

Supporting information

Composite of KLVFF-Transthyretin-Penetratin and Manganese Dioxide Nanoclusters: A Multifunctional Agent against Alzheimer's β -Amyloid Fibrillogenesis

Haitao Lan^{1,†}, Ying Wang^{1,†}, Wei Liu^{2,*}, Xiaoyan Dong¹ and Yan Sun^{1,*}

¹ Key Laboratory of Systems Bioengineering and Frontiers Science Center for Synthetic Biology (Ministry of Education), Department of Biochemical Engineering, School of Chemical Engineering and Technology, Tianjin University, Tianjin 300350, China; lanhaitao@tju.edu.cn (H.L.); 1020207109@tju.edu.cn (Y.W.); d_xy@tju.edu.cn (X.D.)

² Tianjin Key Laboratory of Radiation Medicine and Molecular Nuclear Medicine, Institute of Radiation Medicine, Chinese Academy of Medical Sciences and Peking Union Medical College, Tianjin 300192, China

* Correspondence: liuwei@irm-cams.ac.cn (W.L.); ysun@tju.edu.cn (Y.S.); Tel./Fax: +86-22-27403389 (Y.S.)

† These authors contributed equally to this work.

Table S1. Lag time (T_{lag}) of A β ₄₀ aggregation kinetics calculated from Figure S2.

Sample	T_{lag} (h)
A β only	49.8 ± 5.8
A β + 10 µg/mL TTR	58.5 ± 2.1
A β + 25 µg/mL TTR	81.1 ± 3.0
A β + 50 µg/mL TTR	86.5 ± 2.2
A β + 10 µg/mL KTP	16.8 ± 1.5
A β + 25 µg/mL KTP	26.4 ± 2.5
A β + 50 µg/mL KTP	— —
A β + 10 µg/mL KTP @MnO ₂	21.2 ± 3.1
A β + 25 µg/mL KTP @MnO ₂	68.1 ± 3.5
A β + 50 µg/mL KTP @MnO ₂	— —

Table S2. The content of secondary structure of A β ₄₀ incubated with different inhibitor was calculated by the BeStSeL algorithm (<http://bestsel.elte.hu/>). Others include 3₁₀-helix, bends, π -helix, β -bridge, and irregular/loop.

Secondary structure	A β only	A β only (0 h)	+ 50 µg/mL TTR	+ 50 µg/mL KTP	+ 50 µg/mL KTP@MnO ₂
Helix	22.4	30.3	12.8	11.3	0
Antiparallel β -sheet	20.5	64.8	35.5	44.9	44.5
Parallel β -sheet	57.1	0	36.8	0	0
Turn	0	0	15	13.2	9.7
Others	0	5	0	30.6	45.8

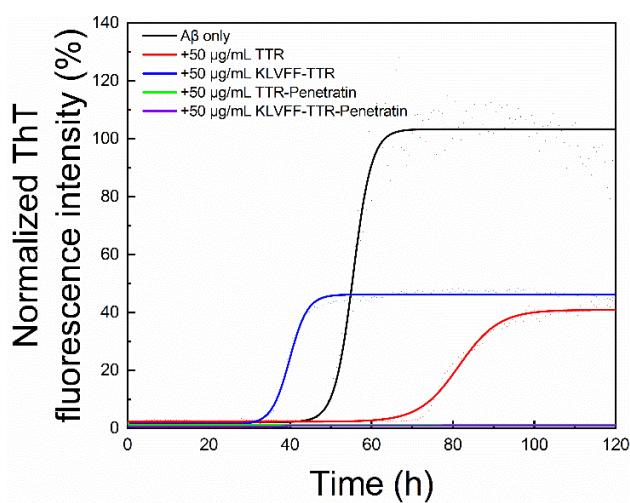


Figure S1. Aggregation kinetics of A β_{40} incubated with TTR-derived proteins.

