sequence A β_{25-28} (GSNK, the C-terminal domain of the nontoxic A β_{1-28} fragment). Striatal ACh release is relatively insensitive to A β peptides.⁸⁶ This regional selectivity indicates that transmitter phenotype expression does not fully explain the susceptibility of specific cell populations to effects of A β . Factors such as the distance over which cholinergic axons project to their terminal fields and regional variation in the expression of A β binding sites may contribute to the differences in cellular responsiveness to A β . However, the sensitivity to A β of

cholinergic neurons in the cortex, hippocampus and striatum matches the pattern of regional vulnerability in AD.

The inhibitory effects of Aβ on ACh release have been confirmed in rat and guinea pig cortical synaptosomes,⁸⁷ rat retinal neurons⁸⁸ and in cholinergic synaptosomes from the electric organ of the electric ray *Narke japonica*.⁸⁹ These effects may be affected by aging. Higher levels of Aβ₁₋₄₀ were observed in the aged rat hippocampus than were found in young adult rats, and the cholinergic neurons of aged, cognitively impaired rats may be more sensitive to Aβ-mediated inhibition of hippocampal ACh release than either cognitively unimpaired, aged rats or young adult rats.⁹⁰ Lee et al⁹¹ reported that inhibition of ACh release by Aβ₂₅₋₃₅ could be reversed by certain ginseng saponins at concentrations that did not by themselves alter ACh release. This effect was insensitive to tetrodotoxin, suggesting a direct interaction of ginseng at the level of the cholinergic synapse.

The cellular mechanisms by which Aβ-related peptides acutely attenuate ACh release from selected brain regions remain unclear. Steps that are critical for ACh synthesis and release, ranging from precursor recruitment to vesicular fusion, could be impaired by Aβ peptides (Table 1, Fig. 2). Turnover of ACh in the cholinergic terminals is regulated so that increased transmitter release is associated with increased synthesis. When brain slices are exposed to submaximal concentrations of depolarizing agents such as K⁺ or veratridine,

ongoing synthesis of ACh keeps pace with transmitter release from the terminals.¹⁰³ ACh synthesis under these conditions depends on the high-affinity uptake of choline from extracellular sources to intracellular acetyl

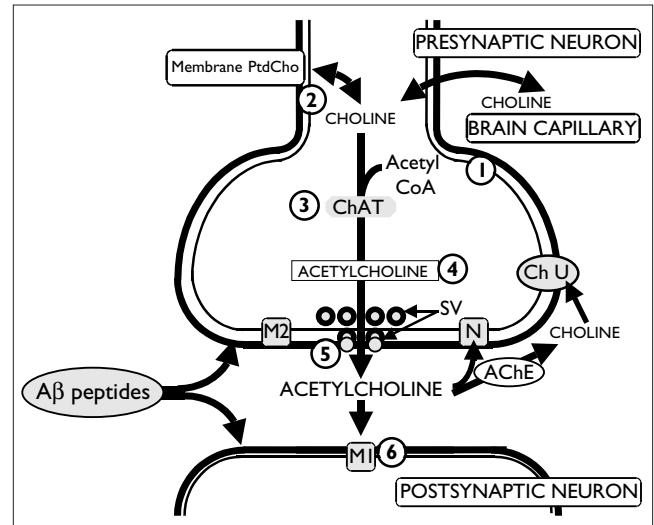


Fig. 2: Targets of Aβ that modulate cholinergic transmission: (1) Aβ reduces activity of pyruvate dehydrogenase, an enzyme that generates acetyl coenzyme A (CoA) from pyruvate; (2) Aβ reduces high-affinity uptake of choline; (3) long-term or high-dose exposure to Aβ reduces activity of the choline acetyltransferase (ChAT) enzyme; (4) Aβ reduces acetylcholine (ACh) content; (5) Aβ reduces ACh release from synaptic vesicles (SV); (6) Aβ impairs muscarinic M1-like signalling. AChE = acetylcholinesterase, Ch U = site of choline uptake, M2 = presynaptic muscarinic M2 receptor, N = presynaptic nicotinic receptor, PtdCho = phosphatidylcholine.

