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Novel drug targets based on metallobiology of Alzheimer's disease

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Abstract

Importance of the field—Increased localization of Zn, Fe, Cu and Al within the senile plaques (SP) exacerbates amyloid beta ($A\beta$)-mediated oxidative damage, and acts as catalyst for $A\beta$ aggregation in Alzheimer's disease (AD). Thus, disruption of aberrant metal-peptide interactions via chelation therapy holds considerable promise as a rational therapeutic strategy against Alzheimer's amyloid pathogenesis.

Areas covered in this review—The complexities of metal-induced genesis of SP are reviewed. The recent advances in the molecular mechanism of action of metal chelating agents are discussed with critical assessment of their potential to become drugs.

What the reader will gain—Taking into consideration the interaction of metals with the metalresponsive elements on the Alzheimer's amyloid precursor protein (APP), readers will gain understanding of several points to bear in mind when developing a screening campaign for ADtherapeutics.

Take home message—A functional iron-responsive element (IRE) RNA stem loop in the 5' untranslated region (UTR) of the APP transcript regulates neural APP translation. Desferrioxamine, clioquinol, tetrathiolmolybdate, dimercaptopropanol, VK-28 and natural antioxidants, such as curcumin and ginko biloba need critical evaluation as AD therapeutics. There is a necessity for novel screens (related to metallobiology) to identify therapeutics effective in AD.

Declaration of interest

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APP; A β ; iron responsive element; metal chelators; screening

1. Introduction

The two predominant pathological features of Alzheimer's disease (AD) are the extracellular amyloid plaques and intracellular neurofibrillary tangles in the brain. The abnormal processing of the amyloid precursor protein (APP) is the initiating event in AD pathogenesis, subsequently causing aggregation of amyloid beta (A β), specifically A β 42. Formation of neuritic plaques instigates the formation of neurofibrillary tangles, composed of hyperphosphorylated microtubule-associated tau protein, and results in significant loss of neurons and synapses leading to cognitive impairment and dementia [1–4].

One of the physiologically relevant environmental factors able to affect the conformation of amyloidogenic proteins/peptides is metal ions. The 'metal hypothesis' of AD proposes that the interaction of APP and its proteolytic product 'AB' with specific metals drives ABpathogenicity [5]. Increased concentrations of Fe, Cu and Zn [6] have been observed within the amyloid plaques (Figure 1) [7] that are released from glutamergic neurons at synapses [5]. The Fe and Zn levels have been reported to reach as high as 1 mM in the vicinity of amyloid plaques [8]. When compared with the surrounding tissue, levels of Zn, Cu, Fe and Ca inside the plaque were higher by 9.09 ± 0.20 -fold, 11.72 ± 1.24 -fold, 5.74 ± 3.31 -fold and 98.40 ± 31.56-fold respectively [9]. Proton-induced X-ray emission, epifluorescence microscopy, immersion autometallography [10] and synchrotron X-ray fluorescence microprobes confirmed 'hot spots' of Fe, Cu, and Zn co-localized with AB in the rims and cores of the senile plaques [6,11,12]. Fe-rich A β plaques could be seen in AD-mouse brains in vivo through MRI and susceptibility-weighted MRI with significant increases in the basal ganglia cortex [13,14]. Furthermore, laser capture micro-dissection coupled with X-ray fluorescence microscopy could determine elemental profiles in Aß amyloid plaques (Figure 1) [7].

Metals play a key role in the aggregation of hyperphosphorylated tau into insoluble paired helical and straight filaments, which are involved in the pathogenesis of AD. Fe and Al were reported to accumulate in neurons within neurofibrillary tangles (NFTs) of AD brain [15], and, notably, tau pathology with hippocampal neurons was exacerbated in copper-exposed triple transgenic (Tg)-AD mice [16].

Thus chronic exposure to metal(s) (mainly Cu, Al, Fe and Zn) accelerates amyloid and tau pathology in AD, and trapping of these metals might be necessary to protect the brain from susceptible degeneration during AD.

2. Metals and AD

2.1 Fe-APP, Aβ and AD

A unique CAGA box, 'amyloid' (+83/+86), present only in the APP gene from amyloid plaque-forming species and absent in genes of APP-like-proteins (APLP1 and APLP2),

plays a key role in APP gene regulation [17]. The functional iron responsive element (IRE)-Type II around the amyloid CAGA in the 146 nt-5'-untranslated region (UTR) of APP mRNA (+ 51 to + 94 from the 5'-cap site) [18] binds to the iron regulatory proteins (IRP1 and IRP2) and controls mRNA translation rate [8]. In normal human brain extracts, the IRP is detected as a double IRE-IRP complex but in two of six AD brain extracts a single IRE-IRP complex with decreased mobility was observed. Alterations in the IRP-IRE interaction possibly through elevated endogenous RNase activity could be the site in the AD brains at which Fe mismanagement occurs and the Fe regulatory system becomes dysfunctional [19].

Fe enhances amyloidogenicity through different mechanisms. Firstly, Fe downregulates furin protein levels that promote non-amyloidogenic α -secretase activity [20]. Secondly or possibly most commonly Fe participates in the production of the reactive oxygen species (ROS) (Figure 2) that upregulate inhibition of matrix metalloprotease and shift aconitase to the IRP1 form.

When Fe is scarce, the IRP1 binds to IREs that regulates translation or stability of IREmRNAs (Figure 3), whereas when Fe is abundant IRP1 forms a [4Fe–4S] cluster and is converted from an RNA-binding form to an aconitase [21]. Unlike IRP1, IRP2 is degraded by the proteasome by a process involving Fe-catalyzed oxidation. IRP-2 colocalizes with redox-active Fe, and change in IRP-2 has been reported to be directly linked to impaired Fe homeostasis in AD [22]. Such changes could modulate APP mRNA translation and APP synthesis in astrocytes [23] and neurons to significantly reduce expression of the neuroprotective secreted APP (APP(s)) [24]. In terms of AD pathology, the APP–IRE–RNA secondary structure might be disrupted in the presence of an adjacent 5' UTR-specific single nucleotide polymorphism that could be genetically linked to increased risk for spontaneous AD [25].

A changed IRP-IRE binding modulates expression of the Fe-storage protein (Figure 3), transferrin and melanotransferrin in brain white matter [26–28] in a way that parallels the pathological lesions in AD [29,30]. Furthermore, genetic evidence implicates synergy between the C282Y allele of the hemochromatosis gene (*HFE*) and the C2 allele of transferrin as risk factors for developing AD [31]. It has also been reported that although each of the HFE variants alone had relatively little effect on Fe status, the combination of either *HFE* C282Y and *HFE* H63D or of *HFE* C282Y and transferrin C2 markedly raised transferrin saturation in those without dementia, but had little effect in those with mature AD [32].

