Supplementary Information

Liquid-Liquid Phase Separation Induces Pathogenic Tau Conformations In Vitro Nicholas M. Kanaan et al.

Supplementary Methods

Fluorescent Labeling of Tau

Purified recombinant Tau was labeled with FITC and Alexa568 to establish the best approach for studying tau liquid droplet formation. First, tau protein (1 mg/ml) was buffer exchanged into 1x phosphate buffered saline using Micro Bio-Spin 6 columns (BioRad, 7326200) per the manufacturer's instructions. The proteins were then labeled with FITC (FITC-Tau, Thermo F6434) or Alexa568 (A568-Tau, Thermo, A20184) following the protocols provided by the manufacturers. Briefly, FITC labeled tau protein was created by adding 20 µl of 1M sodium bicarbonate buffer (from kit) to 200 µl of tau protein, then 10 µl of the reactive FITC dye (10 mg/ml stock in DMSO) was added while gently stirring the sample to mix, followed by incubation at room temperature for 1 hr (protected from light). After labeling, the protein sample was cleaned of free dye and buffer exchanged into PBS (pH 7.2, with 2mM sodium azide) using the spin columns provided following the kit instructions. AlexaFluor568 labeled tau protein was generated by adding 10 μ of 1M sodium bicarbonate buffer (from kit) to 100 μ of tau protein, then 100 µl of this solution was added to reactive AlexaFluor 568 dye and gently inverted to mix, followed by incubation at room temperature for 1 hr (protected from light) with gentle inversion to mix every 10 minutes. After labeling, the protein sample was cleaned of free dye and buffer exchanged into PBS (pH 7.2, with 2mM sodium azide) using the spin columns provided following the kit instructions. The labeling efficiency was calculated at $0.6 - 0.7$ M dye per M of tau protein.

Arachidonic Acid-Induced Tau Aggregation *In Vitro*

In vitro aggregation of Tau, Tau-GFP, FITC-Tau and A568-Tau (all at 2 µM) was induced using arachidonic acid (75 µM) in tau polymerization buffer (100mM NaCl, 10mM HEPES (pH 7.6), 5mM DTT and 0.1mM EDTA) as described in detail previously1. Aggregated tau samples were imaged using standard electron microscopy imaging as detailed below1.

Microtubule Binding Assay and Western Blotting

The effects of labeling tau with GFP or conjugating tau to fluorescent molecules (i.e. FITC and Alexa 568) on microtubule binding *in vitro* was assessed using the Microtubule Binding Protein Spin-Down Assay according to the manufacturer's instructions (Cytoskeleton, BK029). Briefly, microtubules (5 µM tubulin dimers) were generated at 35˚ C for 20 minutes and then 20 µM taxol was added to stabilize the microtubules. The samples tested included microtubules only (no tau) and either unlabeled Tau, Tau-GFP, FITC-Tau or A568-Tau (all at 0.5 µM final concentration) without and with microtubules (in a total reaction volume of 50 µl). These samples were incubated for 30 minutes at room temperature while binding took place and then spun over a sucrose cushion (100 µl, supplemented with taxol) at 100,000 x g at room temperature for 40 minutes. The supernatant was collected for each sample (50 µl, 10 µl of 6x Laemmli buffer was added), the cushion was discarded and the pellet was collected by adding 50 µl 1x Laemmli buffer.

Standard SDS-PAGE and Western blotting analyses as described previously² were used to evaluate results of the microtubule binding assay. Briefly, samples were heated at 95° C for 5 minutes before being separated on 26-well BioRad precast Criterion 4-20% TGX gels for ~30 minutes at 250 V. The supernatant and pellet fractions were run for each sample. The transfer was 50 minutes at 400 mA and then the membranes were blocked in 2% non-fat dry milk for 30 minutes. The blots were then incubated in a mixture of R1 antibody³ (a rabbit polyclonal pan-tau antibody; AB 2832929) diluted 1:10,000 and 5H1 antibody⁴ (a β -tubulin antibody, IgM mouse monoclonal; AB 2832941) diluted 1:15,000 in 2% non-fat dry milk overnight at 4° C. The following day, the blots were rinsed in TBS with 0.1% tween-20 and then incubated for 1 hour in a mixture of goat anti-rabbit 800 (Licor, 926-32211, AB_621843) and goat anti-mouse IgM Alexa Fluor 680 (Jackson ImmunoResearch, 115-625-075, AB_2338934) secondary antibodies each diluted 1:20,000 in non-fat dry milk. The blots were rinsed again and imaged on the Licor system. The signal intensity of the tau bands were quantified and then the signal from the supernatant and pellet fractions were added together to obtain a value representing 100% of the tau signal for each sample set. Then the signals for the supernatant and pellet fractions were expressed as a

