

Amyloid β -Cu²⁺ Complexes in both Monomeric and Fibrillar Forms Do Not Generate H₂O₂ Catalytically but Quench Hydroxyl Radicals[†]

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ABSTRACT: Oxidative stress plays a key role in Alzheimer's disease (AD). In addition, the abnormally high Cu²⁺ ion concentrations present in senile plaques has provoked a substantial interest in the relationship between the amyloid β peptide (A β) found within plaques and redox-active copper ions. There have been a number of studies monitoring reactive oxygen species (ROS) generation by copper and ascorbate that suggest that A β acts as a prooxidant producing H₂O₂. However, others have indicated A β acts as an antioxidant, but to date most cell-free studies directly monitoring ROS have not supported this hypothesis. We therefore chose to look again at ROS generation by both monomeric and fibrillar forms of A β under aerobic conditions in the presence of Cu²⁺ with/without the biological reductant ascorbate in a cell-free system. We used a variety of fluorescence and absorption based assays to monitor the production of ROS, as well as Cu²⁺ reduction. In contrast to previous studies, we show here that A β does not generate any more ROS than controls of Cu²⁺ and ascorbate. A β does not silence the redox activity of Cu^{2+/+} via chelation, but rather hydroxyl radicals produced as a result of Fenton–Haber Weiss reactions of ascorbate and Cu²⁺ rapidly react with A β ; thus the potentially harmful radicals are quenched. In support of this, chemical modification of the A β peptide was examined using ¹H NMR, and specific oxidation sites within the peptide were identified at the histidine and methionine residues. Our studies add significant weight to a modified amyloid cascade hypothesis in which sporadic AD is the result of A β being upregulated as a response to oxidative stress. However, our results do not preclude the possibility that A β in an oligomeric form may concentrate the redox-active copper at neuronal membranes and so cause lipid peroxidation.

Along with the accumulation of extracellular amyloid plaques, Alzheimer's disease (AD)¹ is characterized by marked oxidative injury. Numerous studies have shown evidence of increased levels of oxidative damage in the brain tissue, as well as in the cerebrospinal fluid (CSF) and plasma, of patients with AD. Observed oxidative damage includes protein oxidation (1), DNA and RNA oxidation (2), and also lipid peroxidation (3) (as reviewed in ref 4). Indeed, much of the A β isolated from plaques contains oxidized methionine (Met³⁵) (5, 6). Intriguingly, some studies have shown that oxidative stress occurs before the appearance of amyloid plaques (7, 8).

The principal component of amyloid plaque deposits is a small peptide, amyloid β peptide (A β), of between 39 and 43 residues in length. Soluble A β 40 and A β 42 are found in all humans in the blood plasma, bound principally to serum albumin (9), and the CSF where A β 40 has a concentration

of 5 nM (10). One of the most compelling pieces of evidence for A β being the primary culprit in AD comes from genetic mutations underlying familial AD. These mutations are within A β itself or are associated with increased production of A β 42 in the brain, which promotes more rapid fibril growth (11). Small oligomers or pore-like protofibrils rather than mature amyloid fibers may be the neurotoxic form of A β (12). The mechanism by which A β is toxic to neurons remains unclear, but it is suggested the cause may be ROS production (13, 14), perturbation in Ca²⁺ regulation, or interaction of A β oligomers with the membrane causing cell depolarization.

In vivo, copper ions are a major source of reactive oxygen species (ROS) generation via Fenton–Haber Weiss reactions. Under aerobic conditions oxidation of Cu⁺ ions results in the generation of hydrogen peroxide, superoxide, and hydroxyl radicals (15). The redox-active metal ion Cu²⁺ is found bound to A β in isolated senile plaque cores (5). Total copper content within the brain increases with age, as do oxidative processes. Due to the potentially toxic nature of copper ions they are carefully regulated and compartmentalized. However, the dyshomeostasis of redox-active metal ions occurs in AD, with copper being found in abnormally high concentrations, 340 μ M, in cerebral amyloid deposits (16). Controversially, trace amounts of copper in a cholesterol-high diet were found to induce amyloid β plaques and learning deficits in a rabbit model of Alzheimer's disease (17, 18).

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¹ Abbreviations: A β , amyloid β ; AD, Alzheimer's disease; APP, amyloid precursor protein; BC, bathocuproinedisulfonic acid; 3-CCA, coumarin-3-carboxylic acid; CSF, cerebrospinal fluid; EDTA, ethylenediaminetetraacetic acid; His, histidine; H₂O₂, hydrogen peroxide; 7-OHCCA, 7-hydroxycoumarin-3-carboxylic acid; 8-OHG, 8-hydroxyguanine; LDL, low-density lipoprotein; Met, methionine; Met^{ox}, methionine sulfoxide; Msr, methionine sulfoxide reductase; NMR, nuclear magnetic resonance; ODAD, *o*-dianisidine; 2-oxo-His, 2-oxohistidine; PBS, phosphate-buffered saline; ROS, reactive oxygen species.

As such, antioxidants and copper chelators represent possible therapeutics for AD (19).

The most widely accepted model of Cu^{2+} bound to $\text{A}\beta$ indicates Cu^{2+} ions bind to histidine residues within the N-terminal portion (His⁶, His¹³, His¹⁴) in a square-planar complex (20). The nature of the fourth ligand is still debated, with the N-terminus (20–22), tyrosine (23), and glutamic or aspartic acid (24) proposed. Reported Cu^{2+} affinities for monomeric $\text{A}\beta$ vary widely between micromolar (μM) (25), nanomolar (nM) (20, 26), and the much higher attomolar, $K_d \sim 10^{-18}$ affinity for $\text{A}\beta 42$ (27). Cu^{2+} ions have a plasma concentration of $\sim 15\text{--}20\ \mu\text{M}$ and a lower concentration of $\sim 250\ \text{nM}$ in the CSF (28); however, when released at the synapse concentrations can be as high as $250\ \mu\text{M}$ (29).

Both antioxidant and prooxidant roles for $\text{A}\beta$ have been suggested. As a prooxidant, the neurotoxicity of $\text{A}\beta$ peptides has been linked to oxidative ROS generation (30–32). Neuronal cell viability studies indicate that reactive oxygen species mediate the toxicity of $\text{A}\beta$, with the neurotoxicity of $\text{A}\beta$ and Cu^{2+} being abolished by the addition of catalase or anti- $\text{A}\beta$ antibodies (33) or attenuated through the use of metal chelators (34). In contrast, others have highlighted an antioxidant activity of $\text{A}\beta$ (35, 36). In particular, monomeric $\text{A}\beta 40$ and $\text{A}\beta 42$ were shown to inhibit neuronal death caused by transition metal-induced oxidative damage (37). A dual role for $\text{A}\beta$ has also been suggested (38–40).

A number of groups have published studies monitoring the generation of ROS in the presence of $\text{A}\beta$ and Cu^{2+} in a cell-free system. Early studies suggested spontaneous peptide free radical generation (41). However, this was later disputed by a number of groups (42, 43). It was then suggested that $\text{A}\beta$ in fact required the presence of transition metal ions (e.g., Cu^{2+} and Fe^{3+}) to produce reactive oxygen species via Fenton–Haber Weiss-type reactions (44–47). However, redox potentials recently determined for the Cu – $\text{A}\beta$ complex suggested that this is unlikely (48). Initial studies by Huang et al. suggested that Cu^{2+} -bound $\text{A}\beta$ was capable of generating H_2O_2 without the presence of an additional reducing agent (44, 45). However, subsequent studies by the same laboratory indicated that the initial experiments were flawed, and the method used to detect the H_2O_2 contained a reagent (TCEP) capable of acting as a reducing agent for $\text{A}\beta 40$ and $\text{A}\beta 42$ but not the controls (33). This study highlighted the additional requirement of biological reductants (such as ascorbic acid, cholesterol, L-DOPA, or dopamine) to act as substrates for the sequence of electron transfer reactions (33, 49). A model proposed by Opazo et al. for catalytic hydrogen peroxide production by $\text{A}\beta$ and Cu^{2+} hypothesizes that reduction of the metal ion occurs through the transfer of an electron from the peptide backbone, with the peptide radical then being reconstituted through recruitment of a biological reducing agent (33). However, in contrast more recently studies have indicated H_2O_2 depletion in a Cu^{2+} /ascorbate system by $\text{A}\beta$ (31) and similarly hydroxyl radical depletion (50).