The physiological relevance of Fe to A β is demonstrated by their concurrent accumulation in A β deposits in AD [29,33] as well as in transgenic mouse models [34]. Fe-binding to A β enhances its aggregation and facilitates the oxidative damage in the immediate vicinity of the senile plaques [35–37]. Dysfunction of ferritin with ferroxidase activity [38], particularly abundant in myelinated axons and oligodendrocyte processes, results in an increase of toxic brain Fe²⁺ ions. In support of this, null ferritin heterozygous mutants mice of mixed C57BL6/J × 129SvEv genetic background have also been found to develop oxidative features in the cortex that are reminiscent of AD and Parkinson' disease (PD) [39].

In AD pathobiology, intracellular A β forms a complex with free heme that results in functional heme deficiency [40]. Regulation of cerebral heme biosynthesis is profoundly altered in AD and may contribute towards disease pathogenesis by affecting cell metabolism as well as Fe homeostasis [41]. A significant decrease in loosely bound Fe has been reported in the hippocampal white matter of mild-moderate and severe AD patients with a trend towards increased non-heme iron in the hippocampal gray matter of severe AD [42]. The AD brain has been observed to have 2.5-fold more heme-b and 26% less heme-a compared with controls, with a significant 2.9-fold decrease in the heme-a:heme-b ratio [43]. When the rate of production of A β exceeds the capacity of heme synthesis or there is an increase in free metals, A β reacts with free metals and forms aggregates [43]. The A β -heme complex, being a peroxidase, catalyzes the oxidation of serotonin and 3, 4-dihydroxyphenylalanine by H_2O_2 that leads to oxidative damage to macromolecules and depletes specific neurotransmitters. The intracellular oxidative stress-induction of the astroglial and neuronal hemoxygenase-1 gene through a 'common pathway' leads to pathological brain Fe²⁺ deposition. This further causes enhanced free radical generation [44], mitochondrial insufficiency, enhanced cytochrome c oxidase activity [45] and H₂O₂-generation that releases free radical hydroxyl (OH) via the Fenton reaction: $Fe^{+2} + H_2O_2 \rightarrow Fe^{+3} + OH^- + OH^-$ OH or by virtue of hypervalent iron compounds, as observed in vitro [46,47]. Oxidative stress triggers activation and/or translocation of NF-kB, p53 and c-Jun transcription factors resulting in enhanced apoptosis [48]. This is accompanied by DNA damage, blood-brain barrier (BBB) disruption [49] and age-related myelin breakdown [50,51]. The toxicity of Aß is mediated, at least in part via redox active Fe that precipitates lipid peroxidation and cellular oxidative stress [52].

2.2 Zn and AD

In AD, recurrent episodes of focal high Zn release are observed from the presynaptic vesicles to the postsynaptic neurons of the neocortex, hippocampus and amygdale in a Ca²⁺- and depolarization-dependent fashion [53]. This indicates prior injury with toxic 'floods' of free Zn in the brain as a major risk factor in AD-development [54]. A decreased concentration of astrocytic growth inhibitory factor that chelates Zn [55] causes the highest concentration of Zn in the hippocampus. This is associated with increased extracellular Zn metalloproteinase activities [56] due to cholinergic deafferentation of the hippocampus [57]. The sequestration of Zn in A β -Zn complexes further leads to reduced Zn availability at synaptic terminals and consequent loss of Zn modulatory activity at excitatory synapses [58].

The Zn transporter proteins (ZNTs, ZnT2 - 8), abundantly expressed in A β plaque, and cerebral amyloid angiopathic vesicles [59,60] induce an abnormality in the uptake or distribution of Zn in the AD brain [61]. The ZnT immunoreactions were detected in the amyloid plaques and amyloid angiopathic vessels of brains of APPswe/PS1dE9 transgenic mice. ZnT1 and ZnT4 are extensively expressed in all parts of the plaques. ZnT3, ZnT5, and ZnT6 are expressed most prominently in the degenerating neurites in the peripheral part of the plaques, while ZnT7 is present in the core of the plaques. The amyloid angiopathic vessels showed a strong ZnT3 immunoreactivity [59]. The disruption in neuronal Zn homeostasis triggers ZnT1 expression that caused increased efflux of Zn ions in the

extracellular space. This might involve altered homeostasis of metallothionein and other Znbinding proteins such as a 2 macroglobulin, or pro-inflammatory cytokine polymorphism [62]. ZnT2 – ZnT8 are localized in the intracellular membranes and transport Zn ions into different intracellular compartments when the intercellular level of Zn ions is raised [63]. The synaptic ZnT3 activity promotes cerebral amyloid angiopathy by indirectly increasing exchangeable Zn concentrations in the perivascular spaces of the brain [59,64,65]. The increase in synaptic Zn correlates well with higher levels of insoluble A β and plaque loads in aging females, and these sex differences completely disappear in ZnT3^{-/-} mice, suggesting the role for synaptic Zn in the sex differences in AD [65].

Zn interaction may play an important, evolutionary conserved role in APP function and metabolism. The sequence of APP harboring the Zn binding site (181 – 200 amino acids) is evolutionarily conserved with the coordination sites as Glu-185, Cys-188 and Cys-189, and a saturation binding κ_A of 750 nmol/l [66]. The obligatory motif for the ectodomain Zn binding region on APP is a novel sequence for a Zn binding site, GVEFVCCP that is highly conserved in APLP 1 and APLP2, as well as in the *Drosophila* and *Caenorhabditis elegans* APP-like protein [67]. The Zn binding domain may play a role in regulating the adhesiveness of APP through its control over the Kunitz-type protease inhibitory insert and heparin binding affinity of APP695 [66,68]. Disturbed homeostasis of extracellular Zn²⁺ in AD may interfere with the normal binding of APP to heparin-like molecules such as heparan sulfate moiety of proteoglycans (known to alter protein conformation, and the clearance and processing of bound proteins) [66]. It also modulates the binding of APP to extracellular matrix components such as laminin [69] that controls crucial cell–cell and cell–matrix interactions [70].

Indeed, Zn-binding may also influence APP processing and has been found to specifically inhibit the α -secretase cleavage of APP [71]. Exogenous Zn enhances the synthesis of presenilin in a dose-dependent manner. This further elevates γ -secretase activity that could result in increased production of A β and the formation of more senile plaques, which in turn could trap more Zn [72]. The Zn-dependent transcription factors NF- κ B and specific protein-1 (Sp1) bind to the promoter region of the APP gene, and also inhibit enzymes that degrade APP to non-amyloidogenic peptides and degrade the soluble form of A β [73]. A reduction in Zn-stimulated protein tyrosine kinase activities in AD hippocampus indicates a possible connection of neuronal protein tyrosine kinase activity loss to severe memory and intellectual impairment that is characteristic of AD [74].

