

Supplementary Information

Effects of the toxic metals arsenite and cadmium on α -synuclein aggregation *in vitro* and in cells

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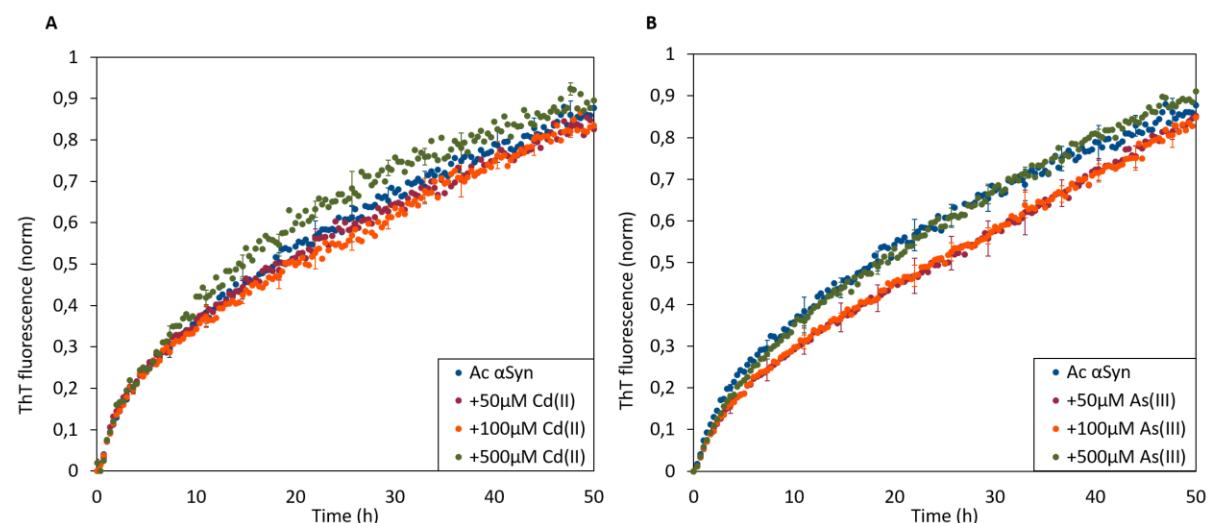


Figure S1 - ThT fluorescence kinetic traces for 50 μM Ac α Syn aggregation (pH 7.4) in the presence of 10% amyloid fibre seeds. The effect of the metals is no longer visible.

Table S1 – Reported values for α Syn amyloid fibril pitch based on AFM and cryo-EM measurements.

Reference	Protein	Method	Pitch
Li Y. et al., 2018 [1]	WT α Syn	CryoEM	119 nm
R. Guerrero-Ferreira et al., 2019 [2]	WT α Syn	CryoEM	108 nm
Li B. et al., 2018 [3]	WT α Syn	CryoEM	rod: 92 nm twister: 46 nm
Ni X. et al., 2019 [4]	Ac α Syn	CryoEM	121 nm
Ni X. et al., 2019 [4]	Ac α Syn1-122	CryoEM	85 nm
Ni X. et al., 2019 [4]	Ac α Syn1-103	CryoEM	65 nm
Zhao K. et al., 2020 [5]	Ac α Syn	AFM	120 nm
Zhao K. et al., 2020 [5]	Ac α Syn E46K	AFM/CryoEM	64 nm
Iyer A. et al., 2016 [6]	Ac α Syn	AFM	115 nm
Boyer D. et al., 2019 [7]	α Syn H50Q	CryoEM	90 nm
Sun Y. et al., 2020 [8]	Ac α Syn A53T	CryoEM	97 nm

Table S2 - Measured values of As(III) and Cd(II) concentrations in ppb, measured using ICP-MS. Samples (1 to 1 protein to metal) were spun down (to get amyloids at the bottom) after no incubation and after 5 days of aggregation, followed by metal analysis of the supernatant. The 'missing' metal from the solution (comparing 0 to 5 days) must have been incorporated into the amyloids.

As ³⁺	Conc. (μM)	Cd ²⁺	Conc. (μM)
AVG As(III) 0 h	50	AVG Cd(II) 0 h	50
AVG As(III) 5 days	20	AVG Cd(II) 5 days	24
Left in solution (%):	40%		49%

