

Neurotoxicity in Alzheimer's disease: is covalently crosslinked A β responsible?

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Abstract Alzheimer's disease is the most common form of dementia in the elderly, and is characterised by extracellular amyloid plaques composed of the β -amyloid peptide (A β). However, disease progression has been shown to correlate more closely with the level of soluble A β oligomers. Recent evidence suggests that these oligomers are covalently crosslinked, possibly due to the interaction of A β with redox-active metal ions. These findings offer new avenues for the treatment and prevention of disease, by modulating metal binding or preventing the formation of neurotoxic A β oligomers.

Keywords Alzheimer · Beta-amyloid · A β · Oligomer · Covalent crosslinks · Metal · Neurotoxicity

Introduction

Alzheimer's disease (AD) is the most common form of dementia in the elderly, comprising more than 50% of reported cases (Small 2000). The major pathological hallmark of AD is protein deposition, particularly the

extracellular accumulation of β -amyloid peptide (A β) in amyloid plaques. Recent research, however, has revealed the surprising finding that, although amyloid plaques are the most obvious pathological feature of AD and A β has been shown to be cytotoxic, plaque burden does not correlate with disease progression. Instead, the severity of the disease correlates with the level of soluble A β oligomers [see Fig. 1, (McLean et al. 1999)]. The stability of these oligomers, coupled with the demonstrated ability of A β to undergo copper-induced redox chemistry, has led to the proposal that they are chemically crosslinked. That is, a covalent bond has formed between monomeric A β units.

At present, AD has a global incidence of 22 cases per 1,000 people in adults over the age of 75. The disease is characterised by neuronal death, oxidative damage, the hyperphosphorylation and intracellular deposition of the microtubule-associated protein tau, and, as noted above, the extracellular deposition of A β (Small 2000).

A β is cleaved from the amyloid precursor protein (APP) by proteases known as secretases (Small 2000). Controversy exists as to the pathogenic process of AD and the role of A β therein, but even those researchers who suggest that A β over-production or reduced clearance is not the primary cause of late-onset AD allow the peptide some role in the pathogenic process (e.g. Maurer and Hoyer 2006; Lee et al. 2007; Webber et al. 2007). Establishing the aetiology of AD has been complicated by the fact that γ -secretase can cleave APP at a number of residues, creating a pool of A β species of 38–46 residues long. The major species, A β_{1-40} , is 40 residues long, but a 42 residue species, A β_{1-42} , is more hydrophobic, has a greater tendency to aggregate, shows increased neurotoxicity, is more redox active and will seed aggregation of other A β species in the brain (Small 2000). A β also displays N-terminal truncations, including the modification of Glu-3 into pyroglutamate, which has been

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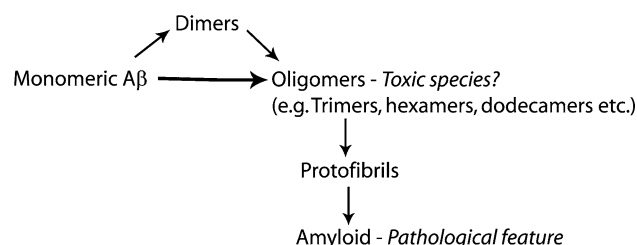


Fig. 1 Diagram of the proposed cascade leading from the formation of monomeric A β to amyloid deposition in AD. Although amyloid is the main pathological feature in AD, soluble oligomers are believed to be the main toxic species. Aggregation is believed to be transition metal dependant

shown to be a major component of amyloid plaques (Hartigaya 2000).

A β oligomers

Soluble A β oligomers are defined as those species that remain in the supernatant following centrifugation at $100,000 \times g$. These oligomers are resistant to degradation into monomers by compounds such as hexafluoroisopropanol (HFIP), urea, sodium dodecyl sulphate (SDS) and formic acid (Podlisny et al. 1995; Walsh et al. 2002; Lesne et al. 2006). A β oligomerisation is essential to the pathology of AD: A β is a normal cellular product, and is only considered harmful when aggregated (Walsh et al. 2005; Cohen et al. 2006). It can assume a number of different aggregation states, and much controversy exists over which of these states is “the” toxic agent in AD. Trimers, hexamers, dodecamers and a host of other soluble assembly states up to the protofibril level have all been implicated as the potential toxic agent (Lambert et al. 1998; Lesne et al. 2006; Townsend et al. 2006; Wu et al. 2006). However, the concept of a single toxic state has been challenged by the observation that different aggregation states induce neurotoxicity by distinct mechanisms or have different cellular effects (Deshpande et al. 2006). Additionally, monomeric A β is capable of adopting a wide variety of conformations depending on its environment, which can result in structural polymorphisms in higher aggregation states (Petkova 2005). This finding suggests that a specific isoform of, for example, trimer, not trimers per se, may be the most toxic A β species. Townsend et al. (2006) demonstrate this by showing that oral administration of *scyllo*-inositol neutralises the synaptotoxic effects of A β trimers without destabilising them or otherwise influencing their profile on Western blots.

