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# The effects of platinum(IV) complexes on A $\beta$ <sub>1–42</sub> aggregation: a synergistic inhibition upon axial coordination†

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Among novel metallodrugs, Pt(IV) complexes have been receiving increasing attention as alternatives to Pt(II) analogues since they exhibit superior kinetic inertness and, hence, fewer undesirable reactions. Pt(IV) complexes have been designed as prodrugs with mechanisms of action finely tuned by properties of the axial ligands. In this context, the insertion of known bioactive molecules as axial ligands to generate multi-target drugs (MTDs) is considered a valuable drug design strategy because of the possible synergistic effects between the metal centre and the organic moiety. By employing the repurposing of drugs in different diseases, in this study, the Pt(IV) complex (OC-6-44)-acetatodiamminedichlorido(4,5-dihydroxy-9,10-dioxo-9,10-dihydroanthracene-2-carboxylato)platinum(IV) was investigated for its ability to modulate the self-aggregation process of the amyloid peptide A $\beta$ <sub>1–42</sub>. Specifically, the complex features 4,5-dihydroxy-9,10-dioxo-9,10-dihydroanthracene-2-carboxylato (rhein) as a ligand that is a natural aromatic molecule already known as a discrete inhibitor of amyloid aggregation. Herein, several biophysical and microscopic assays, such as thioflavin T (ThT) fluorescence, dynamic light scattering (DLS) and scanning electron (SEM) and confocal microscopy, indicated that the Pt(IV) complex can inhibit and disassemble A $\beta$ <sub>1–42</sub> aggregation to a greater extent with respect to rhein alone. This effect is likely due to the formation of  $\pi$ – $\pi$  interactions between the rhein moiety and the side chains of the A $\beta$ <sub>1–42</sub> peptide. This experimental evidence was confirmed by molecular docking studies of monomeric and tetrameric A $\beta$ <sub>1–42</sub>. Overall, the data support the application of Pt(IV) complexes as innovative neurotherapeutics.

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† Electronic supplementary information (ESI) available: Fig. S1: stability overtime of Pt-Ac-rhein and Pt-Ac-OH; Fig. S2: time-courses of ThT fluorescence emission of A $\beta$ <sub>1–42</sub>, in the absence and presence of Pt-Ac-rhein, Pt-Ac-OH and rhein and compounds alone, and upon the addition (indicated by an arrow) of the complexes; Fig. S3: fluorescence emission spectra of Pt complexes and rhein at different times ( $\lambda_{\text{ex}} = 440$  nm); Fig. S4: fluorescence emission spectra of A $\beta$ <sub>1–42</sub> in the absence and presence of compound Pt-Ac-rhein or rhein at different times ( $\lambda_{\text{ex}} = 275$  nm); Fig. S5: ESI-MS spectra of A $\beta$ <sub>1–42</sub> in the absence and presence of Pt-Ac-OH and of Pt-Ac-OH alone; Fig. S6: ESI-MS spectra of A $\beta$ <sub>1–42</sub> in the absence and presence of rhein and of rhein alone; Fig. S7: SEM analysis of A $\beta$ <sub>1–42</sub> alone; Fig. S8: SEM analysis of A $\beta$ <sub>1–42</sub> in the presence of Pt-Ac-rhein; Fig. S9: SEM analysis of A $\beta$ <sub>1–42</sub> in the presence of rhein; Fig. S10: ligand interactions for the best docked complex of Pt-Ac-rhein with the A $\beta$ <sub>1–42</sub> monomer and tetramer; and Fig. S11: best docked pose of Pt-Ac-rhein with the A $\beta$ <sub>1–42</sub> tetramer. Table S1: table of main observed ions relative to the species formed by the A $\beta$ <sub>1–42</sub> alone and mixed with Pt-Ac-rhein; Table S2: docking scores for Pt-Ac-rhein, rhein and Pt-Ac-OH with the A $\beta$ <sub>1–42</sub> monomer and tetramer. See DOI: <https://doi.org/10.1039/d5dt00691k>

## Introduction

Transition metal coordination compounds possess distinctive properties, including variable oxidation states, diverse geometries, coordination numbers, and ligands.<sup>1–3</sup> Over time, a large number of metal complexes have been developed and investigated for their therapeutic uses in cancer, microbial and viral infections, diabetes, and neurodegenerative diseases (NDDs).<sup>4</sup> Research has underscored the importance of coordination chemistry in shaping the metallodrug mechanisms of action (MOAs), which are influenced by the metal's binding preferences and ligand exchanges with functional groups from proteins and oligonucleotides.<sup>5</sup>

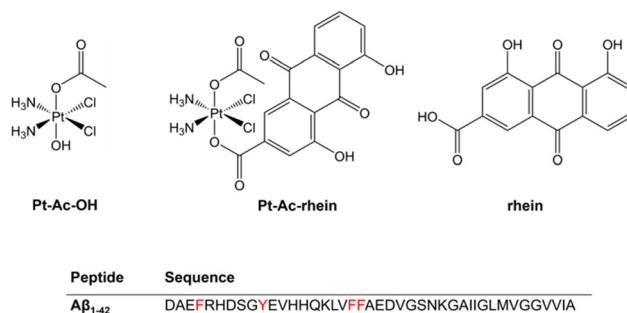
Although NDDs exhibit clinical diversity, they share key pathological characteristics, such as the cellular accumulation of intrinsically disordered proteins and disrupted metal ion homeostasis in the brain.<sup>6</sup> While significant advances have been made in understanding these diseases and developing treatments, to date no definitive cures have been achieved.<sup>7</sup> The abnormal formation of oligomers and fibrils through the self- or hetero-assembly of amyloidogenic systems—such as



amyloid- $\beta$ ,  $\alpha$ -synuclein, huntingtin, tau, and islet amyloid polypeptides—is directly implicated in NDDs like Alzheimer's, Parkinson's, and Huntington's diseases, as well as frontotemporal dementia and type II diabetes.<sup>8</sup>

