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# **Physiological and pathological aspects of A**β **in iron homeostasis via 5'UTR in the APP mRNA and the therapeutic use of iron-chelators** Yael Avramovich-Tirosh, Tamar Amit, Orit Bar-Am, Orly Weinreb and Moussa BH Youdim\*

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#### **Abstract**

Many studies have highlighted the pathological involvement of iron accumulation and iron-related oxidative stress (OS) in Alzheimer's disease (AD). Iron was further demonstrated to modulate expression of the Alzheimer's amyloid precursor holo-protein (APP) by a mechanism similar to that of regulation of ferritin-L and -H mRNA translation through an iron-responsive element (IRE) in their 5' untranslated regions (UTRs). Here, we discuss two aspects of the link between iron and AD, in relation to the recently discovered IRE in the 5'UTR of APP mRNA. The first is the physiological aspect: a compensatory neuroprotective response of amyloid-β protein (Aβ) in reducing iron-induced neurotoxicity. Thus, given that Aβ possesses iron chelation sites, it is hypothesized that OS-induced intracellular iron may stimulate APP holo-protein translation (via the APP 5'UTR) and subsequently the generation of its cleavage product,  $\mathsf{AB},$  as a compensatory response that eventually reduces OS. The second is the pathological aspect: iron chelating compounds target the APP 5'UTR and possess the capacity to reduce APP translation, and subsequently Aβ levels, and thus represent molecules with high potential in the development of drugs for the treatment of AD.

#### **Introduction**

There is increasing evidence that iron accumulation in the brain can cause a vast range of disorders of the central nervous system. It has become apparent that iron progressively accumulates in the brain with age [[1](#page-6-0),2], and that iron-induced oxidative stress (OS) can cause neurodegeneration [3]. Free iron induces OS through its interaction with hydrogen peroxide (Fenton reaction), resulting in increased formation of hydroxyl free radicals. Free radicalrelated OS causes molecular damage that can then lead to a critical failure of biological functions and ultimately cell death [4,5].

In Alzheimer's disease (AD) pathology, iron is significantly concentrated in and around amyloid senile plaques, and neurofibrillary tangles (NFTs), leading to

alterations in the pattern of the interaction between iron regulatory proteins and their iron responsive elements (IREs), and disruption in the sequestration and storage of iron [6,7]. Also, high levels of iron have been reported in the amyloid plaques of the Tg2576 mouse model for AD, resembling those seen in the brains of AD patients [8]. In addition to the accumulation of iron in senile plaques, it was demonstrated that the amount of iron present in the AD neuropil is twice that found in the neuropil of nondemented brains [6]. Further studies have suggested that accumulated iron supports the AD pathology as a possible source of OS-dependent reactive oxygen radicals, demonstrating that neurons in AD brains experience high oxidative load [9-12]. *Post mortem* analysis of AD patients' brains have revealed activation of two enzymatic indicators of cellular OS: heme oxygenase-1 [13] and NADPH oxidase [14]. Also, heme oxygenase-1 was greatly enhanced in neurons and astrocytes of the hippocampus and cerebral cortex of AD subjects, co-localizing to senile plaques and NFTs [15]. A recent study reported that ribosomal RNA provided a binding site for redox-active iron and serves as a redox center within the cytoplasm of vulnerable neurons in AD brain, in advance of the appearance of morphological change indicating neurodegeneration [16]. In addition, other evidence suggests that the metabolism of iron is disrupted in AD. For example, the location of the iron-transport protein transferrin in senile plaques, instead of its regular location in the cytosol of oligodendrocytes, indicated that it becomes trapped within plaques while transporting iron between cells [17]. The mediator of iron uptake by cells, melanotransferrin, and the iron-storage protein ferritin are altered in AD and are expressed within reactive microglial cells that are present both in and around senile plaques [18,19].

Previous studies assessing the effects of certain genes encoding proteins involved in iron metabolism, such as hemochromatosis (*HFE*) and Transferrin (*TF*) genes, on the onset of AD have been contradictory [20,21]. At the biochemical level, iron was demonstrated to facilitate the aggregation of β-amyloid peptide (Aβ) and increase its toxicity [22]. Indeed, the iron chelator deferrioxamine (DFO) prevented the formation of β-pleated sheets of  $Aβ<sub>1-42</sub>$  and dissolved preformed β-pleated sheets of plaque-like amyloid [23]. Also, iron induced aggregation of hyperphosphorylated  $\tau$  (tau), the major constituent of NFTs [24].