The aim of this study is to characterize the anti- and/or prooxidant nature of the $\text{A}\beta$ peptide, in both monomeric and fibrillar form. To date there have been many inconsistencies in the literature on this subject, and the role of $\text{A}\beta$ in relation to the observed oxidative stress in AD is still highly disputed. Cu – $\text{A}\beta$ samples were studied in the presence and absence of a physiological reductant, ascorbate, in a cell-free system.

ROS generation was monitored using absorption and fluorescence assays. Cu^{2+} reduction was monitored using the BC assay, and effects of redox cycling on $\text{A}\beta$ itself were monitored using ^1H NMR.

MATERIALS AND METHODS

Peptide Syntheses and Purification. Peptides representing various fragments of the amyloid β peptide were synthesized by employing solid-phase F-moc chemistry (ABC; Imperial College, London). After removal from the resin and deprotection, the samples were purified using reverse-phase HPLC and characterized using mass spectrometry.

Peptides synthesized were as follows: $\text{A}\beta(1\text{--}42)$, $\text{A}\beta(1\text{--}40)$, $\text{A}\beta(1\text{--}28)$, and $\text{A}\beta(1\text{--}16)$, designated as $\text{A}\beta 42$, $\text{A}\beta 40$, $\text{A}\beta 28$, and $\text{A}\beta 16$ in the text. In addition, the neurotoxic fragment $\text{A}\beta(25\text{--}35)$ was synthesized. $\text{A}\beta 42$ is DAE-FR H D S G Y E V H H Q K L V F F A E D V G S N K G A I-I G L M V G G V V I A.

Full-length $\text{A}\beta$ peptides ($\text{A}\beta 42$ and $\text{A}\beta 40$) were synthesized with the N-terminus left as the native amino group and a carboxyl group at the C-terminus, while $\text{A}\beta$ fragments were synthesized with an amino group at the N-terminus, and the truncated C-terminus was blocked as the corresponding ethyl ester. Although $\text{A}\beta(25\text{--}35)$ is not a naturally occurring peptide, its N-terminus was not acetylated as previous studies into this fragment's toxicity used a free N-terminal amino group.

The peptide concentrations were determined using the extinction coefficient at 280 nm of the single tyrosine residue at position 10, specifically $1280\ \text{M}^{-1}\ \text{cm}^{-1}$. Typically, the freeze-dried peptides contained $\sim 20\%$ moisture by weight.

Solubilization of $\text{A}\beta$ and Amyloid Fibril Formation. Lyophilized peptide stocks were solubilized in water, with additional processes being needed for full-length $\text{A}\beta$ stocks to prevent aggregation/oligomerization. Briefly, $\text{A}\beta 40$ and $\text{A}\beta 42$ were dissolved at $\sim 0.7\ \text{mg/mL}$ concentration at pH 10.5 and kept on ice ($0\text{--}5\ ^\circ\text{C}$) to keep aggregation/oligomerization to a minimum. Samples described in this paper as monomeric $\text{A}\beta$ were freshly prepared at pH 7.4, with ThT and CD measurements suggesting minimal fibrillar content. However, even freshly prepared $\text{A}\beta$ may contain some dimers and low molecular weight oligomers.

Amyloid fibrils of $\text{A}\beta 40$ and $\text{A}\beta 42$ were generated by incubation of full-length $\text{A}\beta$ ($50\ \mu\text{M}$) at $37\ ^\circ\text{C}$ in 10 mM phosphate buffer at pH 7.4 with 300 mM NaCl (PBS) and 0.05% sodium azide. Fibrils were generated over a period of 120 h with gentle agitation on a rocker. Fibril formation was confirmed by a variety of techniques including far-UV circular dichroism (CD), thioflavin T (ThT) fluorescence assay (see Supporting Information Figure S1), seeding experiments, and electron microscopy (EM) imaging.

Materials. Unless otherwise stated, all chemicals were purchased from Sigma-Aldrich at the highest purity available. Deuterium oxide was obtained from GOSS Scientific Instruments Ltd. Water used was of ultrahigh quality ($> 18\ \Omega\ \text{cm}^{-1}$ resistivity) level. All studies were performed under aerobic conditions.

H_2O_2 Detection: *o*-Dianisidine (ODAD) and Peroxidase Assay. A modified version of the *o*-dianisidine (ODAD) and peroxidase assay (51, 52) was utilized to monitor hydrogen peroxide production. A Thermo Labsystems Multiskan

Ascent photometric microplate reader fitted with a 450 nm filter was used to detect hydrogen peroxide with this assay, using 96-well culture plates (Corning Costar). Stock solutions were prepared freshly for each experiment, and each 50 μ L reaction mixture typically contained 10 μ M peptide, 10 μ M CuCl_2 , and 30 μ M ascorbate in 10 mM phosphate buffer, pH 7.4, with 300 mM NaCl (PBS). Samples were left to incubate for 1 h at 37 °C, and then a volume of 250 μ L of assay reagents was added to each reaction mixture. The assay reagents consisted of 240 μ L of ODAD (0.066 mg in 100 mL of phosphate buffer) and 10 μ L of peroxidase (1 mg/mL in H_2O). Following reagent addition, the samples were subjected to a period of shaking (30 s at 60 rpm) using the inbuilt shaker to mix the sample and reagents, and the absorbance readings were measured. An extensive number of different sample/control conditions were tested, with up to 24 repeats being performed of each sample condition and then an average calculated. Absorbance readings of phosphate buffer plus assay reagents were recorded to be used later for baseline subtraction. The net changes in absorbance (ΔA) were determined by multiplying the absorbance by 6 to return it to original concentrations (50 μ L of sample was diluted by 250 μ L of assay reagent) and then subtracting the baseline absorbance readings of phosphate buffer plus reagents. The amount of H_2O_2 produced was quantified by comparing resultant absorbance readings with a linear calibration curve acquired with known concentrations of H_2O_2 (see Supporting Information Figure S2).

Hydroxyl Radical Detection: Coumarin-3-carboxylic Acid (3-CCA) Assay. A variety of different peptide fragments/forms were studied in the presence and absence of ascorbate and CuCl_2 and monitored for hydroxyl radical production. A modified version of the coumarin-3-carboxylic acid (3-CCA) assay (52, 53) was used to quantify hydroxyl radical production. Upon reaction with hydroxyl radicals the 3-CCA forms a fluorescent product, 7-OH-CCA, with characteristic emission at 450 nm and excitation maximum at 388 nm. Samples typically contained 1 mM 3-CCA, 10 μ M peptide, 10 μ M CuCl_2 , and 30 μ M ascorbate, in 40 mM phosphate buffer (pH 7.4). Prior to recording spectra samples were incubated for 1 h at 37 °C. At least three replicates were performed of each sample condition. Emission readings were measured on a Hitachi F-2500 fluorescence spectrophotometer, and fluorescence spectra were collected over the wavelength range 410–550 nm at 37 °C. A very slight baseline fluorescence of 3 AFU was observed from the unreacted 3-CCA reagent. Phosphate buffer plus reagent was used as a baseline sample and subtracted from all other fluorescence values.