Accurately designed metal complexes can stabilize alternative, non-toxic conformations of amyloid proteins by inhibiting fibril formation, or promote disaggregation of existing aggregates.<sup>9,10</sup> Among others, Pt-based compounds have been explored for this purpose: several complexes inhibited amyloid aggregation and modulated their toxicity<sup>11–14</sup> through a predominantly coordinative MOA, targeting as metal-binding sites the histidine residues (in  $A\beta_{1-42/1-40}$ : His<sup>6</sup>, His<sup>13</sup>, and His<sup>14</sup>).<sup>15</sup> Studies demonstrated that the charge, the steric hindrance and aromatic features around Pt(II) exert pivotal roles in tuning the MOAs of these complexes. For example, Pt(II) complexes, containing the 1,10-phenanthroline ligands, modulated  $A\beta_{1-42}$  peptide aggregation and reduced neurotoxicity in mouse hippocampal tissue by coordinating His residues.<sup>16</sup> In recent studies, Pt(II) complexes with  $\beta$ -hydroxy dithiocinnamic esters as ligands have been shown to inhibit the aggregation of different amyloid models such as NPM1<sub>264–277</sub>, the heptapeptide GNNQQNY from the Yeast Prion Protein Sup35p (Sup35p<sub>7–13</sub>) and  $A\beta_{21-40}$  fragments.<sup>17–20</sup> The limited solubility in aqueous solutions of these initial complexes hindered their development into viable therapeutic drugs; consequently, other studies have been focused on the use of water-soluble Pt(II) complexes. The incorporation of the tridentate ligand terpyridine in the square-planar structure of Pt(II) complexes to obtain (*SP*-4-2)-chlorido (2,2':6',2''-terpyridine)platinum(II) chloride (Pt-terpy) aided the inhibition of  $A\beta_{21-40}$  aggregation.<sup>21</sup> Consistently, the role of charge has been further investigated with positive and neutral complexes that demonstrated their ability as efficient inhibitors of  $A\beta_{21-40}$  aggregation, which reduced peptide cytotoxicity in SH-SY5Y neuroblastoma cells.<sup>22</sup> In this *scenario*, two glycoconjugate Pt(II) bipyramidal complexes suppressed amyloid aggregation of the polypeptide fragments  $A\beta_{21-40}$  and  $A\beta_{25-35}$ .<sup>23</sup>

In the search for different interaction modes between amyloid proteins and metal complexes, the peculiar features of Pt(IV) complexes can be explored. These octahedral compounds are actively studied as anticancer drugs because they are more inert than traditional Pt(II) compounds and require activation by reduction *in vivo* in the hypoxic tumor tissue, making them potentially more selective and less toxic, and producing fewer adverse effects than Pt(II) complexes,<sup>24–26</sup> also making them applicable in the amyloid context.<sup>27</sup> Moreover, the octahedral coordination geometry makes the introduction of one or two bioactive ligands possible in the axial positions, producing potentially multi-functional agents combining the MOA of the platinum moiety with that of the bioactive ligands in an additive or, better, synergistic way.<sup>28</sup> In this context, to generate a suitable complex, the well-known cisplatin was combined with a molecule of rhein (4,5-dihydroxy-9,10-dioxo-9,10-dihydroanthracene-2-carboxylic acid or cassic acid) and an acetate group to complete the coordination, thus providing the compound (*OC*-6-44)-acetatodiamminedichlorido(4,5-dihydroxy-9,10-dioxo-9,10-dihydroanthracene-2-carboxylato)plati-



**Fig. 1** Structure of the compounds and sequence of the  $A\beta_{1-42}$  investigated in this study. The aromatic amino acids of the amyloid polypeptide are highlighted in red.

num(IV), nicknamed **Pt-Ac-rhein** (Fig. 1).<sup>29</sup> This complex proved to be similar to or more potent than the parent cisplatin and rhein compounds on two human glioblastoma cell lines (U87-MG and T98G). Importantly, physicochemical and computational studies indicated its superior ability to cross the blood–brain barrier than rhein itself.<sup>29</sup> The rhein molecule alone demonstrates several pharmacological effects, such as anti-inflammatory, anti-fibrosis, anti-oxidant and anticancer abilities, manifested by modulating cell proliferation, apoptosis, migration, and invasion.<sup>30,31</sup> It has been demonstrated that rhein can partially dissociate or antagonize the oligomerization of class A scavenger receptors (SCARA). SCARA is a class of cell surface receptors involved in cell adhesion and uptake of ligands and are expressed on microglia and astrocytes. The first member of the SCARA family, SCARA-1, exhibited high interaction levels around the  $A\beta$  plates in microglial cells. Docking and molecular dynamic simulations indicated that both hydrophobic and H-bonds are responsible for these interactions.<sup>32,33</sup> Importantly, rhein demonstrated a discrete ability to inhibit  $A\beta_{1-42}$  amyloid aggregation, where its carboxylic group is able to interact with Lys<sup>16</sup> and Lys<sup>28</sup> residues and these direct interactions targeted the monomer, dimer, and trimer, further suppressing oligomerization.<sup>34</sup>

