



Published in final edited form as:

Neurobiol Dis. 2020 February ; 134: 104668. doi:10.1016/j.nbd.2019.104668.

A peptide inhibitor of Tau-SH3 interactions ameliorates amyloid- β toxicity

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Abstract

The microtubule-associated protein Tau is strongly implicated in Alzheimer's disease (AD) and aggregates into neurofibrillary tangles in AD. Genetic reduction of Tau is protective in several animal models of AD and cell culture models of amyloid- β (A β) toxicity, making it an exciting therapeutic target for treating AD. A variety of evidence indicates that Tau's interactions with Fyn kinase and other SH3 domain-containing proteins, which bind to PxxP motifs in Tau's proline-rich domain, may contribute to AD deficits and A β toxicity. Thus, we sought to determine if inhibiting Tau-SH3 interactions ameliorates A β toxicity. We developed a peptide inhibitor of Tau-SH3 interactions and a proximity ligation assay (PLA)-based target engagement assay. Then, we used membrane trafficking and neurite degeneration assays to determine if inhibiting Tau-SH3 interactions ameliorated A β oligomer (A β o)-induced toxicity in primary hippocampal neurons from rats. We verified that Tau reduction ameliorated A β o toxicity in neurons. Using PLA, we identified a peptide inhibitor that reduced Tau-SH3 interactions in HEK-293 cells and primary neurons. This peptide reduced Tau phosphorylation by Fyn without affecting Fyn's kinase activity state. In primary neurons, endogenous Tau-Fyn interaction was present primarily in neurites and was reduced by the peptide inhibitor, from which we inferred target engagement. Reducing

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Authors' contributions

TR, JNC, and EDR conceived of the study. TR, JRR, and JNC developed assays and reagents for the study. TR, JRR, and EDR designed experiments, which were performed by TR, JRR, SJT, and ARA, all of whom completed statistical analysis. Data was analyzed by TR, JRR, and EDR. The manuscript was written by JRR and EDR with input from TR, SJT, and JNC. All authors read and approved the final manuscript.

Declaration of Competing Interest

EDR is an owner of intellectual property relating to Tau.

Availability of supporting data

The data sets used and analyzed during this study are available from the corresponding author upon reasonable request.

Consent for publication

Not applicable.

Ethics approval

All experiments were conducted in accordance with the guidelines set forth by the Institutional Animal Care and Use Committee at the University of Alabama at Birmingham.

Appendix A. Supplementary data

Supplementary data to this article can be found online at <https://doi.org/10.1016/j.nbd.2019.104668>.

Tau-SH3 interactions in neurons ameliorated A β toxicity by multiple outcome measures, namely A β -induced membrane trafficking abnormalities and neurite degeneration. Our results indicate that Tau-SH3 interactions are critical for A β toxicity and that inhibiting them is a promising therapeutic target for AD.

Keywords

Amyloid- β ; Oligomer; Tau; SH3; Fyn; Alzheimer's disease

1. Introduction

The microtubule-associated protein Tau canonically stabilizes microtubules (Weingarten et al., 1975) and regulates many cellular processes beyond microtubule dynamics including axonal transport, DNA maintenance, and both pre- and post-synaptic signalling (Wang and Mandelkow, 2016). Mutations in Tau lead to the neurodegenerative disease frontotemporal dementia (FTD) (Rademakers et al., 2004) and Tau inclusions are found in many neurodegenerative disorders including FTD (Hutton et al., 1998), Alzheimer's disease (AD) (Grundke-Iqbal et al., 1986; Ihara et al., 1986), and temporal lobe epilepsy (Tai et al., 2016). Therapies for these diseases are desperately needed and while Tau is an exciting potential target, the question of how to best target Tau remains open.

The discovery that genetic reduction of Tau prevents behavioral and cognitive deficits in an AD mouse model provided an important insight with translational implications (Roberson et al., 2007). This finding has been replicated in multiple AD models (Ittner et al., 2010; Leroy et al., 2012; Meilandt et al., 2008; Nussbaum et al., 2012; Palop et al., 2007a; Roberson et al., 2011) as well as in other disease models including epilepsy (Gheyara et al., 2014; Holth et al., 2013). Tau reduction also ameliorates network hyperexcitability, a potential early driver of AD deficits (Palop and Mucke, 2009), in AD models (Ittner et al., 2010; Roberson et al., 2007; Roberson et al., 2011) and in wild-type mice (Devos et al., 2013; Li et al., 2014). Thus, Tau's role in neurodegeneration may be due to its influence on network hyperexcitability.

Due to the beneficial effects of Tau reduction across diverse models, many studies have examined its underlying mechanisms, seeking to replicate its beneficial effects therapeutically. However, efforts targeting Tau post-translational modifications, Tau aggregation, or microtubule stabilization have not yet been clinically successful (Congdon and Sigurdsson, 2018), highlighting the importance of developing new approaches to target Tau therapeutically (Cummings et al., 2019).

One such approach is inhibiting the interaction between Tau and other AD-related proteins. Tau interacts with several proteins implicated in AD, many of which contain SH3 domains and bind Tau's central proline-rich domain. The proline-rich domain is heavily posttranslationally modified in AD (Gong et al., 2005) and has several PxxP motifs that mediate binding to SH3 domain-containing proteins, including Fyn kinase (Cochran et al., 2014; Ittner et al., 2010; Lee et al., 1998). Fyn is strongly implicated in AD and exogenous A β activates Fyn in dendritic spines, leading to NMDA receptor

(NMDAR) phosphorylation and excitotoxicity in neurons (Ittner et al., 2010; Roberson et al., 2011). Diverse evidence provides a strong premise for therapeutically targeting Tau-SH3 interactions, especially the Tau-Fyn interaction. Manipulating Tau or Fyn produces a converging phenotype: either Tau (Rapoport et al., 2002) or Fyn (Lambert et al., 1998) knockout protects against A β toxicity, and both proteins promote network hyperexcitability (Holth et al., 2013; Kojima et al., 1998; Roberson et al., 2007; Roberson et al., 2011). Tau reduction prevents cognitive deficits caused by Fyn overexpression in an AD mouse model (Roberson et al., 2011). In addition, Tau plays an important role in the dendrite and postsynapse (Ittner and Ittner, 2018) and mice expressing a truncated form of Tau that excludes Fyn from dendrites are protected against A β -induced cognitive deficits and seizure susceptibility (Ittner et al., 2010), showing that these abnormalities may be influenced by Tau-SH3 interactions like that of Tau-Fyn. However, whether Tau-SH3 interactions directly contribute to these effects, as opposed to more indirect mechanisms, remains unknown.

In this study, we sought to determine the effects of inhibiting Tau-SH3 interactions, a novel target for treating AD. Since Tau reduction ameliorates A β toxicity in primary neurons (Rapoport et al., 2002), we used that system to explore a potential mechanism by which Tau reduction prevents A β toxicity. These studies provide insight to the mechanism by which A β toxicity is Tau-dependent and identify Tau-SH3 interactions as a potential therapeutic target for AD.

