

Different Inhibitors of A β 42-Induced Toxicity Have Distinct Metal-Ion DependencyAshley J. Mason,[#] Ian Hurst,[#] Ravinder Malik, Ibrar Siddique, Inna Solomonov, Irit Sagi, Frank-Gerrit Klärner, Thomas Schrader, and Gal Bitan*Cite This: *ACS Chem. Neurosci.* 2020, 11, 2243–2255

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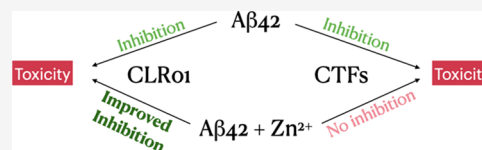


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ABSTRACT: Oligomers of amyloid β -protein (A β) are thought to be the proximal toxic agents initiating the neuropathologic process in Alzheimer's disease (AD). Therefore, targeting the self-assembly and oligomerization of A β has been an important strategy for designing AD therapeutics. In parallel, research into the metallobiology of AD has shown that Zn²⁺ can strongly modulate the aggregation of A β in vitro and both promote and inhibit the neurotoxicity of A β , depending on the experimental conditions. Thus, successful inhibitors of A β self-assembly may have to inhibit the toxicity not only of A β oligomers themselves but also of A β -Zn²⁺ complexes. However, there has been relatively little research investigating the effects of A β self-assembly and toxicity inhibitors in the presence of Zn²⁺. Our group has characterized previously a series of A β 42 C-terminal fragments (CTFs), some of which have been shown to inhibit A β oligomerization and neurotoxicity. Here, we asked whether three CTFs shown to be potent inhibitors of A β 42 toxicity maintained their activity in the presence of Zn²⁺. Biophysical analysis showed that the CTFs had different effects on oligomer, β -sheet, and fibril formation by A β 42-Zn²⁺ complexes. However, cell viability experiments in differentiated PC-12 cells incubated with A β 42-Zn²⁺ complexes in the absence or presence of these CTFs showed that the CTFs completely lost their inhibitory activity in the presence of Zn²⁺ even when applied at 10-fold excess relative to A β 42. In light of these results, we tested another inhibitor, the molecular tweezer CLR01, which coincidentally had been shown to have a high affinity for Zn²⁺, suggesting that it could disrupt both A β 42 oligomerization and A β 42-Zn²⁺ complexation. Indeed, we found that CLR01 effectively inhibited the toxicity of A β 42-Zn²⁺ complexes. Moreover, it did so at a lower concentration than needed for inhibiting the toxicity of A β 42 alone. In agreement with these results, CLR01 inhibited β -sheet and fibril formation in A β 42-Zn²⁺ complexes. Our data suggest that, for the development of efficient therapeutic agents, inhibitors of A β self-assembly and toxicity should be examined in the presence of relevant metal ions and that molecular tweezers may be particularly attractive candidates for therapy development.



KEYWORDS: Amyloid, Alzheimer's disease, metal ions, inhibitor, aggregation, neurotoxicity

■ INTRODUCTION

Amyloid β -protein (A β) self-assembly into neurotoxic oligomers and aggregates is a seminal pathological process in Alzheimer's disease (AD).^{1,2} Although deposition of fibrillar A β aggregates in amyloid plaques originally led to the *Amyloid Cascade Hypothesis*,^{3,4} nonfibrillar A β oligomers are now considered by most researchers to be the true culprit causing the initial disruption of synaptic communication and instigating the cascade of events that leads to AD.^{5–11} The other offending protein self-assembly process in AD, that of hyperphosphorylated tau, is believed to be secondary to the initial A β insults.¹² Therefore, targeting A β oligomers, particularly those of the longer and more toxic form, A β 42, is a popular strategy for therapy development.¹³

In parallel, research into the metallobiology of AD has identified several metal ions that interact with A β , modulate its self-assembly and toxicity, and may be important for the onset and progression of the disease.^{14,15} Metal cations, such as Zn²⁺, Cu²⁺, and Fe³⁺, have been reported to stimulate the deposition of A β and promote the hyperphosphorylation of tau.^{16,17} Much

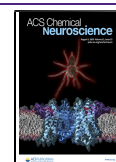
of the research has focused on the effects of Cu²⁺ and Zn²⁺ due to their role in the maintenance of neuronal excitability, their function as cofactors, and their contribution to oxidative stress and inflammation in the AD brain.¹⁸ In addition, both ions have been shown to modulate the oligomerization and aggregation of A β itself.^{19–22}

Zn²⁺ has been of particular interest due to its presence in deposited amyloid plaques and its concentration in glutamatergic neurons in the hippocampus.^{23,24} Interestingly, Zn²⁺ ions have been reported to both accelerate and inhibit A β aggregation and both increase A β neurotoxicity and protect against A β neurotoxicity, depending on the specific conditions used by each group, including the concentration of A β and

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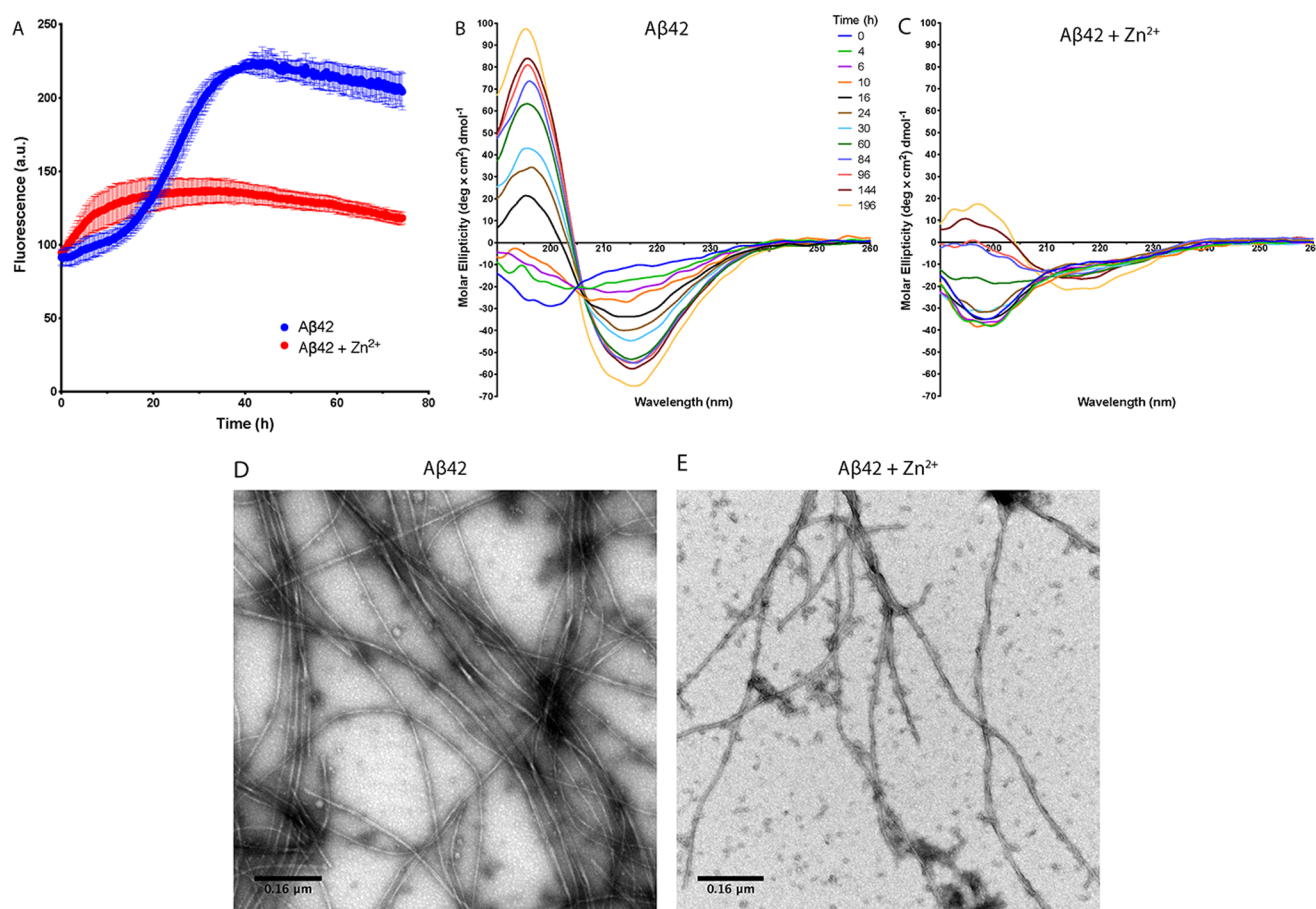


Figure 1. Aβ42 aggregates differently in the absence or presence of Zn²⁺. Twenty micromolar Aβ42 was incubated in the absence or presence of 40 μM Zn²⁺. (A) ThT fluorescence was measured every 15 min in five technical replicates per condition. (B, C) Representative CD spectra in aggregation reactions of Aβ42 (B) and Aβ42-Zn²⁺ (C). (D, E) Representative electron micrographs of Aβ42 (D) and Aβ42-Zn²⁺ (E) after 408 h of incubation. Scale bars represent 0.16 μM.