A direct link between iron metabolism and AD pathogenesis was provided recently by Rogers *et al.* [25], who described the presence of an IRE in the 5' untranslated region (5'UTR) of the amyloid precursor protein (APP) transcript. Thus, APP 5'UTR is selectively responsive to intracellular iron levels in a pattern that reflects irondependent regulation of intracellular APP synthesis. Indeed, iron levels were shown to regulate translation of APP holo-protein mRNA in astrocytes [26] and neuroblastoma cells [25] by a mechanism similar to iron control of the translation of ferritin-L and -H mRNAs via IREs in their 5'UTRs.

This review will discuss two main aspects of the link between iron and AD in relation to the recently discovered IRE in the 5'UTR of APP mRNA. First is the physiological aspect, which considers the neuroprotective response of Aβ in reducing iron-induced neurotoxicity. Thus, given that Aβ possesses iron chelating sites, it may be hypothesized that OS-induced intracellular iron levels stimulate APP holo-protein translation (via the APP 5'UTR) and the subsequent generation of its cleavage product, Aβ, as a compensatory response that eventually reduces OS. Second is the pathological aspect, which considers iron chelator compounds targeting the APP 5'UTR that possess the capacity to reduce APP translation, and subsequent Aβ generation, as molecules with high potential in the development of drugs for the treatment of AD (Figure 1).

### **A**β **generation as a compensatory mechanism**

Conflicting results have been reported in recent literature concerning whether the interaction between Aβ and iron is neurotoxic or neuroprotective [27]. Thus, while *in vitro* studies demonstrate that Aβ can induce OS and neurotoxicity (at high concentrations of aged Aβ), it is apparent from other reports that OS promotes Aβ generation (perhaps as a result of altered metal iron metabolism), presumably as a protective/compensatory response, leading to reduced neuronal OS. Consistent with this, amyloid plaques and NFTs in the cortex were found to be inversely correlated with OS markers (for example, 8-hydroxyguanosine) [28], indicating that oxidative damage is an early event in AD that decreases with disease progression. Indeed, in different APP transgenic mouse lines, the subtle functional deficits occur before the formation of amyloid plaques [29]. In addition, various sources of OS (for example,  $H_2O_2$ , UV, and reactive oxygen species (ROS))



#### Figure 1

Chemical structures of the novel multifunctional iron chelators M30, HLA20 and VK28.

have been demonstrated to increase neuronal Aβ production [30-32]. Moreover, Aβ has been shown to be upregulated by many forms of stress conditions, including apoptosis, ischemia, shortage of energy supply, hypoglycemia, and brain injury [33-35]. The levels of β-secretase were markedly increased by oxidative agents, with consequent augmentation of the levels of carboxy-terminal fragments of APP [36,37], further suggesting that OS may be the cause of Aβ production. Taken together, these studies suggest that Aβ formation may be a response to, rather than a cause of, neurotoxic oxidative challenge.

Previous studies have reported similarities between APP holo-protein and ferritin gene expression, both of which are driven by translational regulatory events [38]. The expression of the APP gene was up-regulated at the translational level by iron and interleukin-1, which was paralleled by the action of 5'UTR sequences that are similar to the 5'UTR sequences in the mRNA coding for the L- and H-subunits of ferritin. Evidently, IRE-dependent pathways govern the post-transcriptional expression of many proteins involved in iron metabolism, in addition to ferritin and transferrin receptor.

Aβ has been characterized as a metalloprotein that binds transition metal ions via three histidines and a tyrosine residue located in the hydrophilic amino-terminal part of the peptide. Thus, given that  $Aβ$  is a metalloprotein that possesses strong chelating properties for transition metal ions [27], it is proposed that Aβ generation under oxidative conditions may be aimed at sequestering metal ions in order to prevent further potentional oxidative damage. Consistent with this, previous studies demonstrated that the injection of Aβ-iron complexes into rat cerebral cortex was less toxic than iron alone [39]. Additionally, Aβ1–40 at 5 μM was found to protect primary neuronal cultures from the neurotoxicity of iron [40,41]. Recent findings demonstrated that three histidine residues in Aβ control the redox activity of iron, indicating that Aβ is likely to be an important antioxidant. Thus, it was shown that Fe3+ catalyzed ascorbate oxidation and hydroxyl radical generation were inhibited in the presence of Aβ1–40 or Aβ1–42 [42].