2. Materials and methods

2.1. Primary hippocampal neurons

All animal procedures were approved by the Institutional Animal Care and Use Committee of the University of Alabama at Birmingham. Timed-pregnant, albino Sprague-Dawley rats (Charles River) were euthanized by isoflurane anesthesia asphyxiation followed by bilateral thoracotomy. Hippocampal tissue from E19 embryos was harvested on ice in cold Hibernate E media (4 °C; Life Technologies A12476–01), and then digested with Papain (20 units/mL, Worthington Biochemical Corporation LK003178) for 10 min at 37 °C. After the incubation, neurons were dissociated by manual trituration to a single-cell suspension in Neurobasal medium (ThermoFisher 21,103,049) supplemented with 1 \times B-27 (ThermoFisher 17,504,044), 2 mM L-Glutamine (ThermoFisher 25,030,081) and 10% premium select fetal bovine serum (Atlanta Biologicals S11550). Neurons were then plated at 30,000 neurons per well (9.0×10^4 neurons per cm²) in 200 μ L plating medium. Neurons were plated in the inner 60 wells of 96-well plates coated overnight at 4 °C with 0.1 mg/mL Poly-D-Lysine (Millipore Sigma P6407) and 0.02 mg/mL laminin (Millipore Sigma L2020) 24–48 h prior to the neuron harvest, with the outer wells containing autoclaved ultrapure water (MilliQ filtered) to prevent evaporation in wells with neurons. Neurons used for the MTT membrane trafficking assay were plated in clear plastic 96-well plates (Fisher 08–772–2C), and those used for the MAP2 neurite degeneration assay were plated in black plastic, clear bottom 96-well plates (Fisher 07–200–565). Cultures were maintained in a 37 °C humidified incubator with 5% CO₂. 24 h after plating, 75% media was exchanged for Neurobasal supplemented with B-27 and L-Glutamine but lacking serum. 5 μ M cytosine β -D-arabino-furanoside (Millipore Sigma C6645) was added at DIV2 to inhibit glial proliferation. 50% media

changes were performed weekly with Neurobasal supplemented with B-27 and L-Glutamine until experiments were started at DIV19–21.

2.2. Antisense oligonucleotide application

At DIV14, neurons were treated with either 1 μ M Tau ASO (sequence: 5-ATCACTGATTTTGAAGTCCC-3), 1 μ M nontargeting control ASO (sequence: 5-CCTTCCCTGAAGTTCCTCC-3), both ASOs were previously published (Devos et al., 2013), or 1 μ M Fyn ASO (sequence: 5-CACAGCCCATTATCCA-3), which was previously published (Girasol et al., 2009), and ordered from IDT. ASO was left on for one week before performing experiments.

2.3. Inhibitor peptides

Peptides were synthesized by Peptide 2.0. Tau-PxxP_{5/6}, rrrqrrkkrgrSRTPSLPTPPTREPCK, consisted of amino acids 209–225 of Tau, with an N-terminal TAT tag (lower case) for cell permeability. Tau-CTD, rrrqrrkkrgrENAKAKTDHGAEIVYKS, consisted of amino acids 380–396 of Tau, also with an N-terminal TAT tag. Peptides were received as lyophilized powder, dissolved in DMSO to 30 mM, then ali-quoted and frozen until use, with a final working concentration of 15 μ M, unless otherwise stated.

2.4. Western blotting

For primary neurons, one week after ASO application, neurons were lysed using RIPA buffer (50 mM Tris, pH 7.5, 150 mM NaCl, 5 mM EDTA, 0.1% SDS, 0.1% Triton X100, 0.5% sodium deoxycholate) with protease inhibitors and centrifuged to remove cell debris. HEK-293 cells were lysed with lysis buffer (50 mM Tris, 150 mM NaCl, 5 mM EDTA, 1% Triton X-100, 0.1% sodium deoxycholate) with protease inhibitor and phosphatase inhibitor can centrifuged to remove cell debris. The samples were diluted with LDS buffer (Thermo Fisher NP0007) and sample reducing agent (final concentration 50 mM dithiothreitol, Fisher B0009), and then heated at 70 °C for 10 min. The samples were then run on 4–12% bis-tris gels (ThermoFisher NW04127BOX) in SDS buffer (25 mM Tris base, 190 mM glycine, 0.1% SDS; pH 8.3) with 20 μ g of protein per lane. The gels were transferred to Immobilon-FL PVDF membranes (Fisher IPFL00010), and the membranes were blocked for 1 h in 50% Odyssey blocking buffer (Fisher NC9125955). The membranes were probed with primary antibody overnight at 4C. The next day, the membranes were incubated at room temperature for 1 h in an IRDye 700 or 800-conjugated secondary antibody (1:20,000 Fisher NC0252291 or Fisher NC0252290) and scanned on an Odyssey Scanner (LI-COR Biotechnology). Bands were quantitated using Image Studio Lite software (LI-COR Biotechnology) and ImageJ. Statistical analysis was performed with GraphPad Prism 8. Antibodies used were: rabbit monoclonal anti-Tau (1:1000, DAKO #A0024), mouse monoclonal antiNeuN (Millipore MAB377), mouse monoclonal anti-pY18 Tau (1:2500 GeneTex 9G3), rabbit polyclonal anti-SFKpY420 (1:1000, Cell Signalling Technology #2010), rabbit polyclonal anti-SFKpY531 (1:1000, Cell Signalling Technology #2015), mouse monoclonal anti-Fyn (1:500, Fyn15, Santa Cruz sc-434, used for HEK-293 lysate), rabbit polyclonal anti-Fyn (1:1000, Fyn3, Santa Cruz sc-16, used for neuron lysate), mouse monoclonal anti-GAPDH (1:5000, Millipore MAB374).

2.5. Amyloid- β oligomer preparation

Lyophilized recombinant amyloid beta 1–42 (A β ; California Peptide 641–15) was dissolved in 1,1,1,3,3,3-Hexafluoro-2-propanol (Millipore Sigma 52517), dried overnight, and then stored in a desiccator at –20 °C until oligomerization. To prepare oligomers, A β was dissolved in fresh, anhydrous DMSO (Millipore Sigma D2650) to 1 mM, vortexed and incubated for 5 min at room temperature, and then diluted to 100 μ M in 1 \times PBS on ice. The A β solution was then sonicated in an ice bath off and on at 30 s increments for 15 min and then left on ice undisturbed to oligomerize for 24 h. Immediately prior to use, A β o was centrifuged at 4 °C for 5 min at 5,000 \times g. Vehicle solution was prepared in identical fashion, beginning with 1,1,1,3,3,3-Hexafluoro-2-propanol evaporation, except lacking A β peptide.

2.6. Modified MTT assay for membrane trafficking

After 24 h exposure to A β o (or vehicle), 100 μ M Tetrazoleum salts (MTT; Millipore Sigma M2128) were applied to neurons and incubated at 37 °C for 60 min. Media from each well was replaced with 15 μ L 1xPBS containing 1.6% Tween[®] 20 (Fisher BP337500) and incubated at room temperature with mild agitation for 5 min. The entire sample volume (referred to as “Tween Soluble MTT”) was then transferred to a 384-well white-plastic, clear bottom plate, and 15 μ L of isopropanol (Millipore Sigma 190764) was added to the plate. The isopropanol containing plates were incubated at room temperature with mild agitation for 5 min, and then (referred to as “Tween Insoluble MTT”) transferred to distinct wells of the same 384-well plate as the associated tween-containing samples. Sample pH was raised by adding 2 μ L of 1 N NaOH to each well to improve formazan absorbance spectrum. The 384-well plates were spun for 2 min at 100 \times g to eliminate any confounding air bubbles, and then absorbance at 590 nm read on a Synergy2 (BioTek) plate reader with absorbance at 660 nm used as the reference wavelength. The ratio of Tween Soluble MTT to Tween Insoluble MTT was calculated on a per sample basis, and then ratios normalized to the vehicle treated group average per experiment.

2.7. MAP2 immunocytochemistry for neurite retraction assay

After 48 h exposure to A β o (or vehicle), neurons were fixed in 1xPBS containing 4% paraformaldehyde and 4% sucrose at 37 °C for 45 min. Neurons were washed 3 \times 5 mins in 1xPBS, blocked in saponin blocking buffer (5% normal horse serum, 5% normal goat serum, 1% bovine serum albumin, and 0.5% saponin in 1xPBS) for 1 h, and then incubated overnight at 4 °C with MAP2 antibody (ThermoFisher, PA1–10005, 1:5000) in saponin blocking buffer. After overnight incubation, neurons were washed 3 \times 5 mins with saponin rinse buffer (0.5% normal horse serum, 0.5% normal goat serum, 0.05% saponin in 1xPBS). Neurons were washed 4x5minutes with saponin rinse buffer, then secondary antibody (Donkey anti-chicken conjugated with Alexa-594, Jackson ImmunoResearch, 702–585-155, 1:2000) applied for 1 hat room temperature in saponin blocking buffer. Neurons were rinsed 4x5minutes with saponin rinse buffer then washed 5x5mins with 1xPBS. Plates were wrapped in parafilm, protected from light, and stored at 4 °C until imaging. Neurons were imaged on the Operetta high-content Imager (PerkinElmer) with a 40 \times objective collecting 4 adjacent fields per well. Intact neurite length was measured in an unbiased, automated manner using Harmony software (PerkinElmer). Intact neurites have relatively

smooth MAP2 labeling, and the neurite detection algorithm excludes neurites that have high contrast along their path (e.g. produced by blebbing). See Supplementary Fig. 3 for workflow. For analysis, total intact neurite length from each condition was normalized to vehicle-treated control wells from each plate and compared by Two-Way ANOVA followed by Dunnett's post hoc test with correction for multiple comparisons.