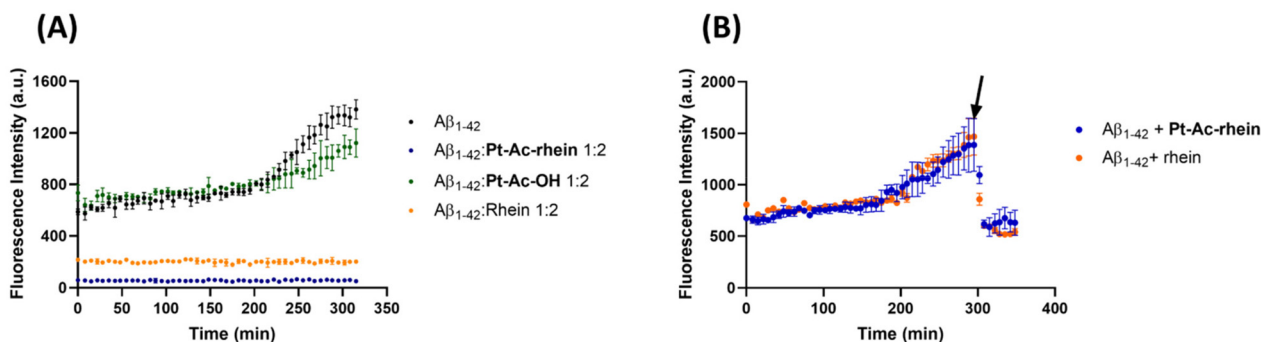
In light of the above considerations, the ability of **Pt-Ac-rhein** to interfere with the aggregation of  $A\beta_{1-42}$  was investigated *via* a range of spectroscopic and microscopic techniques, as well as by molecular dynamics (MD) methods. The complex **Pt-Ac-OH**, as the precursor of **Pt-Ac-rhein**, and rhein were included in the study in order to distinguish between the effects of the separate moieties.

## Results and discussion

### Modulation of $A\beta_{1-42}$ aggregation

Prior to performing amyloid-related assays, the stability of **Pt-Ac-rhein** and **Pt-Ac-OH** was evaluated by registering UV-vis absorption spectra at different times and at three concentrations (Fig. S1†). **Pt-Ac-OH** was stable at all tested concentrations, up to 24 h (Fig. S1B†). Regarding **Pt-Ac-rhein**, the almost complete superimposition of the spectra at 25 and





**Fig. 2** Overlay of time-courses of ThT fluorescence emission intensity of (A)  $A\beta_{1-42}$ , in the absence and presence of **Pt-Ac-rhein**, **Pt-Ac-OH** and **rhein**, and (B) upon the addition (indicated by an arrow) of the complexes at a 1:1 molar ratio to pre-formed amyloid aggregates. The results are representative of three independent measurements.

50  $\mu\text{M}$  indicated great stability of the complex, while at 100  $\mu\text{M}$ , a certain variability of absorbance suggested potential aggregation over time (Fig. S1A<sup>†</sup>).

To analyze the effects of these compounds on the self-recognition process of  $A\beta_{1-42}$ , thioflavin T (ThT) binding assay was employed,<sup>35–37</sup> and experiments were carried out by adding both compounds at the beginning of aggregation and once oligomers had become pre-formed (Fig. 2). The time course profiles of the ThT fluorescence of  $A\beta_{1-42}$  alone and in the presence of compounds are reported in Fig. 2A. The  $A\beta_{1-42}$  peptide alone exhibited a typical sigmoidal profile presenting a  $t_{1/2}$  value of  $\sim 250$  min. The effects of the presence of **Pt-Ac-rhein** were analyzed at two different peptide:complex molar ratios (1:1 Fig. S2A<sup>†</sup> and 1:2 Fig. 2A) and, in both cases, the complex showed a clear inhibitory effect reducing the maximum fluorescence value at 150 a.u. with an average inhibition of  $\sim 90\%$ . A similar, but less marked, effect was observed in the presence of **rhein** with a fluorescence value of 255 a.u. and 20% inhibition (Fig. 2A). **Pt-Ac-OH** showed a negligible effect, presenting ThT profiles almost superimposable on those of the peptide alone at both tested ratios (Fig. 2A and S2A<sup>†</sup>). None of the compounds exhibited interference with ThT (Fig. S2A<sup>†</sup>). Control emission spectra of compounds alone,  $\lambda_{\text{ex}} = 440$  nm, are reported in Fig. S3.<sup>†</sup>

Having assessed the suppressive abilities of compounds toward amyloid aggregation starting from monomers, their effects on pre-formed oligomers were evaluated by monitoring the ThT signals after the addition of compounds to  $A\beta_{1-42}$  aggregates in a 1:1 peptide:compound molar ratio (Fig. 2B). Upon the addition of the compounds, both **Pt-Ac-rhein** and **rhein** caused a decrease in ThT fluorescence intensity, with an inhibitory effect of  $\sim 60\%$ . Conversely, the addition of **Pt-Ac-OH** had no effect on the pre-formed aggregates (Fig. S2A<sup>†</sup>).

The lack of effect of **Pt-Ac-OH** led us to speculate on a potential aromatic effect among the anthracene portion of rhein and aromatic residues of the polypeptide sequence as Phe at positions 4, 19 and 20, and Tyr at position 10. To gain insights into this aspect, the fluorescence emission of  $A\beta_{1-42}$ , upon excitation at 275 nm, in the presence and absence of the

investigated compounds, was recorded. As displayed in Fig. S4,<sup>†</sup> the  $A\beta_{1-42}$  alone shows a progressive decrease in its fluorescence intensity at 303 nm over time, as previously observed.<sup>41</sup> The presence of rhein and in particular of **Pt-Ac-rhein** already causes a significant reduction of this signal at  $t = 0$  h, suggesting involvement of aromatic residues in the interaction with the compounds and thus important contributions of  $\pi$ - $\pi$  bonds in the inhibition of aggregation observed.