Zn²⁺, the ratio between these concentrations, and the solution conditions.^{24–29} Indeed, our own work showed that Zn²⁺ accelerated formation of nonfibrillar, yet β-sheet rich, aggregates of Aβ40 and that the toxicity of Aβ40-Zn²⁺ complexes depended on the method of their addition to cultured neurons.³⁰

In the AD brain, high concentrations (~1 mM) of Zn²⁺ have been found in amyloid plaques along with elevated levels in the hippocampus and serum of patients.³¹ Zn²⁺ concentrations up to 300 μM have been measured in synaptic vesicles^{32,33} and during synaptic transmission, and direct imaging of Zn²⁺ using fluorescent probes has shown a release of 10–30 μM of Zn²⁺ into the synaptic cleft.³⁴ It is predominantly under these circumstances that free Zn²⁺ is thought to associate with Aβ^{32,35–37} and induce the formation of neurotoxic assemblies.^{32,38} Previously, we found that Aβ40-Zn²⁺ complexes were β-sheet-rich, nonfibrillar, quasi-spherical aggregates that suppressed spontaneous Ca²⁺ transients and were toxic to primary hippocampal neurons when added at small concentrations in several portions, but not when added at a higher concentration in the beginning of the experiment.³⁰ These findings led us to ask how the presence of Zn²⁺ might impact the interaction of Aβ assembly inhibitors/modulators with Aβ and whether such inhibitors maintained their inhibitory activity in the presence of Zn²⁺.

Therapy development efforts targeting Aβ–metal-ion interactions have focused on chelators, such as clioquinol, a hydroxyquinoline-derived drug, which was shown to prevent cognitive decline in Zn²⁺-transporter knockout mice,^{39,40} and its successor, PBT2, which was found to lower Aβ levels in the CSF of patients with AD.⁴¹ More recently, several compounds have been reported to have both metal-ion chelating activity and act as Aβ assembly inhibitors regardless of metal binding.^{42–49} However, the latter activity may arise from the weak nature of the forces mediating Aβ oligomerization, which is easily modulated by many small molecules nonspecifically,¹³ potentially through the formation of colloids.⁵⁰

Previously, we reported that certain C-terminal fragments (CTFs) of Aβ42 interacted with the full-length peptide, modulated its oligomerization, and inhibited its toxicity.^{51–54} In particular, out of 15 peptides with the general formula Aβ(x–42), where x = 28–39, Aβ(31–42) and Aβ(39–42) had notably high inhibitory activity. In addition, an Aβ40 CTF that originally was tested as a control, Aβ(30–40), was found to be a potent inhibitor of Aβ42-induced toxicity.⁵⁵ Thus, we decided to use these CTFs as a test case and determine how their inhibitory activity might be affected by the presence of Zn²⁺. We studied the effect of the CTFs on the conformational change kinetics and morphology of Aβ42-Zn²⁺ complexes and on their toxicity in cell culture. We show that the CTFs' effects on Aβ42 β-sheet formation, morphology, and cytotoxicity are

dramatically altered by the interactions of A β 42 with Zn²⁺. In light of these results, we also tested another inhibitor of A β self-assembly and toxicity, the molecular tweezer CLR01, and found that its activity also was affected by the presence of Zn²⁺, but in a distinct way.

RESULTS AND DISCUSSION

Zn²⁺ Alters A β 42 Assembly and Toxicity. Because the CTFs were originally conceived as potential inhibitors of A β 42, here we characterized first the aggregation and toxicity of A β 42 in the presence of Zn²⁺. In our previous study of A β 40-Zn²⁺ complexes, the biophysical characterization of the system was done primarily in a 3-(*N*-morpholino)propanesulfonic acid (MOPS) buffer.³⁰ An interesting finding was that the addition of Zn²⁺ reduced the increase in thioflavin T (ThT) fluorescence even though circular dichroism (CD) spectroscopy showed that A β 40-Zn²⁺ complexes were rich in β -sheet. Therefore, we used both techniques here again to assess the temporal conformational changes in A β 42 in the presence of Zn²⁺. However, initial attempts to characterize the temporal changes of A β 42-Zn²⁺ complexes in MOPS buffer using CD spectroscopy yielded low signal-to-noise ratio spectra, because MOPS strongly absorbs light below 200 nm.⁵⁶ For this reason, we decided to change the buffer from MOPS to phosphate buffer (PB). Using PB, we found that reproducibility was low and data interpretation was difficult when using a concentration of A β 42 below 20 μ M. Therefore, although lower concentrations were used in our toxicity assays, the investigation of A β 42 aggregation and conformational change in the presence of Zn²⁺ and the effect of the CTFs on this process was done using 20 μ M A β 42. To be able to compare these experiments with the toxicity assays, the A β 42/Zn²⁺ concentration ratio was kept at 1:2, respectively, in all experiments.

In the absence of Zn²⁺, A β 42 showed a two-phased sigmoidal curve, consisting of a short lag phase of \sim 3 h, a relatively slow increase up to \sim 14 h, and a faster increase reaching a plateau at \sim 42 h (Figure 1A). This rate of ThT fluorescence increase was slow relative to other reports in the literature, including from our own group, even when using lower concentrations of A β 42.^{24,27,57,58} In our experience, many factors influence the rate of the increase, including the peptide source and the exact preparation method, including the steps used for disaggregation and solubilization, peptide concentration, buffer type and concentration, salt type and concentration, pH, temperature, agitation method, and agitation rate. As explained above, we chose the conditions for these experiments to allow meaningful comparisons with the conditions in which Zn²⁺ and/or CTFs were added to the reaction mixture, taking into account that precise quantitative comparison with the multiple previous reports in the literature may not be possible due to the different conditions used in each study.

In contrast to A β 42 alone, in the presence of 2 equiv of Zn²⁺, no lag phase was observed. Rather, the ThT fluorescence increased rapidly during the first 8 h and reached a maximum at \sim 16 h. However, the maximum fluorescence was substantially lower than the value for A β 42 alone (Figure 1A), in agreement with the behavior of A β 40 in the presence of Zn²⁺ reported previously.³⁰

CD spectroscopy analysis of the A β 42 conformation in the absence of Zn²⁺ showed a gradual transition from a predominantly statistical coil spectrum at $t = 0$, characterized

by a minimum at 198–200 nm, to a primarily β -sheet spectrum at 196 h, displaying the typical maximum at 195–196 nm and minimum at 215–218 nm (Figure 1B). In the presence of Zn²⁺, an overall transition from statistical coil to β -sheet also was observed (Figure 1C), but with two important differences. First, unlike the gradual conformation change observed in the spectra of A β 42 alone already at early time points, in the presence of Zn²⁺ the spectra remained largely unchanged for the first 24 h, suggesting a delay in the conformational transition. Second, the amplitude of the spectra in the presence of Zn²⁺ was substantially reduced compared to the spectra of A β 42 alone, suggesting overall smaller changes in the presence of Zn²⁺. These data suggested that, upon addition of Zn²⁺ to A β 42, A β 42-Zn²⁺ complexes formed rapidly, and although they changed over time, the change involved relatively minor conformational rearrangements.

Interestingly, the rapid increase in ThT fluorescence (Figure 1A) disagreed with the small change in conformation observed by CD during this time (Figure 1B), suggesting that, in the presence of Zn²⁺, ThT might bind rapidly to A β 42-Zn²⁺ complexes without formation of substantial β -sheet conformation. The binding may involve a ternary complex formed by the protein, the metal ion, and the dye molecules, yet it is an atypical phenomenon that does not actually report on the formation of a cross- β structure or of amyloid fibrils, as would be expected in a typical ThT-fluorescence experiment. In agreement with this interpretation, the overall magnitude of the change in fluorescence during the reaction was only \sim 30% compared to A β 42 alone. An alternative explanation might stem from formation of insoluble aggregates that bind ThT but have little impact on the CD spectrum. We did not observe particulate matter in the solutions in either experiment, but we cannot rule out the formation of insoluble, ThT-positive aggregates that were too small to detect by the naked eye.