The aggregation of the A β peptide appears to be driven by interaction between A β and metal ions, particularly copper and zinc, for which the peptide has a high binding affinity (Atwood et al. 2000). This observation accounts for the

tendency of amyloid plaques to be focused on synapses, where bioavailable metals are present at elevated concentrations ($[Cu^{2+}] \sim 15 \mu M$; $[Zn^{2+}] \sim 300 \mu M$). One form of metal coordination that can drive peptide aggregation is the formation of histidine bridges, where the imidazole side-chain of a His residue in a Cu-bound A β can also coordinate to the copper of a second A β peptide [see Fig. 2, (Smith et al. 2006)]. This forms a metal coordination site similar to the active site of SOD1, an enzyme involved in the neutralisation of superoxide radicals. Metal-induced aggregation is generally non-covalent and can be reversed by the addition of chelators (Cherny 1999). However, coordination of metals, particularly copper, by A β can also lead to the formation of covalent bonds (Atwood 2004; Barnham et al. 2004). Copper coordination induces redox chemistry, which leads to the oxidative modification of A β . One such modification is the formation of covalent dityrosine crosslinks. Another is the oxidation of the methionine at residue 35 of A β , which has been shown to profoundly affect the oligomerisation and toxicity of the peptide (Bitan 2003).

Proposed mechanisms of toxicity: dityrosine, membrane binding and radical formation

A detailed mechanism for the formation of dityrosine A β adducts is proposed in Barnham et al. (2004) and summarised in Fig. 3. In short, when A β binds copper (coordinated to histidine residues at positions 6, 13 and 14 and the tyrosine at position 10), it is capable of activating oxygen in the presence of a reducing substrate such as ascorbate, dopamine or thiols such as glutathione, and catalyses the reduction of oxygen to H_2O_2 . The final step of this reaction involves the transfer of a hydrogen atom of the side-chain hydroxyl group of Tyr-10 to the nascent H_2O_2 , forming a tyrosyl radical. While the A β /Cu catalyst can be regenerated by further reaction with the original reducing substrate, two tyrosyl radicals can also react to form dityrosine. Thus, the two peptides become covalently linked. Further support for the toxicity of these dityrosine-based oligomers has been shown by investigating a mutant peptide of A β where

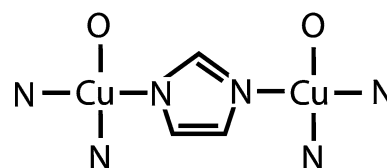


Fig. 2 Illustration of Cu^{2+} coordination by A β , where the metal is believed to be coordinated by His-6, His-13 and His-14 and Tyr-10. The imidazole side chain of a His residue bridges between two Cu^{2+} atoms to form dimeric A β