#### ESI-MS analysis of adducts between complexes and $A\beta_{1-42}$

To evaluate direct interaction between  $A\beta_{1-42}$  and the investigated compounds, native electrospray ionization mass spectrometry (ESI-MS) was used.  $A\beta_{1-42}$  was incubated with the compounds (at 1:1 molar ratio) (Fig. 3);  $A\beta_{1-42}$  and Pt(IV) compounds alone were analyzed as references. As can be observed, in the presence of **Pt-Ac-rhein**, two additional peaks, with respect to  $A\beta_{1-42}$  alone, were found at  $m/z$  1719.12 and 1290.76 a.m.u. (Table S1<sup>†</sup>). These peaks were attributed to the formation of a non-covalent adduct between  $A\beta_{1-42}$  and one **Pt-Ac-rhein** molecule.

Conversely, in the presence of **Pt-Ac-OH** (Fig. S5<sup>†</sup>), no adduct formation was observed, nor was there any formation in the presence of rhein (Fig. S6<sup>†</sup>). Because of the observed inefficacy of the **Pt-Ac-OH** complex to modulate  $A\beta_{1-42}$  aggregation, it was not included in subsequent experiments. The absence of any adduct in the case of **Pt-Ac-OH** is likely due to the lack of potential of aromatic interactions between the complex and the polypeptide.

#### Effects of Pt(IV) complexes on amyloid $A\beta_{1-42}$ aggregation: DLS assay

Dynamic light scattering (DLS) was employed to evaluate the effects of compounds on the size of  $A\beta_{1-42}$  aggregates (Fig. 4). Correct autocorrelation of the native  $A\beta_{1-42}$  was achieved after 24 h with a diameter of  $\sim 155$  nm (Fig. 4A) as already reported.<sup>38,42,43</sup> Conversely, the presence of **Pt-Ac-rhein** (1:1 molar ratio) anticipated autocorrelation at  $t = 0$  h, by stabilizing greater oligomers centered at  $\sim 1130$  nm (Fig. 4B). Smaller diameters, compatible with the peptide alone, are also



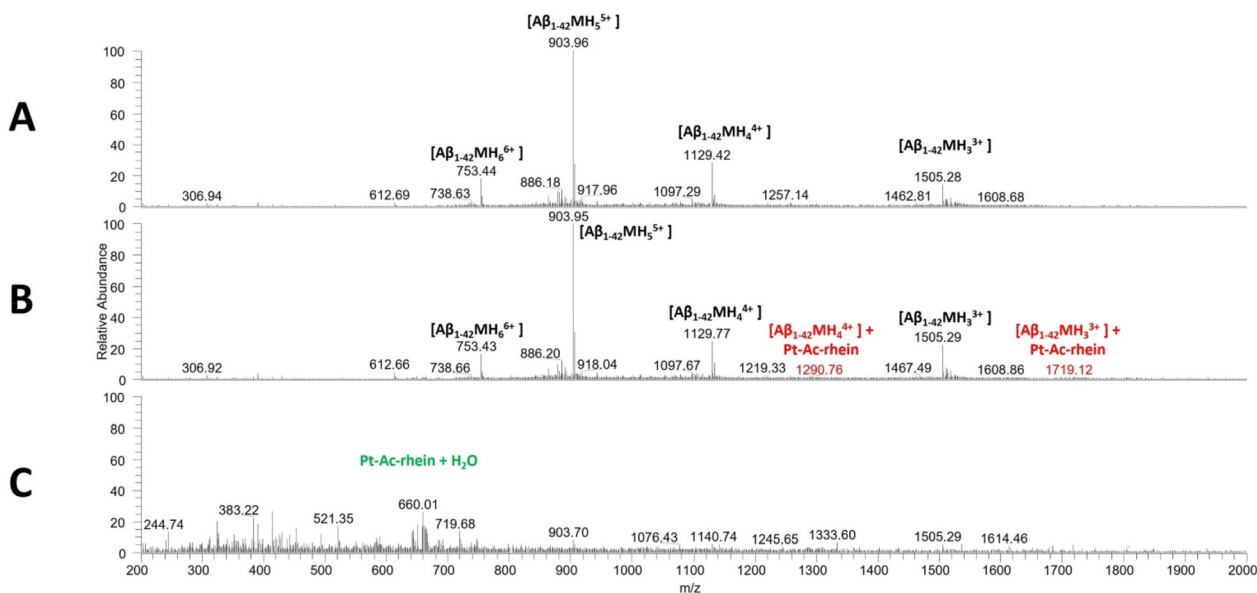


Fig. 3 ESI-MS spectra of  $A\beta_{1-42}$  (A) in the absence and (B) presence of Pt-Ac-rhein, and (C) the spectrum of the latter compound alone.

observed. Potential differences in the surface charges of monomers and/or small oligomers can lead, in the presence of Pt-Ac-rhein, to the formation of oligomers that are larger than those formed by  $A\beta_{1-42}$  alone. The small variation in the sizes of oligomers formed over time falls within the error of the measurement. DLS data suggest that the adduct formed by Pt-Ac-rhein and  $A\beta_{1-42}$  stabilizes oligomeric states, which are unable to further oligomerize.<sup>38,39</sup> Samples derived from the incubation of  $A\beta_{1-42}$ :rhein at 1:1 ratio, rhein and Pt-Ac-rhein alone did not correlate (data not shown). The inability to correlate the  $A\beta_{1-42}$ :rhein sample does not imply lack of interaction of rhein with  $A\beta_{1-42}$ , but instead indicates that this interaction was insufficient to stabilize definitive sizes of amyloid aggregates.