2.8. HEK-293 transfection and inhibitor application

HEK-293 cells were passaged by removing media and washing with 1xPBS, then applying 0.25% trypsin (Fisher 25–200–056) to cells for 3–5 min at 37 °C to dissociate them from the plate. Trypsin was inactivated by diluting it 1:10 in DMEM (Fisher MT10013CV) supplemented with 10% premium select fetal bovine serum and 1% Pen Strep (ThermoFisher 15140122) and the cells were plated to 10% confluency in 500 μ L supplemented DMEM onto 12 mm No 1 glass coverslips (Carolina Biological 633029) coated overnight at 4 °C with 0.1 mg/mL Poly-D-Lysine and 0.02 mg/mL laminin in a 24-well plate (Fisher 08–772–1). After 24 h, each well was transfected with 125–500 ng of each plasmid (human Tau tagged with mKate2 and human Fyn tagged with Myc) and 2 μ L of Promega Fugene (Fisher) in serum-free DMEM. The following day, 5 μ L of 3 mM Tau-PxxP_{5/6} or Tau-CTD (Peptide 2.0) in 10% DMSO in serum-free DMEM was added to each well as appropriate to bring the final concentration to 15 μ L of peptide. Twenty-four hours after peptide application, cells were fixed in 1xPBS containing 4% paraformaldehyde and 4% sucrose at room temperature for 30 min, then washed 3 \times 5 minutes in 1xPBS and stored at 4 °C in the dark until PLA was performed, or lysed for Western blot analysis.

2.9. HEK-293 proximity ligation assay and image analysis

Duolink In Situ Fluorescence kit (Millipore Sigma DUO92014) was used for PLA. Fixed HEK-293 cells on coverslips were permeabilized for 10 min with 0.25% Triton X-100 (Fisher BP151–500) in 1xPBS at room temperature then blocked in 5% normal goat serum in 1xPBS for 1 h at room temperature. Coverslips were then incubated overnight at 4 °C with Tau5 (1:500) and Fyn3 (Santa Cruz, sc-16, 1:250) antibodies in 1% NGS in 1xPBS in a humidity chamber to prevent evaporation. The next day, coverslips were washed 3 \times 5 minutes in 1xPBS then incubated for 1 h at 37 °C in 8 μ L Duolink In Situ PLA Probe Anti-Rabbit PLUS (Millipore Sigma DUO92002) and 8 μ L Anti-Mouse MINUS (Millipore Sigma DUO92006) in 24 μ L 1% NGS in 1xPBS per reaction. Coverslips were washed 2 \times 5 minutes in Duolink In Situ Wash Buffer A at room temperature then incubated for 1 h at 37 °C in 8 μ L Duolink In Situ 5 \times ligation buffer and 1 μ L ligase in 31 μ L ultrapure DNase/RNase-free distilled water (ThermoFisher 10977015) per reaction. Coverslips were washed 2 \times 2 minutes in Duolink In Situ Wash Buffer A at room temperature then incubated for 100 min at 37 °C in 8 μ L Duolink In Situ 5 \times amplification buffer and 0.5 μ L polymerase in 31.5 μ L ultrapure DNase/ RNase-free distilled water per reaction. Coverslips were washed 2 \times 10 minutes in Duolink In Situ Wash Buffer B at room temperature then washed for 1 min in 1% Duolink In Situ Wash Buffer B and for 5 min in 1xPBS. To view Fyn, coverslips were incubated for 1 h at room temperature in secondary antibody (Goat anti-rabbit conjugated with Alexa-647, ThermoFisher, A-21245, 1:1000) in 1% NGS in 1xPBS, then washed 3 \times 5 minutes in 1xPBS. Coverslips were mounted with Duolink In Situ Mounting Medium with DAPI (Sigma) and stored at 4 °C while protected from light until

imaging. Fluorescent images were taken using an epifluorescent microscope at 60× with four channels: DAPI (nuclei), FITC (PLA), TRITC (Tau-mKate2), and Cy5 (Fyn). Seven images per slide were obtained then analyzed using ImageJ. Because it is challenging to disentangle cells contributing from outside of the field of view, we measured PLA density per field-of-view and took multiple images of each coverslip to accurately measure PLA density for each coverslip. For each of the channels, we subtracted the background and adjusted brightness/contrast equally across treatments. Since Tau filled the cells most evenly, we measured cell area as the area covered by Tau fluorescence by thresholding the TRITC channel. To quantify PLA puncta, we thresholded the FITC channel to view puncta and ran the ImageJ particle analyzer, specifying size and circularity to exclude any non-punctate signal. We then divided the PLA puncta by the area covered by Tau for each image to measure PLA density and normalized to the values for untreated slides from that experiment.

2.10. Primary neuron PLA

Neurons were grown on coverslips until DIV20, then treated with 15 μ M Tau-PxxP_{5/6} or vehicle control for 24 h and fixed in 4% PFA with 4% glucose in 1× PBS. The same Duolink In Situ Fluorescence kit was used as for the HEK-293 PLA with the only difference being primary and secondary antibodies used. Anti-Tau (1:1000, DAKO #A0024) and anti-Fyn (1:250, Santa Cruz SC-434) primary antibodies were used, and Goat anti-rabbit conjugated with Alexa-594, (ThermoFisher, A-11037, 1:1000) secondary antibody was used to view Tau. Imaging and quantification was completed using the same methods described for the HEK-293 PLA.

2.11. Statistics

Sample sizes were determined based on power calculations to provide 80% power to detect a difference of 20% with 30% standard deviation at an alpha of 0.05 using the effect size from preliminary studies or from literature. All statistical tests were performed using Graphpad Prism 7, and appropriate statistical tests were used based on the number and types of groups for each experiment. Amelioration of A β toxicity was defined as a significant interaction between main factors of peptide concentration and A β by two-way ANOVA, with a significant difference between vehicle and A β at 0 μ M peptide and no significant difference between vehicle and A β at 15 μ M peptide by Dunnett's post hoc.

3. Results

3.1. Tau reduction ameliorates A β toxicity

We first replicated the beneficial effects of Tau reduction in a primary hippocampal neuron model of A β toxicity. We applied a previously published Tau antisense oligonucleotide (ASO) (Devos et al., 2013), which after 1 week of exposure reduced Tau levels to about half of normal levels (Fig. 1A,B). Six days after treatment with either the Tau ASO or a nontargeting control (NTC) ASO (Devos et al., 2013), we applied 2.5 μ M synthetic A β (characterized in Supplementary Fig. 1) or their vehicle control for 24 h. To measure A β toxicity, we used a modified MTT assay that provides a measure of membrane trafficking (Izzo et al., 2014; Liu and Schubert, 1997). Briefly, neurons metabolize MTT into formazan, which can either remain in soluble intracellular vesicles or be exocytosed, forming insoluble

crystals on the cell surface. A β o increases exocytosis of formazan, thus decreasing the ratio of soluble vs insoluble formazan (see Supplementary Fig. 2). Tau reduction ameliorated this A β o-induced dysfunction, as the Tau ASO protected neurons from A β o-induced membrane trafficking abnormalities (Fig. 1C). Importantly, Tau reduction did not compromise viability of neurons, measured by either NeuN levels (Fig. 1D) or MTT assay (Fig. 1E). These results add to the body of work demonstrating that Tau reduction does not impair neuronal survival.