Morphological examination of each preparation by transmission electron microscopy (TEM) at intermediate time points (Supporting Information Figure S1) and at the end of the aggregation reaction (Figure 1D,E) showed that, during the reaction, oligomers and fibrils were observed, whereas at the end of the aggregation reaction, abundant fibrils were present in both cases. However, the abundance was higher in samples containing A β 42 alone, in which some fibrils appeared thicker and comprised multiple thin filaments. Quasi-globular structures, presumably oligomers, also were observed but were relatively sparse in samples of A β 42 alone. In the samples containing A β 42-Zn²⁺ complexes, the fibrils were thinner on average than in those containing A β 42 alone, and twisted fibrils appeared to comprise mainly two intertwined threads. The abundance of quasi-globular structures and amorphous aggregates in these samples was substantially higher than in samples of A β 42 alone (Supporting Information Figure S1 and Figure 1D,E).

To measure the toxicity of A β 42 in the absence or presence of Zn²⁺, we employed a strategy developed in our previous study of A β 40-Zn²⁺ complexes.³⁰ In that study, we found that, if 10 μ M A β 40-Zn²⁺ complexes were added to primary rat hippocampal neurons all at once, they did not cause measurable toxicity 48 h later, likely because under these conditions the formation of toxic structures was transient and too short before the complexes transformed into larger, nontoxic aggregates.³⁰ In contrast, if the sample was divided into four equal portions, each of which was prepared freshly and added 12 h apart, more than 80% of the cells were found

to be dead at 48 h using a neuron-specific enolase assay. Here, we used differentiated PC-12 cells and the lactate dehydrogenase (LDH) assay to measure cell viability. First, we confirmed that upon addition of $A\beta_{42}$ - Zn^{2+} complexes in one portion to the cells, little toxicity was observed after a 48 h incubation ($3.8 \pm 2.0\%$ cell death, Figure 2) compared to $A\beta_{42}$ alone

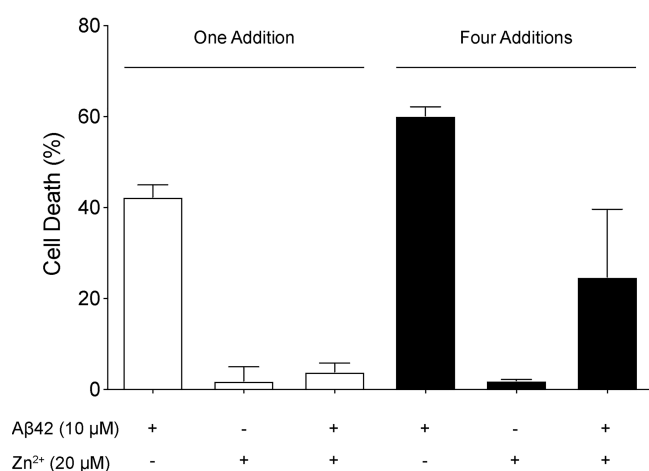


Figure 2. $A\beta_{42}$ - Zn^{2+} complexes are toxic when added in small portions. $A\beta_{42}$ was prepared in the absence or presence of Zn^{2+} , and the same total amount was added to differentiated PC-12 cells either at time 0 or in four equal portions at times 0, 12, 24, and 36 h. Cell death was measured using the LDH assay at 48 h. Each column is an average of three or four biological replicates, each done in six technical replicates. The error bars represent the standard error of measure (SEM).

($42.3 \pm 2.7\%$ cell death). Zn^{2+} on its own was not toxic under these conditions ($1.9 \pm 3.1\%$ cell death). Therefore, we adopted the previously established protocol,³⁰ added a fresh aliquot equal to a quarter of each sample at $t = 0, 12, 24$, and 36 h, and measured cell viability at 48 h. Under these

conditions $A\beta_{42}$ showed increased toxicity ($60.1 \pm 2.0\%$ cell death), Zn^{2+} was not toxic ($1.9 \pm 0.3\%$ cell death), and $A\beta_{42}$ - Zn^{2+} complexes gave somewhat variable results, but overall were ~ 6 times more toxic than when added in one portion ($24.7 \pm 14.9\%$ cell death, Figure 2). Therefore, we used the four-portion paradigm in all the toxicity-inhibition experiments.

CTFs Have Distinct Effects on the Self-Assembly of $A\beta_{42}$ - Zn^{2+} Complexes.

To test the effect of $A\beta(39-42)$, $A\beta(31-42)$, and $A\beta(30-40)$ on the aggregation of $A\beta_{42}$ - Zn^{2+} complexes, we incubated mixtures of 20 μM $A\beta_{42}$, 40 μM Zn^{2+} , and 100 μM of each CTF at 37 °C with agitation and measured the change in ThT fluorescence and CD spectra during the aggregation reactions. In control ThT experiments, the CTFs were examined on their own or in the presence of Zn^{2+} at the same concentrations. Compared to $A\beta_{42}$ - Zn^{2+} alone (Figure 3A), the ThT signal of $A\beta_{42}$ - Zn^{2+} in the presence of $A\beta(39-42)$ increased initially at a similar rate, whereas the final fluorescence was $\sim 25\%$ lower (Figure 3A, blue curve). In contrast, a markedly different behavior was observed in the presence of $A\beta(31-42)$ or $A\beta(30-40)$. In both cases, the initial fluorescence was ~ 5 times higher than in the presence of $A\beta_{42}$ - Zn^{2+} alone or $A\beta_{42}$ - Zn^{2+} in the presence of $A\beta(39-42)$, suggesting a rapid formation of quaternary complexes comprising $A\beta_{42}$, Zn^{2+} , the CTF, and ThT. This interpretation was supported by the observation that Zn^{2+} had a minimal impact on the behavior of the CTFs on their own (Figure 3B). $A\beta(39-42)$ and $A\beta(31-42)$ showed essentially no change in fluorescence during the experiment, regardless of the presence of Zn^{2+} . In contrast, the fluorescence of $A\beta(30-40)$ showed a two-step increase followed by a slow decrease after 24 h. In the presence of Zn^{2+} the initial increase in ThT fluorescence was accelerated and was followed by a decrease up to ~ 10 h and then a plateau (Figure 3B).

In the mixtures with $A\beta_{42}$, $A\beta(31-42)$ caused a rapid increase in the ThT fluorescence signal, which peaked at ~ 18 h and then declined steadily until the end of the experiment

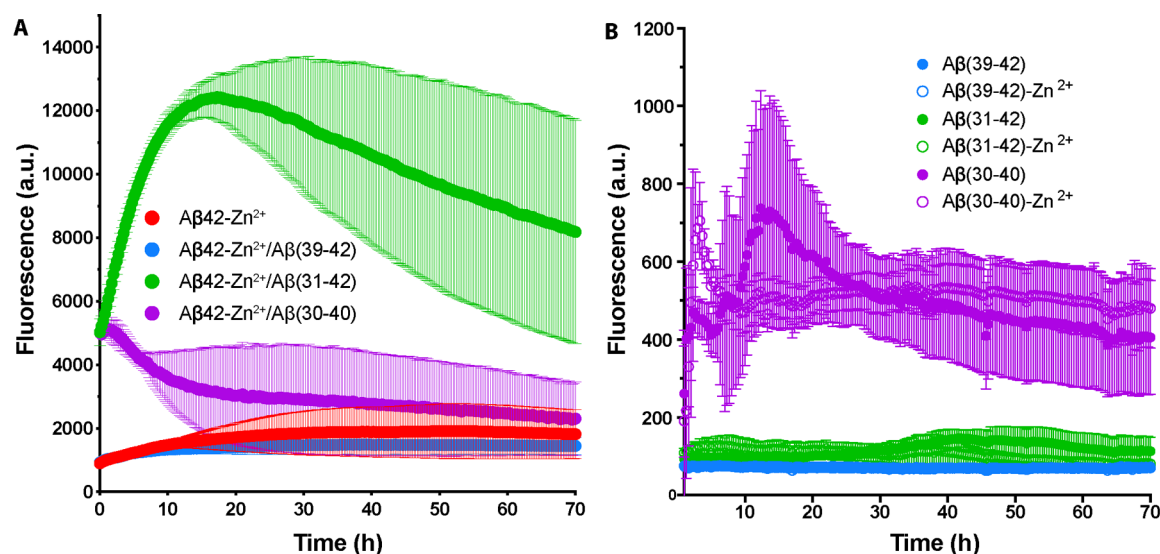


Figure 3. Time-dependent ThT fluorescence of $A\beta_{42}$ - Zn^{2+} complexes in the presence of CTFs. (A) $A\beta_{42}$ (20 μM) and 40 μM Zn^{2+} were incubated in the absence or presence of 100 μM of each CTF, and the change in ThT fluorescence was monitored every 15 min and averaged for five technical replicates per condition. (B) The CTFs in the absence of $A\beta_{42}$ were monitored at the same concentrations under the same conditions (though using a different plate reader, reflected by the different scales of the arbitrary fluorescence units on the y-axis of the two panels). Error bars represent the standard deviation (SD).