#### Effects of metal complexes on the morphologies of amyloid fibers

Potential modifications of the morphology of  $A\beta_{1-42}$  aggregates induced by Pt-Ac-rhein and rhein alone were studied by using the technique of scanning electron microscopy (SEM). Samples, prepared by mixing at a 1:1 molar ratio, were analyzed across four independent experiments, with the images from one representative experiment being shown in Fig. 5. Additional data from the other independent experiments are presented in Fig. S7–S9.† As expected, after 6 h of stirring, the  $A\beta_{1-42}$  peptide alone provides well-defined fibers with an average length of  $\sim 1040 \mu\text{m}$  and a diameter of  $\sim 13 \mu\text{m}$  (Fig. 5 upper panel, Table 1). In the presence of Pt-Ac-rhein, a significant reduction in fiber diameter ( $\sim 8 \mu\text{m}$ ) and a change in their morphology were evident; indeed, the fiber appeared folded back several times, making measurement of its length impossible (Fig. 5B–B’). In the presence of rhein, a fiber with a diameter comparable to that of the peptide alone was observable, but with a length approximately 7 times shorter. The amyloid

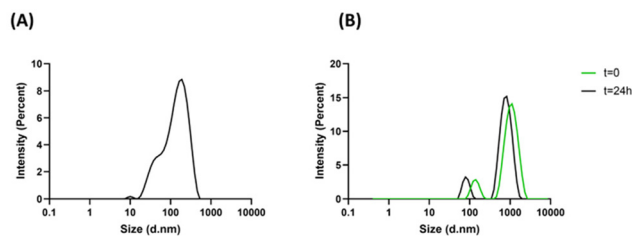


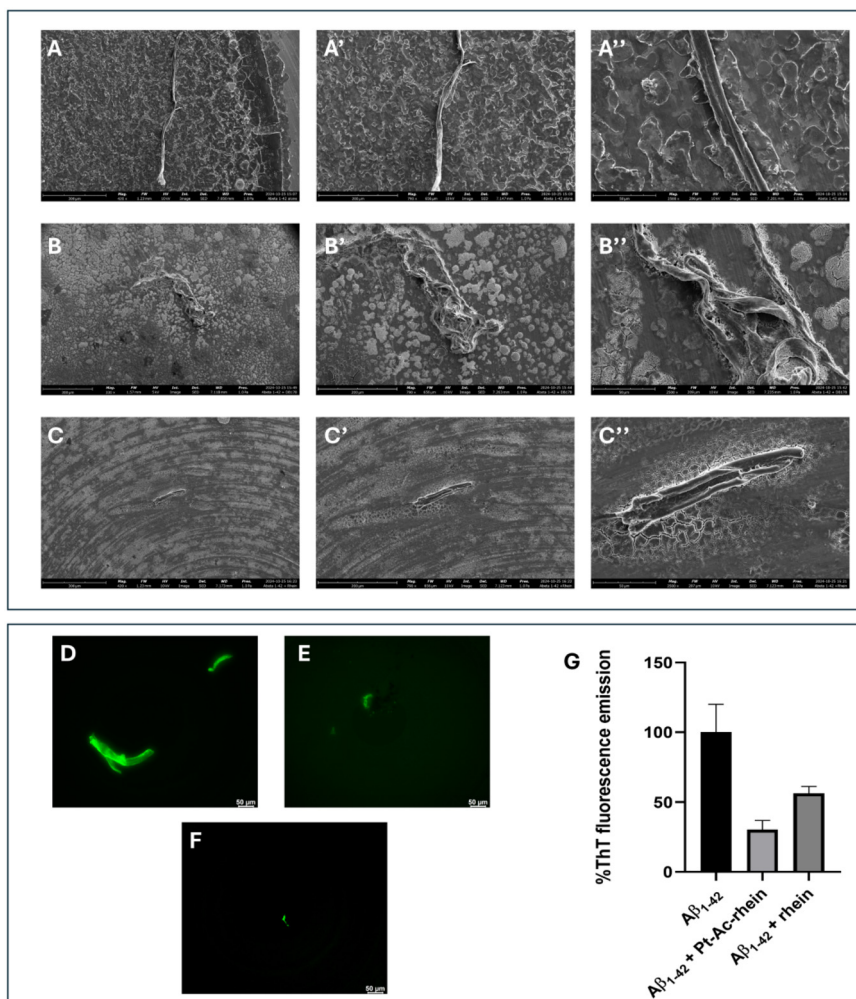
Fig. 4 Overlay of size distribution by DLS measurements of the native  $A\beta_{1-42}$  peptide at  $t = 24$  h (A) and with the complex Pt-Ac-rhein (B) at an  $A\beta$ : complex mole ratio of 1:1, at  $t = 0$  (green line) and  $t = 24$  h (black line).

character of the observed aggregates in SEM was corroborated by confocal microscopy analysis (Fig. 5, lower panel). In the case of  $A\beta_{1-42}$  alone and with rhein, fibers, although different in size, demonstrated incorporation of ThT, confirming their amyloid character (Fig. 5, lower panel, D–G). Conversely, for  $A\beta_{1-42}$  with Pt-Ac-rhein, the low fluorescence intensity observed (Fig. 5, lower panel, E–G) confirmed the amorphous nature of the observed aggregates.

#### Modelling studies of the interaction of Pt-Ac-rhein with $A\beta_{1-42}$

Pt-Ac-rhein was docked into monomeric and tetrameric  $A\beta_{1-42}$ , along with rhein and Pt-Ac-OH, as controls. To reflect the intrinsically disordered nature of  $A\beta_{1-42}$ , docking was carried out using 10 different conformations of the monomer.<sup>40</sup> As reported in Table S2,† Pt-Ac-rhein forms stable adducts in 1:1 complexes, each of which are significantly more stable than equivalent forms involving either rhein or Pt-Ac-OH. It is apparent that the conjugation of rhein with the Pt(IV) center gives rise to markedly greater interaction with  $A\beta_{1-42}$  than rhein alone. Pt-Ac-rhein is more strongly bound to the  $A\beta_{1-42}$





**Fig. 5** Upper panel: scanning electron microscopy images of Aβ<sub>1-42</sub> alone (A) and in the presence of Pt-Ac-rhein (B) and rhein (C) at a magnification of 330× (300 μm scale bar, A, B, and C), 790× (200 μm scale bar, A', B', and C') and 2500× (50 μm scale bar, A'', B'', and C''). Lower panel: confocal microscopy images ( $\lambda_{\text{ex}} = 440 \text{ nm}$  and  $\lambda_{\text{em}} = 480 \text{ nm}$ ) of Aβ<sub>1-42</sub> alone (D) and in the presence of Pt-Ac-rhein (E) and rhein (F) at a magnification of 10×. (G) Histogram showing the percentage of ThT fluorescence intensity from confocal analysis. The SEM images are representative of four independent experiments (the images from other experiments are shown in Fig. S7–S9†).