Cu^+ Detection: Bathocuproinedisulfonic Acid (BC) Assay. For monitoring the reduction of Cu^{2+} ions the specific Cu^+ chelator bathocuproinedisulfonic acid (BC) was used, by observing the characteristic purple color produced by the complex $\text{Cu}^+(\text{BC})_2$ ($\epsilon_{483\text{nm}} = 12250 \text{ M}^{-1} \text{ cm}^{-1}$) which has an absorbance maximum at 483 nm (52, 54). Typically, a 1 mL sample containing 10 μ M $A\beta$, 10 μ M CuCl_2 , 10 mM phosphate buffer, pH 7.4, and 300 mM NaCl (PBS) was allowed to incubate with 360 μ M BC. Absorbance readings were monitored at two time intervals, after 60 s and 60 min, with samples being incubated and recorded at 37 °C. The capacity of BC to detect Cu^+ is rapid (less than 30 s).

UV-visible absorption spectra were collected using a Hitachi U-3010 spectrophotometer. An array of control samples were recorded for baseline readings to be subtracted from resultant spectra. Three replicates were performed for each sample condition. This assay uses optical absorption and therefore may be influenced by light scatter from turbidity in the solution which should be considered. So, further control samples of peptide plus CuCl_2 , and peptide plus BC, were performed to check absorbance contributions from light scatter. We see from the controls of $A\beta$ that the addition of BC (or CuCl_2) alone to $A\beta_{40}$ does not cause an increase in apparent absorption due to light scatter (see Figure 3b), while the less soluble $A\beta_{42}$ fragment does produce slight light scatter indicated by very weak absorbance of ~ 0.003 at 483 nm. However, the effect is small and does not alter the interpretation of the data. Phosphate buffer plus BC reagent was therefore used as the baseline condition to be subtracted.

^1H Nuclear Magnetic Resonance (NMR) Spectroscopy. Experiments were carried out on a Bruker Avance spectrometer, operating at a 600 MHz ^1H frequency. Water was suppressed using the excitation sculpting with gradients method. NMR experiments were recorded at either 30 or 37 °C, pH 7.4. Samples contained 100 μ M peptide incubated and monitored over time for up to 60 h with varying amounts of CuCl_2 and ascorbate. A lack of NaCl and agitation mean that fibrillization is minimal over the period. Alternatively, samples of $A\beta_{25-35}$ at 140 μ M were incubated at 4 °C and then monitored after a 7 day incubation. NMR samples contained 10% D_2O and 90% H_2O or 100% D_2O . The pH of NMR samples was recorded before and after acquisition. An exponential line broadening of 3.0 Hz was applied to the data before Fourier transformation. Chemical shifts were typically referenced to water, or formate, which produces a singlet at 8.495 ppm. Typically, ^1H NMR spectra were obtained with either 256 or 512 transients with a recycle time of ~ 4 s. For analysis purposes, individual spectra were summed to compare the results of spectra acquired over ~ 40 min duration with results acquired over several hours of incubation. Assignments of His ring protons and other resonances are based on 2D TOCSY (total correlation spectroscopy) studies and the use of His/Ala analogues described elsewhere (20, 55).

RESULTS

Influence of $A\beta$ on H_2O_2 Production by Copper and Ascorbate. In the course of $\text{Cu}^{2+/+}$ redox cycling via Fenton–Haber Weiss reactions, the reactive oxygen species H_2O_2 is generated. Thus detection of H_2O_2 is an effective method of monitoring Fenton–Haber Weiss reactions. The ability of $A\beta$ to generate H_2O_2 in the presence of Cu^{2+} ions was studied using the ODAD and peroxidase assay, monitoring the absorbance value at 450 nm.

Reaction mixtures of CuCl_2 (10 μ M) with/without various $A\beta$ peptide fragments (10 μ M) were incubated at 37 °C for 60 min prior to measurement under aerobic conditions. The recorded absorbance measurements for these samples were converted into concentration of H_2O_2 , based on H_2O_2 calibration curve data (see Supporting Information Figure S2). Figure 1 shows the results of a selection of control and peptide samples. As expected, the buffer or Cu^{2+} ions on their own do not generate any H_2O_2 . A solution of ascorbate

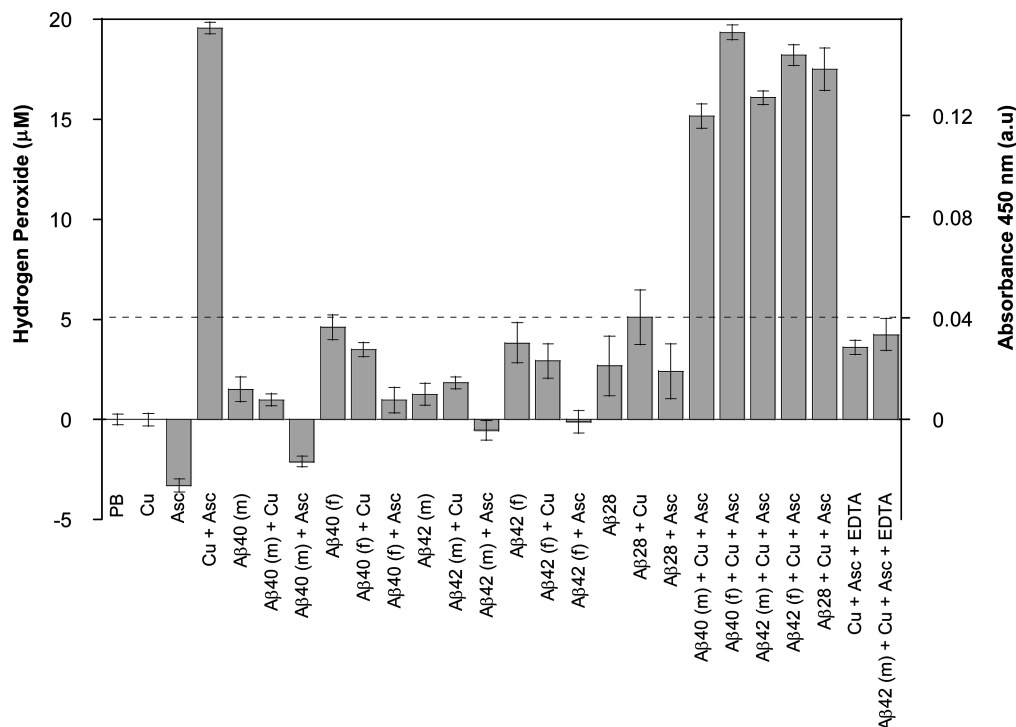


FIGURE 1: H_2O_2 production by Cu^{2+} and ascorbate in the presence of monomeric and fibrillar $\text{A}\beta$, monitored using ODAD and peroxidase assay. $\text{A}\beta_{28/40/42}$ (10 μM) and/or CuCl_2 (10 μM) and/or ascorbate (30 μM) in various combinations, all in 10 mM phosphate buffer, pH 7.4. Mean values are shown in both absorbance at 450 nm and hydrogen peroxide concentration on the y-axes. Samples were incubated for 60 min at 37 °C prior to reagent addition. Plot shows the average ODAD assay result for samples, based on 18 or 24 replicates, with standard error bars plotted. Dashed line represents background noise level observed in the assay (m = monomeric, f = fibrillar).