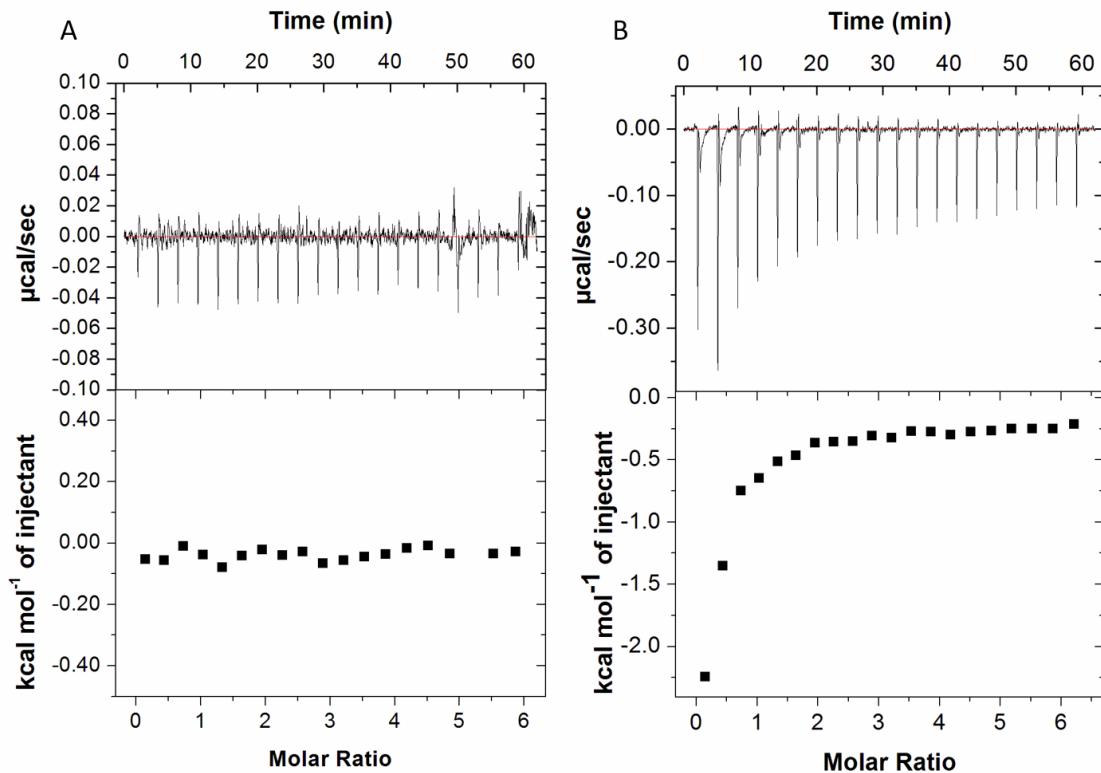


Figure S2 - ITC data of 50 μM monomeric Ac α Syn with addition of A) arsenite (NaAsO₂) and B) cadmium (CdCl₂) at a final ratio of 1:6 monomers to metal ions. There's no quick binding to be detected for As(III). There is a significant heat exchange with the addition of Cd(II) that may suggest a weak binding to Ac α Syn assembled species.

References

1. Li, Y.; Zhao, C.; Luo, F.; Liu, Z.; Gui, X.; Luo, Z.; Zhang, X.; Li, D.; Liu, C.; Li, X. Amyloid fibril structure of α -synuclein determined by cryo-electron microscopy. *Cell Research* **2018**, *28*, 897-903, doi:10.1038/s41422-018-0075-x.
2. Guerrero-Ferreira, R.; Taylor, N.M.; Arteni, A.-A.; Kumari, P.; Mona, D.; Ringler, P.; Britschgi, M.; Lauer, M.E.; Makky, A.; Verasdonck, J.; et al. Two new polymorphic structures of human full-length alpha-synuclein fibrils solved by cryo-electron microscopy. *eLife* **2019**, *8*, doi:10.7554/elife.48907.
3. Li, B.; Ge, P.; Murray, K.A.; Sheth, P.; Zhang, M.; Nair, G.; Sawaya, M.R.; Shin, W.S.; Boyer, D.R.; Ye, S.; et al. Cryo-EM of full-length α -synuclein reveals fibril polymorphs with a common structural kernel. *Nature Communications* **2018**, *9*, doi:10.1038/s41467-018-05971-2.
4. Ni, X.; Mcglinchey, R.P.; Jiang, J.; Lee, J.C. Structural Insights into α -Synuclein Fibril Polymorphism: Effects of Parkinson's Disease-Related C-Terminal Truncations. *Journal of Molecular Biology* **2019**, *431*, 3913-3919, doi:10.1016/j.jmb.2019.07.001.
5. Zhao, K.; Li, Y.; Liu, Z.; Long, H.; Zhao, C.; Luo, F.; Sun, Y.; Tao, Y.; Su, X.-D.; Li, D.; et al. Parkinson's disease associated mutation E46K of α -synuclein triggers the formation of a distinct fibril structure. *Nature Communications* **2020**, *11*, doi:10.1038/s41467-020-16386-3.
6. Iyer, A.; Roeters, S.J.; Schilderink, N.; Hommersom, B.; Heeren, R.M.A.; Woutersen, S.; Claessens, M.M.A.E.; Subramaniam, V. The Impact of N-terminal Acetylation of α -Synuclein on Phospholipid Membrane Binding and Fibril Structure. *Journal of Biological Chemistry* **2016**, *291*, 21110-21122, doi:10.1074/jbc.m116.726612.
7. Boyer, D.R.; Li, B.; Sun, C.; Fan, W.; Sawaya, M.R.; Jiang, L.; Eisenberg, D.S. Structures of fibrils formed by α -synuclein hereditary disease mutant H50Q reveal new polymorphs. *Nature Structural & Molecular Biology* **2019**, *26*, 1044-1052, doi:10.1038/s41594-019-0322-y.
8. Sun, Y.; Hou, S.; Zhao, K.; Long, H.; Liu, Z.; Gao, J.; Zhang, Y.; Su, X.-D.; Li, D.; Liu, C. Cryo-EM structure of full-length α -synuclein amyloid fibril with Parkinson's disease familial A53T mutation. *Cell Research* **2020**, *30*, 360-362, doi:10.1038/s41422-020-0299-4.