Here, we analyzed the capacity of Aβ to reduce ironinduced cell death in Chinese hamster ovary (CHO) cells stably transfected with the APP 'Swedish' mutation (CHO/ΔNL), which express elevated levels of holo-APP in cell lysate and Aβ1–40 and Aβ1–42 peptides in the medium compared to controls (Figure 2). Indeed, as demonstrated in Figure 3, CHO/ΔNL cells provided significant increased protection against iron-mediated cell toxicity compared to control CHO cells, further suggesting that Aβ significantly reduced the neurotoxicity of iron. Based on these findings, we hypothesize that the IRE of the 5'UTR



#### **Figure 2**

APP expression in CHO cells. **(A)** A representative western blot analysis of cellular holo-APP in CHO cells and CHO cells stably transfected with the APP 'Swedish' mutation (CHO/ΔNL). Cellular holo-APP was detected in cell lysates with 22C11 antibody (directed to the APP amino terminus). The loading of the lanes was normalized to levels of β-tubulin. **(B)** Aβ was detected in the medium of CHO and CHO/ ΔNL cells by immunoprecipitation and western blotting with monoclonal antibody 6E10 (which recognizes an epitope within residues 1–17 of Aβ domain). **(C)** Aβ1–40 and Aβ1– 42 levels were measured using standard sandwich ELISA (BioSource, Camarillo, California, USA).

of the APP transcript may be linked to the compensatory response of Aβ that helps neurons cope with altered iron homeostasis. Figure 4 illustrates the main physiological iron homeostatic mechanisms, including the IRE in the 5'UTR of the APP mRNA as a potential target involved in the compensatory response of Aβ. First, following moderate OS conditions and abnormal iron metabolism, APP synthesis is enhanced via the APP-5'UTR; second, as Aβ is a cleavage product of APP, the increase in APP level will be accompanied by elevation of Aβ generation; and third, Aβ peptides may have compensatory/neuroprotective properties as a result of their ability to trap free iron [27]. In addition to known iron regulation targets (for example, transferrin receptor, ferritin), the formation of Aβ peptides may be considered a novel compensatory response that reduces moderate OS damage.

#### **Iron chelation for the treatment of AD**

Under extreme pathological conditions – that is, at some threshold level of ROS generation – it appears that the main role of Aβ switches from neuroprotective to dyshomeostatic in terms of cerebral biometals and APP/Aβ/ metal-redox complexes, leading to a vicious cycle of increased ROS production and Aβ generation. Chelation has the potential to prevent iron-induced ROS, OS, and Aβ aggregation and, therefore, chelation therapy may be considered a valuable therapeutic strategy for AD. In fact, intramuscular administration of DFO, a potent iron chelator, slowed the clinical progression of AD dementia [43]



### Figure 3

Comparison of viability between CHO and CHO/ΔNL cells following  $FeSO<sub>4</sub>$  treatment. CHO and CHO/ $\triangle$ NL cells were incubated in the absence (Control) or presence of different concentrations of  $FeSO<sub>4</sub>$  (5–20  $\mu$ M) for 24 h. Viability of the cell cultures was determined by the MTT (3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyl tetrazolium bromide) method. The data are expressed as percent of Control. Representative curves from four independent experiments are shown. \**P* < 0.05; \*\**P* < 0.01 versus CHO cells.

and some success has also been achieved with clioquinol, another metal-complexion agent [44,45]. However, clioquinol is highly toxic and DFO has poor blood-brain barrier penetration.

The identification of an IRE in the 5'UTR of the APP transcript led to a novel therapeutic approach aimed at reducing amyloidosis by FDA pre-approved drugs targeted to the IRE in the APP mRNA 5'UTR [38]. For example, the APP 5'UTR-directed drugs DFO (Fe3+ chelator), tetrathiomolybdate (Cu2+ chelator), and dimercaptopropanol (Pb2+ and Hg2+ chelator) were found to suppress APP holo-protein expression and lower Aβ secretion [38,46,47]. In addition, the bi-functional molecule XH-1, which contains both amyloid-binding and metal-chelating moieties, was shown to reduce APP expression in SH-SY5Y cells and attenuate cerebral Aβ in PS1/APP transgenic mice [48]. Additional drug classes were also reported to suppress the APP 5'UTR and limit APP expression, including antibiotics, selective serotonin reuptake inhibitors, and other selective receptor antagonists and agonists [47].