(no copper) causes a slight reduction in absorbance from baseline levels due to interactions with the assay reagents. Monomeric $\text{A}\beta_{40}$, in the absence of Cu^{2+} , gives a slight H_2O_2 signal ($1.5 \pm 0.6 \mu\text{M}$); however, this is clearly not caused by trace amounts of Cu^{2+} because addition of 1 mol equiv of Cu^{2+} (10 μM) to $\text{A}\beta$ produces the same amount of H_2O_2 ($1.0 \pm 0.4 \mu\text{M}$). The addition of a reducing agent, ascorbate, to $\text{A}\beta_{40}$ (in the absence of CuCl_2) does not generate H_2O_2 either. Similarly, fibrillar $\text{A}\beta_{40}$, and monomeric and fibrillar $\text{A}\beta_{42}$, also showed no significant difference in absorbance at 450 nm with or without the presence of Cu^{2+} ions. This indicates that the small absorbance at 450 nm observed here is not the result of Fenton's cycling of Cu^{2+} to Cu^+ . The very weak absorbance values are simply an artifact of turbidity in this optically based assay. $\text{A}\beta_{28}$ plus Cu^{2+} , which we later show does not generate Cu^+ ions and therefore cannot perform Fenton's cycling, also produces a slight absorbance at 450 nm comparable to $\text{A}\beta_{40}$ and $\text{A}\beta_{42}$. A cutoff at which turbidity in the assay can affect apparent absorbance is shown in Figure 1 set at 0.04 absorbance unit.

These experiments indicate that $\text{Cu-A}\beta_{40}$ and $\text{Cu-A}\beta_{42}$ in their monomeric or fibrillar form do not generate H_2O_2 . The only samples to produce H_2O_2 (15–20 μM) are those that contain Cu^{2+} with physiological levels of ascorbate, irrespective of the presence or absence of $\text{A}\beta$.

Further studies using the ODAD and peroxidase assay focused on testing the additional effect of a physiological reducing agent, ascorbate. As before, samples were monitored at 450 nm after 60 min incubation at 37 °C but this time contained a mixture of ascorbate (30 μM) with/without CuCl_2 (10 μM) with/without various $\text{A}\beta$ peptide fragments (10 μM). Note, extracellular levels of ascorbate are typically between 30 and 100 μM (15). The nonpeptide sample of

ascorbate and CuCl_2 showed production of $\sim 19.5 \mu\text{M}$ H_2O_2 after 60 min at 37 °C (see Figure 1). When we directly compare this to samples that additionally contain $\text{A}\beta$ peptides, it is evident that the amount of detectable H_2O_2 is slightly reduced in the presence of $\text{A}\beta$ rather than increased. The exact degree of H_2O_2 depletion varies with peptide fragment, with detected H_2O_2 ranging between 78% and 99% compared to that of CuCl_2 with ascorbate on its own. It is evident from Figure 1 that the monomeric forms of full-length $\text{A}\beta$ ($\text{A}\beta_{40}$ and $\text{A}\beta_{42}$) show a greater capacity to decrease detectable H_2O_2 compared to the fibrillar forms, 78–83% H_2O_2 for monomeric $\text{A}\beta$ compared to the Cu^{2+} and ascorbate sample, while fibrillar forms show 94–99% H_2O_2 . The shorter more soluble $\text{A}\beta_{28}$ fragment also demonstrates a capacity to deplete H_2O_2 generated (see Figure 1) comparable to full-length $\text{A}\beta$.

These data indicate that in the presence of a physiological reductant (ascorbate) the $\text{A}\beta$ peptides will not increase the amount of H_2O_2 generated, compared to the ascorbate and CuCl_2 sample alone. This was a surprise as there have been reports that the $\text{Cu-A}\beta$ complex generates H_2O_2 in the presence of an additional reductant (33). The latter studies were performed in the presence of EDTA, so for this reason we repeated these types of experiments with some additional controls. We see from Figure 1 that the presence of EDTA (50 μM) dramatically decreases the amount of H_2O_2 generated by the ascorbate (30 μM) and CuCl_2 (10 μM) mixture to baseline levels. It has been suggested that the addition of $\text{A}\beta$ to this mixture of ascorbate, CuCl_2 , and EDTA increased H_2O_2 production (33); however, we found no such difference between ascorbate, CuCl_2 , and EDTA with or without the presence of $\text{A}\beta$ (10 μM). Indeed, both samples were below the cutoff for the baseline levels as shown in Figure 1.

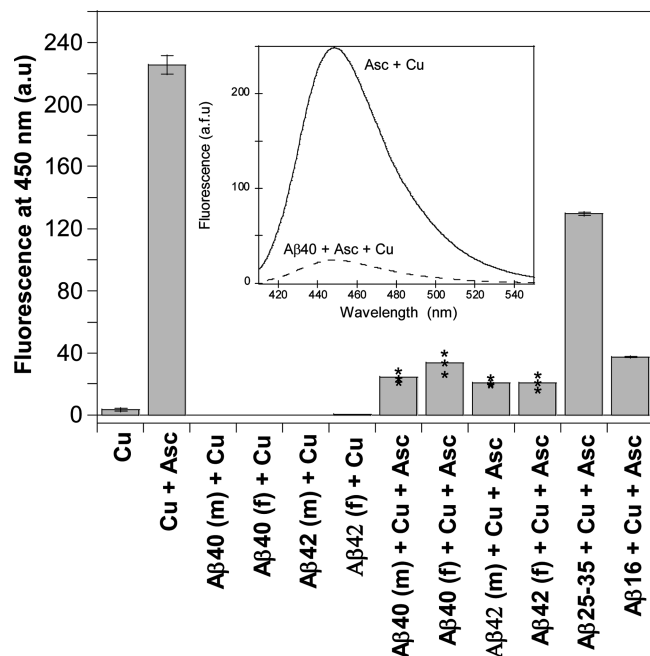


FIGURE 2: Hydroxyl radical production by Cu²⁺ and ascorbate in the presence of monomeric and fibrillar A β , monitored using 3-CCA assay. A β 16/28/25–35/40/42 (10 μ M) and/or CuCl₂ (10 μ M) and/or ascorbate (30 μ M) in various combinations, with 1 mM fluorophore 3-CCA. All are in 40 mM phosphate buffer, pH 7.4. Plot shows the average 3-CCA result for samples, based on three or more replicates of each sample condition. Samples were incubated for 60 min at 37 °C (m = monomeric, f = fibrillar).

The latter studies were performed at a 1:1 ratio of peptide to Cu²⁺. Further studies testing above a stoichiometric ratio of peptide to Cu²⁺ (i.e., 1:2), with Cu²⁺ loaded as Cu(Gly)₂, showed similar behavior. In the presence of A β the amount of H₂O₂ generated was depleted compared to the ascorbate and Cu(Gly)₂ control (see Supporting Information Figure S3).

Influence of A β on Hydroxyl Radical Production by CuCl₂ and Ascorbate. Next, the redox properties of the Cu–A β complex were studied by monitoring hydroxyl radical production. Hydroxyl radicals form an important part of redox chemistry; they are highly reactive and therefore highly toxic to cells. The quantitative detection of hydroxyl radicals was carried out using the 3-CCA fluorescence assay, monitoring the product 7-OHCCA fluorescing at 450 nm. Reaction mixtures were incubated for 1 h at 37 °C. Sample conditions were CuCl₂ (10 μ M) and A β peptide fragments (10 μ M), with/without ascorbate (30 μ M) at pH 7.4 under aerobic conditions.

As expected, the copper and ascorbate mixture produced appreciable amounts of hydroxyl radicals, as seen by the intense fluorescence signal (Figure 2). In contrast, incubation of CuCl₂ with A β 40 or A β 42, both monomeric and fibrillar, did not generate any fluorescence signal. It is clear that CuCl₂ with A β does not generate any hydroxyl radicals, in agreement with the previous H₂O₂ assay, and that reactive oxygen species are not generated in the absence of ascorbate.