**Table 1** SEM analyses: average diameter and length of fibers obtained for Aβ<sub>1-42</sub> in the presence and absence of Pt-Ac-rhein and rhein. Values represent the mean from four independent experiments: one representative experiment is shown in Fig. 5 and the others are presented in Fig. S7–S9†

	Average diameter (μm)	Average length (μm)
Aβ <sub>1-42</sub>	13 ± 2	1040 ± 3.0
Aβ <sub>1-42</sub> :Pt-Ac-rhein	7.4 ± 2	Could not be evaluated
Aβ <sub>1-42</sub> :rhein	18.6 ± 0.5	146 ± 0.9

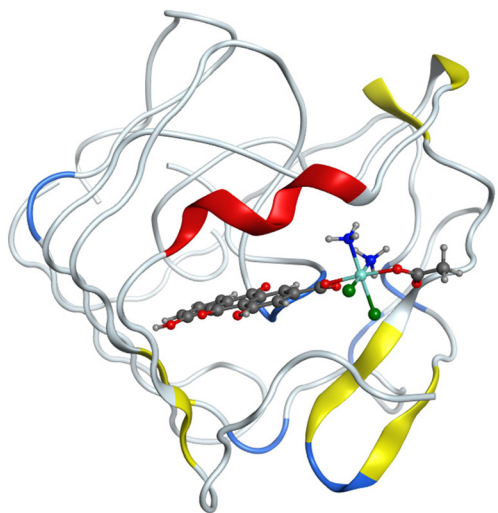
monomer than rhein alone with a ChemPLP score of ~10 on average.

In the 1 : 1 Pt-Ac-rhein complex, intermolecular interactions occur largely within the N-terminal region of Aβ<sub>1-42</sub>. This is particularly evident in the complex indicated with monomer 8 in Table S2† (Fig. 6), in which strong polar interactions

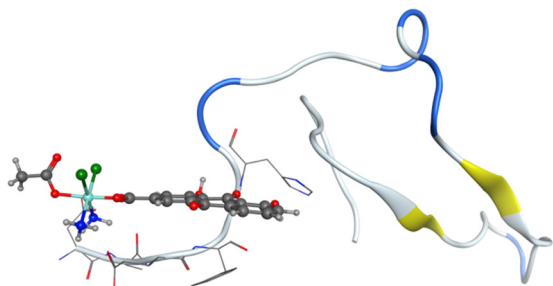
between coordinated -NH<sub>3</sub> groups and sidechains of Asp<sup>1</sup> and Glu<sup>3</sup> as well as backbone carbonyl of Ala<sup>2</sup> are evident. In addition, stacking interactions between the rhein moiety and Phe<sup>4</sup> and His<sup>6</sup> are observed (Fig. S10†). Thus docking provides evidence that the combination of a polar Pt(IV) complex with an extended aromatic ligand forms an excellent combination for binding to monomeric Aβ<sub>1-42</sub>.

Docking was also performed on 5 representative clusters as tetramers of Aβ<sub>1-42</sub> reported by Nguyen *et al.*<sup>41</sup> (Table S2†). Again, it is apparent that Pt-Ac-rhein binds significantly more strongly than either of rhein or Pt-Ac-OH alone. As in the monomer case, the most stable 1 : 4 complex located by docking is stabilized by polar interactions between coordinated -NH<sub>3</sub> groups and acidic residues in Aβ<sub>1-42</sub>, particularly Glu<sup>3</sup> and Glu<sup>22</sup> of chain A. Hydrophobic interactions also form between the rhein moiety and Phe<sup>4</sup>, Phe<sup>19</sup> and Phe<sup>20</sup> of chain A (Fig. 7 and S10†): Pt-Ac-rhein slots between helical and sheet structures





**Fig. 6** Best docked pose of Pt-Ac-rhein with the A $\beta_{1-42}$  monomer. Pt-Ac-rhein is shown as a ball-and-stick model, and residues Asp<sup>1</sup>, Ala<sup>2</sup>, Glu<sup>3</sup>, Phe<sup>4</sup> and His<sup>6</sup> are shown as stick models. The peptide is colored yellow for  $\beta$ -strands, blue for turns/bends, and white for random coils.



**Fig. 7** Best docked pose of Pt-Ac-rhein with the A $\beta_{1-42}$  tetramer. Pt-Ac-rhein is shown as a ball-and-stick model. The peptide is colored red for the helix, yellow for  $\beta$ -strands, blue for turns/bends, and white for random coils.

formed by chain A on the surface of the tetramer. A surface representation of the receptor colored on the basis of lipophilicity (Fig. S11†) shows how this binding pose includes hydrophilic pockets formed by Glu residues, as noted above, while the rhein group slots into a hydrophobic pocket formed by Phe residues. As in the monomer case, Pt-Ac-rhein is predicted to bind more strongly than rhein alone with a ChemPLP score of  $\sim 10$  on average, with Pt-Ac-OH alone being even less strongly bound. This is in agreement with the high lipophilicity of Pt-Ac-rhein, measured in an *n*-octanol/water system and expressed as distribution coefficient *D*, to compare neutral Pt-Ac-rhein ( $\log D = 1.85$ ) with ionizable rhein ( $\log D = -0.27$ ).<sup>29</sup>