Human A β specifically and saturably manifests high-affinity ($\kappa_a = 107$ nM) binding [75] (concomitant with Zn-induced A β aggregation) with Zn in less than a millisecond [76] at a Zn²⁺: A β stoichiometry of 1:1 [77], and induces tinctorial amyloid formation [61]. A low-affinity binding with the second Zn ion at 2:1 stoichiometry seems to have a moderate effect on peptide conformation [77]. The 1 – 16 region represents a possible initiation site for the entire A β transconformation, and the minimal A β fragment required for Zn binding [77]. The Zn is coordinated i) at the N-terminal hydrophilic region of A β with the imidazole side chain of His6, His13, His14 [78–80], ii) with Asp1 either with the N-terminus or/and the carboxylate group and iii) with the carboxylate side chain of Glu11 [80–82]. The dissociation constant (K_d) of Zn for the fragment A β -peptide 1 – 28 (measured by

fluorescence study) and that for $A\beta$ -peptide 1 – 40 (through NMR) had values in the μ M range at pH 7.2 and 286 K. Zn also has a second, weaker binding site involving residues between 23 and 28 [79]. Moreover, the apparent dissociation constant ($K_{d, app}$) of Zn binding to all forms of A β (soluble A β , A β fibrils, or Zn-induced A β aggregates) is in the low μ M range (1 – 20 μ M) [80,83]. Isomerization of Asp7 or the substitution of Asp7 with Asn (in Tottori–Japan mutation with higher susceptibility of spontaneous conversion to iso-aspartate) in A β promotes Zn-induced oligomerization of the A β –Zn binding domain [84]. The Zn-A β binding prevents the formation of the typical amyloid fibrils, the inhibitory effect (IC5₀ =1.8 μ M) being three times stronger than that for Cu(II). It further induces the accumulation of large unorganized aggregates of smaller non-fibrillar pathogenic forms of A β [76,85]. The promotion of A β 28 aggregation by Zn is based on the transformation of the partially α -helical conformer (intermediate) towards the A β -sheet amyloid [80] structure by destabilization of the α -helix in the intermediate [86].

With regard to the tau-protein-mediated AD pathology, the Zn-binding protein S100 β has been identified as an interacting partner with tau. S100 β -tau binding, promoted by Zn, may represent a key pathway for neurite development, possibly through S100 β modulation of tau phosphorylation and/or functional stabilization of microtubules and process formation [87]. Interestingly, S100 β -tau interaction may be disrupted by hyperphosphorylation and/or imbalances in Zn metabolism and this may contribute to the neurite dystrophy associated with AD. Zn has also been reported to enhance tau binding as an important factor in the internalization of S100 β [87].

2.3 Cu and AD

The possibility that Cu may contribute to AD pathology is suggested by the ceruloplasmin fragmentation that indicates improper Cu transport, together with the 'free' Cu rise in AD [88]. Cu^{2+} levels in the brain increase with age in transgenic Tg2576 mice [89], and Cu along with ROS homeostasis are compromised in AD patients [6,89–93]. Even a small increase in the serum free Cu can be of significance, particularly over a long period of time [94]. The free Cu can cross the BBB in living patients and supply the brain with a continuous flux of noxious redox Cu. Cu²⁺ levels in the AD neuropil are 400% higher than in the neuropil of a healthy brain [6], and oxidative damage [95] involving Fenton cycling is the probable source of ROS [96].

Neurotoxicity of A β with Cu-induced dityrosine crosslinking of A β 1 – 28, A β 1 – 40, and A β 1 – 42 [97] has been linked to H₂O₂ production [98], with the generation of highly toxic hydroxyl radical species [99]. The oxidative coupling is initiated by interaction of H₂O₂ with a Cu²⁺ tyrosinate that induces A β aggregation under mildly acidic conditions (e.g., pH 6.8 – 7.0) [100].

Hyperlipidemia is a significant risk factor of an interaction between free copper and A β [94,101], and Cu alters the structure of lipid rafts through flotillin-2 lipid raft association that inhibits APP endocytosis. The primary products of lipid peroxidation are phospholipid hydroperoxides that are degraded by free Cu²⁺ in the presence of ascorbic acid to yield hydroxy-2-nonenal, a toxic factor in the pathogenesis of AD [102]. It has also been seen that total cellular Cu²⁺ is associated inversely with lipid raft Cu²⁺ levels, so that under

intracellular Cu²⁺ deficiency A β –Cu complexes are more likely to form [103]. Diffusible A β oligomers concentrate Cu²⁺ in a toxic redox-active state at the membrane that in turn causes further oxidative stress and upregulation of A β . The A β accumulation causes altered kinase and phosphatase activities leading to neurofibrillar tangles of tau protein and dementia [104].

APP has many of the features of a Cu transporter, and the Cu²⁺ binding site at residues 135 – 155 at a $K_{d APP}$ of approximately 10 nM can promote the reduction of bound Cu²⁺ leading to increased oxidative stress in neurons [105,106]. The APP-Cu binding domain consists of an α -helix (residues 147 – 159) packed against a triple-stranded β -sheet (residues 133 – 139, 162 – 167 and 181 – 188) through Cy- rich disulfide bonds [107]. The metal ligands are His147, His151, Tyr168 and two water molecules that are arranged in a square pyramidal geometry [108]. There is a clear link between APP processing and Cu²⁺ homoeostasis in the brain with the 24-residue peptide of the C-terminal domain of β -secretase 1 (BACE1) binding with high affinity to a single Cu¹⁺ atom through Cys residues. Overproduction of BACE1 reduces superoxide dismutase (SOD)1 activity in cells through the interaction of cytoplasmic domain of APP with the Cu²⁺ chaperone for SOD 1 [109]. Perturbations to APP metabolism and in particular, its secretion or release from neurons may alter Cu homeostasis [110] resulting in increased A β accumulation and free radical generation [111].

It is known that monomeric A β peptides bind Cu(II) ions *in vitro* [112], with concomitant acceleration of A β aggregation and precipitation [100,113,114]. The oxidative damage to A β amino acid side chains was profound in tests performed *in vitro*, and some of the modified chains, for example, 2-oxo-His, were also found in amyloid plaques [115].

Cu may participate in oxidative stress through redoxcycling between its + 2 and + 1oxidation states to generate ROS that is governed by the binding mode. All of the ligands of the high affinity Cu(II) site are contained in the A β 16 N-terminal domain [112]. Two complexes (components I, at lower pH and component II, at higher pH), distinguishable by conventional 9 GHz electron paramagnetic resonance (EPR) spectroscopy are present near physiological pH values. Using pulse EPR techniques such as electron spin echo envelope modulation and hyperfine sublevel correlation spectroscopy (HYSCORE) Shin and Saxena proposed three His bound at pH 7.4 [116]. On the other hand Drew et al. proposed the binding of two His and Asp1 by A β 16 N terminus and side chain carboxylate group at pH 6.3 – 6.9, and of three His and the carbonyl group of Ala2 at pH 8 [117,118]. Dorlet et al. further used a wide range of advanced EPR techniques in conjunction with specific isotopic labeling that were able to directly detect and thus identify the ligands of the Cu (II) ion coordinated to $A\beta 16$ in the component I and component II forms. They proposed that in component I, the two His, the NH2 terminus, and the carbonyl group from Asp1 composed the equatorial coordination plane, while the side-chain carboxylate group of Asp1 occupied an axial position. In component II, the equatorial ligands were the NH2 terminus, the amide and carbonyl groups of Ala2 (due to deprotonation of the amide nitrogen atom of A2 upon pH increase), and one His. The side-chain carboxylate group of Asp1 was in an apical position [119]. The component I of Dorlet et al. [119] was in close agreement with Drew et al. [117,118] but component II differed significantly.