We then went on to study the effect of ascorbate on the Cu–A β system. Incubation of CuCl₂ and A β with ascorbate for 1 h generates appreciable fluorescence signal, although the intensity is ~10–20% that of CuCl₂ and ascorbate on its own. In particular, both monomeric and fibrillar full-length A β (A β 40 and A β 42) show a marked reduction in fluorescence, with ~90% depletion in the amount of hydroxyl

radicals detected compared to the CuCl₂ and ascorbate without A β .

Fragments of A β (A β 16 and A β 28) were also studied, and the depletion (~85%) of hydroxyl radicals detected with these fragments is comparable with the full-length peptide (A β 40 and A β 42) results. However, A β (25–35), that contains Met³⁵ but not the histidines of the Cu²⁺ binding site, is much less effective at reducing hydroxyl radicals detected and only shows ~40% reduction relative to the CuCl₂ and ascorbate sample.

It is clear from the detection of hydroxyl radicals that A β does not silence redox cycling of the Cu²⁺ ions in the presence of a physiological reductant (ascorbate). Interestingly, from the ODAD assay it is clear the amount of H₂O₂ generated by CuCl₂ and ascorbate is only slightly reduced in the presence of A β by between 1% and 22%. In contrast, the 3-CCA assay, which monitors hydroxyl radicals, suggests that the appreciable amounts of hydroxyl radicals generated by CuCl₂ and ascorbate are substantially decreased by the presence of A β by ~80–90%.

Rather than A β acting to redox silence Cu²⁺ ions, it appears to quench the hydroxyl radicals generated through Fenton–Haber Weiss reactions involving CuCl₂ and ascorbate. This quenching results in appreciable oxidation of A β itself (described later) as observed by ¹H NMR and mass spectrometry.

As with CuCl₂-containing samples, experiments using Cu(Gly)₂ as the source of Cu²⁺ ions also showed a marked reduction (~80%) in detectable hydroxyl radicals in the presence of A β (i.e., A β 40 plus Cu(Gly)₂ and ascorbate samples) compared to the Cu(Gly)₂ and ascorbate control (see Supporting Information Figure S4).

Cu²⁺ Reduction to Cu⁺ by A β . Key reactions associated with oxidative stress are the redox cycling of Cu^{2+/+} ions; this process is a major source of reactive oxygen species *in vivo* via the Fenton–Haber Weiss reactions (15). To examine any inherent ability of the A β peptide to reduce Cu²⁺ to Cu⁺, both fragments and full-length A β in monomeric and fibrillar form were studied using the BC assay. The Cu⁺(BC)₂ complex produces a characteristic purple color with an absorption maximum at 483 nm, allowing any Cu⁺ production to be quantitatively detected. In Figure 3 the reduction of Cu²⁺ by three different A β fragments is shown. The absorbance band at 483 nm was monitored after 60 s (Figure 3a) and 60 min (Figure 3b) incubations at 37 °C under aerobic conditions. Control samples in the presence of BC (360 μ M) and CuCl₂ (10 μ M), in which no A β fragments were present, are also shown for both incubation times. We see from Figure 3 that BC alone is able to reduce very small amounts of Cu²⁺ itself, as has previously been noted (56). Contributions to Cu²⁺ reduction from the BC alone were shown to be ~1.0 μ M after 60 s and ~2.2 μ M after 60 min. The effect of this background reduction of Cu²⁺ by BC alone is highlighted in the data, shown in Figure 3.

From the 60 s sample incubations, it is evident that the shorter, more soluble fragment A β 28 actually shows less Cu⁺ production than the nonpeptide sample, detecting ~0.8 μ M Cu⁺ compared to the ~1.0 μ M by BC alone (see Figure 3a). A similar result was observed for other shorter fragments tested, including A β 16 and A β (25–35) (data not shown) with these fragments showing Cu²⁺ reduction less than or equivalent to that by BC alone. In contrast, both the full-

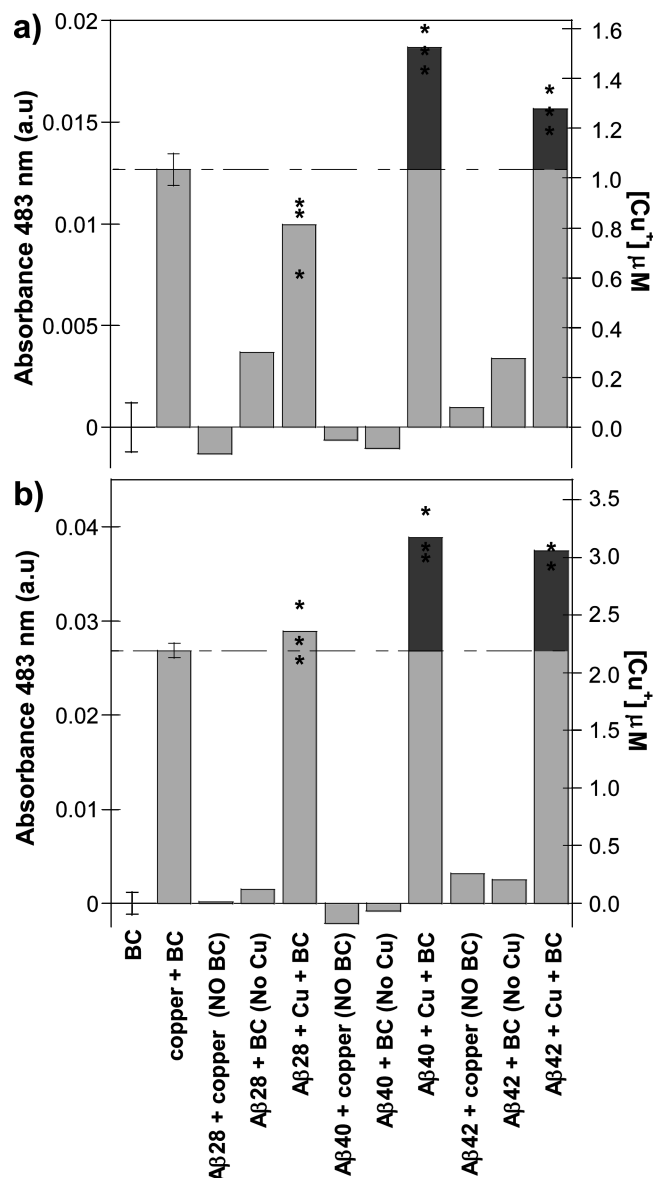


FIGURE 3: Cu²⁺ reduction by Aβ, monitored using the BC assay. Incubations contained Aβ28/40/42 (10 μM) and/or CuCl₂ (10 μM) and/or BC (360 μM). Dashed line represents Cu⁺ reduction level by BC alone. Results are shown in both absorbance at 483 nm and Cu⁺ concentration on the y-axes. Plot shows the average BC assay results, based on three replicates, at two different incubation periods: (a) 60 s incubation at 37 °C and (b) 60 min incubation at 37 °C.

length fragments, Aβ40 and Aβ42, produced slightly greater amounts of Cu²⁺ reduction to Cu⁺ than BC alone after the 60 s incubation at 37 °C. Levels of Cu⁺ ions detected in the presence of full-length peptides are ~1.5 and ~1.3 μM (Aβ40 and Aβ42, respectively) compared to the ~1.0 μM reduced by BC alone. This suggests that under these conditions Aβ possesses some intrinsic ability to convert Cu²⁺ to Cu⁺ by donation of an electron. With longer incubation times (60 min at 37 °C) (see Figure 3b), the Cu²⁺ reduction in the presence of Aβ40 and Aβ42 is more apparent. Specifically, the data in Figure 3b show that full-length Aβ reduces ~3.2 and ~3.1 μM (Aβ40 and Aβ42, respectively) compared to the ~2.2 μM reduced by BC alone after the longer incubation time.