3.2. Tau-PxxP_{5/6} inhibits Tau-SH3 interactions

To determine whether the beneficial effects of Tau reduction would be replicated by inhibiting Tau-SH3 interactions, we developed a cell-permeable Tau-SH3 interaction inhibitor peptide (Tau-PxxP_{5/6}) based on a 17-amino acid peptide spanning the 5th and 6th PxxP motifs of the proline-rich domain of human Tau protein (amino acids 209–225). This region of Tau is the primary binding site for Fyn and may mediate other Tau-SH3 interactions. We added a 10-amino acid sequence derived from the transactivator of transcription (TAT) of human immunodeficiency virus for cell permeability.

We have previously shown that the resulting Tau-PxxP_{5/6} peptide inhibits Tau-Fyn interactions in a cell-free AlphaScreen (Cochran et al., 2014). To determine whether Tau-PxxP_{5/6} inhibits the Tau-Fyn interactions in a cell-based system, we utilized proximity ligation assay (PLA) to measure the amount of Tau-Fyn interaction in cells. If two proteins interact and the employed antigen-targeting domains are within 40 nm of each other, PLA leads to a fluorescent punctum, which can be imaged by immunocytochemistry or immunohistochemistry. We transiently transfected HEK-293 cells with plasmids both for mKate2tagged Tau and for Fyn (Fig. 2A) and verified that the PLA signal was only seen in cells transfected with both Tau and Fyn and treated with primary antibodies (Supplementary Fig. 3). Treatment of cells with Tau-PxxP_{5/6} for 24 h decreased Tau-Fyn PLA signal by almost half (Fig. 2B), measured as the number of PLA puncta per cell area. As a negative control to rule out a non-specific peptide effect, we also designed a cell-permeable TAT peptide (Tau-CTD) spanning 17 amino acids from Tau's C-terminal domain (residues 380–396), a region that does not contain PxxP motifs or mediate Tau's SH3 interactions. As expected, Tau-CTD did not change Tau-Fyn PLA density (Fig. 2B). Tau-PxxP_{5/6} did not change Tau or Fyn levels in HEK-293 cells (Fig. 2C), demonstrating that the PLA results are not due to reducing protein levels.

3.3. Tau-PxxP_{5/6} reduces Tau phosphorylation by Fyn without changing Fyn kinase activity

Fyn phosphorylates Tau at Y18, and this phosphorylation has been implicated in NMDAR-dependent glutamate toxicity (Miyamoto et al., 2017), so we determined whether Tau-PxxP_{5/6} reduced the phosphorylation of Tau Y18 by Fyn. It is likely that Tau-Fyn interactions facilitate subsequent Tau phosphorylation by Fyn, so we expected that inhibiting their interaction via the SH3 domain would decrease phosphorylation of Tau at Y18 by Fyn. To test this, we transiently transfected HEK-293 cells with Tau and Fyn, then determined the effect of 24 h exposure to Tau-PxxP_{5/6} on tau phosphorylation measured by Western blot (Fig. 3A). As expected, Tau-PxxP_{5/6} reduced Y18 phosphorylation of Tau by Fyn (Fig. 3B).

This effect likely results from reduced Tau-Fyn interaction but could be due to direct Fyn inhibition by Tau-PxxP_{5/6}. To distinguish between these possibilities, we examined Fyn's regulatory phosphorylation status. Fyn autophosphorylation at Y420 is associated with active Fyn kinase and assessing phosphorylation at this site is a commonly used measure of Fyn kinase activity (Knox and Jiang, 2015). Importantly, we found that Tau-PxxP_{5/6} did not change phosphorylation status of Fyn at Y420, providing no evidence for a change in Fyn activity (Fig. 3C). As positive controls, the Fyn kinase inhibitor PP2 decreased pY420 (Supplementary Fig. 4), and ethanol, which activates Fyn kinase activity (Wang et al., 2007), increased pY420 (Supplementary Fig. 5). Together, these results support the hypothesis that Tau-PxxP_{5/6} inhibits Tau-Fyn interaction without changing the activity of Fyn kinase.

3.4. Tau-PxxP_{5/6} inhibits endogenous Tau-Fyn interaction in neurons

Since Tau-PxxP_{5/6} inhibits the Tau-Fyn interaction when the proteins are overexpressed in HEK-293 cells, we next used PLA to determine if Tau-PxxP_{5/6} inhibits endogenous Tau-Fyn interaction in neurons, achieving the intended pharmacodynamic effect and allowing us to infer target engagement. This also provided the opportunity to visualize the amount and localization of endogenous Tau-Fyn interaction in neurons (Fig. 4A). Interestingly, the interaction predominantly occurred throughout the neurites rather than in the soma and appeared adjacent to, but not directly within, Tau-positive neuritic shafts (see inset). This localization is consistent with the possibility that the Tau-Fyn interaction may occur in dendritic spines, which would support the hypothesis that Tau and Fyn interact at the postsynapse in dendrites, allowing A β to transduce its signal through Fyn, leading to neuronal dysfunction.

Treating DIV20 neurons with Tau-PxxP_{5/6} for 24 h decreased endogenous Tau-Fyn PLA density by about two-thirds (Fig. 4B,C). Lower magnification images of neurons treated with Tau-PxxP_{5/6} demonstrate that the peptide did not affect the morphology of neurons (Supplemental Fig. 6). Importantly, Tau-PxxP_{5/6} did not change Tau or Fyn levels in primary neurons (Supplementary Fig. 7), and reducing Tau or Fyn with ASOs reduced Tau-Fyn PLA (Supplementary Fig. 8). These results show that Tau and Fyn interact endogenously in neurons, that this interaction predominately occurs in neurites, and that Tau-PxxP_{5/6} inhibits endogenous Tau-Fyn interactions.

3.5. Inhibiting Tau-SH3 interactions ameliorates A β toxicity

With validated tools in hand for inhibiting Tau-SH3 interactions in neurons, we next asked if inhibiting Tau-SH3 interactions ameliorated A β toxicity in primary neurons, mimicking the beneficial effects of Tau reduction. We assessed A β toxicity using two orthogonal measures: membrane trafficking abnormalities and neurite degeneration.

First, we tested whether inhibiting Tau-SH3 interactions prevented A β -induced membrane trafficking abnormalities. We pretreated DIV20 neurons with Tau-PxxP_{5/6} or its vehicle control for 90 min before applying 2.5 μ M A β or their vehicle control for 24 h and utilizing the same modified MTT assay described earlier (Fig. 5A). Tau-PxxP_{5/6} ameliorated A β -induced membrane trafficking abnormalities in a dose-dependent manner, while the negative control peptide, Tau-CTD, did not (Fig. 5B).

Second, we tested another A β -induced abnormality that is ameliorated by Tau reduction (Rapoport et al., 2002), A β -induced neurite degeneration. A β causes neurite degeneration and blebbing, which can be quantified using immunofluorescent labeling of MAP2 followed by imaging in an unbiased manner using a high-content, automated approach (see Supplementary Fig. 9). As before, we pretreated DIV20 neurons with Tau-PxxP_{5/6} or its vehicle control for 90 min before applying 2.5 μ M A β or their vehicle control (Fig. 6A). Forty-eight hours after A β application, we fixed the neurons, labeled MAP2, then quantified intact neurite length. Tau-PxxP_{5/6} ameliorated neurite blebbing and degeneration while Tau-CTD did not (Fig. 6B,C). Together, these findings provide evidence that Tau-SH3 interactions contribute to A β toxicity in neurons.