Since there is a functional IRE in the 5'UTR of APP mRNA, we further investigated the efficacy of the iron chelator drugs M30, HLA20 and VK28 to modulate the translation of a luciferase reporter gene driven by APP 5'UTR sequences. The assay was performed essentially as described in Reznichenko *et al*. [56]. The pGALA construct was generously provided by JT Rogers (Massachusetts General Hospital, Boston, MA, USA). Human U-87-MG glioma cells, selectively chosen for their high transfection efficiency, were grown in flasks (100 mm2) and transfected with 7 μg of DNA from the parental vector pGL-3 or pGALA constructs and co-transfected with 3 μg of DNA from a construct that expresses green fluorescent protein (GFP) to standardize for transfection efficiency. The pGALA consists of a pGL-3 backbone to which the APP 5'UTR sequences (containing the IRE) were inserted in front of the luciferase gene start codon and the complete APP 3'UTR sequences immediately downstream of the luciferase, to provide the natural arrangement of the APP gene 5' and 3'. After 12 h, the cells were split equally into 96-well plates and grown without (control) or with the iron chelator drugs M30, HLA20 or VK28 for 48 h. Cell viability was established by a fluorescent microscopic examination of each well, quantified for GFP activity at 480/509 nm wavelength (using an automatic Wallac-1420 multilabel counter) and then lysed to determine luciferase activity. For statistical analysis one-way ANOVA followed by Student's *t*-test was performed. Quantitative values are mean ± standard error of the mean (n = 3). †*P* < 0.05, \**P* < 0.01 versus control was considered significant.

The concept of metal chelators for clinical use in neurological disorders that could remove excess iron in the brain recently led our group to develop non-toxic, lipophilic, and brain permeable iron chelators for neurodegenerative diseases. The novel iron chelator VK28 (varinel) [49] and the multi-functional drugs HLA20 and M30 (Figure 1) [50], which possess the potent iron chelating activities and neuroprotective properties of N-propargylamine, were recently shown to induce a significant down-regulation of membrane-associated holo-APP levels in the mouse hippocampus and in human SH-SY5Y neuroblastoma cells, presumably by chelating intracellular iron pools [51]. Indeed, the iron chelator drugs VK28, HLA20, and M30 (Figure 1) were found to suppress translation of a luciferase reporter mRNA via the APP 5'UTR sequence (Table 1) [51]. Furthermore, M30 markedly reduced the levels of the amyloidogenic Aβ in the medium of CHO cells stably transfected with the APP 'Swedish' mutation (CHO/ $\Delta$ NL) [51]. In addition, naturally occurring polyphenols – for example, EGCG ((-)-epigallocatechin-3-gallate)*a*nd curcumin – might be used as another novel and promising therapeutic approach for treating AD. Both compounds have well characterized antioxidant



### **Figure 4** Production of the contract of the c

Schematic representation of physiological iron homeostasis mechanisms, including Aβ generation as a compensatory process that reduces OS damage and the pathological mechanisms of iron-induced neurodegeneration in AD and its prevention by iron chelators.

and metal-chelating (iron and copper) activities [52-54] and have been demonstrated to exert neuroprotective activity against a variety of neurotoxic insults, as well as to regulate APP processing and Aβ burden in cell culture and *in vivo* [55]. EGCG treatment led to a reduction of Aβ levels in CHO/ΔNL [56], murine neuron-like cells (N2a) transfected with the APP 'Swedish' mutation and primary neurons derived from 'Swedish' mutant APP-over-expressing mice [57]. *In vivo*, EGCG significantly reduced cerebral Aβ levels concomitant with reduced Aβ amyloid plaques in TgAPPsw transgenic mice overproducing Aβ [57].

Our recent studies have shown that prolonged administration of EGCG to mice induced a reduction in holo-APP levels in the hippocampus [58]. In SH-SY5Y cells, EGCG significantly reduced both the mature and full-length cellular holo-APP without altering APP mRNA levels, as shown by two-dimensional gel electrophoresis, suggesting a post-transcriptional action [56]. Indeed, we demonstrated that EGCG reduced the translation of a luciferase reporter gene fused to the APP mRNA 5'UTR, which includes the APP IRE [56,58]. The observation that the alteration in APP following treatment with EGCG was blocked by exogenous iron provides further support to the implication of the metal-chelating property of EGCG in the regulation of iron homeostasis-associated proteins. A recent study reported the development of a high-throughput screen (library of 110,000 compounds from the Laboratory for Drug Discovery on Neurodegeneration) to identify APP mRNA 5'UTR-directed compounds that may be developed into therapeutic agents for AD [59]. Table 2 summarizes the effects of various compounds  $(1-50 \mu M)$ that were demonstrated to limit APP 5'UTR-conferred translation on holo-APP and Aβ levels.





Arrows indicate decreased levels. SSRI, selective serotonin reuptake inhibitor.