Cu²⁺ reduction by fibrillar Aβ40 and Aβ42 was studied to compare with the reducing capacity demonstrated by the monomeric forms of the peptide. Fibrillar full-length Aβ40

and Aβ42 also showed some Cu²⁺ reduction capabilities beyond the intrinsic ability of BC, which was comparable to monomeric Aβ. However, the extent of increase was variable and compromised by turbidity in the assay for these fibrillar samples.

Samples of Aβ were incubated at 37 °C with Cu²⁺ for 1 h with or without the presence of BC. Samples incubated for 1 h in the absence of BC, after which the BC was allowed to react for just 60 s, did not show any additional Cu²⁺ reduction beyond that seen in the 60 s incubations. It appears that BC drives the reduction of Cu²⁺ to Cu⁺ more readily in the presence of Aβ.

It is clear then that fragment size of the Aβ peptide has an influence on its Cu²⁺ reduction capabilities. The full-length fragments (Aβ40 and Aβ42) have the ability to promote the reduction of Cu²⁺ in the presence of BC, whereas the shorter fragments Aβ16 and Aβ28, which lack the hydrophobic C-terminus and the highly oxidizable Met³⁵, do not promote reduction of Cu²⁺ ions.

We were also interested in the effect of ascorbate on Cu–Aβ samples. Using the BC assay we saw complete and rapid reduction of all the Cu²⁺ present irrespective of the presence or absence of Aβ. It is clear the Aβ does not redox silence Cu²⁺ ions.

Oxidative Modification of Aβ by a Cu²⁺/Ascorbate System. In addition to monitoring the effect of Aβ on ROS production by a copper and ascorbate, we also examined the effects of this ROS generation on Aβ itself. Full-length Aβ42 was studied using ¹H NMR spectroscopy and monitored over a period of 60 h in the presence of CuCl₂ and ascorbate (or H₂O₂) as shown in Figure 4. In the absence of ascorbate, the paramagnetic effect of small amounts of Cu²⁺, 0.05 mol equiv, on the Aβ42 histidine resonances and other signals is minimal, with only slight broadening observed (see Supporting Information Figure S5). Incubation of Cu–Aβ in the absence of ascorbate showed no oxidation of the peptide. Furthermore, upon the addition of ascorbate the effect on the line width or intensity of most of the Aβ resonances was not marked. Even after 60 h incubation the total loss of Aβ signals was ~35%. This is attributed to gradual precipitation/fibrillization of some Aβ. However, the histidine and methionine resonances were specifically perturbed.

Figure 4a shows ¹H NMR spectra of Aβ42 in the aromatic region, comparing Aβ42 in the absence of CuCl₂ (gray top spectra) and Aβ42 incubated with CuCl₂ (0.05 mol equiv) and ascorbate (10 mol equiv) at various incubation times. The εCH and δCH resonances of all three histidines (His⁶, His¹³, and His¹⁴) gradually lose their intensity; after 60 h the histidine resonances are considerably diminished. There is a shift to low field due to a drop in pH of 0.1 unit. With greater amounts of ascorbate (20 mol equiv) almost complete loss of histidine resonances in the ¹H NMR spectra occurs within ~1 h of the addition of ascorbate to the Cu–Aβ complex. In addition to loss of the histidine signals, two new resonances at 6.25 and 6.29 ppm are observed with CuCl₂ and ascorbate incubation, and an enlargement of this signal is shown as an insert in Figure 4a. This new resonance can be assigned to the δH (C4H) of the 2-oxohistidine species (57). The 2-oxohistidine species results from oxidation of the histidine imidazole side chain to 2-oxoimidazole, with CεH being converted to a carbonyl group, CεO (58). The new signals shown in Figure 4a are most likely a combination

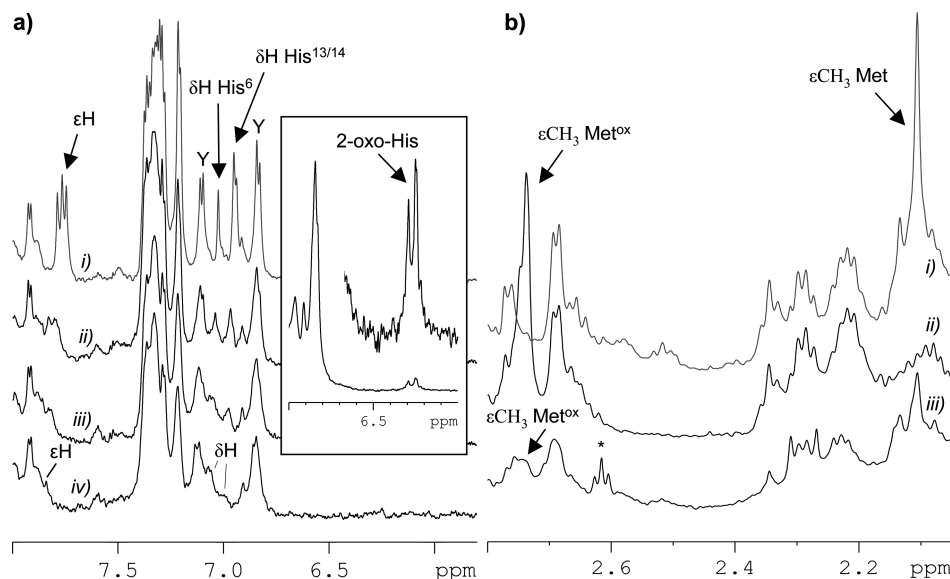


FIGURE 4: 1D proton NMR spectra of A β 42 in the presence of CuCl₂ and ascorbate. (a) Aromatic region: (i) Apo A β 42 (100 μ M) in 10 mM phosphate buffer (pH 7.4) (top gray spectra), (ii–iv) a sample of A β 42 with 5 mM (0.05 mol equiv) CuCl₂ and 1 mM (10 mol equiv) ascorbate in 10 mM phosphate buffer, pH 7.4 at 37 $^{\circ}$ C. Incubation times shown are (ii) \sim 20 h, (iii) \sim 40 h, and (iv) 60 h post addition of CuCl₂ and ascorbate. Insert shows the summed spectra over the complete 60 h incubation at 37 $^{\circ}$ C in the region 7.0–6.0 ppm and a zoomed view showing the 2-oxo-His signal. Y indicates Tyr¹⁰ ring protons. (b) Aliphatic region: (i) Apo A β 42 (100 μ M) in 10 mM phosphate buffer (pH 7.4) at 37 $^{\circ}$ C (top gray spectra), (ii) A β 42 plus H₂O₂ (2.5 mM) after 60 h incubation at 37 $^{\circ}$ C, and (iii) A β 42 plus 5 mM (0.05 mol equiv) CuCl₂ and 1 mM (10 mol equiv) ascorbate after 60 h at 37 $^{\circ}$ C. * indicates ascorbate oxidation product.

of the three histidines (His⁶, His¹³, and His¹⁴) in their oxidized form. Although the loss of signal for the unoxidized histidine is almost complete, the intensity of this 2-oxo-His signal is considerably less than that of other resonances for A β , \sim 10% the intensity of, for example, the Tyr¹⁰ 2,6 ring protons. It is notable that the Tyr ring protons are largely unperturbed, suggesting that under these conditions dityrosine cross-linking is limited, although higher Cu–A β ratios have been shown to mediate dityrosine cross-linking (59).