4. Discussion

This study demonstrates that an inhibitor of Tau-SH3 interactions ameliorates A β toxicity in primary rat neurons, supporting the idea that inhibiting Tau-SH3 interactions is a potential mechanism by which Tau reduction is protective. We developed several novel tools for studying Tau-SH3 interactions. One is a proximity ligation assay for the most well studied Tau-SH3 interaction, Tau-Fyn. In neurons, endogenous Tau-Fyn interactions were observed primarily along neurites adjacent to Tau-positive neuritic shafts. The second new tool is a cell-permeable peptide inhibitor, Tau-PxxP_{5/6}, that competitively inhibits Tau-SH3 interactions in neurons. Blocking Tau-SH3 interactions using this inhibitor reduced Tau-Fyn interactions measured by PLA and ameliorated multiple measures of A β toxicity.

The Tau-SH3 interaction inhibitor also reduced Tau phosphorylation by Fyn, without directly reducing Fyn activity. This finding provides further support for the ability of Tau-PxxP_{5/6} to reduce Tau-Fyn interactions, although it is not clear that Fyn phosphorylation of Tau is the mechanism by which Tau-PxxP_{5/6} reduces A β toxicity. Our data indicates that Tau-PxxP_{5/6} decreases Fyn phosphorylation of Tau and prevents A β toxicity, but we cannot conclude that the former causes the latter. It remains possible that other downstream effects of inhibiting the Tau-Fyn interaction contribute to the prevention of A β toxicity by Tau-PxxP_{5/6}.

One question that will need to be addressed in ongoing studies is which particular Tau-SH3 interactions are most critical. The field has focused on Fyn as a prototypical SH3-containing binding partner for Tau (Lee, 2005) and considerable evidence suggests a role for Tau-Fyn interactions in AD. First, Tau directly binds Fyn's SH3 domain (Cochran et al., 2014; Lau et al., 2016; Lee et al., 1998; Usardi et al., 2011) and both Tau phosphorylation and Tau mutations increase Tau-Fyn interaction (Bhaskar et al., 2005; Cochran et al., 2014). Second, Tau and Fyn have converging phenotypes both in vitro and in vivo. In primary neurons, genetic knockout of either Tau (Rapoport et al., 2002) or Fyn (Lambert et al., 1998) ameliorates A β toxicity. In vivo, reduction of either Tau (Ittner et al., 2010; Leroy et al., 2012; Meilandt et al., 2008; Nussbaum et al., 2012; Palop et al., 2007b; Roberson et al., 2007; Roberson et al., 2011) or Fyn (Chin et al., 2004) prevents A β -induced dysfunction, and Tau reduction decreases susceptibility to network hyperexcitability (Devos et al., 2013; Roberson et al., 2011). Third, Fyn overexpression exacerbates A β -induced neuronal dysfunction (Chin et al., 2005) and Tau reduction prevents these Fyn-dependent

effects (Roberson et al., 2011). Finally, Tau regulates trafficking of Fyn to the postsynaptic density and preventing this Tau-mediated trafficking of Fyn ameliorates A β -induced cognitive deficits, NMDAR activation, and premature mortality in mice (Ittner et al., 2010). Altogether, these findings combined with the results observed here indicate that the Tau-Fyn interaction may have an important role in AD pathogenesis.

Other Tau-SH3 interactions besides Tau-Fyn might also play a role. One recent report suggested that the Tau-Fyn interaction may not be critical for glutamate-induced excitotoxicity (Miyamoto et al., 2017). This would not be inconsistent with the observations here, particularly because the mechanisms of A β - and glutamate-induced toxicity are different. In addition, that study (Miyamoto et al., 2017) used an overexpression system and manipulated the 7th PxxP binding site, which is likely not the primary Fyn binding site in Tau (Cochran et al., 2014; Lau et al., 2016; Usardi et al., 2011). One group recently completed computational modeling of known Tau-SH3 interactions to determine which of Tau's PxxP motifs has the highest affinity for each SH3 domain (Wang et al., 2019). Interestingly, they found that of all 180 PxxP-SH3 combinations they modeled, Tau's 5th PxxP motif and Fyn's SH3 domain had the highest binding affinity, which highlights the relevance of targeting Tau's 5th/6th PxxP motifs, as we did with Tau-PxxP_{5/6}. It is also important to emphasize that other Tau-SH3 interactions besides Tau-Fyn could mediate the effects we observed. Known Tau-SH3 interactions with clear potential relevance to A β -induced abnormalities include Tau's interaction with the AD genetic risk factor BIN1 (Lambert et al., 2013) and the post-synaptic density protein PSD95 (Mondragón-Rodríguez et al., 2012). BIN1 regulates propagation of Tau pathology (Calafate et al., 2016) and its *Drosophila* homolog modulates Tau toxicity (Chapuis et al., 2013). Tau's proline-rich domain binds directly to BIN1's SH3 domain and phosphorylation of Tau regulates this binding (Sottejeau et al., 2015), highlighting a potential role for the Tau-BIN1 interaction in Tau-mediated AD deficits. A recent report used Tau-BIN1 PLA to identify modulators of Tau-BIN1 interaction (Sartori et al., 2019) and found that phosphorylation of BIN1 modulates the interaction and that there is altered BIN1 phosphorylation in AD brain. Of note from the in silico study of SH3-Tau-PxxP interactions (Wang et al., 2019) was that the strongest binding for BIN1's SH3 domain in Tau's proline-rich domain was the 5th PxxP motif. Additionally, PSD95 interacts via its SH3 domain with Tau at the synapse, forming a complex with Fyn and NMDARs (Mondragón-Rodríguez et al., 2012), and manipulations that disrupt this Tau/Fyn/PSD95 complex prevent network hyperexcitability and A β toxicity in vivo (Ittner et al., 2016). This complex allows Fyn to phosphorylate NMDARs which could lead to increased Ca²⁺ influx and excitotoxicity (Liu et al., 2019). Thus, multiple different Tau-SH3 interactions could contribute to A β -induced neuronal dysfunction.

Another remaining question is whether A β promotes increased Tau-SH3 interactions as a mechanism leading to neuronal deficits, or whether Tau-SH3 interactions are permissive of A β toxicity without A β -induced increases. Either possibility is consistent with the findings here that blocking Tau-SH3 interactions reduces A β -induced neuronal dysfunction. On one hand, there is some evidence supporting the idea that A β could promote Tau-SH3 interactions. A β causes Fyn-mediated local translation of Tau in dendrites (Li and Gotz, 2017) and causes Tau translocation into dendritic spines

(Frandemiche et al., 2014), either of which could increase the amount of Tau-SH3 interactions if there are unbound SH3-containing proteins available to bind Tau. However, it is unlikely that there are enough unbound SH3-containing proteins in the dendrites to have A β -induced local Tau translation or translocation significantly alter the amount of Tau-SH3 interactions, and it is unclear how this would induce A β toxicity.

On the other hand, and perhaps more likely, Tau-SH3 interactions could be permissive of A β toxicity, even if A β does not lead directly to more or stronger Tau-SH3 interactions. Considering the prototypical SH3-containing protein Fyn, extracellular A β activates Fyn when it is anchored to the post-synaptic density so that Fyn phosphorylates its targets. One study suggested that Tau traffics Fyn to dendrites and anchors it to the post-synaptic density (Ittner et al., 2010). Based on this model, inhibiting Tau-SH3 interactions, specifically the Tau-Fyn interaction, could prevent activation of Fyn by A β in the postsynapse, preventing signal transduction leading to A β toxicity. Going forward, it is important to determine if the Tau-Fyn interaction is indeed the critical Tau-SH3 interaction required for A β toxicity and which downstream pathways are activated to cause neuronal dysfunction. Possibilities of key downstream pathways include phosphorylation of Tau or the NMDAR, or activation of the AD risk gene Pyk2, another phosphorylation target of Fyn (Qian et al., 1997) that is implicated in A β -induced dysfunction in vivo (Salazar et al., 2019).