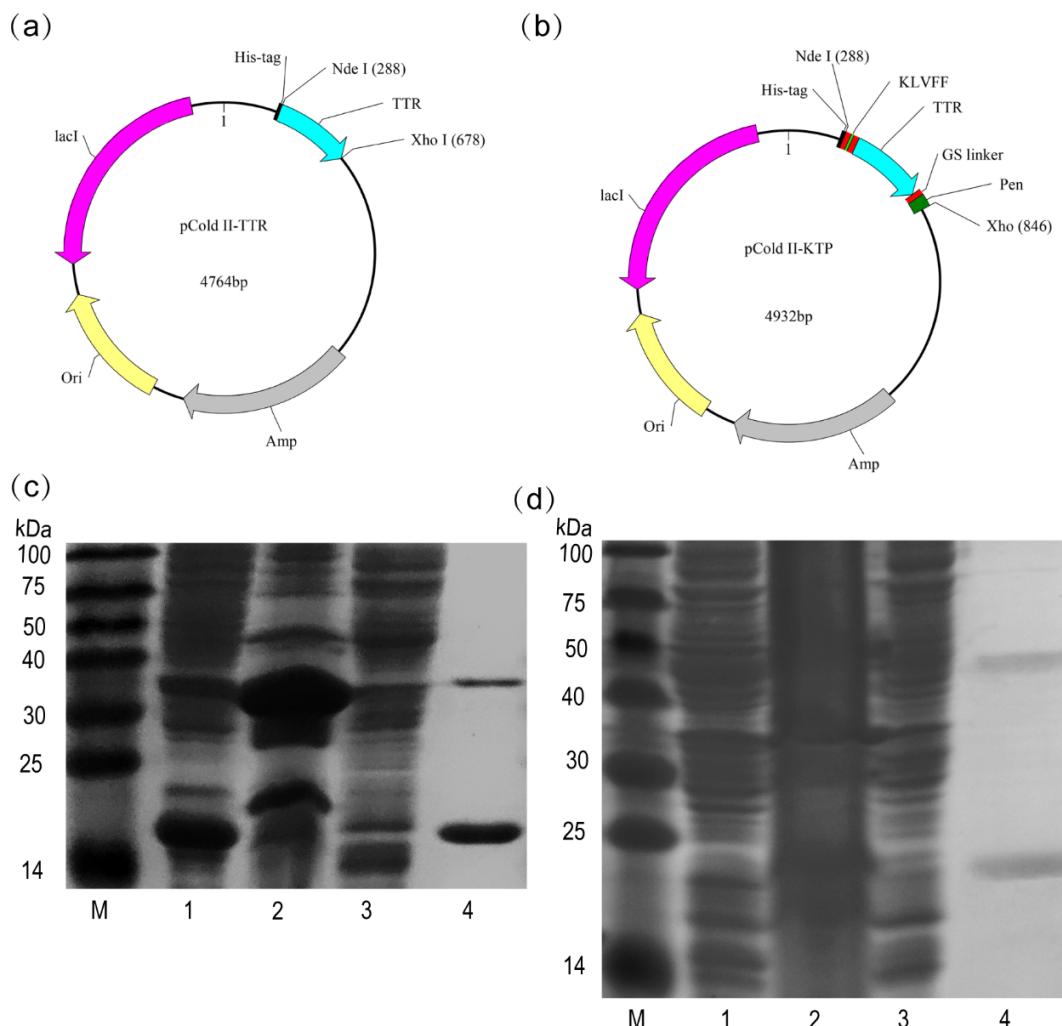


Figure S2. (a)-(b) Construction of pCold II-TTR and pCold II-KTP expression vectors. (c) SDS-PAGE of TTR. Lanes: M, protein marker; 1, supernatant of cell lysate; 2, precipitate of cell lysate; 3, washing eluent; 4, purified TTR. (d) SDS-PAGE of KTP. Lanes: M, protein marker; 1, supernatant of cell lysate; 2, precipitate of cell lysate; 3, washing eluent; 4, purified KTP.