X-ray absorption (XAS) and EPR spectroscopy revealed that Cu(I) is ligated by the imidazole group of His13 and His14 in a linear coordination environment in A β and Cumediated oxidative damage of A β occurs over multiple redox cycles [120,121]. X-ray absorption fine structure spectroscopy and CO-binding studies demonstrated the preference of Cu I ions for two-coordinate geometry in binding to A β through a contiguous His13 – His14 motif [122]. Further, NMR studies revealed that structure is retained, even in the presence of three His residues (His6, His13, His14) [121] and additional potential donors (Tyr10, Asp7, Glu11, Ser8, backbone carbonyl O, amide N) [122].

The Cu interacts with A β to form simple reversible 1:1 complex [123], and at times a 2:1 complex [124]. At the Cu²⁺/peptide molar ratios > 0.3 A β coordinates a second Cu²⁺ atom in a highly cooperative manner [125]. But EPR spectroscopy indicated that both Cu have axial, Type II coordination geometry, square-planar or square-pyramidal, with nitrogen and oxygen ligands [124]. The EPR parameters are consistent with a Type 2 Cu²⁺ center with three nitrogen donor atoms and one oxygen donor atom in the coordination sphere of Cu²⁺, and this coordination in retained during organization of A β monomers into fibrils [126].

The rate constant for the reaction of superoxide with $Cu^{2+}-A\beta$ has been found to be much slower than that with SOD, and the His residues of $A\beta \ 1 - 42$ control the redox activity of transition metals present in senile plaques [127]. The addition of Cu^{2+} to $A\beta$ in a negatively charged lipid environment caused a conformational change from β -sheet to α -helix, accompanied by peptide oligomerization and membrane penetration [128]. Bufferindependent conditional K_d for Cu(II)- $\alpha\beta_{40}$ complex at pH 7.4 is equal to 0.035 µmol/1 [129]. The dissociation constant of the Cu($\alpha\beta$) complexes ranges from µM to pM values [113], with a preference for the region between 100 pM to 1 nM [79,80].

The modulation of oxidative stress related to Cu-dysfunction may also be one of the mechanisms that make apolipoprotein E4 gene a risk factor for AD [130–132].

2.4 AI and AD

High consumption of Al from drinking water has been epidemiologically reported to be a risk factor for AD [133–135]. Al neurotoxicity as a factor for AD onset or AD-like pathology has been observed in Al-dust exposed workers [136]. Although the role of Al in the etiology of AD remains controversial, energy-dispersive X-ray spectroscopy combined with transmission electron microscopy has detected Al colocalized with A β peptides in the cores of SP located in the hippocampus and the temporal lobe that facilitates iron-mediated oxidative reactions [137]. A β peptide (1 – 42) and Al have been found to induce helical transitions in supercoiled DNA, as a first step to AD neuropathology [138]. It further disturbs heme metabolism [139], perturbs neuronal [Ca²⁺]i homeostasis and mitochondrial respiration with enhanced A β accumulation and neurodegenerative damage [140].

A study in Tg2576 mice indicated that the Al-dietary supplementation increased soluble and insoluble A β levels and the levels of an isoprostane marker of oxidative stress in the hippocampus and iso-cortex [141]. This situation was associated with increased total number of proliferating neuronal cells in the dentate gyrus of hippocampus compatible with an accelerated neurodegeneration [142] and significant Al deposit in the cortex [143]. Al has

also been suggested to interact with plasmin proteases involved in the degradation of $A\beta$ and promotion of α -cleavage of the APP [144].

The NFTs observed in human neurons always developed in conjunction with cytoplasmic Al, suggesting that Al played an important role in their formation [145]. Al in brain bulk was observed to be colocalized with NFTs [146] probably as aluminosilicates [147]. In rat and mouse neurons Al was found to be accompanied by transition of both phosphorylated and unphosphorylated form of NFT peptides from α -helix to β -pleated sheet [148,149]. The NFTs were prone to dissolution by desferrioxamine (DFO) or EDTA [150].

3. Metal chelators as therapeutic agents

Disruption of aberrant metal–peptide interactions via metal chelation could play a crucial role in overcoming Alzheimer's amyloid pathogenesis. The strategy of using chelators to block intracellular APP expression, A β fibrilization and A β -dependent metal oxidative stress-induced neurotoxicity could well produce a major new therapeutic effect on AD progression.

3.1 5-chloro-7-iodo-8-hydroxyquinoline (CQ)

CQ (Figure 4A) has selective affinities for Cu and Zn and form relatively stable complexes [151]. Being hydrophobic CQ molecule crosses the BBB and inhibits homeostatic defects of brain metal ion metabolism in APP transgenics it redistributes Cu from plaques (as observed through NMR spectroscopy) to the cells, disaggregates the metal-ion-induced aggregates of Ap1 – 40 through metal chelation, retards the fibril growth along with Zn^{2+} [152] and dissolves A β with about 30 times greater efficiency and speed [153] than A β vaccine therapy [154]. CQ might work by a combined action that facilitates disaggregation of the Zn-mediated A β collections [100,155], while also inhibiting Cu- or Fe-mediated H₂O₂ production with increased A β clearance [96,156]. A paradoxical increase in Cu and Zn in CQ-treated APP2576 mice might be explained by CQ preventing Cu²⁺ and Zn²⁺ from sequestering with extracellular A β , and then diverting metal ions for uptake into metal-ion-deficient brain tissue [157]. The resolubilized A β may either be removed into the blood or degraded by intracellular uptake and hydrolysis. As observed in *Caenorhabditis elegans* and in mice, CQ chelates mitochondrial enzyme clock abnormal protein 1 that results in slowing down of the aging process in AD [158].

CQ has been shown to reduce APP translation and A β production *in vitro* with reduced APP-5'UTR levels in APP Tg2576 [159]. Oral treatment with CQ caused 49% reduction in cerebral A β deposition and approximately 30 times greater reduction in absolute A β in APP 2576 Tg mice with its dissociation (1.45% elevated) into soluble form of A β [156] without affecting APP processing. It also showed a decrease in the amyloid plaque surface area without appreciable effects on weight loss or mean survival, and improved motor activity, alertness and general health. CQ was effective in liberating A β from postmortem brain samples of AD patients and may be the first credible drug candidate based on the amyloid hypothesis of AD [157]. A Phase II double-blind clinical trial on the effects of oral CQ has shown reduced cognitive decline and decreased plasma A β 42 levels in moderate to severe AD patients [160].