Our results support the idea of targeting Tau-SH3 interactions as a novel Tau-directed therapeutic strategy for AD. These data provide preclinical evidence supporting the potential efficacy of inhibiting Tau-SH3 interactions; however, considering the potential of a novel strategy also requires evaluating safety and tractability. While further studies are clearly needed, there is evidence to suggest that inhibiting Tau-SH3 interactions would be safe. Even complete removal of Tau, which de facto would prevent all Tau-SH3 interactions, does not cause dramatic abnormalities in animals (Dawson et al., 2001; Fujio et al., 2007; Harada et al., 1994; Tucker et al., 2001; van Hummel et al., 2016), and even the rather subtle changes in Tau knockout mice are not seen in Tau heterozygous mice with partial Tau reduction (Ikegami et al., 2000; Lei et al., 2014; Li et al., 2014; Morris et al., 2013), which better represent the incomplete block that would be provided by Tau-SH3 interaction inhibitors. In fact, Tau reduction with antisense oligonucleotides did not have apparent adverse effects in mouse models (Devos et al., 2013; DeVos et al., 2017) and human clinical trials using this approach have been underway since 2017, as yet without reported adverse effects. In terms of tractability, inhibiting protein-protein interactions was once thought to be an unrealistic strategy for small molecule therapies, but advances in drug discovery and design have mitigated these concerns (Arkin et al., 2014). We have already conducted high-throughput screening for Tau-SH3 interaction inhibitors, using a Tau-Fyn interaction assay, and identified multiple series of chemically tractable hits (Cochran et al., 2014). Much is left to learn, but the data presented here provide initial preclinical support for the concept of targeting Tau-SH3 interactions as a therapeutic strategy for AD.

5. Conclusions

Tau reduction ameliorates A β -induced dysfunction, but there is a need to identify therapeutic strategies that reproduce this effect. Here, we tested the hypothesis that

Tau-SH3 interactions contribute to A β toxicity and can be targeted therapeutically. We developed a cell-based target engagement assay and a peptide inhibitor that reduces Tau-SH3 interactions. We used PLA to measure endogenous Tau-Fyn interactions throughout the neurites of primary neurons and determined that the peptide inhibitor reduces endogenous Tau-Fyn interactions in neurons. We then used this peptide inhibitor to determine that inhibiting Tau-SH3 interactions ameliorates A β toxicity, highlighting the therapeutic potential of inhibiting Tau-SH3 interactions to treat AD.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgements

We thank Yuliya Voskobiynik for valuable discussions and input throughout the project and for help with data visualization. We also thank the Herskowitz and Arrant labs for valuable discussion and Kelly Chen for assistance with protocol development for immunostaining. Southern Research and the UAB Center for Biophysical Sciences and Engineering graciously provided access to their imaging equipment.

Funding

This work was supported by NIH grants R01NS075487, RF1AG059405, UL1TR001417, and T32NS061455, BrightFocus Foundation grant A2015693S, the Alabama Drug Discovery Alliance, the UAB Center for Clinical and Translational Sciences, and a Weston Brain Institute advisor fellowship.

Abbreviations

AD	Alzheimer's disease
Aβ	amyloid- β
Aβo	A β oligomer
SH3	Src-homology 3 domain
FTD	Frontotemporal dementia
NMDAR	NMDA receptor
ASO	antisense oligonucleotide
NTC	nontargeting control ASO
MTT	3-(4,5-dimethylthiazol-2-yl)-2,5-diphenyltetrazolium bromide
TAT	transactivator of transcription
PLA	proximity ligation assay
DIV	days in vitro

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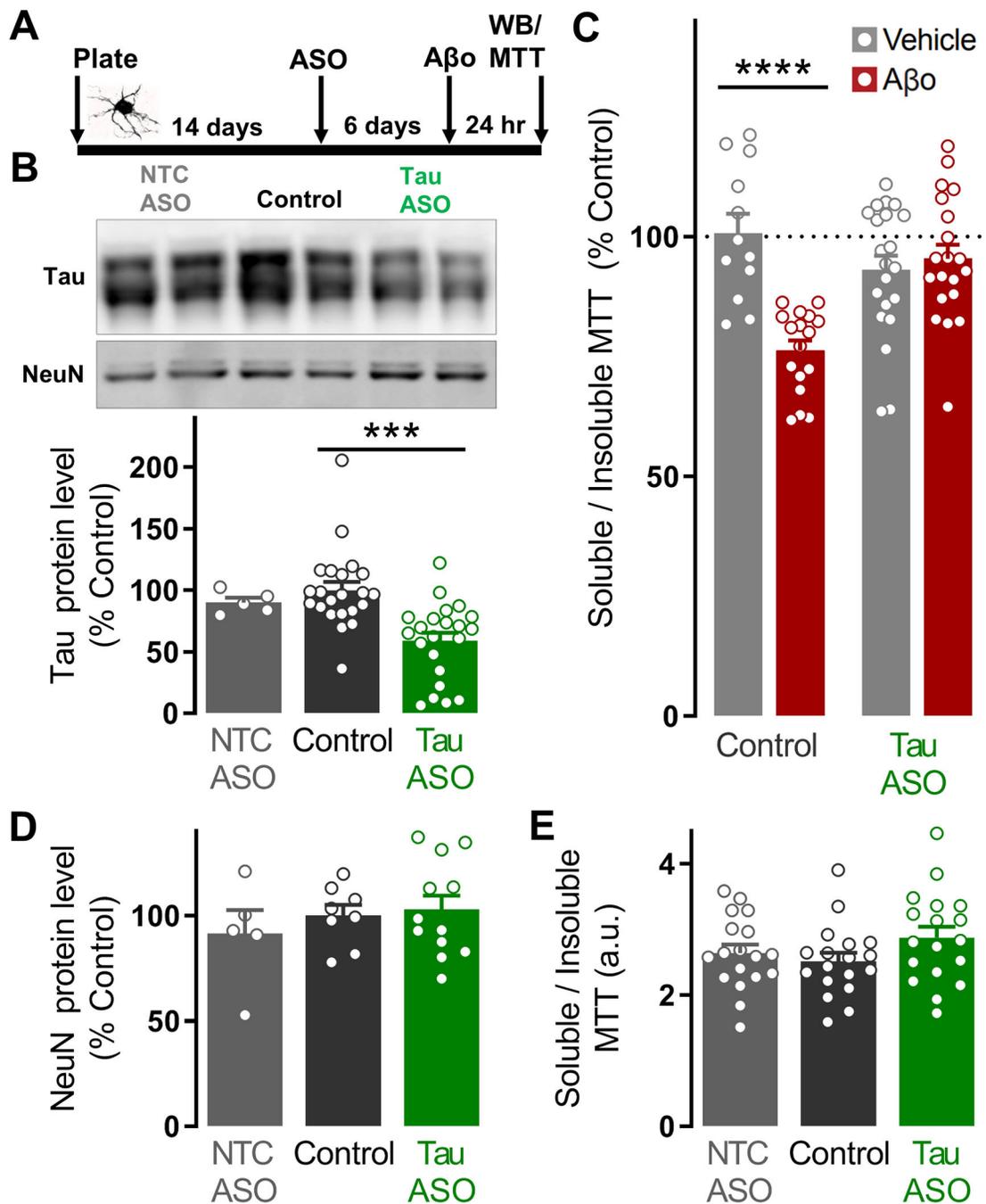


Fig. 1. Tau reduction ameliorates amyloid- β -induced membrane trafficking abnormalities. **A:** Timeline of ASO treatment and modified MTT experiments. **B:** Representative Western blot images of lysate from hippocampal neurons from rats treated for one week (DIV14–21) with a previously published antisense oligonucleotide for Tau or nontargeting control ASO (Devos et al., 2013). The Tau ASO caused a 41% reduction in Tau while a nontargeting control (NTC) ASO did not reduce Tau (ANOVA, $F(2, 64) = 10.23$, $p = .0002$, $n = 5-22$ wells per group). **C:** At DIV20, 2.5 μ M A β o or vehicle equivalent was applied. After 24

h, A β -induced membrane trafficking abnormalities were measured using a modified MTT assay (full description in the methods section and Supplementary Fig. 2). Tau reduction with the ASO ameliorated A β -induced membrane trafficking abnormalities (Two-way ANOVA interaction for factors A β and ASO $F(1,66) = 18.9$, $p < .0001$, $n = 12-21$ wells per group from $N = 2$ rats, 4-12 wells per group per plate with 1 plate per rat). Differences indicated: **** $p < .0001$ by Dunnett's post hoc test with correction for multiple comparisons relative to vehicle control. D: Neither ASO affected neuronal survival, as measured with NeuN protein levels (ANOVA, $F(2, 22) = 0.52$, $p = .60$, $n = 5-12$ wells per group). E: Neither ASO affected neuronal survival, as measured with the MTT assay (ANOVA, $F(2, 51) = 1.66$, $p = .42$, $n = 18$ wells per group). All Panels: Bars indicate mean \pm SEM. Differences indicated: *** $p < .001$ by Dunnett's post hoc test with correction for multiple comparisons relative to CTL.