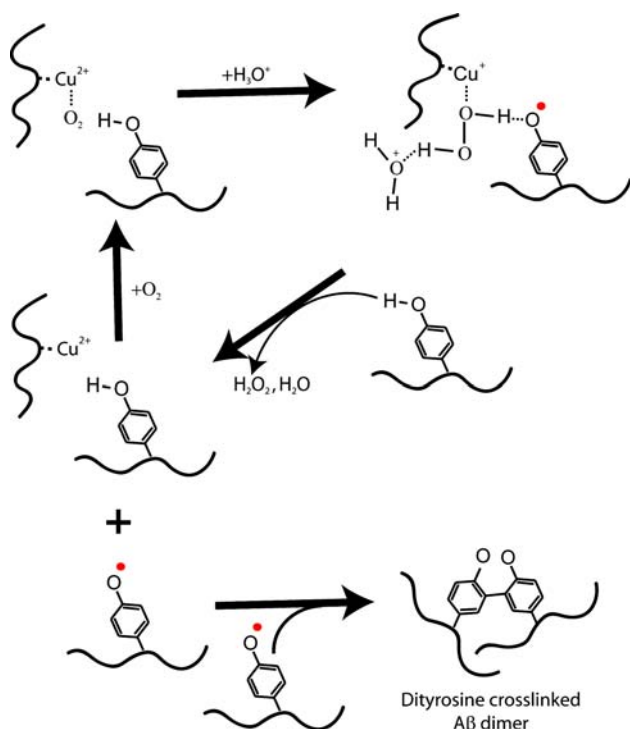


Fig. 3 Simplified diagram of the A β /Cu-catalysed production of H₂O₂ from O₂ and ascorbate. This reaction can generate tyrosyl radicals which can crosslink to form dityrosine crosslinked A β dimers

alanine is substituted for the tyrosine at position 10 (Y10A). The mutant A β peptide is incapable of generating dityrosine crosslinks, and although it produced significant amounts of H₂O₂ and was shown to aggregate with a similar propensity as the wildtype A β , Y10A did not induce any neurotoxicity (Barnham et al. 2004).

It has also been suggested A β toxicity is due to the catalytic formation of reactive oxygen species (ROS) from membrane-associated peptides. Experiments with an M35V mutant A β peptide (methionine to valine at position 35) have shown increased toxicity despite levels of H₂O₂ production similar to wild type A β . This is believed to be due to increased membrane binding (Ciccotosto et al. 2004). In vivo, AD brains display a high level of oxidative stress and ROS-mediated injury. ROS-mediated damage is detected in the brain primarily as an increase in lipid peroxidation, and increased levels of malondialdehyde, 4-hydroxynonenal, 8-hydroxydeoxyguanosine, protein carbonyls and, most importantly, increased dityrosine levels (Hensley et al. 1998).

The intricate relationship between A β , copper, radical and ROS formation and neurotoxicity has led to the development and use of compounds to modulate metal homeostasis and so prevent neurodegeneration. The metal ligand clioquinol (5-chloro-7-iodo-quinolin-8-ol) has been successfully used in vitro, in animal models and in small clinical trials. Clioquinol and similar compounds have been

shown to inhibit the in vitro generation of H₂O₂ by A β , and reverse the aggregation of the peptide both in vitro and from human brain post-mortem specimens. In animal models, oral administration of clioquinol markedly reduced the accumulation of A β (Cherny et al. 2001). Further metal-protein attenuating compounds are in development.

Future directions

Despite this progress, the chemical nature of the toxic oligomeric A β species is still unknown, the exact nature of inter-molecular crosslinking and its method of formation has not been established. Whether this crosslinking is due to the cellular processing of APP and A β or to the intrinsic chemical nature of the peptide remains to be determined. As noted above, A β can be induced in vitro to form dityrosine crosslinks via a mechanism involving metal-catalysed redox chemistry. It is interesting to note that, of the mammalian proteins with intra-molecular crosslinks described in the literature (for example, cytochrome c oxidase, catalase and lysyl oxidase), many involve copper-based enzymes or enzymes associated with reactive oxygen species [reviewed in (Okeley and van der Donk 2000)]. A β , of course, is also associated with both ROS production and copper. However, although data supporting dityrosine formation in vivo have been published (Hensley et al. 1998), it has not yet been shown that dityrosine A β adducts can be isolated from in vivo sources. It has also been suggested that the oligomers may be linked by an N^ε-(γ -glutamyl)-lysine bond mediated by tissue transglutaminase. High levels of this enzyme have also been described in Huntington's disease and Parkinson's disease brains, and tissue transglutaminase is also activated during apoptosis (Ho et al. 1994; Nemes et al. 2001; Citron et al. 2002). If this is the case, clioquinol and other MPACs may be effective treatments due to secondary effects on A β production, such as metal-based modulation of secretase activity. Alternately, A β monomers may become crosslinked by other reactive cellular products, such as carbohydrates or aldehydes, including those produced by the effects of ROS created by A β /Cu complexes. Understanding how A β oligomerisation occurs is an essential step in understanding and preventing AD, so further research into the peptide's chemistry is essential.

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