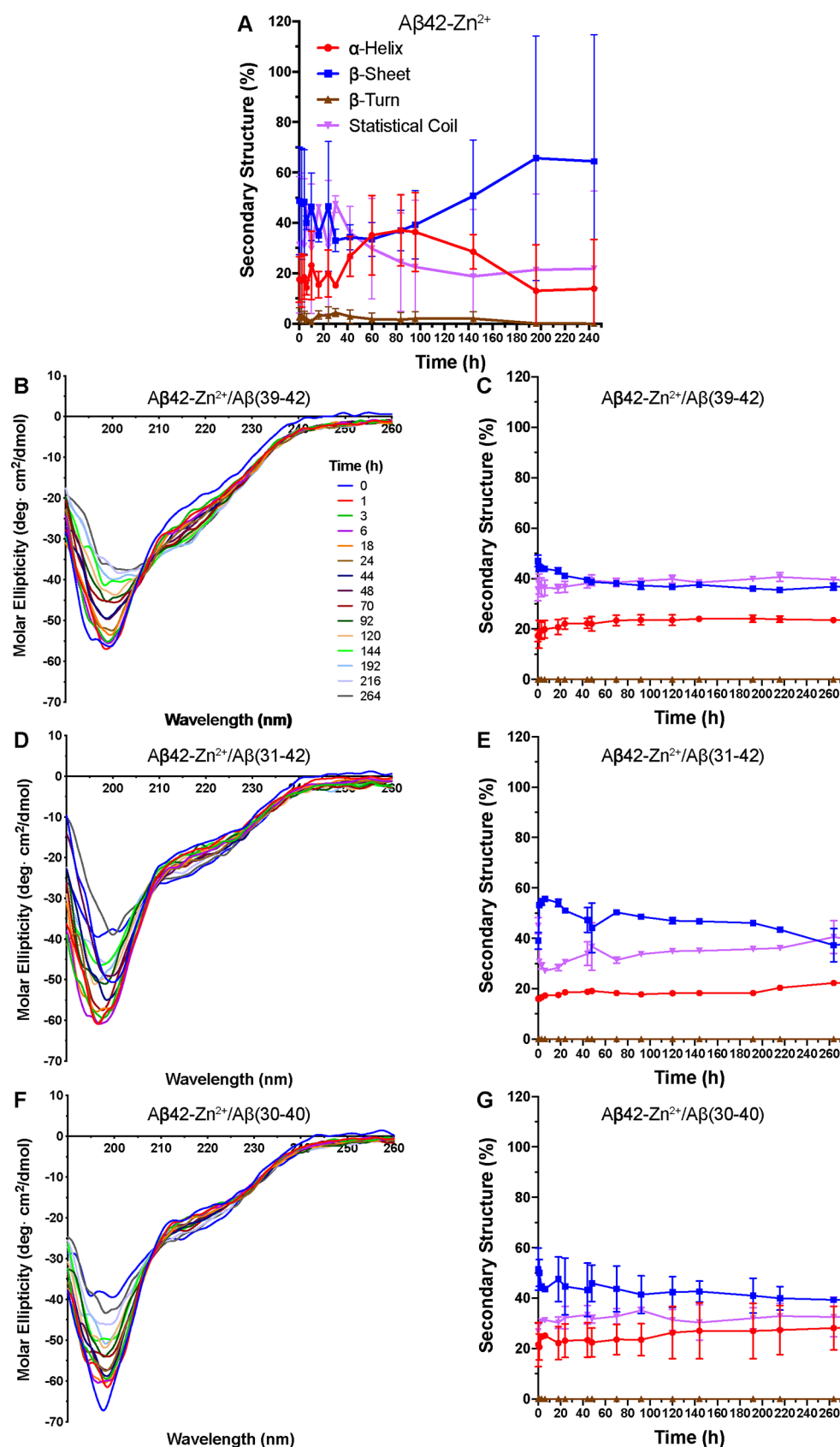


Figure 4. CD-spectroscopy analysis of $A\beta 42-Zn^{2+}$ complexes in the presence of CTFs. Twenty micromolar $A\beta 42$ and $40 \mu M Zn^{2+}$ were incubated in the absence or presence of $100 \mu M$ of each CTF, and the change in CD spectrum was monitored at the times indicated in (B), which apply also to (D, F). (A) Deconvolution of the CD spectra shown in Figure 1C. (C, E, G) Deconvolution of the spectra shown in (B, D, F), respectively.

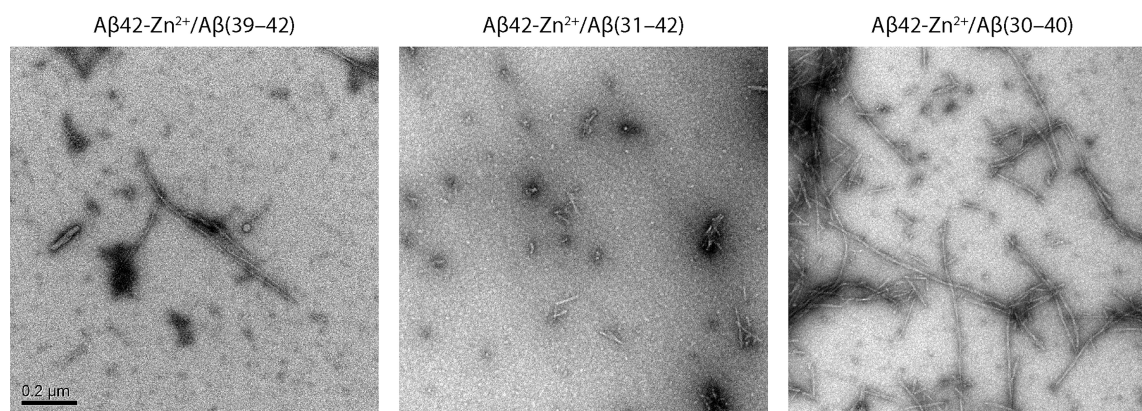


Figure 5. Morphological analysis of $A\beta_{42}$ - Zn^{2+} complexes in the presence of CTFs. Twenty micromolar $A\beta_{42}$ and $40\ \mu M\ Zn^{2+}$ were incubated in the absence or presence of $100\ \mu M$ of each CTF. The morphology of the mixture was assessed using EM after 17 d of incubation. Representative electron micrographs are shown. The scale bar in the left image represents $0.2\ \mu m$ and is applicable to all the images.

(Figure 3A). In the presence of $A\beta(30-40)$ (Figure 3A), the fluorescence increased slightly in the first 90 min, then declined rapidly for the next ~ 16 h, followed by a slower decline until the end of the experiment, at which point, the fluorescence was only a little higher than that of $A\beta_{42}$ - Zn^{2+} alone. Comparison of Figures 3A and B suggested that this behavior might have been influenced by the CTF itself and might not have reflected formation of a complex involving $A\beta_{42}$ and $A\beta(30-40)$, though formation of a complex could not be ruled out (the experiments in Figures 3A and B were done using different plate readers, and therefore the comparison is qualitative rather than quantitative). Interestingly, in the presence of $A\beta_{42}$, in both the cases of $A\beta(31-42)$ and $A\beta(30-40)$, the behavior of the different replicates was highly consistent for the first 8 h of the experiment but then diverged substantially (Figure 3A). Nonetheless, despite the relatively low reproducibility among replicates between 8 and 72 h, the distinction among the behavior of the different mixtures was clear. These peculiar behaviors supported the idea that in these experiments ThT fluorescence either reflected formation of complexes that were different from the typical binding of the dye molecules to cross- β structures of amyloid fibrils or, in the case of $A\beta(30-40)$, the ThT-fluorescence curve was dominated by the behavior of the CTF itself.