Table 1: In-vitro effects of Aβ-related peptides on cholinergic neurotransmission and toxicity

Effect	Peptide fragment	Concentration	Model	References
Short-term				
Decrease in choline uptake	Aβ _{1-42, 1-40, 1-28, 25-35}	pmol/L–μmol/L	Cortical and hippocampal synaptosomes	84,85
Decrease in ACh release	Aβ _{1-42, 1-40, 1-28, 25-35}	pmol/L–μmol/L	Cortical and hippocampal slices, cortical and electric organ synaptosomes, retinal neurons	84,86–91
Decrease in whole-cell currents and increase in excitability	Aβ _{1-42, 25-35}	nmol/L–μmol/L	Dissociated cells from the diagonal band of Broca	92
Long-term				
Decrease in PDH activity	Aβ ₁₋₄₂	nmol/L	Primary septal cultures	93
Decrease in ChAT activity	Aβ _{1-42, 1-40, 1-28, 25-35}	nmol/L–μmol/L	SN56 cell line and primary septal cultures	94,95
Decrease in ACh content	Aβ _{1-42, 1-28, 25-35, 25-28}	pmol/L–nmol/L	SN56 cell line and primary septal cultures	93,94,96
Disruption of muscarinic M1-like receptor signalling	Aβ _{1-40, 25-35}	nmol/L–μmol/L	Primary cortical cultures	97
Induction of tau phosphorylation	Aβ _{1-42, 1-40, 25-35}	μmol/L	SN56 cell line and primary septal cultures	95,98
Induction of toxicity	Aβ _{1-42, 1-40, 25-35}	μmol/L	SN56 cell line, RN46A cell line and primary septal cultures	95,98,99

Note: Aβ = β-amyloid peptide; ACh = acetylcholine; ChAT = choline acetyltransferase; PDH = pyruvate dehydrogenase.

CoA and ChAT. The availability of choline is a rate-limiting determinant of ACh biosynthesis, whereas ChAT activity is not.¹⁰³ Under short-term treatment conditions, picomolar–nanomolar concentrations of A $\beta_{1-40/1-42}$ do not affect ChAT activity in tissue homogenates or in slice preparations from the hippocampus, cortex or striatum.⁸⁴ Similarly, Zambrzycka et al¹⁰⁴ reported that soluble A β_{25-35} did not acutely affect ChAT activity in the adult or aged rat brain. Nanomolar A β_{1-42} (but not A β_{1-40}) can acutely regulate the phosphorylation of the ChAT enzyme in IMR32 neuroblastoma cells expressing human ChAT.¹⁰⁵ The significance of this phosphorylation regarding regulation of cholinergic transmission remains unclear.

Temperature-dependent high-affinity [³H]choline uptake is decreased by 20 minutes' preincubation with A β . This effect is particularly marked in tissues from the hippocampus and cortex, mirroring the effect of A β on ACh release in these regions.⁸⁴ Acute incubation of hippocampal synaptosomes with low nanomolar A β_{1-40} suppresses depolarization-induced high-affinity choline uptake as well as [³H]hemicholinium-3 ([³H]HC-3) binding.⁸⁵ Detailed analysis of these data indicate that changes in the transport are predominantly the result of an alteration of V_{max} , whereas the changes in specific binding probably involve alterations of both B_{max} and K_D . Micromolar concentrations of A β_{1-40} decrease high-affinity choline uptake and the [³H]HC-3 binding under basal conditions in a time-dependent manner.⁸⁵ These results indicate that A β can affect acute ACh release, at least in part, by regulating high-affinity choline uptake. The potential involvement of A β in the intracellular transport of ACh and the fusion of ACh vesicles with the presynaptic membrane remains to be investigated.

A variety of receptors (e.g., receptors for advanced glycation end products [RAGE], the class A scavenger receptor [SR], the 75-kd neurotrophin receptor [p75^{NTR}] and serpin-enzyme complex receptors) interact with A β in vitro.¹⁰⁶⁻¹⁰⁹ These interactions have attracted attention both for the insights they may provide into linking amyloid accumulation to neurodegeneration and as potential targets for drug design. Recent evidence suggests that A β_{1-42} can also bind with high affinity to the α_7/α -BgTx nicotinic receptor and with lower affinity to $\alpha_4\beta_2$ /cytisine nicotinic receptors (but not muscarinic) in the rat and guinea pig hippocampus and cerebral cortex.¹¹⁰ This is supported by whole-cell patch-clamp studies that showed that nanomolar