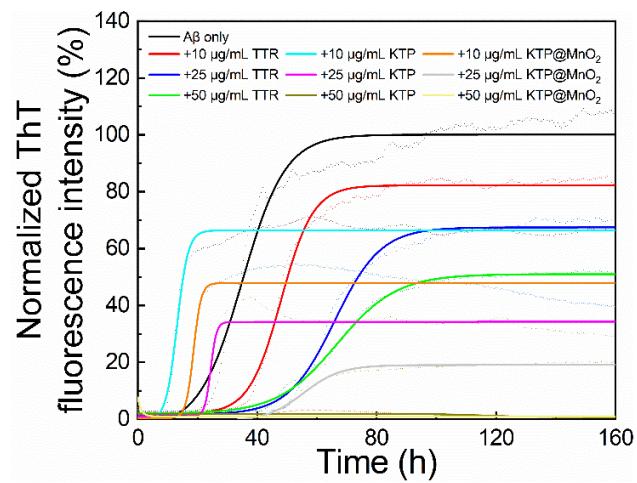


Figure S3. Aggregation kinetics of A β_{40} incubated with different concentrations of TTR, KTP or KTP@MnO₂.

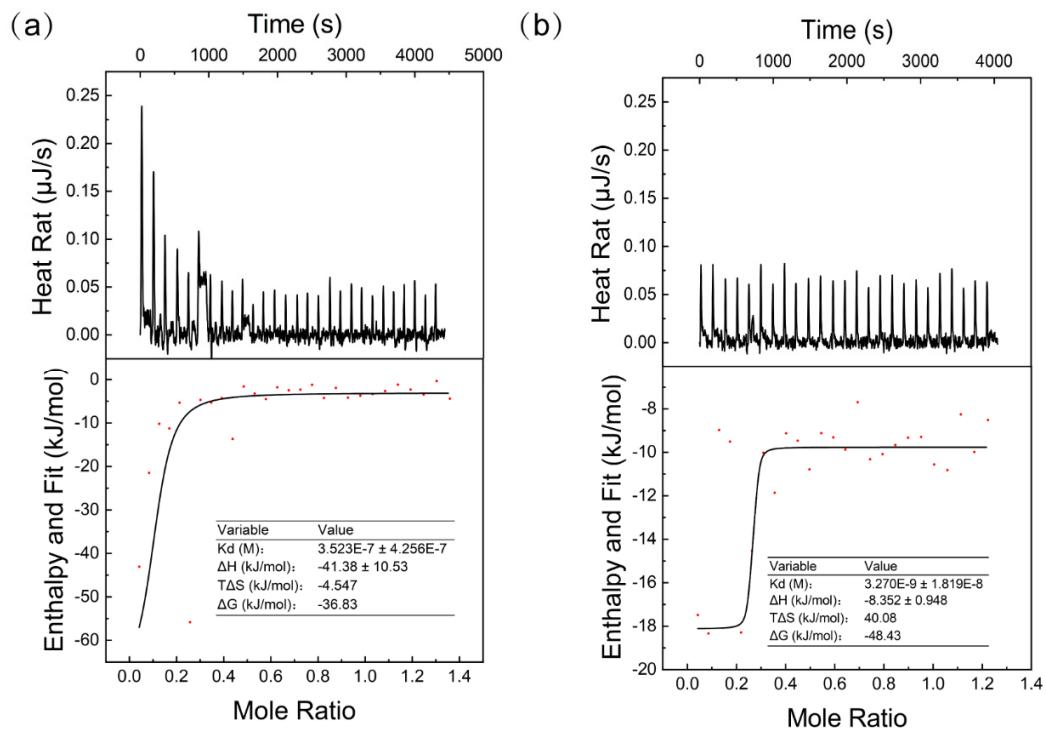


Figure S4. ITC binding isotherm for the titration of (a) TTR and (b) KTP to A β_{40} .