To further assess the change in secondary structure of $A\beta_{42}$ - Zn^{2+} complexes in the presence of CTFs, we examined the temporal changes in CD spectra of the same mixtures (prepared together but incubated without ThT) and deconvoluted the spectra to gain additional insight. CD spectra of the CTFs themselves have been reported previously⁵³ and therefore were not recorded here. Supporting the interpretation that the ThT data reflected atypical complexes rather than cross- β structures, unlike their distinct behavior in the ThT-fluorescence experiments, all three CTFs showed roughly similar attenuation of the conformational change in CD spectroscopy (Figure 4B,D,F). In all cases, the initial spectra were characteristic of a statistical coil. Over 11 d of incubation, relatively small changes were observed in the presence of the CTFs. The magnitude of the minimum at 195–198 nm decreased, and the molar ellipticity at the typical ~ 215 nm minimum characteristic of a β -sheet slightly increased, yet those changes were relatively minor, and the final spectra suggested that the peptides ($A\beta_{42}$ and CTF) were

still largely unstructured in these complexes. Again, dissimilarity between the ThT and CD data might reflect the formation of ThT-positive insoluble aggregates that did not contribute to the CD spectra and were too small to detect by the naked eye.

To quantify the change, we deconvoluted the spectra using the Savitzky-Golay algorithm. In the absence of CTFs, $A\beta_{42}$ - Zn^{2+} complexes showed a somewhat variable behavior (Figure 4A; the data are an average of three experiments, including the one presented in Figure 1C). The conformational transition involved a temporary formation of an α -helical intermediate between 30 and 90 h, which decreased at later time points, concomitant with an increase in β -sheet content and a decline in statistical coil, as described previously for $A\beta$ in the absence of Zn^{2+} .⁵⁹ In the presence of the CTFs, the variability between experiments was substantially lower, and the deconvolution showed only minor conformational changes during the reactions. In the presence of $A\beta(39-42)$, which itself is expected to be unstructured,⁵³ the mixture comprised $\sim 40\%$ each statistical coil and β -sheet, and $\sim 20\%$ α -helix, with minimal changes in these fractions for the duration of the experiment (Figure 4C). In the presence of $A\beta(31-42)$, similar initial fractions were observed at $t = 0$ h. In the next 3 h, the abundance of β -sheet conformation increased to $\sim 55\%$, and that of statistical coil decreased to $\sim 35\%$, possibly reflecting formation of β -sheet in the CTF itself.⁵³ Then, this trend was slowly reversed during the next 261 h, until the β -sheet and statistical coil fractions were again $\sim 40\%$ each. The α -helix fraction remained at $\sim 20\%$ throughout the entire reaction (Figure 4E). $A\beta(30-40)$ caused formation of $\sim 50\%$ β -sheet immediately at $t = 0$ h, followed by a slow decrease to $\sim 40\%$ at 264 h, which was accompanied by minor increases in statistical coil, from 27% to 32%, and α -helix, from 21% to 28% (Figure 4G). Although the numbers generated by the deconvolution algorithm may not be precise, they provide an approximation of the contribution of each secondary structure component, supporting the conclusion that to a large extent the CTFs attenuated the conformational transition observed in $A\beta_{42}$ - Zn^{2+} complexes (Figures 1C and 4A).

Morphological analysis of the $A\beta_{42}$ - Zn^{2+} mixtures in the presence of each CTF at the end of the aggregation reaction showed mixtures of fibrillar and nonfibrillar structures (Figure 5). In all cases, the fibrils were shorter and less abundant than in mixtures of $A\beta_{42}$ - Zn^{2+} complexes formed in the absence of

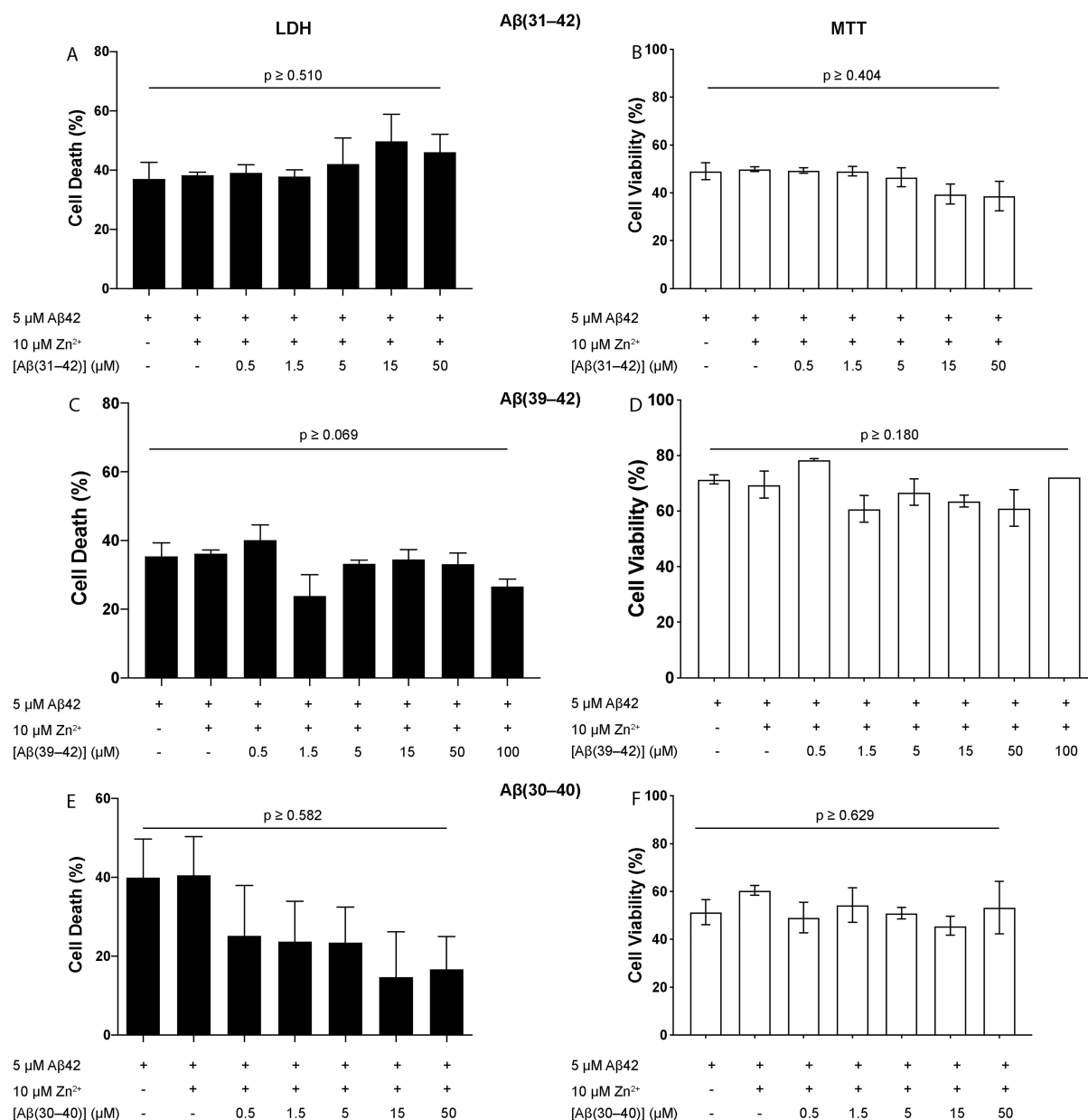


Figure 6. Evaluation of the cytotoxicity of Aβ42-Zn²⁺ complexes in the presence of CTFs. Five micromolar Aβ42 and 10 μM Zn²⁺ were prepared in the absence or presence of increasing concentrations of each CTF and added to differentiated PC-12 cells in four equal portions at times 0, 12, 24, and 36 h. (A, C, E) Cell death was measured at 48 h using the LDH-release assay. (B, D, F) Cell viability was measured at 48 h using the MTT-reduction assay. The data represent three biological replicates, each done with six technical replicates. Error bars indicate SEM. The *p*-values were calculated by a one-way ANOVA with a post hoc Tukey test for multiple comparisons. The value indicated in each panel is for the smallest *p*-value comparing any two columns.