A $\beta_{1-40}/A\beta_{1-42}$ can specifically and reversibly block the α_7 nicotinic receptor current in cultured primary hippocampal neurons. The impairment of this current was noncompetitive, independent of voltage and mediated through the extracellular N-terminal length of the α_7 subunit.¹¹¹ In hippocampal slice preparations from rats that were 13–18 days old, Pettit et al¹¹² reported that A β_{1-42} can reversibly inhibit carbachol-induced α_7 and non- α_7 nicotinic receptor currents.¹¹² The broader range of effects in the slice study could relate to the concentration of A β peptide or to differences in cellular connectivity in the slice paradigm. The effects of A β on the nicotinic currents are consistent with receptor involvement in A β -mediated inhibition of ACh release. In support of this notion, the inhibitory effects of A β_{1-40} on cortical ACh release could be restored by the addition of a α_7 agonist, such as nicotine and epibatidine, but not by the α_4/β_2 nicotinic receptor agonist cytosine.¹¹³ Further studies are needed to define the precise role of the α_7 nicotinic receptor in regulating the inhibitory effects of A β peptides on endogenous ACh release.

In addition to affecting hippocampal and cortical terminals, A β peptide can act at the level of the cell body to increase neuronal excitability.⁹² Application of A $\beta_{1-42/25-35}$, 1 $\mu\text{mol/L}$, to acutely dissociated rat neurons from the diagonal band of Broca decreased whole-cell voltage-sensitive currents in cholinergic neurons that were identified by single-cell reverse-transcriptase polymerase chain reaction.⁹² This reduction was associated with changes in several K⁺ currents, including the Ca²⁺-activated K⁺ currents (Bk or Ic), the delayed rectifier current (Ik) and transient outward current (Ia), but not in calcium or sodium currents. The responses were blocked by tyrosine kinase inhibitors, suggesting that A β induces phosphorylation-dependent cascades to alter these currents.⁹² A β effects on whole-cell currents can be replicated by human amylin (a 37-amino-acid pancreatic peptide that is deposited in the islet cells of patients with type 2 diabetes mellitus) and are not additive with those elicited by amylin and can be blocked by AC187, which is a specific amylin-receptor antagonist. These data raise the intriguing possibility that the amylin receptor may mediate certain effects of A β on basal forebrain cholinergic neurons.¹¹⁴

Long-term effects of A β peptide on cholinergic neurons

A 2-day exposure to picomolar–nanomolar concentrations of A β_{1-42} , A β_{1-28} , A β_{25-35} and, to a lesser extent,

A β_{25-28} decreased intracellular ACh concentration in the cholinergic hybrid SN56 cell line (mouse septal neurons \times neuroblastoma), without causing toxicity (Table 1, Fig. 2). The decrease in intracellular ACh could be attributed to reduced biosynthesis, because it was accompanied by a reduction in activity of ChAT but not of AChE. The decrease could be prevented by a co-treatment with trans-retinoic acid, a compound that increases ChAT mRNA expression in SN56 cells, or by co-administration of tyrosine kinase inhibitors.^{50,94,96} Inhibition of DNA synthesis or treatment with antioxidants did not alter the long-term effects of A β on ACh concentrations. Thus, gene transcription and free radical production are likely not involved in mediating the effect of A β on this cholinergic SN56 cell line.⁹⁶ In keeping with these results, treatment of rat septal neurons with nanomolar concentrations of A β_{1-42} decreased ACh production and reduced activity of the acetyl-CoA biosynthesizing enzyme pyruvate dehydrogenase (PDH), without affecting ChAT activity or neuronal survival. The decreased PDH activity likely results from A β activation of the tau protein kinase I/glycogen synthase kinase-3 β (GSK-3 β), which can phosphorylate and inactivate PDH.⁹³ This study suggests that long-term exposure to A β peptide may impair ACh processing by reducing the availability of acetyl CoA.

In addition to regulating ACh synthesis, solubilized A β peptide can disrupt transduction of the muscarinic M1-like receptor signal.⁹⁷ A 4-hour exposure to nanomolar–micromolar A β_{1-40} reduced carbachol-induced GTPase activity in rat cortical cultured neurons without affecting muscarinic receptor ligand binding parameters. At higher concentrations, similar treatment with A β attenuated muscarinic M1 receptor signalling by decreasing intracellular Ca²⁺ and the accumulation of inositol phosphates Ins(1)P, Ins(1,4)P₂, Ins(1,4,5)P₃ and Ins(1,3,4,5)P₄.⁹⁷ Exposure of rat cortical cultured neurons to nanomolar A β_{1-42} /A β_{25-35} inhibits the carbachol-induced, but not glutamate-induced, increase in intracellular Ca²⁺ and Ins(1,4,5)P₃, indicating that selective disruption of muscarinic signalling is another means by which A β can affect the function of cholinergic neurons.¹¹⁵