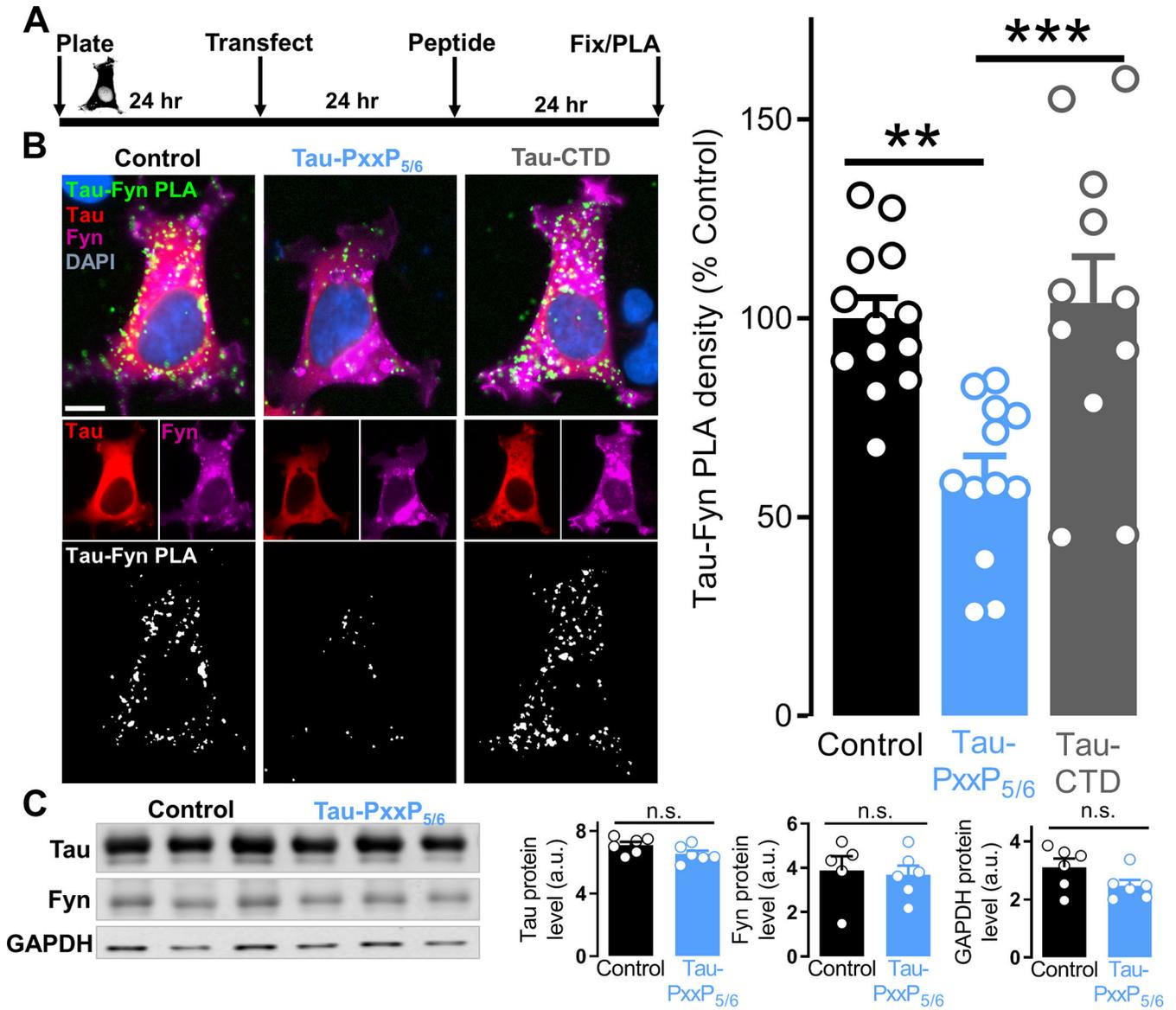


Fig. 2. Tau-PxxP_{5/6} inhibits the Tau-Fyn interaction in cells. **A:** Timeline of Tau-Fyn PLA experiments in HEK-293 cells. **B:** Representative immunofluorescent images of HEK-293 cells treated with Tau-PxxP_{5/6} (rrrqrkkrgRSRTPSLPTPPT-REPKK) or Tau-CTD (rrrqrkkrgENAKAKTDHGAEIVYKS) after Tau-Fyn proximity ligation assay (PLA) was performed (full description in the methods section). Green puncta represent sites of Tau-Fyn interaction, red is Tau-mKate2, and magenta is immunofluorescently labeled Fyn. Scale bar = 10 μm. Quantification of Tau-Fyn PLA density demonstrates that Tau-SH3 inhibitor Tau-PxxP_{5/6} reduced Tau-Fyn interaction by 40% while negative control peptide Tau-CTD did not (ANOVA, $F(2, 33) = 10.14$, $p = .0004$, average of 5–7 images per coverslip from $n = 11–13$ coverslips from $N = 6$ distinct passages, see Methods for details of quantification and analysis). Differences indicated: ** $p < .01$, and *** $p < .001$ by Dunnett's post hoc test with correction for multiple comparisons relative to control. **C:** Western blot of lysate from