## Experimental

### Synthesis of peptides

The A $\beta_{1-42}$  polypeptide was purchased from NovoPro Bioscience Inc. (Shanghai, China). The peptide was treated

with 1,1,1,3,3,3-hexafluoro-2-propanol (HFIP) to guarantee a monomeric state, lyophilized and stored at  $-20$  °C until use.<sup>42</sup>

### Synthesis of metal compounds

4,5-Dihydroxy-9,10-dioxo-9,10-dihydroanthracene-2-carboxylic acid or rhein (Tokyo Chemical Industry) and all the other chemicals (Alfa Aesar; Thermo Fisher Scientific or Sigma Aldrich; Merck) were used without further purification. Cisplatin was prepared according to Dhara's method.<sup>43</sup> Complex (OC-6-44)-acetatodiamminedichloridohydroxidoplatinum(IV), Pt-Ac-OH, was prepared by treating cisplatin (100 mg) with 50% w/w oxygen peroxide (1 mL) in acetic acid (40 mL) as previously described.<sup>44,45</sup> Complex (OC-6-44)-acetatodiamminedichlorido(4,5-dihydroxy-9,10-dioxo-9,10-dihydroanthracene-2-carboxylato)platinum(IV), Pt-Ac-rhein, was prepared as reported in the literature.<sup>29</sup> Briefly, Pt-Ac-OH (0.100 g) was reacted with rhein (0.091 g) in the presence of 1-[bis(dimethylamino)methylene]-1*H*-1,2,3-triazolo[4,5-*b*]pyridinium 3-oxide hexafluorophosphate (HATU, 0.145 g) and *N,N*-diisopropylethylamine (50  $\mu$ L) in anhydrous DMF (3 mL) overnight. Upon centrifugation of the mixture, the volume of the supernatant was reduced using a rotary evaporator. The oily residue was triturated with a dichloromethane–diethyl ether mixture to give a yellow powder, which was washed with diethyl ether, 1% formic acid and cold water, and finally dried under nitrogen flow (58% yield).

### UV-vis absorption spectroscopy

The solution stability of Pt-Ac-rhein and Pt-Ac-OH was evaluated by recording their UV-vis absorption spectra in 10 mM phosphate buffer at pH 7.4 over 24 hours. To record the spectra, the compounds were dissolved in DMSO (stock solutions 50 mM) and then added to the selected buffers to reach a final concentration of 25, 50 or 100  $\mu$ M. The DMSO final concentration was 0.2% v/v. UV-vis absorption spectra were recorded on BioDrop Duo UV-visible spectrophotometers (Cambridge, United Kingdom), at room temperature, using 1 cm path length cuvettes and the following parameters: 350–600 nm range, 200 nm min<sup>-1</sup> scanning speed, and 2.0 nm bandwidth.

### Fluorescence spectroscopy

The ThT emission assay was carried out on an EnVision 2105 fluorescence reader (PerkinElmer) using black plates (96 well) while stirring. Measurements were recorded every 8 min ( $\lambda_{ex}$ : 440 nm and  $\lambda_{em}$ : 485 nm). Assays were performed at 25 °C employing a peptide concentration of 50  $\mu$ M (buffer composition: 50 mM NaCl and 20 mM phosphate buffer, pH 7.4), using a ThT final concentration of 5  $\mu$ M, at different molar ratios with metal complexes (stock solutions of 50 mM in DMSO, final DMSO concentration of 0.2% v/v). For disaggregation experiments, the fibrillation of A $\beta_{1-42}$  was carried out by mixing the peptide at the desired concentration with ThT, in 50 mM NaCl and 20 mM phosphate buffer, at pH 7.4, and the solution was placed in the black plates, kept under agitation and moni-



tored; metal complexes were added when maximum fibrillation was reached.

Intrinsic fluorescence was carried out using 50  $\mu\text{M}$   $\text{A}\beta_{1-42}$  mixed with the compounds at a 1 : 1 molar ratio (peptide : compound), in a quartz cuvette with an optical path length of 10 mm, on a Jasco FP 8300 spectrofluorometer ( $\lambda_{\text{ex}} = 275 \text{ nm}$ ).

### ESI-MS analysis

The solution of  $\text{A}\beta_{1-42}$  at a concentration of 50  $\mu\text{M}$  in 15 mM ammonium acetate (AMAC) buffer at  $\text{pH} = 7.0$  was incubated with **Pt-Ac-rhein** and **Pt-Ac-OH** in a peptide to metal compound molar ratio of 1 : 1. The solutions were diluted 10 times with 15 mM AMAC and then analyzed using an LTQ XL Ion Trap mass spectrometer equipped with an electrospray ionization (ESI) source, operating at a needle voltage of 3.5 kV and temperature of 320  $^{\circ}\text{C}$ , combined with a complete Ultimate 3000 HPLC system, including a pump MS, autosampler, and photodiode array (all from Thermo Fisher Scientific). Spectra of the complexes alone were recorded as controls.

### DLS assays

DLS measurements were performed using a Zetasizer Nano S DLS device from Malvern Instruments (Malvern, Worcestershire, UK) equipped with a 633 nm laser, at a backscatter angle of 173 $^{\circ}$ , controlled with a Peltier thermostat system while using a plastic micro cuvette.  $\text{A}\beta_{1-42}$  at a concentration of 50  $\mu\text{M}$  in 50 mM NaCl and 20 mM phosphate buffer ( $\text{pH} 7.4$ ) was maintained under conditions of stirring at 25  $^{\circ}\text{C}$  either alone or at a 1 : 1 peptide-to-metal complex molar ratio on a Multi Reax Vortexer 115 V (Heidolph Instruments GmbH & Co., Schwabach, Germany). Size distribution values were determined by intensity measurements in automatic mode at regular time-intervals over a period of 10 min for each measurement. Thirteen acquisitions were recorded, each of 10 seconds in duration.