In silico techniques that incorporate chelating properties into well-known intercalation compounds have designed new multifunctional agents for application in AD. The combination of main features of thioflavin-T, with strong affinity for fibrillar amyloid proteins, and CQ led to the three compounds 2-(2-hydroxyphenyl) benzoxazole, 2-(2-hydroxyphenyl)-benzothiazole, and 2-(2-aminophenyl)-1H-benzimidazole. They increased lipophilicity and suitability for crossing the BBB, obeyed Lipinski's rules for pharmacokinetic properties, enhanced antioxidant properties and increased affinity for A β fibrils [161].

The second-generation 8-hydroxy quinoline analogue PBT2 is found to be superior to and a safer ionophore than CQ or CQ-PBT1 [162]. It promoted the transport and clearance of Cu, Zn and Fe across cell membranes, and prevented formation of A β aggregates through enhanced soluble A β formation. It also showed increased activities of the matrix metalloproteases such as neprilysin, insulin degrading enzyme and tissue plasminogen activator. PBT2 accelerated the degradation of soluble interstitial A β and reduced AD-like neuropathology and cognitive dysfunction in Tg models of AD [163]. AD patients (78 in number) who were treated with PBT2 (250 mg) in a Phase IIa double-blind, randomized and placebo-controlled trials showed a dose-dependent reduction in cerebrospinal fluid level of A β 42 [164]. They demonstrated improvement in two executive function component tests of the neuropsychological test battery and in the executive factor z-scores without any serious adverse events [164,165].

DP-109, a diester derivative of BAPTA (1,2-bis(2-aminophenyloxy) ethane-N,N,N',N'-tetra acetic acid), selectively chelated transition metals such as Zn, Cu and Fe within membrane compartments [166]. DP-109 could be similar to CQ in attenuating A β plaque deposition, inflammation and neuronal damage in hippocampus. It reduced cerebral amyloid angiopathy and increased soluble:insoluble A β 40:42 ratio. DP-109 effectively crossed the BBB neuronal and vesicular membranes to quench free Zn in synaptic vesicles to reach the brain of Tg2576 mice. However, with DP-109, possible chelating effects on other metals cannot be ignored [166].

1-(benzimidazole-2-ylmethyl)-1,4,7-triazacyclononane and 1,4-bis(benzimidazole-2-ylmethyl)-1,4,7-triazacyclonone), quite similar to CQ exhibited radical-scavenging potential with metal-protein-attenuating ability. The imidazole in the both compounds held sites that could be modified for preventing A β aggregation [167].

However, in contrast to previous studies, it has also been observed that CQ promoted the *in vitro* aggregation/fibrillogenesis of human A β in the presence of Cu and Zn with decreased viability in neuroblastoma cells [168]. There was an increased lethality of APP transgenics upon CQ treatment, which could be rescued by a co-treatment with Cu, confirming toxicity of CQ. Moreover, the exposure to Cu with CQ led to a modest but significant increase in cerebral Cu levels, most probably due to enhanced transport of CQ–Cu complexes with a secondary role as a chelator. Most conspicuously, until 1975 there were 10,000 cases of subacute myelo-optic neuropathy (SMON) associated with CQ administration. SMON resembled an accelerated form of sub acute combined degeneration due to vitamin B₁₂ deficiency [156,169,170].

3.2 DFO

DFO (Figure 4B) has become the mainstay of iron- and Al-chelating therapy [171] (later on the beneficial effect was suggested to be due to the chelating of Cu and Zn also [172]) with probable inhibition of free radical formation and inflammation [173]. DFO either prevented the formation of β -pleated amyloidal fibrils or it initiated the dissolution of febrile amyloidal plaques [150]. In a clinical trial that involved the treatment of AD patients, the administration of DFO led to a significant reduction in the rate of decline of daily living skills as assessed by both group means and variances (p < 0.04) [171].

DFO is non-oral, rapidly metabolized and relatively unstable molecule, and is administered via painful intramuscular injections [172]. It is a highly costly drug with the need for parenteral administration. Systemic metal ion depletion (anemia) as a side effect of DFO are important reasons to search for an orally effective, cheaper and less toxic chelating agent than DFO [172].

3.3 Bifunctional metal chelators

New oral Fe chelators, 5-[*N*-methyl-*N*-propargylaminomethyl]-8-hydroxyquinoline (M-30) (Figure 4C) and the 8-hydroxyquinoline derivative, VK28 (Figure 4D), are being developed for the treatment of AD and other neurodegenerative diseases [174,175]. The neuroprotective activity of propargylamines led to the development of several novel bifunctional iron chelators, (5-[4-propargylpiperazin-1-ylmethyl]-8-hydroxyquinoline) HLA-20, M30 and M30A from the prototype brain-permeable iron chelator, VK-28. The chelators had cholinesterase and monoamine oxidase B inhibitory activities and retained the *in vitro* and *in vivo* neuroprotective activity of rasagiline. The *N*-propargyl moiety of the anti-PD drugs rasagiline (Azilect, Teva) and selegiline have been found to be the most effective that exerted Fe chelation potency, served as radical scavengers and inhibited Fe-induced membrane lipid peroxidation features [176].

Fe chelators of low molecular weight, minimal toxicity and satisfactory lipophilicity have added a new facet in the etiology of AD therapy. Fe-binding drugs may also stabilize hypoxia-inducible factor that in turn would transactivate the expression of established protective genes, including VEGF, erythropoietin, aldolase and p21.

The metal-complexing bifunctional molecule, XH1 (Figure 4E), ([(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]- $\{2-[(2-\{[(4-benzothiazol-2-yl-phenylcarbamoyl)methyl]-carboxymethyl-amino]-ethyl]-amino]-acetic acid, which crosses the BBB and holds the potential of being an amyloid-targeting metal chelator for AD treatment [177] contains two identical amyloid-binding and one metal-chelating moiety that specifically targets amyloid. It i) binds to A<math>\beta$ 1 – 40 peptide putatively, ii) decreases Zn(II)-induced A β aggregation *in vitro*, iii) specifically reduces APP protein expression in human SH-SY5Y neuroblastoma cells and iv) attenuates cerebral A β amyloid pathology in PS1/APP transgenic mice without inducing apparent toxicity and behavior disturbances [177].