CTFs. The shortest and thinnest fibrils appeared to form in the presence of Aβ(31-42) (Figure 5B), whereas the longest and most abundant fibrils formed in the presence of Aβ(30-40) (Figure 5C). These fibrils were still shorter than those formed by Aβ42 in the presence of Zn²⁺ in the absence of CTFs (Figure 1E), whereas in the presence of Aβ(39-42) an intermediate effect was observed. These results suggest that the C-terminal residues of Aβ42, namely, Ile⁴¹-Ala⁴², which are missing in Aβ(30-40), contribute to the interference of the CTFs with fibril formation. The relatively low abundance of the fibrils agreed, in general, with the CD spectroscopy data, though the differences in morphology among the mixtures containing different CTFs were more pronounced than the

differences in conformational transition as reflected by the CD spectra.

CTFs Do Not Inhibit the Toxicity of Aβ42-Zn²⁺ Complexes. We then used the LDH-release and 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide (MTT)-reduction assays to assess if the addition of Zn²⁺ to Aβ42 would impact the inhibitory activity of CTFs. In our previous work, Aβ(31-42) and Aβ(39-42) were shown to inhibit the toxicity of 5 μM Aβ42 with half-maximal inhibition (IC₅₀) values of 20 ± 4 and 47 ± 14 in the LDH-release assay,⁵¹ whereas Aβ(30-40) yielded IC₅₀ = 29 ± 4 μM in the LDH assay under similar conditions.⁵⁴ Remarkably, in the presence of Zn²⁺ (i.e., complexes comprising 5 μM Aβ42 and 10 μM Zn²⁺), all three CTFs lost completely their inhibitory

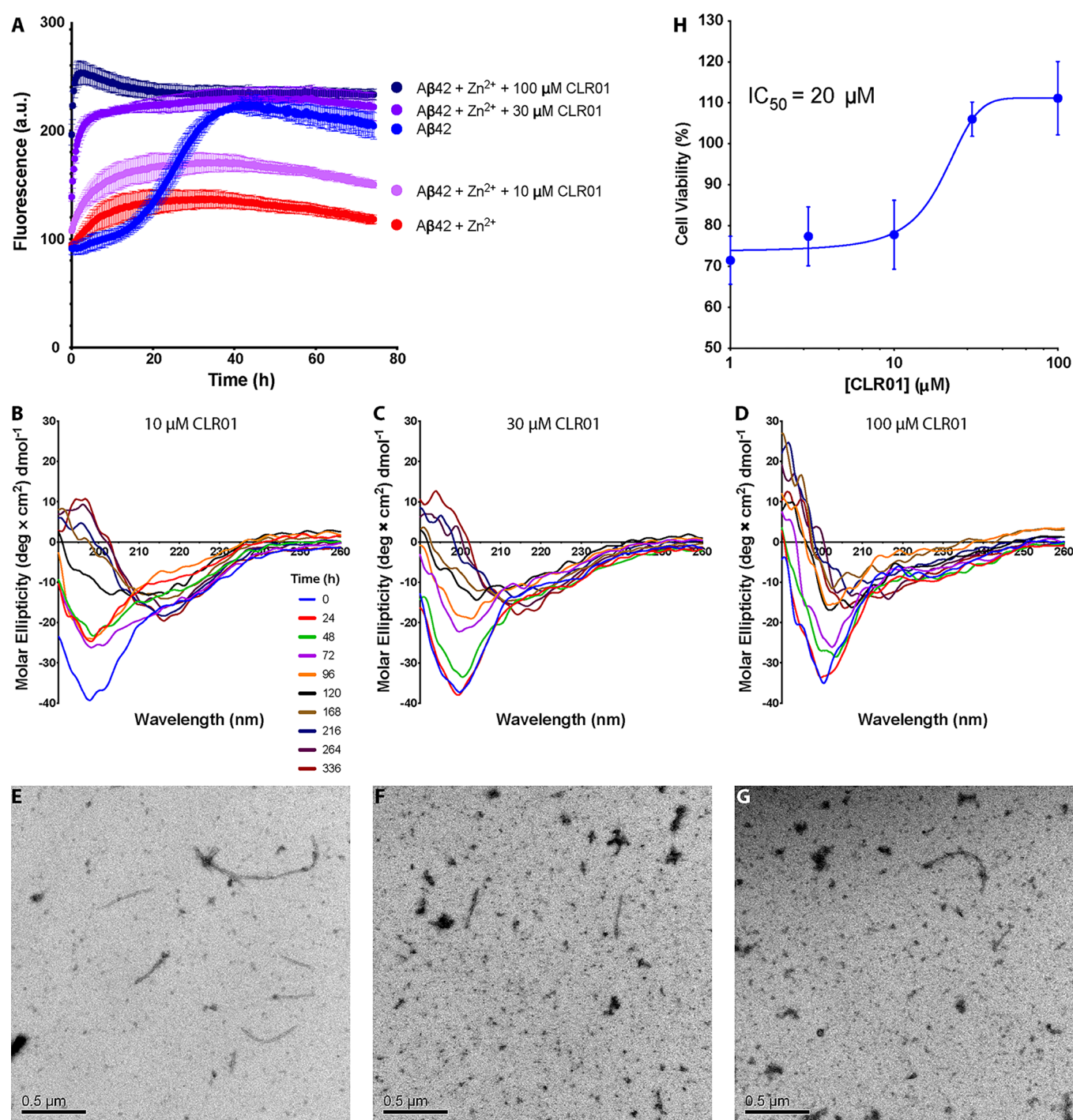


Figure 7. Effect of CLR01 on secondary structure, morphology, and toxicity of $A\beta 42$ - Zn^{2+} complexes. (A) Twenty micromolar $A\beta 42$ and 40 μM Zn^{2+} were incubated in the absence or presence of 10, 30, or 100 μM of CLR01, and the change in ThT fluorescence was monitored every 15 min and averaged for five technical replicates per condition. Error bars represent SD. (B–D) CD spectra of aggregation reactions, similar to those in (A) and monitored at the times indicated in (B), which apply also to (C, D). (E–G) Electron micrographs of the same reaction mixtures used for the corresponding CD experiments at the end of the reaction. (H) Five micromolar $A\beta 42$ and 10 μM Zn^{2+} were dissolved in the absence or presence of increasing concentrations of CLR01 and added to differentiated PC-12 cells in four equal portions at times 0, 12, 24, and 36 h. Cell viability was measured at 48 h using the MTT assay. Each data point represents three or four biological replicates, each done with six technical replicates. IC_{50} was calculated using Prism 8.4.0 by nonlinear fitting with four-parameter variable slope. The program did not allow calculating the error of the IC_{50} value due to the magnitude of the experimental error.

activity even when used at 10-fold excess (20-fold excess in the case of $A\beta(39-42)$, Figure 6), suggesting that the structures of the $A\beta 42$ - Zn^{2+} complexes and/or their interaction with the cells were distinct from those of $A\beta 42$ oligomers formed in the absence of Zn^{2+} .

CLR01 Modulates $A\beta 42$ Assembly and Inhibits Its Toxicity in the Presence of Zn^{2+} . In light of these results, we asked whether other inhibitors acting by different mechanisms could still inhibit the formation or toxicity of the $A\beta 42$ - Zn^{2+} complexes and tested whether a similar loss of

inhibition would be observed for the molecular tweezer CLR01. CLR01 is a broad-spectrum inhibitor,^{60–62} which previously has been shown to prevent the conformational change of A β 42 in ThT-fluorescence and CD-spectroscopy assays and inhibit the toxicity of A β 42 in differentiated PC-12 cells with IC₅₀ = 52 \pm 18 μ M in the MTT-reduction assay.⁵⁷ CLR01 is a particularly interesting test case for inhibiting the toxicity of A β 42-Zn²⁺ complexes, because it could target both A β 42 and Zn²⁺ simultaneously. Its binding to Zn²⁺ was reported by Wilch et al., who discovered that CLR01 bound Zn²⁺ with a relatively high affinity, K_d = 5 μ M, as opposed to a number of other metal ions, including Fe³⁺, Cu²⁺, Cs⁺, Ca²⁺, and Mg²⁺, for which CLR01 had low or no affinity.⁶³ Interestingly, binding of CLR01 to Zn²⁺ appeared to be cooperative, rather than competitive, with binding to Lys. ¹H NMR spectroscopy and fluorescence titrations suggested that CLR01 chelated Zn²⁺ by both phosphate groups while maintaining binding to a Lys side-chain inside its cavity.⁶³ However, how simultaneous binding to Lys and Zn²⁺ affects the affinity of CLR01 for either ligand is not known.