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percentage of the total tau signal. This experiment was repeated twice with similar results.

Generation and Characterization of MTBR Monoclonal Antibody

We generated a mouse monoclonal antibody against the MTBRs, which was used here to probe the MTBR-GFP proteins on immunoblots. Mice were housed with a normal light cycle at room temperature and humidity with food and water provided *ad libitum*. An adult (10 week-old) female tau knockout mouse (B6.129X1-Mapttm1Hnd/J; Jackson Labs, 007251) was immunized with a tau peptide conjugated to keyhole limpet hemocyanin (KLH-CSTENLKHQPGGGKVQIVYKPVDLSKVT, Genscript) and antibody hybridomas (fusion with SP2/o myeloma cells, Kanaan lab) were generated and screened using methods similar to those described previously by our group⁵. A stable hybridoma line producing an antibody, referred to as the MTBR3 antibody (a mouse IgG1 isotype, AB_2832994), was obtained and subcloned at least 3 times. The MTBR3 antibody displayed strong reactivity against the immunizing peptide (without KLH) and was validated for reactivity against full-length 4R (2N4R) and 3R (2N3R) human tau isoforms using ELISA titer assays as described 5 (Supplementary Fig. 5a). The MTBR3 antibody, was further validated for its utility with immunoblots (the purpose of its use here) using recombinant full-length human 4R and 3R proteins. The ELISA titer and recombinant protein experiments were repeated twice with similar results. All rodent uses/procedures conducted in compliance with federal, state and institutional guidelines and were approved by the Michigan State University Institutional Animal Care and Use Committee (protocol 01/17-004-00).

Supplementary Figures

Supplementary Fig. 1. Strategy for studying tau liquid-liquid phase separation. a, Representative blots from a microtubule binding assay to measure binding of Tau, Tau-GFP, FITC-Tau and Alexa568(A568)-Tau proteins. **b,** Quantitation of tau immunoblots

of soluble (S) and pelleted (P) fractions from a representative experiment indicate that Tau and Tau-GFP display the same degree of effective microtubule binding as assessed by their co-pelleting with microtubules (MT), while crosslink labeling of tau with either FITC or A568 impairs microtubule binding. Representative data from one experiment presented, this experiment was repeated two independent times with similar results. **c,** Transmission electron micrographs of Tau and Tau-GFP aggregates (2 µM) formed, which included small globular oligomers and short, intermediate and long filaments typically found with arachidonic acid-induced aggregation *in vitro*. In contrast, FITC-Tau and A568-Tau (2 µM) do not form normal tau aggregate species, but instead form large clumps of protein. Scale bar is 200 nm. These experiments were repeated three independent times. **d, e,** Despite the lack of normal microtubule binding and *in vitro* aggregation properties of FITC-Tau and A568-Tau, both of these labeled proteins undergo liquid droplet formation (4 μ M) when incubated with PEG. Scale bar is 10 μ m. These experiments were repeated three independent times. **f,** Multiple independent preparations of recombinant Tau-GFP proteins (n = 3) produced the same results confirming that tau phase separation was not due to abnormalities with a single protein preparation. Scale bar is 10 µm. **g,** Incubation of GFP alone (8 µM) with 10% PEG did not produce liquid droplet formation indicating GFP does not undergo phase separation at 2-4 times the amount of tau-GFP constructs used in this study. Scale bar is 10 µm. Source data for panels **a**, **b** provided as a Source Data file.