Due to the attached carbohydrate moiety responsible for increased tissue specificity at a physiological pH, and due to the phenolic moieties as suitable antioxidants the

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tetrahydrosalens such as $(N,N'-bis[(5-\beta-D-glucopyranosyloxy-2-hydroxy)benzyl]-N, N'$ $dimethyl-ethane-1,2-diamine (H2GL1) and N,N'-bis[(5-\beta-D-glucopyranosyloxy-3-tert$ butyl-2-hydroxy)benzyl]-N,N'-dimethyl-ethane-1, 2-diamine (H2GL2)) were developed $[178]. Both H2GL1 and H2GL2 were found to reduce Zn-and Cu-induced a<math>\beta$ 1 – 40 aggregation *in vitro*, with a higher affinity for Cu over Zn with H2GL1 displaying better coordinating ability at physiological pH [179]. The moderate affinity of H2GL1 and 2 for metal ions at physiological pH might obviate the toxicity commonly associated with chelating therapy [179].

3-hydroxy-4-pyridinones that contain phenol groups for antioxidant functionality are further elaborated with pendant glucose moieties for improved BBB targeting. Glycosidase removal of the carbohydrate substituents gives ligands that are ready to passivate excess metal ions, especially Cu and Zn, in the brain [180,181].

Curcumin and other polyphenols are anti-inflammatory and antioxidant agents, and structurally interfere with A β aggregation and metal dyshomeostasis [182]. The use of synthetic activity to rescue protein-aggregate-mediated cellular toxicity led to the synthesis of apocyclen attached to selective A β recognition motifs (KLVFF or curcumin). The resultant complexes interfere with A β aggregation and degrade A β into fragments, preventing H₂O₂ formation and toxicity in neuronal cell culture [183].

3.4 Al chelators

Tacrine, ascorbate plus Feralex-G have also been found to be particularly effective in removing Al(III) from the nuclear matrix in addition to enhancing cholinergic transmission [184]. Feralex-G disaggregated compacted paired helical filaments isolated from aged human brain [185]. Feralex could also dissociate binding of Al and Fe with hyperphosphorylated Tau of AD [185]. Simultaneous administration of two Al chelators, DFO and tacrine as a palliative treatment for AD patients has also been taken into consideration. Administration of *N*-(2-hydroxyethyl) EDTA, a potential antidote for Al overload, in combination with citric acid has been proposed as a chelation therapy for AD [186].

3.5 Nanoparticle chelator delivery

Chelator-delivery with the help of nanoparticles has been suggested to be a significantly improved method of chelation therapy with higher efficacy, reduced toxicity and substantial tissue-specific targeting. Covalent conjugates of Fe chelators with nanoparticles, such as prototype nanoparticle–chelator conjugate, could attenuate the lipophilic character of the chelator. They provided better BBB permeability and safe treatment with higher chelator bioavailability in AD, without affecting metal binding ability of chelators. The nanoparticle–chelator conjugates could effectively inhibit A β aggregate formation and, thereby, protect human brain cells from A β -related toxicity. Using *in vitro* studies, it was shown that chelator–nanoparticle system complexed with Fe. When incubated with human plasma it preferentially adsorbed apolipoprotein E and apolipoprotein A-I that would facilitate transport of chelators and chelator–metal complexes in both directions across the BBB. The

system thus provided safer and more effective chelation treatment in AD and other neurodegenerative diseases [187–190].

4. Targeting neuronal signalling

A novel neurotherapeutic approach involved the activation of neuronal cell signaling mechanisms using metallo-complexes. The metal ligand-Cu complexes as with CQ or PBT2 or alternative metal-complexes such as Cu/Zn-bis(thiosemicarbazone) (Figure 5A) complex entered brain cells and upregulated MMP. This involved activation of ERK signaling to cleave the monomeric A β (Figure 6) [90]. In another situation, inhibition of PI3K and C-JNK prevented depletion of A β by the metallo-complexes [191].

Lipoic acid (LA) (Figure 5B) or dihydrolipoic acid (DHLA) exhibited the ability to chelate metal ion with the disulfide group of dithiolane ring [192]. They reduced liposomal peroxidation [193] and activated the pro-survival PI3K and ERK1/2 signaling pathways. LA/ DHLA protected cultured hippocampal neurons against A β and Fe/H₂O₂ [194] and improved learning-memory in a Tg2576 mouse model for AD [195,196].

5. Natural antioxidants

Curcumin (Figure 7A) molecules chelated cations with the diketone and pairs of phenol and methoxy groups, reduced oxidative stress and prevented amyloid aggregation. Its two molecules bind the Cu and Fe on A β at $K_{d1} \sim 10 - 60 \mu$ M and $K_{d2} \sim 1.3 \mu$ M (for binding of the first and second curcumin molecules, respectively) [197,198]. But the antioxidants like curcumin and ginkgo (common name kaempferol) (Figure 7B) extracts fail to reach the target site in sufficient amounts during oxidative stress, and hence are best as prophylactics to complex the metals before the stress cascade starts [199].

Human epidemiology confirms that tea extract contains nutrients endowed with possible prospective neurobiological-pharmacological actions. They are beneficial to human health due to the natural antioxidant, catechin polyphenol constituent (–)-epigallocatechin-3-gallate (EGCG) (Figure 7C) [200]. EGCG targets APP 5'UTR and decreases A β levels [169] and plaques in the cingulate cortex, hippocampus, and entorhinal cortex. It was found to promote non-amyloidogenic α -secretase proteolytic pathway in 'Swedish' mutant APPsw and in Tg2576 mice with a disintegrin and metalloprotease (ADAM)-10 activation. EGCG administration also markedly suppressed sarkosyl-soluble phosphorylated tau isoforms with significant behavioral improvements [201,202].

Moderate consumption of wine and increased intake of fruits and vegetables is associated with a lower incidence of AD [203–205]. The antioxidant, resveratrol (a bioactive compound in red wine), through the protein kinase pathway could lower A β peptide formation, and promote A β degradation in 15-week-old male APP/PS1 transgenic mice (B6C3-Tg (APPswe, PSEN1dE9)85Dbo/J), The Jackson Laboratory [206].

Antioxidants, such as α -tocopherol (Vitamin E) and ascorbic acid had modest benefits in elderly subjects [207]. Gossypin (3,3',4',5,7,8-hexahydroxyflavone 8-glucoside) [208] and ginko biloba [209] protected cortical cell cultures from A β -induced toxicity. They exhibited

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neuroprotective effects in several mouse models and maintained and improved cognitive function in AD patients [208]. It was found that 6-hydroxy melatonin administration proved successful in reducing Fe²⁺-induced neurotoxicity, lipid peroxidation and necrotic cell damage in the rat hippocampus *in vivo* [210].

A pilot study examining the efficacy of the Cu-chelating agent D-penicillamine (treatment for Wilson's disease) in AD patients showed decrease in oxidative damage with depletion of bio-available Cu and increased excretion of Cu in the urine [211]. But, the role of Dpenicillamine in antioxidant balance remains inconclusive, possibly due to the action of vitamin B_6 that was administered to placebo- and D-penicillamine-treated patients [212]. Further studies with larger cohorts of AD patients are needed to elucidate the real therapeutic efficacy of D-penicillamine.