The impact of 10, 30, or 100 μ M CLR01 on temporal changes of ThT fluorescence in a solution of complexes made of 20 μ M A β 42 and 40 μ M Zn²⁺ showed a rapid increase in ThT fluorescence without a lag phase, similar to A β 42-Zn²⁺ in the absence of CLR01 (Figure 7A). The slope and final fluorescence value increased dose-dependently with CLR01 concentration. This behavior was distinct from the effect of CLR01 on the behavior of A β 42 in the ThT-fluorescence assay in the absence of Zn²⁺, which was a typical inhibition, that is, a longer lag time, slower increase, and reduced final fluorescence at a 1:1 concentration ratio and near quantitative inhibition (no increase in fluorescence) at 3- or 10-fold excess CLR01.⁵⁷ However, the rapid increase in ThT fluorescence without a lag phase, observed here upon addition of CLR01 to A β 42 in the presence of Zn²⁺, was similar to the effect of CLR01 on ThT fluorescence in the presence of other amyloidogenic proteins for which the rapid increase in fluorescence correlated with formation of nonfibrillar structures,^{57,64} suggesting that, in this case as well, the fluorescence increase reported on formation of nonamyloid complexes. In support of this interpretation, CD spectroscopy showed that the conformational transition in the presence of low concentrations of CLR01, 10 or 30 μ M (Figures 7B, C, respectively), resembled that of A β 42-Zn²⁺ complexes (Figure 1C) rather than that of A β 42 alone (Figure 1B). In the presence of 100 μ M CLR01, although a gradual change in the CD spectrum suggested a slow conformational change, the spectra were consistent with a statistical coil at all times and showed no formation of β -sheet. Thus, we conclude that, despite the sharp and rapid increase in ThT fluorescence in mixtures containing A β 42, Zn²⁺, and CLR01, the fluorescence does not reflect formation of a cross- β structure. In agreement with this interpretation, the morphologies observed in the samples at the end of 14 d of incubation consisted of a mixture of short, thin fibrils and oligomer-like structures when CLR01 was present at a substoichiometric concentration (Figure 7E), whereas mostly oligomers and very rare fibrillar structures were observed in the presence of 30 or 100 μ M CLR01 (Figures 7F and G, respectively).

Evaluation of the toxicity of A β 42-Zn²⁺ complexes in the presence of increasing CLR01 concentrations using the MTT reduction assay showed that, unlike the CTFs, CLR01 effectively inhibited the toxicity of A β 42 in the presence of Zn²⁺. Dose–Response analysis showed that the IC₅₀ of CLR01

in this assay was 20 μ M (Figure 7H). This value was \sim 2.5 times lower than the IC₅₀ for CLR01 inhibition of A β 42 in the absence of Zn²⁺, suggesting not only that CLR01 is a better inhibitor of the toxicity of A β 42-Zn²⁺ complexes than the CTFs but also that its ability to bind Zn²⁺ may act cooperatively with binding to Lys for inhibition of A β 42-induced toxicity.

Our study of peptides derived from the C-terminus of A β 42 and the molecular tweezer CLR01 show that, although the two types of compounds inhibit the toxicity of A β 42 on its own in cell culture with half-maximal inhibition in the micromolar range, the CTFs completely lose their ability to inhibit the toxicity in the presence of Zn²⁺, whereas CLR01 not only remains an effective inhibitor but even gains inhibitory activity.

Biophysical investigation established that, under the conditions we used, the presence of Zn²⁺ altered the self-assembly of A β 42 into formation of a mixture of fibrils and oligomers (Figure 1E), in which the β -sheet content was lower (Figure 1C) than in fibrils of A β 42 in the absence of Zn²⁺ (Figure 1B). As we reported previously for A β 40,³⁰ Zn²⁺ altered dramatically the interaction of ThT with A β 42. A rapid, relatively small increase in the fluorescence, without a lag phase, was observed in mixtures of A β 42 and Zn²⁺, but it apparently reflected the formation of atypical ternary complexes in which presumably the ThT molecules were rather flat, leading to the observed increase in their fluorescence. This was evidenced by the absence of substantial β -sheet structure in the CD spectra at the corresponding time points. Similar to A β 40-Zn²⁺ complexes, the A β 42-Zn²⁺ complexes did not show toxicity when applied to cultured cells all at once, presumably due to rapid aggregation into benign structures.³⁰ However, when the same total amount of A β 42-Zn²⁺ complexes was added in four portions over 36 h, their toxicity was substantially higher, likely because, under these conditions, the concentration of the complexes did not allow formation of large, nontoxic aggregates.

The behavior of the ThT fluorescence of A β 42-Zn²⁺ complexes in the presence of the inhibitor compounds further supports the conclusion that this fluorescence did not reflect formation of β -sheet-rich fibrils. The only inhibitor that slightly reduced the fluorescence was A β (39–42) (Figure 3A), which previously had been shown not to affect the ThT increase upon fibrillation of A β 42 in the absence of Zn²⁺.⁵⁸ In contrast, A β (30–40), A β (31–42), and CLR01 all caused an immediate increase in the ThT fluorescence (Figures 3A and 7A) followed by gradual changes that did not correlate with the degree of β -sheet formation evidenced by CD spectroscopy (Figures 4 and 7B–D). We conclude that ThT fluorescence is not a reliable method for measurement of β -sheet formation or amyloid fibrils by A β 42 (or A β 40³⁰) in the presence of Zn²⁺.

In contrast to the ThT fluorescence, the CD spectra did not seem to suffer from artifacts and reported on changes in the secondary structure of the A β 42-Zn²⁺ complexes in a manner that appeared consistent with the morphological evaluation and toxicity measurement. As deconvolution algorithms rely on sampling of known structures and can lead to overfitting of the data, we focus the discussion on the spectral changes themselves. In the presence of fivefold excess of the CTFs (Figure 4B–G) or CLR01 (Figure 7B–D), little conformational change was observed over multiple days of incubation, suggesting that the peptides remained predominantly unstructured. Previously, little to no change was observed in solutions of A β (39–42) or A β (30–40) incubated on their own at \sim 200

μM , suggesting that these peptides did not form β -sheets, whereas $A\beta(31-42)$ incubated at $\sim 60 \mu\text{M}$ gradually formed β -sheet-rich fibrils over 48 h.⁵³ Here, despite being used at a higher concentration, $100 \mu\text{M}$, little conformational change was observed over time for all the CTFs, supporting the interpretation that they formed relatively stable ternary complexes with $A\beta 42$ and Zn^{2+} . The larger amplitude of the spectra in the presence of the CTFs, where the value of the minimum at 195 nm was $60-70 \text{ deg} \times \text{cm}^2 \text{ dmol}^{-1}$, in comparison to those in the presence of CLR01, where the minimum was $\sim 40 \text{ deg} \times \text{cm}^2 \text{ dmol}^{-1}$, likely reflects the contribution of the CTFs themselves. This behavior suggested that all the inhibitors reduced the tendency of the $A\beta 42\text{-Zn}^{2+}$ complexes to form fibrils. Indeed, morphological analysis showed shorter, thinner, and less-abundant fibrils in the presence of the CTFs (Figure 3) than in their absence (Figure 1E). This effect was stronger in the presence of $A\beta(31-42)$ and $A\beta(39-42)$ than in the presence of $A\beta(30-40)$, suggesting that the Ile⁴¹-Ala⁴² dipeptide contributed to the disruption of $A\beta 42\text{-Zn}^{2+}$ fibril formation. Inhibition of fibril formation by CLR01 was stronger than by the CTFs, even at low concentrations (cf. Figure 5 and Figure 7E–G). In the presence of all the inhibitors, the suppression of fibril formation appeared to increase the presence of oligomer-like structures.