A β -mediated toxicity, tau phosphorylation and cholinergic neurons

Long-term exposure to A β peptides can induce toxicity in a variety of cell lines, as well as in primary rat and

human cultured neurons. The toxicity of the peptide, unlike its neuromodulatory effects, is related to its ability to form insoluble aggregates.^{29,30} However, recent evidence suggests that the most detrimental forms of A β peptides are the soluble oligomers and that the insoluble amorphous or fibrillar deposits represent a less harmful inactivated form of the peptide.¹¹⁶ Some neuronal phenotypes, such as γ -aminobutyric acid (GABA)-ergic and serotonergic neurons, appear resistant to A β toxicity, and various cell lines differ in their degree of sensitivity.^{99,117} Differentiated SN56 cholinergic cell lines are a susceptible line for toxicity studies, and when exposed to A β_{1-40} these cells exhibit retraction of neurites, cell shrinkage and death.⁹⁸ When treated with ciliary neurotrophic factor, the RN46A cell line develops a cholinergic phenotype and is highly sensitive to A β peptides. In contrast, stimulation of RN46A differentiation with brain-derived neurotrophic factor yields an A β -insensitive cell population with a serotonergic transmitter phenotype.⁹⁹ Prolonged exposure of rat primary septal culture neurons to micromolar A β peptides induces both cell death and a concomitant decrease in ChAT activity.⁹⁵ Collectively, these results suggest that cells that express cholinergic transmitter phenotype are vulnerable to the toxic effects of the A β peptide.

The mechanisms by which A β induces cholinergic cell death may involve alteration in intracellular calcium and/or the production of toxic and inflammatory mediators such as nitric oxide, cytokines and reactive oxygen intermediates.^{118–120} Studies of a variety of cell lines and primary cultured neurons suggest that A β toxicity might be mediated either by interaction with a hydroxysteroid dehydrogenase enzyme or by plasma membrane RAGE, SR, p75^{NTR} or α_7 nicotinic receptors.^{107–110} A role for the death domain of p75^{NTR} in A β -induced cell death was observed in neuroblastoma (SK-N-BE) cells expressing full-length or truncated forms of p75^{NTR}.¹²¹ Studies of transfected neuroblastoma (SK-N-MC) cells indicate that expression of the α_7 nicotinic receptor may also have a critical role in the degeneration by facilitating internalization and accumulation of A β_{1-42} into neurons.¹²² Given the marked expression of p75^{NTR} and of the α_7 nicotinic receptor in the cholinergic basal forebrain, their role in cholinergic cell death bears further investigation.

Prolonged exposure to micromolar A β_{1-40} can increase choline conductance from PC12 cells.¹²³ Should this also occur in cholinergic neurons, severe choline

depletion could result as a consequence of decreased choline uptake and its increased leakage from neurons. Under conditions of choline depletion, cholinergic neurons can use choline from membrane phosphatidylcholine to synthesize ACh. A β -induced alteration in intracellular choline levels might thereby lead to an autocannibalistic process in which membrane turnover is disrupted to sustain neurotransmission.¹²⁴

Tau phosphorylation can also contribute to the vulnerability of neurons by destabilizing microtubules and impairing axonal transport.^{98,125–127} Aggregated A β induces the phosphorylation of tau protein in SN56 cholinergic cell lines.⁹⁸ Studies of cultured rat septal neurons have indicated that aggregated A β increases levels of tau and especially those of phosphorylated tau.⁹⁵ Phosphorylated tau immunoreactivity could be detected primarily in the distal axons of untreated cells, whereas staining was evident in axons, soma and dendrites of neurons exposed to A β .⁹⁵ Hyperphosphorylated tau protein can lead to neuronal death via disruption of the cytoskeletal network,^{16–18} it is likely that the increase in tau phosphorylation plays some role in A β -induced cell death.