The second marked change in the ¹H NMR of A β 42 is shown in part of the aliphatic region (Figure 4b). In the apo (top) spectra, the characteristic singlet from ϵ CH₃ of Met³⁵ in A β is evident at 2.10 ppm. Addition of H₂O₂ (2.5 mM) causes loss of this signal and the appearance of a new singlet peak in the ¹H spectra at 2.74 ppm with similar intensity. This new resonance (and shoulder) can be assigned to the *R*- and *S*-forms of ϵ CH₃ in methionine sulfoxide (Met^{ox}) (C α H–C β H₂–C γ H₂SOC₆H₃) (60), in which an oxygen has been added to the sulfur atom in the methionine side chain. We note that addition of H₂O₂ alone has no effect on the histidine resonances. Furthermore, it is notable that incubations of A β with Cu²⁺ in the absence of ascorbate shows no marked oxidation of His or Met residues.

Incubation with CuCl₂ (0.05 mol equiv) and ascorbate (10 mol equiv) causes a gradual decrease in intensity of the Met³⁵ ϵ CH₃ signal over \sim 60 h. The appearance of a new ϵ CH₃ signal for the Met^{ox} species is slight, and after 60 h a small peak, \sim 10% the intensity of the apo unoxidized Met³⁵, is detected. Although the unoxidized Met³⁵ signal is almost completely lost, this is not accompanied by the appearance of a Met^{ox} signal of equal intensity. In contrast, studies with the A β (25–35) fragment under similar conditions of CuCl₂ and ascorbate showed a significant peak for the Met^{ox} (see Supporting Information Figure S6).

We looked for the appearance of a signal that may correspond to methionine sulfone (C α H–C β H₂–C γ –

H₂SO₂C₆H₃), a product rarely formed in biological systems resulting from further oxidation of methionine sulfoxide (61), but none were apparent. 2-Oxo-His can also be further oxidized to asparagine/aspartic acid, but these products are also not observed. The weak intensities of the 2-oxo-His and Met^{ox} suggest possible oxidation of histidine and methionine to other unidentified species. A β can form oligomeric species whose line width may make them difficult to detect by NMR. However, the intensities and line widths of the rest of the A β spectra remain largely unaffected; even after 60 h incubation, two-thirds of the oxidized A β is detected in the ¹H NMR spectra.

DISCUSSION

Cu–A β Does Not Possess a Prooxidant Activity in a Cell-Free System. Although oxidative injury is an established characteristic of AD (4), the exact role that the A β peptide plays is disputed. Some claim that the neurotoxicity of A β is due to generation of ROS in the presence of the A β –Cu complex (30, 32, 33). While others point to an antioxidant role for A β (35–37, 40), until now however, studies directly measuring ROS in a cell-free system in the presence of A β and Cu²⁺ ions have typically suggested that A β has prooxidant activity (33, 44, 45, 49). It is now clear from our ODAD and peroxidase and 3-CCA ROS detection assays that A β 40 and A β 42 in either their fibrillar or monomeric state do not generate either H₂O₂ or hydroxyl radicals in the presence of Cu²⁺ ions alone. Even in the presence of an additional reducing agent, ascorbate, the peptide has no ability to enhance reactive oxygen species production above and beyond that of control samples. Indeed, levels of detectable hydrogen peroxide are the same or less than controls, and hydroxyl radicals were appreciably diminished by the presence of A β peptide, in the CuCl₂ and ascorbate samples.

Our findings are in complete contrast to *in vitro* studies published by Opazo et al. in which they state that A β 42 with Cu²⁺ generates hydrogen peroxide catalytically by recruitment of a biological reducing agent (e.g., ascorbate) as a substrate (33). Variation between the methodology used by Opazo et al. and that described here includes the presence of EDTA to bind free Cu²⁺ during incubations. For this reason, additional sample incubations containing EDTA were performed using the hydrogen peroxide assay; however, we still did not observe any catalytic production of hydrogen peroxide by A β plus Cu²⁺ and ascorbate. The amount of H₂O₂ described to be generated by A β and Cu²⁺ was small in the Opazo et al. study, but no control for Cu²⁺ with ascorbate and EDTA was reported. In our study the level of H₂O₂ recorded for the control of Cu²⁺ and ascorbate with EDTA was comparable to A β 42 with Cu²⁺, ascorbate, and EDTA; both were very low and within the noise level of the study. The Opazo et al. study used Cu(Gly)₂ as the source of Cu²⁺ ions, and it has been suggested that more hydroxyl radicals are generated by Cu(Gly)₂ and ascorbate with the addition of A β (62); however, we have found no such difference.

A β Acts as a Sacrificial Antioxidant by Quenching Hydroxyl Radicals. The antioxidant-like activity of the A β peptide can be directly correlated to observed metal catalyzed oxidation of A β . The loss of ¹H NMR signals for unoxidized histidine and methionine is almost complete. 2-Oxo-His and Met^{ox} ¹H NMR resonances are also observed, although they only account for about ~10% of the total A β detected. Methionine and histidine residues are particularly prone to metal-catalyzed oxidation (63), and 2-oxo-His is considered a useful biological marker under conditions of oxidative stress (64). Pronounced oxidation of A β has been observed in plaques with the Met³⁵ residue oxidized to Met sulfoxide (5, 6). Other studies by Schoneich et al., using HPLC-MS/MS, have shown that His¹³ and His¹⁴ in the A β sequence are targeted first for oxidation and converted to 2-oxo-His (65).

These results suggest that the antioxidant activity of A β is mediated by a type of free radical scavenging, rather than through metal chelation into a redox-inactive state. We have shown that the hydroxyl radicals generated by Fenton–Haber Weiss reactions of CuCl₂ and ascorbate are largely (80–90%) quenched by A β . The highly reactive hydroxyl radicals are very short-lived, and since copper is bound to A β , the radical species are produced in close proximity to the peptide, readily oxidizing A β .

Similar behavior is observed with the Cu²⁺ transport protein serum albumin, and this is believed to be the mechanism by which albumin acts as a principle antioxidant within blood plasma, accounting for 70% of the free radical trapping activity of serum (66). In addition, similar behavior is observed for Cu²⁺ bound to the prion protein and suggests that the prion protein functions as an antioxidant at the synaptic cleft during release of Cu²⁺ ions (52).

It has been proposed that proteins with surface-exposed methionine residues can act as antioxidants. In conjunction with the repair enzyme methionine sulfoxide reductase (Msr) the oxidation/reduction of methionines can function as a sink for ROS (67). Importantly, studies of Msr activity in the brains of AD patients revealed that the enzyme's activity was reduced, compared to control subjects (68). The methionine residue within A β regulates oxidation products;

specifically, histidine residues in the peptide were only oxidized in the absence of the methionine (69).

Cu⁺ Generation. Our measurements of Cu²⁺ reduction capability by assorted A β peptides shows variation depending on the length of the A β fragment used. In particular, the smaller, more soluble copper binding fragment (A β 28) showed no ability to reduce Cu²⁺ to Cu⁺. Additionally, no Cu²⁺ reduction was observed when the fragments A β (25–35) and/or A β 16 were coincubated with CuCl₂. These results are consistent with previous reports (44, 45, 70). An early study suggested Cu⁺ generation by A β (25–35) (47); however, this observation has never been reproduced.

In contrast, we show that both monomeric and fibrillar full-length A β peptides (A β 40 and A β 42) possess some intrinsic Cu²⁺ reduction capability, in agreement with Huang et al. (44, 45). Although our Cu⁺ assay results did not show a distinction in reduction ability between the two full-length peptides or the fibrillar state, Cu²⁺ reduction by A β 40/42 may involve the Met³⁵ residue and hydrophobic tail not present in A β 28.

Although full-length A β (monomeric or fibrillar) is capable of reducing small amounts of Cu²⁺, this is not reflected by production of hydrogen peroxide or hydroxyl radicals. The lack of H₂O₂ and hydroxyl radicals indicates that the Cu²⁺ reduction is not significant in terms of ROS generation. The reducing capacity of A β seems to require the presence of the Cu⁺ chelator BC to drive the reaction by chelating the Cu⁺ generated. Furthermore, our ¹H NMR studies with Cu–A β 42 suggest that no reactive oxygen species are generated in the absence of ascorbate.