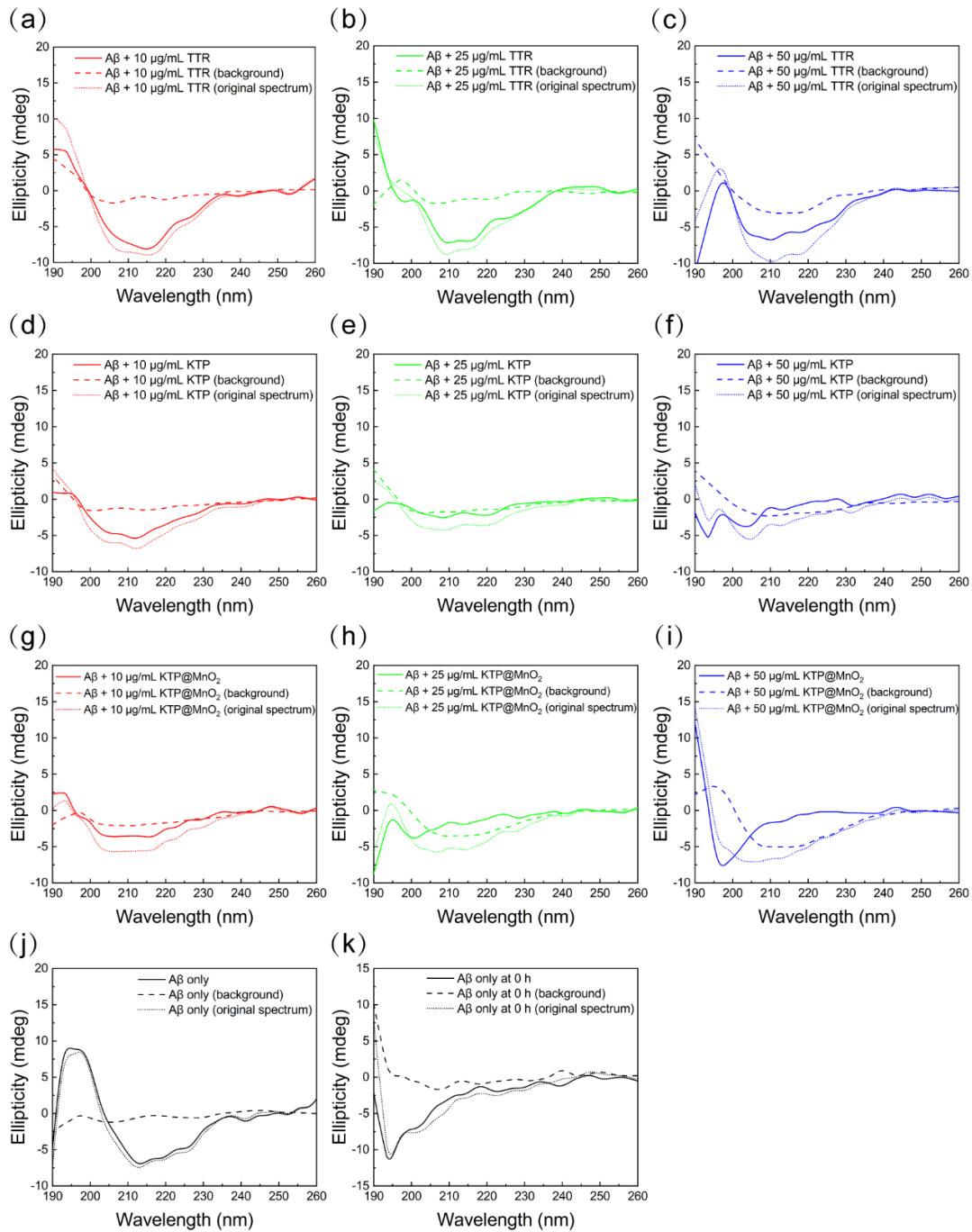


Figure S5. CD spectra of A β_{40} incubated with (a) 10 $\mu\text{g/mL}$ TTR, (b) 25 $\mu\text{g/mL}$ TTR, (c) 50 $\mu\text{g/mL}$ TTR, (d) 10 $\mu\text{g/mL}$ KTP, (e) 25 $\mu\text{g/mL}$ KTP, (f) 50 $\mu\text{g/mL}$ KTP, (g) 10 $\mu\text{g/mL}$ KTP@MnO₂, (h) 25 $\mu\text{g/mL}$ KTP@MnO₂, and (i) 50 $\mu\text{g/mL}$ KTP@MnO₂. (j) CD spectra of A β_{40} after incubation. (k) CD spectra of A β_{40} without incubation. (Background is an A β -free solution, and the spectrum corresponding to the solid line is the original spectrum minus the background spectrum).