How A β might induce the phosphorylation of the tau protein is unclear. Reactive oxygen species and the lipid peroxidation product 4-hydroxynonenal may be involved in A β neurotoxicity and cross-linking of tau proteins.¹²⁸ However, A β might also affect tau phosphorylation by directly increasing relevant kinase activity or by decreasing phosphatase activity.^{98,125,129–131} Activation of GSK-3 β ^{127,130,132} and mitogen-activated protein (MAP) kinase¹²⁹ induces tau protein phosphorylation and cell death in a variety of cultured neuron paradigms, and prolonged exposure of rat septal cultured neurons to micromolar A β peptide has been shown to induce tau phosphorylation by activating MAP kinase and GSK-3 β .⁹⁵ Various kinases phosphorylate tau at discrete sites, and it is likely that the phosphorylation of tau protein in cholinergic neurons is regulated by multiple kinases, including MAP kinase and GSK-3 β . Thus, it is important to explore both the biochemical potential of additional tau kinases, such as cyclin-dependent kinase 5, PKC and calcium-calmodulin kinase, to phosphorylate tau^{17,18} and the particular cellular expression of these kinases by cholinergic neurons.

Tau phosphorylation can be regulated by cholinergic agonists, and control of tau hyperphosphorylation by muscarinic receptor activation may provide a side benefit of cholinomimetic therapeutics. Muscarinic

agonists, carbachol and AF 102B, attenuate tau phosphorylation in cultured PC12 cells stably transfected with muscarinic m₁ receptors.¹³³ On the other hand, activation of the nicotinic receptor by nicotine and epibatidine increased the levels of phosphorylated as well as nonphosphorylated tau in SH-SY5Y human neuroblastoma cells.¹³⁴ The mechanisms by which muscarinic m₁ or nicotinic receptor activation modify tau phosphorylation remain unclear. These activities probably involve alteration of protein kinase/protein phosphatase systems.⁷⁵

In-vivo effects of A β peptide on cholinergic neurons

Attempts have been made to measure the impact of intracerebroventricular or local administration of A β on the cholinergic system under in-vivo conditions. Several studies have reported that A β peptides can induce cholinergic hypofunction when administered to the brain.^{50,100} Injection of A β _{25–35/1–40} into the rat medial septum causes a reduction in ACh release from the hippocampus in the absence of toxicity.¹³⁵ Using a similar approach, Harkany et al³⁹ demonstrated that A β _{1–42} is toxic to cholinergic neurons, as indicated by the reduction in ChAT-immunoreactive cell bodies in the basal forebrain and fibres in the cerebral cortex. Other studies have shown that infusion of A β into the lateral ventricles of adult rats impairs performance on learning and memory tasks in a manner similar to the effect of cholinergic inhibition.^{38,40,100} Local injection of preaggregated A β _{1–42} into the nucleus basalis magnocellularis (NBM) produces congophilic deposits and a strong inflammatory response, characterized by activation of astrocytes and microglia and by induction of microglial p38MAP kinase activity.¹³⁶ These changes were accompanied by a decrease in the number of cholinergic neurons around the congophilic amyloid deposit and hypofunction of the cortical cholinergic system.¹³⁶ Clearly, the influence of these astrocytic and microglial responses must be considered in assessing the in-vivo effects of A β peptides on cholinergic function.

The cholinergic system in APP, PS1 and APP/PS1 transgenic mice

Over the past few years, the central cholinergic system has been examined extensively in a variety of mutant APP, PS1 or APP/PS1 transgenic mouse lines, all of which exhibit elevated A β levels.^{137–146}

In mice that express the hAPP_{V642I} London mutant transgene, a selective decrease was found in the size of medial septal cholinergic neurons, but not in NBM cholinergic neurons. At 17–22 months of age, this line exhibits both reorganization of AChE-positive fibres in the hippocampus and dystrophic AChE-positive fibres around amyloid plaques in the cortex.¹³⁸

Cerebral amyloidosis was found to cause a significant cholinergic fibre loss and severe disruption of the neocortical cholinergic fibre network in aged APP23 mice that expressed the hAPP_{KM670/671NL} Swedish mutant transgene.¹³⁷ Although the cholinergic neurons of the medial septum and vertical limb of the diagonal band of Broca were smaller in APP23 transgenic mice than in nontransgenic controls, the number and volume of ChAT-positive neurons in the NBM complex were not affected. Hippocampal cholinergic fibre density in APP23 mice has yet to be reported.¹³⁷