A relatively new concept deals with possibility of using Zn compounds (used for Wilson's disease treatment [213]) that could induce or maintain a state of Cu-malabsorption. Successful clinical trials with Zn could lead to safe, inexpensive and effective biological anti-Cu agent in AD [214,215].

6. Expert opinion

Over the last few decades, researchers have provided a wealth of information on the underlying nature of AD therapy. Although the amyloid-based pathogenic mechanisms that result in the onset and progression of AD are yet to be clearly understood, there are several growing bodies of evidence to support a central role for bio-metals in many critical aspects of the illness. The APP and $A\beta$ still remain the current focus of AD research that is associated with their integral function in iron metabolism and homeostasis of metal stores in the brain. The deposition of metals in the plaque of AD patients and the demonstration of metal dependent translation of APP 5[']UTR mRNA have distinctly indicated the involvement of metals in amyloid-associated characteristic pathological feature of AD. However, further research is necessary to fully understand the complex and interdependent pathways of biometal homeostasis and amyloid metabolism in AD.

In this review we have provided an update on the development of potential therapeutic agents for AD based on the modulation of metal bioavailability. The compounds could target the metal binding sites on APP and A β , deprive the biological systems of metal ions, or promote metal uptake into cells, and thus inhibit A β :metal-mediated redox activity.

The metal chelating drug, CQ, was well tolerated and did appear to produce some modest benefits in AD patients that could support the proof of concept in humans that drugtargeting metal-A β interactions can have a significant beneficial effect on the progression of AD. But the study only involved a small number of patients and required further observations in larger groups of patients.

The metal chelating drugs DFO (Figure 4B), tetrathiolmolybdate (Figure 4F), and dimercaptopropanol (Figure 4G) [47,216] have shown a significant effect on A β metabolism *in vitro* and/or *in vivo*. A transfection-based screen of a library of FDA drugs to identify compounds that limited APP luciferase reporter expression translated from the APP 5'UTR

in neuroblastoma cells (SY5Y) cells identified the leads, paroxetine (PaxiITM) and dimercaptopropanol. The compound limited A β peptide secretion from lens epithelial cells (B3 cells), the former probably through chelation of or change in distribution of interacellular iron [217]. Tetrathiolmolybdate (Cu²⁺ chelator), dimercaptopropanol (Pb²⁺ and Hg²⁺ chelator) [217] and XH1 (Figure 4E), with A β -binding and metal-chelating moieties [177], suppressed APP holoprotein expression and A β secretion; tetrathiolmolybdate also showed excellent efficacy in animal models [218]. In addition to these compounds, phenserine, which is a novel and highly selective acetylcholinesterase inhibitor, is being tested for the treatment of AD [219]. Phenserine was extremely efficient in blocking translation under conditions of intracellular Fe chelation with DFO, suggesting that the anticholinesterase operated through an Fe-dependent pathway at the APP 5'-UTR site. The FDA-preapproved drugs had the major advantage of being pharmacologically fully characterized, with respect to the toxicity, half-life, capacity for oral administration, and capacity to cross the BBB.

However, a major problem associated with the widespread clinical use of the available metal-complexing agents is their poor target specificity and consequent clinical safety. The long-term use of these agents is likely to perturb the homeostasis of many biometals and normal physiological functions of essential metal-requiring biomolecules. Thus, the development of metal passivating agents with necessary water solubility, efficacy, minimum toxicity and specific targeting is essential for new effective therapies for AD.

New screening approaches targeting APP 5' UTR could thus be very useful in identifying novel metal-complexing agents from world-wide drug libraries. We have performed such a high-throughput screen (HTS) of 110,000 compounds obtained from the library of the Laboratory for Drug Discovery on Neurodegeneration that yielded several non-toxic specific inhibitors of APP mRNA 5' UTR-driven luciferase in the stable SH-SY5Y transfectants [220]. The identified compounds could be expected to hold therapeutic promise as metal chelators at least for those retaining long-term bio-activity. Use of transgenic models of AD could be the next and required step for testing our APP-directed compound hits. We could test the efficacy of lead APP 5' UTR-directed drugs to limit amyloid burden in CRND8 mice wherein the human APP-695 transgene (London/Swedish double mutations in APP-695 cDNA) is expressed under the transcriptional control of the prion protein gene promoter and the translational control of the natural APP 5' UTR [221].

It has been observed that intracellular modifiers of levels of Zn [159] and Fe [222] adjust α secretase to limit A β peptide. Thus, the screening of chemical compounds or antioxidants that could selectively target and promote metal-dependent catalysis of ADAM-protease and the decrease in A β levels could serve as a strategy for identifying anti-AD drugs [220,223,224]. Signaling events involved in the non-amyloidogenic, metalloprotease- α secreatse activity could be the site of pharmacological intervention in AD. Activators of PKC, adenylate cyclase/protein kinase A system, phoslpholipase C (PLC) and the MAPKsignaling system could also claim to be potential and new anti-amyloid agents [225].

The selective expression of human metal-binding protein biomarker S100A7 in the brain of transgenic mice resulted in significant promotion of α -secretase activity. Furthermore, the

promotion of S100A7 expression in the brain selectively promoted α -secretase activity in the brain of AD [225]. Thus upregulating S100A7 in the brain or cerebrospinal fluid could be developed as another strategy for promoting of 'non-amyloidogenic' α -secretase/ADAM-10 mediated responses.

Although development of drugs that target abnormal metal accumulation of $A\beta$ is in process, we still need to explore new screening approaches and technologies to identify novel therapeutics that could promote neuroprotective signalling pathways in AD with minimum adverse effects.

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Article highlights

- Metal dependent amyloid formation: Amyloid precursor protein (APP) is a Cu-, Zn- and Fe- binding protein, and these metals clearly provide one of the pathological requirements for polymerization of amyloid-beta (Aβ) peptide. Fe (perhaps Cu/Zn) controls iron-responsive element (IRE)-iron regulatory protein (IRP) binding to APP mRNA to reset and increase its translation rate thus worsening metal-associated pathological lesions in Alheimer's disease (AD). Al is also a long-known risk factor for AD.
- Chelation therapy: Metal chelation serves as a therapeutic strategy for treatment of AD. Clioquinol inhibits binding of Cu and Zn to Aβ and promotes Aβ clearance. Desferrioxamine (DFO) has been shown to be therapeutic after intramuscular injection. A new range of bifunctional metal chelators (M-30, VK-28, HLA-20, XH1, curcumin and polyphenols) are being assessed for AD treatment. Simultaneous administration of two chelators is being taken into consideration. Nanoparticle chelator delivery should significantly improve the efficacy and reduce the toxicity of chelation therapy.
- *Neuroprotective signaling:* A promising future lies in the use of metallocomplexes to trigger neurotherapeutic signaling pathways with subsequent inhibition of tau phosphorylation and Aβ generation.
- *Anti-oxidants:* Curcumin, kaempferol, epigallocatechin-3-gallate (EGCG), resveratrol, vitamin E and D-penicillamine are beneficial to counteract the toxicity of the redox interaction between metals and Aβ peptide.
- *Future directions:* The challenge of a transfection-based screening approach will be to find new drugs that suppress APP holoprotein translation and thereby limit Aβ peptide. Use of transgenic models of AD is the next and required step for testing the novel APP directed lead compounds.