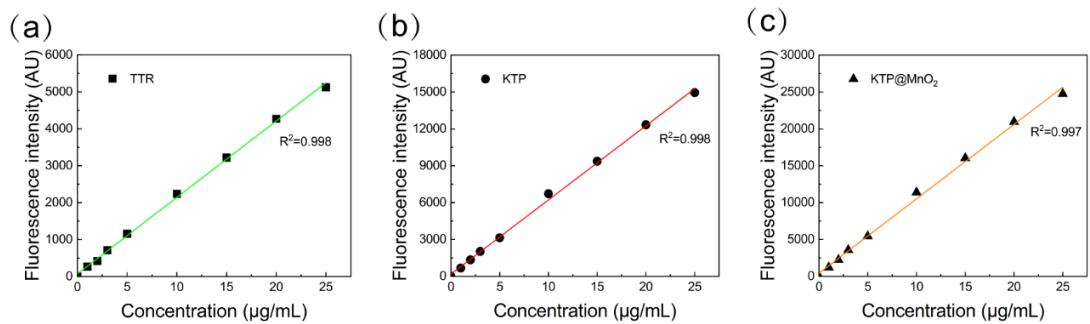


Figure S6. Standard curves of Cy5 fluorescence intensity of (a) TTR, (b) KTP, and (c) KTP@MnO₂.

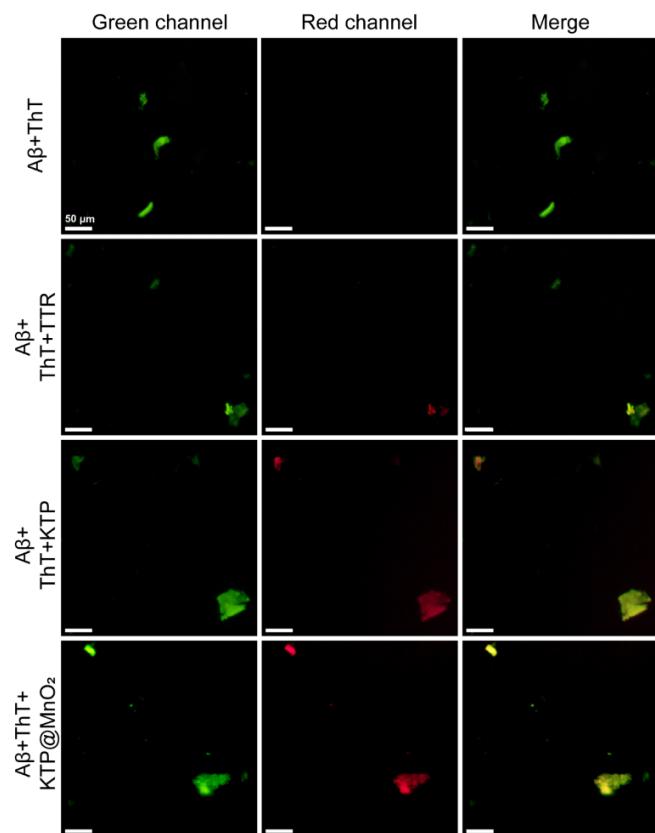


Figure S7. *In vitro* targeting capability of different inhibitors. Aβ plaques were stained with Aβ-specific probe ThT (Green channel) and incubated with Cy5-labelled inhibitors (Red channel). Scale bars, 50 µm.

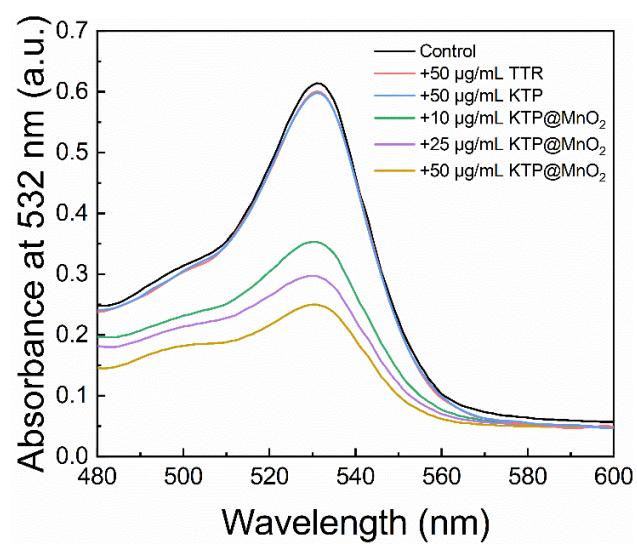


Figure S8. ·OH scavenging ability of different inhibitors.