In Support of an Antioxidant Role for A β . The studies shown here indicate that both the monomeric and fibrillar A β peptide act as an antioxidant by reaction with (quenching sacrificially) reactive oxygen species to prevent their harmful effects elsewhere, and this is supported by a range of studies. There is evidence that A β can act as an antioxidant for lipoproteins in both CSF and plasma (71). Other studies have also demonstrated A β to possess antioxidant function, with Zou et al. reporting on the ability of monomeric A β 40 and A β 42 to inhibit neuronal death (37). These monomeric species were shown to mimic the role of antioxidant scavengers, such as catalase and vitamin E, under the same conditions, preventing neuronal death caused by transition metal-induced oxidative damage. This neuroprotective activity of A β was shown to be lost on formation of oligomeric and aggregated forms of the peptide. Kontush et al. suggested that only at low concentrations (0.1–1.0 nM) did A β inhibit autooxidation of CSF lipoproteins and plasma low density lipoprotein (LDL) (71). Similarly, an earlier study by Chan et al. has reported an antiapoptotic action only at low A β concentrations (40).

A number of studies suggest A β is upregulated as a response to oxidative stress (72). It has been shown that β - and γ -secretases (BACE1 and presenilins) that cleave A β from its precursor protein (APP) are upregulated as a response to oxidative stress (73–76). A β levels are therefore possibly increased as an antioxidant defense mechanism. Studies on human neuroblastoma cells showed increased levels of A β , through enhancement of the amyloidogenic pathway of APP, in response to the addition of hydrogen peroxide (77). Increased production of A β has also been identified in response to injuries such as head trauma (78–80).

A β 28 has been shown to have neurotrophic activity, by enhancing neuronal survival (81). An inverse correlation between amyloid deposits and 8-OHG (8-hydroxyguanosine) levels, a marker for oxidative damage, also suggests that A β may be produced in a compensatory response to increased oxidative stress, with oxidative damage actually reducing with senile plaque formation (82, 83). The neuroprotective nature of A β is highlighted with a negative correlation between amyloid deposits and observed oxidative damage, neuronal loss, and cognitive impairment (84–86). Similar inverse correlation between A β deposits and oxidative damage was detected in patients with Down's syndrome. Interestingly, A β deposition follows rather than precedes increased 8-OHG levels, and strikingly 8-OHG levels decline to control levels after plaques form (87).

ROS-Induced Toxicity Mediated by Cu–A β Is Not Ruled Out. It is important to note that a so-called prooxidant role for A β is not ruled out by our observations. ROS-mediated cellular toxicity of A β has been demonstrated by a number of groups (14). However, we have clearly shown that A β does not generate any more ROS than free Cu²⁺ ions or weakly chelated Cu(Gly)₂ irrespective of the concentration of Cu²⁺ ions or the monomeric/fibrillar state of A β . However, this does not preclude an oligomeric diffusible form of A β binding Cu²⁺ ion and then trafficking and concentrating the Cu²⁺ ions in a redox-active state to the neuronal cell surface.

Thus the conflicting observations that A β can promote lipid peroxidation (31) while others suggest A β acts as a neuronal antioxidant (35–37) may well reflect the form of A β . This is supported by the concentration dependence of these studies, where they only exhibit antioxidant properties at low (monomeric) levels of A β (40, 71). The hydrophobic nature of A β peptide means it has a strong affinity for the cell membrane, but it is likely that monomeric A β will interact with the cell membrane differently from oligomeric or fibrillar forms of A β . Monomeric A β may protect neurons by binding potentially toxic Cu²⁺ ions released at the synapse and so quench the hydroxyl radicals, while oligomeric A β could concentrate the redox-active form of copper at the membrane and so cause lipid peroxidation and eventually cell death.

The prooxidant properties of Cu–A β are clearly not due to the ability of A β to catalytically generate H₂O₂ and/or hydroxyl radicals. Rather it is A β 's ability to chelate Cu²⁺ in a redox-active form and then concentrate Cu²⁺ ions at the membrane surface.

CONCLUSION

The amyloid cascade hypothesis indicates that in cases of familial AD it is the upregulation of A β 42 due to mutations in APP or the presenilins that causes the cascade of events leading to dementia (11). We suggest that, for sporadic AD, A β is also upregulated as a response to oxidative stress. Interestingly, oxidative stress increases with age and is linked to head trauma and the inflammatory response (78, 79), both of which are risk factors associated with sporadic AD. Furthermore, Cu²⁺ levels (a major source of ROS) in the brain increase with age, and Cu²⁺ homeostasis is compromised in AD patients (16, 88).

Our studies, in a cell-free system, show no evidence that A β acts as a prooxidant in a Cu²⁺/ascorbate system; to the

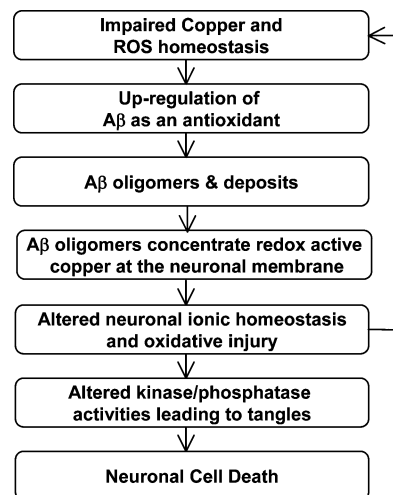


FIGURE 5: Modified amyloid cascade hypothesis. Sporadic AD is caused by upregulation of A β as a response to impaired copper and ROS homeostasis, both of which are linked with aging. Failure to clear increased levels of A β causes oligomerization and diffuse plaque formation. Diffusible oligomers could concentrate Cu²⁺ in a toxic redox-active state at the membrane. This in turn causes further oxidative stress and upregulation of A β . Further A β accumulation causes altered kinase and phosphatase activities leading to neurofibrillar tangles of tau protein and dementia.

contrary, we observe the opposite with A β exhibiting antioxidant-like properties. Thus rather than causing the oxidative stress observed in the early stages of sporadic AD (7, 8), the upregulation of A β (73) could be a protective response to it; initially, A β actually reduces oxidative stress (37). However, in the long term, the upregulation of A β then causes the neurotoxic effects of A β oligomers and the accumulation of amyloid plaques. A β oligomers could act to concentrate Cu²⁺ ions at the membrane surface and allow ROS generation (31), leading to neuronal cell death, which causes further marked oxidative stress and the further upregulation of A β as shown in Figure 5.

Thus the age of onset of sporadic AD may reflect an individual's ability to maintain ROS homeostasis within the brain. Antioxidant therapeutics may therefore be the most effective treatment of sporadic AD (which accounts for 85% of AD cases) and could delay or prevent the onset of sporadic AD. Therapeutics that attempt to solubilize plaques may actually release the more neurotoxic oligomeric form of A β and so may be counterproductive. Thus far, antioxidant therapies such as vitamin E have shown only limited efficacy (89).

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SUPPORTING INFORMATION AVAILABLE

Six figures showing (1) confirmation of fibril formation using a variety of methods, (2) ODAD and peroxidase, hydrogen peroxide detection assay calibration, (3) hydrogen peroxide production with higher Cu²⁺ concentrations, using

CuGly₂, (4) hydroxyl radical detection using Cu²⁺ ions complexed with glycine, (5) 1D proton NMR spectra of A β 42 in the presence of CuCl₂, and (6) 1D proton NMR spectra of A β 25–35 in the presence of CuCl₂ and ascorbate. This material is available free of charge via the Internet at <http://pubs.acs.org>.

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