This box summarizes key points contained in the article.



Figure 1. Synchrotron X-ray fluorescence (XRF) microprobe images of human Alheimer's disease AD plaque.

Elemental profiles (S, Fe, Cu, and Zn) in a typical Alzheimer's amyloid beta (A β) amyloid plaque. The cryo-sectioned (10 µm thickness) AD brain tissues were stained with 0.1% Thioflavin-T for amyloid plaques. The amyloid plaque-bearing human brain tissues were procured by laser capture microdissection (LCM) (Arcturus Pixcell IIE platform) and mounted on Si₃N₄ membrane grids (2.0 × 2.0 mm). Guided by the optical amyloid plaque images, the samples were excited with incident synchrotron X-ray of 10 keV for elemental K α characteristic emission lines. Elemental profiles (S, Fe, Cu, and Zn) were obtained using synchrotron scanning X-ray fluorescence microscopy (µ-XRF) at the Advanced Photon Source of the Argonne National Laboratory. Red depicts the hottest spot of the metals in plaques. (The significance of sulfur (S) element may reflect its high abundance in proteinaceous elemental composition and as an indicator for amyloid plaque-associated oxidative stress since protein *S*-glutathionylation is a salient feature of oxidative stress). Reproduced from [7].



Figure 2. The proteolytic processing of amyloid precursor protein (APP) to produce A β that coordinates the metal ions (M: Zn, Cu and Fe) to induce aggregation, and generation of ROS. APP (695, 751, 770 amino acid isoforms that predominate in brain) can be processed in the plasma membrane as it travels from the intracellular origin to extracellular matrix through the non-amyloidogenic route that involves cleavage at the α -secretase site at amino acid 17 of the 40 – 42 amino acid A β domain resulting in two fragments, sAPP α and a C-terminal fragment (CTF α). Further proteolysis of the CTF α fragment by α -secretase generates the non-amyloidogenic peptide p3 and a C-terminal fragment CTF γ . When APP escapes processing at the α -site, it undergoes β -secretase cleavage at the beginning of A β domain, resulting in a C-terminal fragment CTF β and sAPP β . Next, the resultant β -stub becomes the substrate for γ -secretase cleavage, culminating in extracellular A β secretion. The hypermetallated (by Zn, Fe and Cu) state of A β as a consequence of age-dependent elevations in tissue metal concentrations can induce A β aggregation. H₂O₂ can initiate a number of oxidative events, including Fenton reactions to form toxic hydroxyl radicals and calcium dysregulation, and subsequent reactive oxygen species (ROS) generation.

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Figure 3. Model for the iron-induced change of iron regulatory protein (IRP) interaction with the APP/ferritin iron-responsive element (IRE) to modulate APP/ferritin translation.

The APP IRE is homologous with the canonical L-and H-ferritin IRE mRNA stem-loop that binds the iron regulatory proteins (IRP1 and IRP2), and modulates translation of ferritin to control intracellular iron homoeostasis [226]. Iron influx increases ferritin mRNA translation by releasing IRP1–IRP2 binding to the 5' cap site of IRE stem–loop. The iron-induced change of IRP1 interaction with the APP-IRE activates either 5' cap translation or internal 40S ribosome entry and the onset of APP protein synthesis [227]. The IRE of APP interacts with IRP1, whereas the canonical H-ferritin IRE RNA stem-loop binds to IRP2 in neural cell lines, in human brain cortex tissue and human blood lysates. The canonical H-ferritin IRE RNA stem-loop binds also to IRP2. The APP mRNA acute box domain, as for H-ferritin mRNA, is located immediately upstream of the start codon and may well also interact with RNA poly(C)-binding proteins, CP-1 and CP-2 and control cytokine-induced APP mRNA translation.

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Figure 4. Important metal-chelators characterized as suppressors of APP and A β aggregation. Chemical structures of **A**. Clioquinol, which inhibits Zn, Cu- or Fe-mediated oxidative stress and reduces clinically observed AD-induced cognition; **B**. Desferrioxamine, intracellular Fe³⁺ chelator that suppresses APP translation without changing α -secretase activity; **C** & **D** M-30 and VK-28. M-30 being derived from a prototype iron chelator, VK-28; both being developed for anti-amyloid efficacy and α -secretase co-activation; **E**. Bifunctional XH1: [(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(2-{[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(2-{[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(2-{[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(2-{[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(2-{[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(2-{[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(2-{[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(2-{[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(2-{[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(2-{[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(2-{[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(2-{[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(2-{[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(2-{[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(2-{[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(2-{[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(4-benzothiazol-2-yl-phenylcarbamoyl)-methyl]-{2-[(4-benzot

amino)-acetic acid, with amyloidtargeting metal chelating property; **F**. Tetrathiomolybdate, which shows excellent efficacy in animal AD models and is presently under clinical trial. **G**. Dimercaptopropanol, which has a significant effect on A β metabolism *in vitro* and/or *in vivo*.



Lipoic acid Molecular weight; 206.32

Figure 5. Metallo-complexes that target neuronal signaling.

Chemical structures of **A**. Cu-bis(thiosemicarbazone) that reduces tau phosphorylation through PI3K and ras/raf signalling; **B**. Lipoic acid that chelates metal ion and promotes prosurvival signaling pathways.



Figure 6. Proposed metal ligand action that targets neuronal cell signaling in treatment of AD. Metal-free ligands (L) such as CQ or PBT2 may bind with Cu of the A β peptide-Cu complex resulting in dissolution of A β into Cu-free monomers. The metal ligand–Cu complexes or alternative metal complexes such as Cu-bis(thiosemicarbazone) then enter cells, activate PI3K followed by sequential phosphorylation of AKT and glycogen synthase kinase beta (GSK3 β) that inhibits tau phosphorylation [228]. The complex-mediated activation of ras/raf signalling activates ERK, upregulates MMP activity, which cleaves the monomeric A β .

Adapted from [90].



Figure 7. Important metal-chelators characterized as natural antioxdants for AD. Chemical structures of A. curcumin, a polyphenol that binds Fe and Cu on A β and prevents amyloid aggregation; B. Ginko biloba, inhibits a free radical scavenger that reduces clinically observed AD-induced cognition; C. (–)-epigallocatechin-3-gallate (EGCG) decreased A β levels and plaques via promotion of α -secretase activity.