SUPPLEMENTARY INFORMATION

The effects of Platinum (IV) complex on A β_{1-42} aggregation: a synergistic inhibition upon axial coordination

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Figure S1. Stability overtime of A) **Pt-Ac-rhein** and B) **Pt-Ac-OH**, in 10 mM phosphate buffer at pH 7.4 (0.2% DMSO, v/v). 25, 50 and 100 μ M.



Figure S2: Overlay of time-courses of ThT fluorescence emission intensity of A) $A\beta_{1-42}$, in the absence and in the presence of **Pt-Ac-rhein**, **Pt-Ac-OH** and rhein and compounds alone, and B) upon the addition (indicated by an arrow) of the complexes at 1:1 ratio to preformed amyloid aggregates. The results are representative of two independent measurements



Figure S3. Fluorescence emission spectra at different times of Pt complexes and rhein (λ_{ex} = 440 nm).



Figure S4. Fluorescence emission spectra at different times of $A\beta_{1-42}$ in the absence and presence of compound **Pt-Ac-rhein** or rhein. (λ ex = 275 nm). As insets intensity at 303 nm versus time. A β_{1-42} at 50 μ M with the compounds at 1:1 molar ratio



Figure S5. ESI-MS spectra of: A) $A\beta_{1-42}$ alone, B) $A\beta_{1-42}$ in presence of **Pt-Ac-OH** (at 1:1 molar ratio) and C) **Pt-Ac-OH** alone.



Figure S6. ESI-MS spectra of: A) $A\beta_{1-42}$ alone, B) $A\beta_{1-42}$ in presence of rhein (at 1:1 molar ratio) and C) rhein alone.



Figure S7. Scanning electron microscopy of $A\beta_{1-42}$ alone at a magnification of 330x (300 µm scale bar, A, B, C), 790x (200 µm scale bar, A', B', C') and 2500x (50 µm scale bar, A'', B'', C''). These data represent three of the four independent experiments performed.



Figure S8. Scanning electron microscopy of $A\beta_{1-42}$ in the presence of **Pt-Ac-rhein** at a magnification of 330x (300 µm scale bar, A, B, C), 790x (200 µm scale bar, A', B', C') and 2500x (50 µm scale bar, A'', B'', C''). These data represent three of the four independent experiments performed.



Figure S9. Scanning electron microscopy of $A\beta_{1-42}$ in the presence of rhein at a magnification of 330x (300 µm scale bar, A, B, C), 790x (200 µm scale bar, A', B', C') and 2500x (50 µm scale bar, A'', B'', C''). These data represent three of the four independent experiments performed.



Figure S10. Ligand interactions for best docked complex of Pt-Ac-Rhein with A β_{1-42} monomer and tetramer.



Figure S11. Best docked pose of **Pt-Ac-rhein** with $A\beta_{1-42}$ tetramer. **Pt-Ac-rhein** is shown as ball-andstick, receptor is shown as a surface coloured by lipophilicity: green represents lipophilic areas, pink potential hydrogen bonding sites, and blue polar regions.

Table S1: Table of main observed ions relative to the species formed by the $A\beta_{1-42}$ alone and mixed with **Pt-Ac-rhein**. Experimental and theoretical mass and charge were reported for each adduct.

	Description	m/z (charge)		Theoretical m/z
		Peptide	+metal complex	meoretical m/2
		1505.28 (+3)	1505.29 (+3)	1505.71
	Αβ ₁₋₄₂	1129.42 (+4)	1129.77 (+4)	1129.54
		903.96 (+5)	903.95 (+5)	903.83
Aβ ₁₋₄₂ : Pt-Ac-rhein		753.44 (+6)	753.43 (+6)	753.36
	Aβ ₁₋₄₂ + Pt-Ac-rhein	-	1719.12 (+3) 1290.76 (+4)	1719.71 1290.03

Table S2: Docking ChemPLP scores for Pt-Ac-rhein, rhein and Pt-Ac-OH with $A\beta_{1-42}$ monomer and tetramer

	Pt-Ac-rhein	rhein	Pt-Ac-OH
Monomer 1	-78.0	-72.7	-51.8
Monomer 2	-83.4	-66.5	-62.1
Monomer 3	-86.3	-69.0	-64.1
Monomer 4	-79.3	-73.4	-57.8
Monomer 5	-76.2	-67.0	-54.7
Monomer 6	-75.5	-69.3	-53.6
Monomer 7	-80.0	-89.0	-56.9
Monomer 8	-106.8	-62.7	-57.1
Monomer 9	-79.7	-67.2	-60.1
Monomer 10	-79.3	-80.3	-51.5
Mean monomer	-82.5	-71.7	-57.0
Tetramer 1	-91.4	-72.1	-60.6
Tetramer 2	-81.0	-75.8	-63.1
Tetramer 3	-78.4	-75.5	-60.6

Tetramer 4	-102.4	-93.9	-60.3
Tetramer 5	-78.5	-65.4	-57.4
Mean tetramer	-86.3	-76.5	-60.4