Finally, a new, additional aspect of iron chelator compounds in the etiology of AD therapy is related to their ability to abort anomalous cell cycle re-activation in postmitotic degenerating neurons. Indeed, during the last few years accumulating evidence for an activated cell cycle in the vulnerable neuronal population in AD has suggested a crucial role for cell cycle abnormalities in AD pathogenesis. Therefore, therapeutic interventions targeted towards ameliorating mitotic changes would be predicted to have a positive impact on AD progression. Previous studies have shown that the re-activation of the cell cycle is an obligatory component of the apoptotic pathway evoked by  $\overrightarrow{AB}$  [60-62]. Recently, we have found that M30 (0.1) μM) significantly reduced the percentage of neurons in S phase (approximately 50%), while increasing their relative cell number again in  $G_0/G_1$  phase (approximately 1.4fold) and lowered apoptotic levels after exposure to Aβ25–35 (25 μM) in primary cultures of rat cortical neurons. In support, the novel iron chelator drugs were previously shown to induce cell cycle arrest; M30 and HLA20 increased the number of PC12 cells in  $G_0/G_1$ , and decreased the cell number in S phase, as well as the proportion of cells in the  $G_2$  phase, further indicating that both compounds inhibited cell progress beyond the  $G_0/$  $G_1$  phase.

We recently presented a novel neuroprotective target for iron chelators with regard to the aberrant cell cycle reentry of postmitotic neurons in AD. Accordingly, similar to cancer drug therapy, a newly therapeutic strategy for neurodegenerative diseases is currently directed at interfering with mitogenic signaling and cell cycle progression to ameliorate cell death. Because iron chelators have been shown to affect critical regulatory molecules involved in cell cycle arrest and proliferation [55], a therapeutic intervention with these compounds is assumed to have a profound

impact on neuron preservation and AD progression. Indeed, our studies revealed that the multi-functional iron chelators M30 and HLA20 [51], as well as EGCG [63], induced differentiation features in neuroblastoma and PC12 cells, including cell body elongation, stimulation of neurite outgrowth, and up-regulation of the growth associated protein-43 (GAP-43). Taken together, the data suggest that iron chelators may be considered potential therapeutic agents in AD, targeting early cell cycle anomalies, and re-establishing the lost synaptic connection in the injured neuronal cells.

#### **Conclusion**

The presence of an IRE stem-loop in the APP transcript suggests that this ubiquitous membrane-associated protein, as well as its cleavage product, Aβ, may have a significant role in iron homeostasis, as already exemplified by other iron-associated proteins. Thus, regarding the physiological pathway, Aβ production may be considered a compensatory or neuroprotective response that reduces OS damage. In the pathological aspect, novel therapeutic strategies may comprise iron chelating agents targeted to the IRE in the 5'UTR of the APP mRNA, specifically preventing iron-induced toxicity and over-production of Aβ. We have recently designed and synthesized several novel antioxidant/iron chelators with an 8-hydroxyquinoline moiety, and demonstrated their capacity to lower the expression of APP and the generation of Aβ. These latest findings implicate the therapeutic potential of our drugs as iron-chelator candidates targeting the regulation of APP/Aβ in AD.

In addition, considering a recent report describing a putative IRE in the 5'UTR of Parkinson's disease related αsynuclein mRNA [64], a parallel can be drawn between APP and  $\alpha$ -synuclein both in the physiological and pathological aspects with respect to iron regulation. Indeed, it can be predicted that this RNA structure may have the potential to function as a post-transcriptional regulator of α-synuclein protein synthesis by age-related iron and redox pathophysiology upstream of neurodegeneration.

#### **List of abbreviations used**

Aβ: β-amyloid; AD: Alzheimer's disease; APP: amyloid precursor protein; CHO: Chinese hamster ovary; DFO: deferrioxamine; EGCG: (-)-epigallocatechin-3-gallate; IRE: iron responsive element; NFT: neurofibrillary tangle; OS: oxidative stress; ROS: reactive oxygen species; UTR: untranslated region.

#### **Competing interests**

The authors declare that they have no competing interests.

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**[for neuroprotection in Alzheimer's, Parkinson's, and other](http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?cmd=Retrieve&db=PubMed&dopt=Abstract&list_uids=15653345) [neurodegenerative diseases.](http://www.ncbi.nlm.nih.gov/entrez/query.fcgi?cmd=Retrieve&db=PubMed&dopt=Abstract&list_uids=15653345)** *Bioorg Med Chem* 2005, **13:**773-783.

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