

Supplementary Information for:

Repurposing doxycycline for synucleinopathies: remodelling of α -synuclein oligomers towards non-toxic parallel beta-sheet structured species

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1. Analysis of doxycycline binding to α -synuclein by NMR

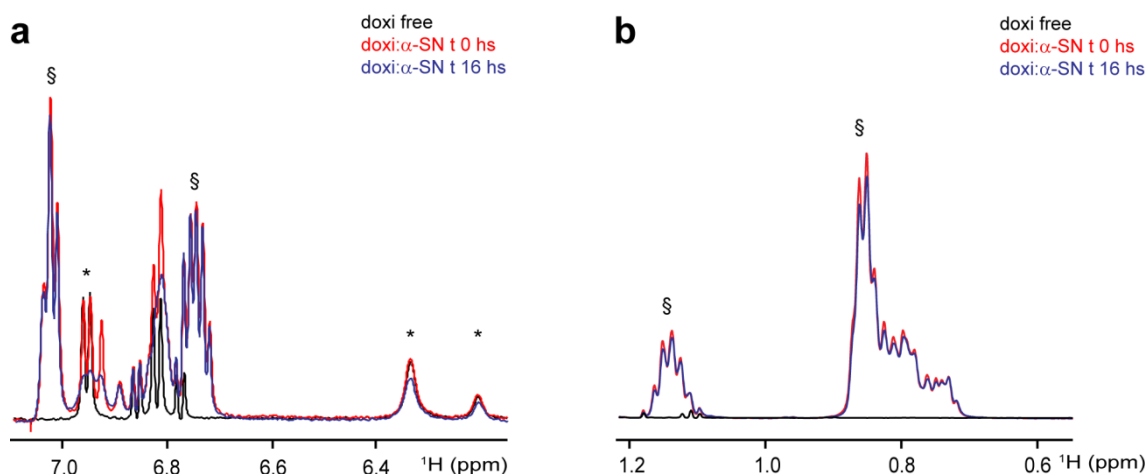


Figure S1. Analysis of doxycycline binding to α -synuclein by NMR. ^1H 1D NMR spectra in the aromatic (a) and methyl region (b) of 200 μM doxycycline alone (black line) and upon addition of 100 μM monomeric α -synuclein (red line) or α -synuclein that was aged for 16 hs (blue line). Before preparing the mixtures, the α -synuclein samples were spun down to remove insoluble particles that might have formed during this time frame. Asteriks denote isolated doxycycline ^1H NMR signals that broaden upon binding to α -synuclein oligomers or protofibrils. The symbols § indicate isolated clusters of α -synuclein ^1H NMR signals. We did not detect major changes in α -synuclein signals between time 0 and 16 hs indicating that slow-tumbling, highly ordered structures such as amyloid fibrils or large protofibrils were not formed at great extents. NMR spectra were acquired at 25°C. Samples were dissolved in 20 mM HEPES supplemented with 150 mM NaCl and 10% D_2O .

2. Dose:response effect of doxycycline on α -synuclein aggregation

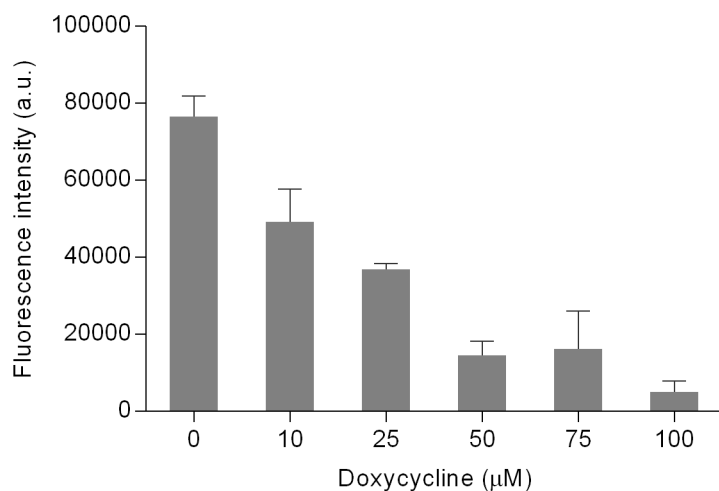


Figure S2. Dose-response effect of doxycycline on the α -synuclein fibrillation process. A solution containing 70 μM α -synuclein in 20 mM HEPES buffer, 150 mM NaCl, pH 7.4 was incubated at 37°C under orbital agitation for 48 h in the presence of 0 μM , 10 μM , 25 μM , 50 μM , 75 μM and 100 μM doxycycline as described in Methods. On each sample Thioflavin T was added to a final concentration of 25 μM and the formation of cross-beta structure was estimated by the increase in fluorescence at $\lambda_{\text{exc}}=450$ nm, $\lambda_{\text{em}}=482$ nm.