Supplementary Fig. 2 Comparison of each tau construct preparation. a, Unlabeled Tau, P301L, AT8, Tau-GFP, P301L-GFP and AT8-GFP recombinant proteins are comparable to one another as assessed by Coomassie stained SDS-PAGE gel analysis. 3 µg of each protein was loaded per lane. **b**, GFP alone, Tau-GFP, the Nterminus with the microtubule binding repeats (NTMT-GFP; amino acids 1-380), the MTs with the C-terminus (MTCT-GFP; amino acids 225-441), the NT alone (NT-GFP, amino acids 1-224), the MTs alone (MT-GFP; amino acids 225-380), and the CT alone (CT-GFP; amino acids 381-441) recombinant proteins are comparable to one another as assessed by Coomassie stained SDS-PAGE gel analysis. 2 µg of each protein was loaded per lane. Source data for panels **a**, **b** provided as a Source Data file.

Supplementary Fig. 3. **Formation of pathological tau conformations and oligomers is caused by molecular crowding. a,** Representative dot blots of Tau, P301L and AT8 incubated without (-PEG, monomeric tau) or with PEG (+PEG, liquid droplets) for 4 hours. Blot was probed for total tau (R1 antibody) and PAD-exposed tau (TNT2 antibody). **b,** Quantitation of dot blots show significant increases in PAD exposure upon LLPS of Tau, P301L and AT8 proteins (n = 3 independent experiments, data are mean \pm SD, two-way ANOVA with Sidak post-hoc test; Interaction: $F_{(2, 12)}$ = 0.5928, p = 0.5682; Crowding Factor: $F_{(1, 12)} = 84.83$, p < 0.0001; Tau Factor: $F_{(2, 12)} =$ 0.5449, p = 0.5936). **c,** Representative dot blots of Tau, P301L and AT8 incubated without (–PEG) or with PEG (+PEG) for 4 hours. Blot was probed for total tau (R1 antibody) and oligomeric tau (TOC1 antibody). **d,** Quantitation of dot blots show significant increases in oligomeric tau species upon LLPS of Tau P301L and AT8 (n = 3 independent experiments, data are mean ±SD, two-way ANOVA with Sidak post-hoc test; Interaction: $F_{(2, 12)} = 0.4173$, p = 0.6681; Crowding Factor: $F_{(1, 12)} = 84.26$, p <

0.0001; Tau Factor: $F_{(2, 12)} = 0.8712$, $p = 0.4433$). Source data for panels **a-d** provided as a Source Data file.

Supplementary Fig. 4. Image analysis of tau liquid droplets. a, Images of tau GFP fusion proteins were acquired on a confocal microscope system. All images within an experiment were acquired using identical confocal settings (scan speed, resolution, magnification, laser intensity, gain, and offset). Scale bar is 10 µm. **b,** Images were analyzed by applying threshold functions in ImageJ software that identified the phase separated tau liquid droplets and excluded the background of the image. All droplets within the threshold limits were analyzed for total area (μm^2) , average size of individual droplets (μ m²) and mean fluorescence intensity of individual droplets (arbitrary units).

Supplementary Fig. 5. Characterization of MTBR3 antibody. A novel antibody against the MTBR domain of tau was generated by immunizing tau knockout mice with a peptide containing sequences between amino acids 262 and 319 of human tau protein. **a,** The MTBR3 antibody was titered in an ELISA using full-length human 4R tau isoform (2N4R) and 3R tau isoform (2N3R) proteins (representative experiment that was replicated 2 times with similar results, data analyzed using nonlinear regression). **b,** To confirm the MTBR3 antibody reacts with recombinant tau on immunoblots, 2N4R and 2N3R proteins were used (monomeric, Mono and aggregated tau samples, Agg). Tau was confirmed using the polyclonal R1 tau antibody. Note the MTBR3 antibody reacts with both tau isoforms in each assay, but displays a higher affinity for 3R proteins (representative experiment that was replicated 2 times with similar results). Source data for panels **a**, **b** provided as a Source Data file.

Supplementary Table 1. Resources used in this study.

Supplementary Table 2. Mutagenesis and PCR Primers.

Note: All primers from Integrated DNA Technologies, Inc. and all restriction enzymes from ThermoFisher Scientific.

Supplementary References

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