### Scanning electron microscopy

$\text{A}\beta_{1-42}$  alone (50  $\mu\text{M}$ ) and in the presence of compounds **Pt-Ac-rhein** and **rhein** (1 : 1 peptide : complex molar ratio) in 10 mM phosphate buffer,  $\text{pH} 7.4$ , were morphologically analyzed after 6 h of aggregation using field-emission SEM (Phenom\_XL, Alfatest, Milan, Italy). The aggregation was achieved by maintaining the solutions under stirring conditions at 25  $^{\circ}\text{C}$  on a Multi Reax Vortexer 115 V (Heidolph Instruments GmbH & Co., Schwabach, Germany) for 6 h. After this time,  $\sim 50 \mu\text{L}$  of solution was drop-cast on an aluminum stub and this was dried under vacuum to prepare each sample. For 75 s, a thin layer of gold was sputtered at a current of 25 mA. Following the introduction of the sputter-coated samples into the specimen chamber, micrographs were obtained using a secondary electron detector (SED) at an accelerating voltage of 10 kV. The **Pt-Ac-rhein** (50  $\mu\text{M}$ ) and **rhein** (50  $\mu\text{M}$ ) compounds were analyzed as controls.

### Confocal microscopy

Confocal microscopy analysis was performed using a Leica X Mica Microhub.  $\text{A}\beta_{1-42}$  (50  $\mu\text{M}$ ), either alone or mixed with **Pt-Ac-rhein** or **rhein** at 1 : 1 molar ratio, was first incubated with 50  $\mu\text{M}$  ThT in 50 mM NaCl and 20 mM phosphate buffer ( $\text{pH} 7.4$ ) at room temperature for 6 h, under conditions of stirring. After incubation, the fluorescence of the formed fibers was analyzed using the excitation wavelength of ThT at 440 nm, with the detection occurring at 483 nm. Fluorescence images were acquired using a 10 $\times$  objective with the pinhole set to its maximum opening. The percentages of the ThT fluorescence intensity signals were analyzed using the image processing program ImageJ.

### Computational methods

Geometries of **Pt-Ac-rhein**, **rhein** and **Pt-Ac-OH** were fully optimized at the B3LYP-G3BJ level with a def2-SVP basis set without any symmetry constraint, and confirmed as minima by harmonic frequency calculation. Single point calculations at the optimal geometry were then performed at the B3LYP/def2-TZVP level to obtain atomic charges using the Merz-Kollman scheme. Each structure, as determined, was then docked into receptors consisting of different conformations of monomeric or tetrameric  $\text{A}\beta_{1-42}$  taken from the literature. Docking employed the PLANTS program,<sup>46</sup> with the entire structure of the monomer or tetramer taken as the receptor.

## Conclusion

In recent years transition metal complexes have garnered significant attention for their potential applications in the diagnosis, imaging, and treatment of NDDs.<sup>47</sup> Very few studies have reported on metal complexes bearing Pt(IV) coordination as modulators of amyloid aggregation. In this study, to exploit potential synergistic effects, a molecule of natural origin that exhibited discrete inhibitory effects on  $\text{A}\beta_{1-42}$  was included as an axial ligand into an octahedral Pt(IV) species to obtain the complex named **Pt-Ac-rhein**. This complex was investigated to determine its effects toward amyloid aggregation in comparison with the parent metal complex lacking the **rhein** ligand, **Pt-Ac-OH**, and the **rhein** molecule alone (Fig. 1). ThT and aromatic assays indicated that **Pt-Ac-rhein** was able to drastically suppress aggregation and disassemble pre-formed high-level oligomers. DLS studies suggested a MOA involving the stabilization of large oligomers induced by the presence of **Pt-Ac-rhein** and these were unable to further form amyloid fibrils, as indicated by SEM and confocal images. Mass spectrometry showed that **Pt-Ac-rhein** directly interacts with  $\text{A}\beta_{1-42}$  for the formation of a non-covalent adduct between the complex and the amyloid polypeptide without exchange of ligands around Pt(IV), as expected. The presence of **rhein** in the coordination sphere of Pt(IV) allows for  $\pi$ - $\pi$  interactions with aromatic residues of the amyloid sequence, as shown by docking simulations with both monomeric and tetrameric forms of  $\text{A}\beta_{1-42}$ . These interactions presumably stabilize the formation of



adducts with A $\beta_{1-42}$  as the monomer as well as small oligomers, hampering further aggregation. The superior inhibitory effect observed for **Pt-Ac-rhein** with respect to rhein alone could be ascribed to the precise geometrical position of rhein in the context of the metal complex as an axial ligand in the octahedral geometry around Pt(IV), which can enhance  $\pi$ - $\pi$  interactions and/or allow the stabilization of soluble oligomers of A $\beta_{1-42}$ .

Unfortunately, the intrinsic cytotoxicity of **Pt-Ac-rhein** observed in human glioblastoma cell lines, with IC<sub>50</sub> values in the low micromolar range, hampered any cytotoxicity assay to evaluate potential rescue of cell viability due to the suppression of amyloid aggregation.<sup>29</sup> However, to the best of our knowledge, this is the first investigation of a Pt(IV) complex incorporating rhein as the axial ligand specifically in relation to its anti-amyloidogenic properties. In conclusion, this study provides critical information on the inhibition of amyloid fibrillation by Pt(IV) metal complexes with unprecedented impact on the biomedical value of clinical metal-based MTDs against amyloid diseases.

## Author contributions

Funding acquisition, conceptualization, and original draft: D. M.; supervision, review and editing: D.M. and M.R.; investigation and formal analysis: S.L.M., D.F., E.G., and J.A.P.

## Data availability

The data supporting this article have been included as part of the ESI.†

## Conflicts of interest

There are no conflicts to declare.

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