In another study, hAPP_{KM670/671NL} mutant mice demonstrated an upregulation in the density of cholinergic synapses in the frontal cortex, parietal cortex and the hippocampus, whereas PS1_{M146L} transgenic mice showed no changes in either the size or density of cholinergic synapses. When crossed to yield hAPP_{KM670/671NL}/PS1_{M146L} double transgenic mice, extensive amyloid plaques were found to be associated with decreased density and size of cholinergic synapses in the frontal cortex and hippocampus.¹³⁹ One recent study showed a selective increase in immunostaining for p75^{NTR} (a marker of basal forebrain cholinergic neurons) in the medial septum of 12-month-old hAPP_{KM670/671NL} or PS1_{M146L} single transgenic mice, but not in hAPP_{KM670/671NL}/PS1_{M146L} double transgenic mice. Staining of p75^{NTR}-immunoreactive fibres in the hippocampus was more robust in single transgenic mice, relative to nontransgenic controls, whereas double transgenic mice displayed less intense p75^{NTR} fibre staining.¹⁴⁰ Whether the increased immunostaining in singly transgenic mice indicates a trophic effect on the cholinergic neurons as a consequence of either hAPP_{KM670/671NL} or PS1_{M146L} gene overexpression remains to be investigated. However, a separate study revealed no differences between hAPP_{KM670/671NL} mice and nontransgenic controls in ChAT activity, AChE activity, vesicular ACh transporter binding or high-affinity choline uptake sites in the cortex, hippocampus, striatum or cerebellum, at multiple times up to 23 months of age.¹⁴¹

Densities of M1/[³H]pirenzepine, M2/[³H]AF-DX 384 or α_7 nicotinic/[¹²⁵I] α -BgTx receptor binding sites

in all brain regions of mutant PS1_{L286V} transgenic and wild-type PS1 transgenic mice are comparable with those found in nontransgenic controls.¹⁴² In hAPP_{KM670/671NL} mutant mice, a decrease in M1/[³H]pirenzepine and $\alpha_4\beta_2$ nicotinic/[³H]cytisine, but not M2/[³H]AF-DX 384, receptor binding was evident in the hippocampus and cortex compared with nontransgenic controls.¹⁴⁶ However, in other studies, elevated hippocampal α_7 nicotinic receptor levels have been reported in hAPP_{K670N/M671L} single and 2 lines (i.e., hAPP_{K670N/M671L}/PS1_{A246E} and APP_{KM670/671NL+V717F}/PS1_{M146L+L286V}) of double transgenic mice.^{143,145}

In sum, increased expression of A β peptides produces a range of effects on the cholinergic systems of mutant APP, PS1 or APP/PS1 transgenic mice. Establishing which of these effects are robustly related to the type of pathogenic mutation, the level of transgene expression or to the intensity of amyloid deposits remains a work in progress.

Significance of amyloid interactions with cholinergic neurons

A β -related peptides are produced constitutively by brain cells and are found in the picomolar–nanomolar range in the cerebrospinal fluid of healthy individuals.^{11,31–33} These concentrations of A β can have a neuromodulatory role in the regulation of normal cholinergic functions through their negative effects on ACh biosynthesis and release. ACh may in turn reciprocally regulate APP synthesis and processing. For example, lesions of the basal forebrain cholinergic neurons or transient inhibition of cortical ACh release could elevate local APP synthesis,^{71,147–149} whereas agonist-induced activation of muscarinic m₁ and m₃ receptor subtypes increases the secretion of soluble APP derivatives and reduces the production of amyloidogenic A β peptides.^{71–75,150} These results suggest a mechanism whereby normal cholinergic innervation participates in the nonamyloidogenic maturation of APP via the α -secretase pathway, whereas the amyloidogenic A β -related peptides depress the activity of cholinergic neurons. A shift in the balance between these activities may be a key factor in the early targeting of cholinergic neurons in AD. Insults that reduce cholinergic transmission, increase A β generation or reduce peptide clearance may enhance vulnerability of neurons to direct toxicity of A β peptide^{11,29,30} or to choline limitation.^{84,85,100,102,123,124} Because A β deposits precede any other

lesions in the brains of individuals with AD,^{27,28} it is likely that amyloid-induced tau phosphorylation plays a critical role in neuronal loss. This is supported by some in-vivo studies in which intrathecal administration, or transgene-delivered expression of A β peptides, was shown to induce a loss of neurons, or a change in presynaptic cholinergic markers, within selected brain regions.^{38-41,137-139} The selective interactions of A β with basal forebrain cholinergic neurons provide candidate mechanisms that may contribute, at least in part, to the vulnerability of these neurons and their projections in AD. It remains to be determined whether changes in cholinergic transmission alter APP processing pathways so as to further AD pathology. If so, appropriate cholinomimetic therapeutics might be expected both to provide symptomatic benefit and to abrogate AD pathogenesis. Although as yet unproven, this potential mechanism will drive further research into clarifying the precise interactions of amyloid with the elements of cholinergic neurotransmission.

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