In previous studies of the CTFs and CLR01 as inhibitors of $A\beta 42$ self-assembly and toxicity, we proposed that all the inhibitors stabilized oligomeric structures (we termed them hetero-oligomers in the case of the CTFs⁵¹), which unlike the oligomers formed by $A\beta 42$ itself, were nontoxic in multiple cell-culture systems.^{51,52,54,55,57,65,66} Here, we discovered that although both the CTFs and CLR01 appeared to suppress fibrillogenesis of $A\beta 42\text{-Zn}^{2+}$ complexes, thereby increasing the relative abundance of nonfibrillar oligomers, the oligomers formed in the presence of the three CTFs maintained their toxicity (Figure 6), whereas those formed in the presence of CLR01 were nontoxic at concentrations of at least $30 \mu\text{M}$ (Figure 7H). This opposite effect of the addition of Zn^{2+} on the ability of the two types of inhibitors to suppress $A\beta 42$ toxicity is remarkable and suggests that $A\beta$ inhibitors evaluated in cell culture should be tested not only against the peptide itself but also against its complexes with Zn^{2+} and potentially with other metal ions. Our data also support further development of CLR01, because CLR01 inhibits the toxicity of $A\beta 42$ both in the absence and in the presence of Zn^{2+} . CLR01 is also an effective inhibitor of tau aggregation and spreading^{67,68} making a particularly attractive lead compound for development of AD therapy.

METHODS

All buffers were prepared using deionized water from a Milli-Q water purification system (Millipore) and analytical grade reagents. NaOH, 3-(*N*-morpholino)propanesulfonic acid (MOPS), 1,1,1,3,3,3-hexafluoro-2-propanol (HFIP), and ZnCl_2 were purchased from Sigma-Aldrich. For cell culture, Dulbecco's modified Eagle's medium (DMEM) (ThermoFisher Scientific) was supplemented with 10% heat-inactivated horse serum (HS) (ThermoFisher Scientific), 5% fetal bovine serum (FBS, Invitrogen), and 2% penicillin-streptomycin (P/S, ThermoFisher). Synthetic human $A\beta 42$ was from Anaspec. $A\beta(39-42)$ and $A\beta(30-40)$ were either prepared and purified in-house as described previously⁶⁹ or obtained from New England Peptides. $A\beta(31-42)$ was obtained from New England Peptides. CLR01 was prepared as described previously⁷⁰ and purified as a sodium salt.

Sample Preparation for Biophysical Characterization. $A\beta 42$ was treated with HFIP to remove pre-existing aggregates as described previously.⁷¹ The resulting dry peptide film was dissolved in a 60 mM NaOH solution comprising 10% of the final volume and then diluted to 50% of the final volume using Milli-Q water, either alone or in the presence of Zn^{2+} . $A\beta 42\text{-Zn}^{2+}$ mixtures were examined in the absence or presence of the CTFs or CLR01. The mixture solutions were sonicated for 1 min using a bath sonicator, and 50% of the final volume 20 mM sodium phosphate buffer (PB), pH 7.4, was added so that the final PB concentration was 10 mM . $A\beta 42$ concentration was confirmed by UV spectroscopy ($\epsilon = 1280 \text{ M}^{-1} \text{ cm}^{-1}$).

Sample Preparation for Cell Culture. $A\beta 42$ was pretreated with HFIP as described previously.⁷¹ The resulting peptide film was dissolved in 60 mM NaOH comprising less than 0.2% of the desired volume in the wells and sonicated for 1 min. Samples then were diluted to a stock concentration of $83 \mu\text{M}$ using differentiation media (0.5% FBS, 2% P/S in DMEM), and this solution then was added to the wells to yield a final concentration of 5 or $10 \mu\text{M}$ $A\beta 42$. In our experience, different batches of $A\beta 42$ often have vastly different toxicity. Therefore, we pretest each batch for its toxicity and use it at a concentration that yields 20–50% reduction of cell viability. CTFs and CLR01 solutions were prepared in a similar manner in differentiation media. Each CTF was tested at the following concentrations: 0.5, 1.5, 5, 15, and $50 \mu\text{M}$. Thanks to its higher aqueous solubility,⁵³ $A\beta(39-42)$ also was tested at a maximal concentration of $100 \mu\text{M}$. CLR01 was tested at 1, 3, 10, 30, and $100 \mu\text{M}$. One or two microliters of a 1 mM ZnCl_2 solution was added to each well, as appropriate, to yield a final concentration of 10 or $20 \mu\text{M}$ Zn^{2+} .

Thioflavin T Fluorescence Assay. ThT fluorescence was recorded as described previously.³⁰ Fresh, $100 \mu\text{L}$ samples of $20 \mu\text{M}$ $A\beta 42$ and $20 \mu\text{M}$ ThT in 10 mM PB, pH 7.4, were prepared in a 96-well plate in the absence or presence of $40 \mu\text{M}$ Zn^{2+} and $100 \mu\text{M}$ of each CTF or 10, 30, or $100 \mu\text{M}$ CLR01. In control experiments, the CTFs were incubated in the absence or presence of Zn^{2+} at the same concentrations in the absence of $A\beta 42$. The samples were incubated inside BioTek Synergy HT or HTX Multi-Detection Microplate Readers at 37°C with orbital shaking at 307 rpm, and ThT fluorescence was measured using excitation: 420/50 nm and emission: 485/20 nm every 15 min for the indicated times.

Circular Dichroism Spectroscopy. Secondary structure analysis by CD was performed as described previously³⁰ with some modifications. Briefly, fresh, $300-450 \mu\text{L}$ samples of $20 \mu\text{M}$ $A\beta 42$ were prepared in the absence or presence of $100 \mu\text{M}$ of each CTF or CLR01 in 10 mM PB, pH 7.4. During preparation, a 1 mM Zn^{2+} solution was added to half of the samples to obtain a final concentration of $40 \mu\text{M}$ of Zn^{2+} . The samples were incubated at 37°C with rotational agitation, and CD spectra were recorded at different time points ranging from 0 (immediately after preparation) to 11 d. CD spectra were recorded using a J-810 Jasco spectropolarimeter and analyzed using SM2 software. The data were smoothed using the Savitzky-Golay algorithm⁷² set to a convolution width of 25. The spectra were deconvoluted using Bestel software.⁷³ Each deconvolution curve is an average of two to three experiments.

Electron Microscopy. The same samples used in the CD experiments were also used for EM. The samples were incubated at 37°C with agitation and aliquots were taken for TEM examination at different time points up to 17 d. Ten microliter aliquots from each solution were deposited on carbon-coated copper grids and stained with 1% uranyl acetate. These samples were analyzed using a JEM1200-EX electron microscope (JEOL) operated at 120 kV. Images were recorded using a wide-angle BioScan 600W $1\text{k} \times 1\text{k}$ digital camera and saved using digital micrograph acquisition software (Gatan).

Cell Viability Assays. Cell viability assays were conducted as described previously⁵¹ with the following changes: Rat pheochromocytoma (PC-12) cells were cultured in DMEM, containing 10% heat-inactivated HS, 5% FBS, and 1% P/S at 37°C in an atmosphere of 5% CO_2 . Only cells with a passage number between 9 and 15 were

utilized for these experiments to minimize a potential phenotype drift due to a high passage number.⁷⁴ Cells were plated at a density of 20 000 per well. Peptide solutions containing 5 or 10 μM A β 42, 2 mol equiv of ZnCl_2 (i.e., 10 or 20 μM , respectively), and increasing concentrations of CTFs or CLR01 were added to the cells. As has been reported previously,³⁰ to observe the toxicity of A β 42-Zn²⁺ complexes, fresh samples of each preparation, including A β 42, A β 42-Zn²⁺, or A β 42-Zn²⁺ in the presence of CTFs or CLR01 were added to the cells every 12 h (i.e., at $t = 0, 12, 24$, and 36 h). At these time points, 12.5 μL of media were removed from each well, and 12.5 μL of fresh samples were added to each well. At $t = 48$ h, cell viability was determined using the lactate dehydrogenase release assay and/or the 3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide reduction assay.

■ ASSOCIATED CONTENT

Supporting Information

The Supporting Information is available free of charge at <https://pubs.acs.org/doi/10.1021/acscchemneuro.0c00192>.

Morphological evaluation of A β 42 incubated in the absence or presence of Zn²⁺ (PDF)

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G.B. designed the research. A.J.M., I.H., R.M., and I.S. performed the research. F.G.K. and T.S. provided unique resources. A.J.M., I.H., and G.B. wrote the manuscript. I. So, I. Sa, F.G.K., and T.S. provided critical reading and discussion of the manuscript.

Author Contributions

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Notes

The authors declare no competing financial interest.

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