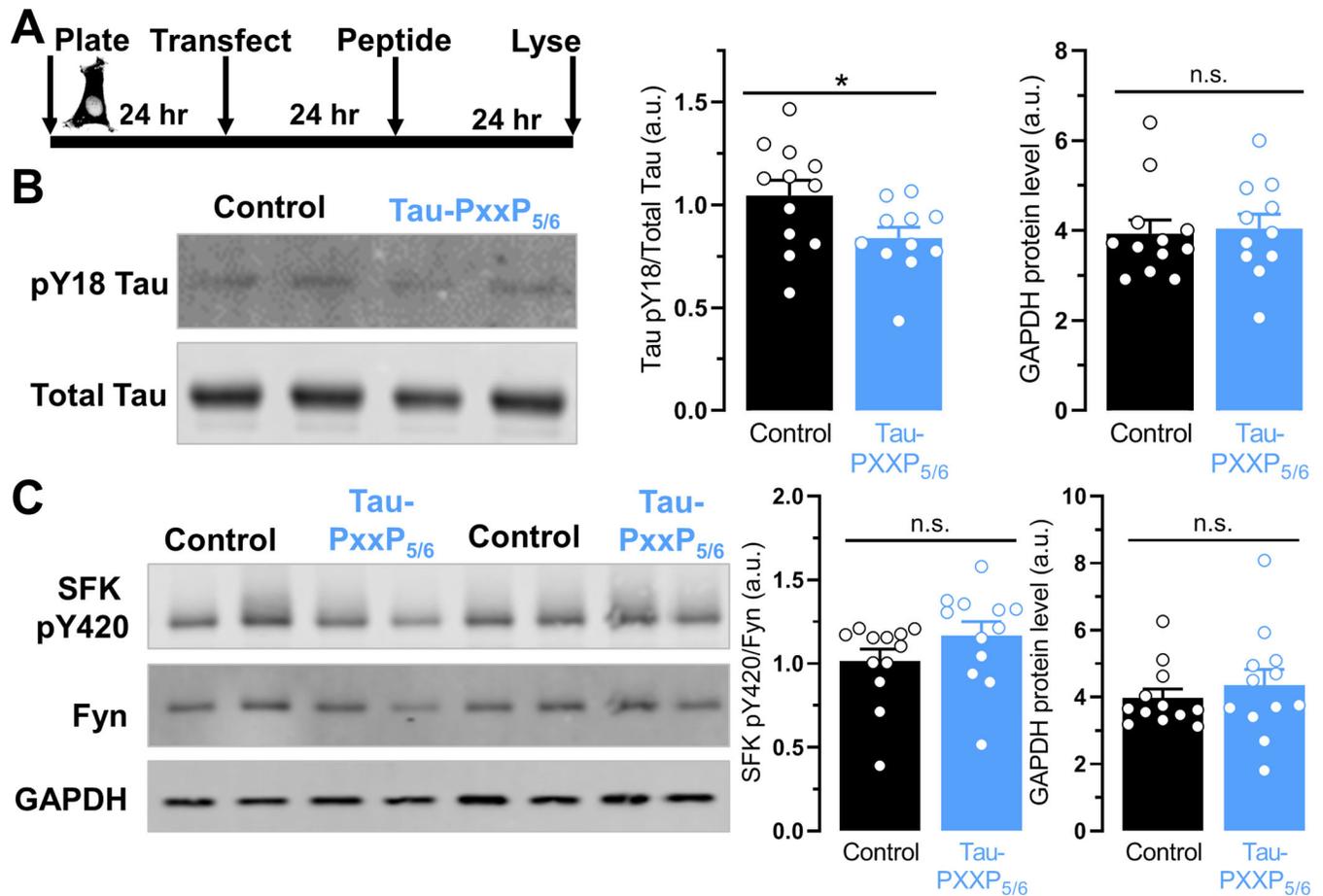
HEK-293 cells demonstrates that Tau-PxxP_{5/6} did not change Tau, Fyn, or GAPDH levels (Student's *t*-test, $t(10)=1.86, p=.092$; $t(9)=0.28, p=.79$; $t(10)=1.78, p=.10$, respectively, $n=5-6$ wells per group).

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**Fig. 3.**

Tau-PxxP_{5/6} reduces phosphorylation of Tau by Fyn without changing Fyn kinase activity. **A:** Timeline of Tau and Fyn phosphorylation experiments in HEK-293 cells. **B:** Representative Western blot images of lysate from HEK-293 cells treated with Tau-PxxP_{5/6} and probed for Tau PY18 and total Tau. Tau-PxxP_{5/6} reduced phosphorylation of Tau pY18 by Fyn by 19.75% (Student's *t*-test, $t(21) = 2.23$, $p < .037$, average of 2 technical replicates of $n = 11$ – 12 wells per group) without affecting GAPDH protein level (Student's *t*-test, $t(21) = 0.25$, $p = .80$, average of 2 technical replicates of $n = 11$ – 12 wells per group). **C:** Representative Western blot images of lysate from HEK-293 cells treated with Tau-PxxP_{5/6} and probed for Src family kinase (SFK) pY420, Fyn, and GAPDH. Tau-PxxP_{5/6} did not affect Fyn pY420 (Student's *t*-test, $t(22) = 1.40$, $p = .17$, $n = 12$ wells per group) or GAPDH (Student's *t*-test, $t(22) = 0.71$, $p = .48$, $n = 12$ wells per group).

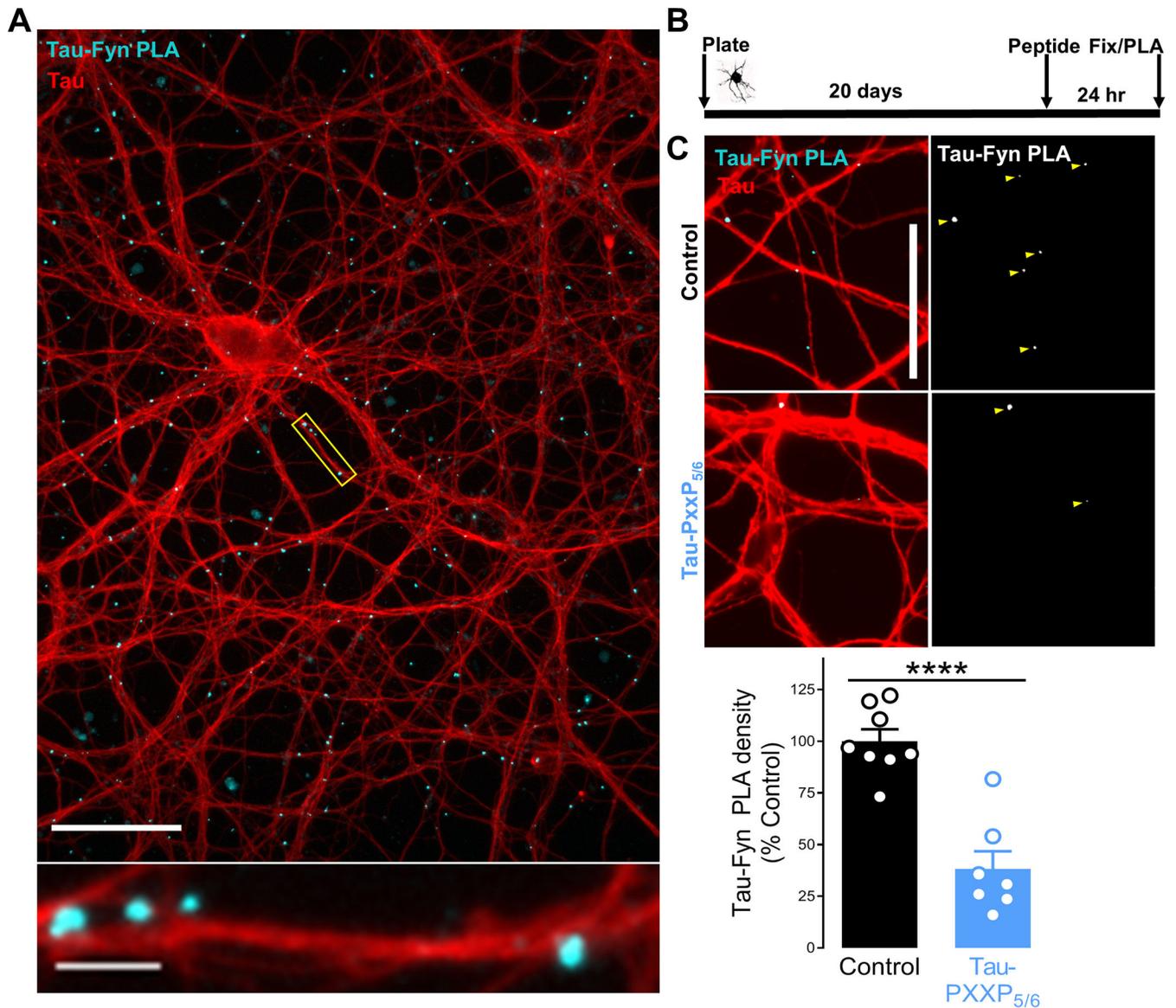


Fig. 4. Tau-PxxP_{5/6} inhibits endogenous Tau-Fyn interaction in neurons. **A:** Representative immunofluorescent image of DIV21 primary neurons after Tau-Fyn PLA. Cyan puncta represent sites of Tau-Fyn interaction and red is immunofluorescently labeled Tau. Scale bar = 50 μ m. *Inset:* magnification of a neurite showing Tau-Fyn PLA (cyan) adjacent to Tau-positive neurites (red). Scale bar = 5 μ m. **B:** Timeline of endogenous Tau-Fyn PLA inhibition experiments. **C:** Representative images of endogenous Tau-Fyn PLA after Tau-PxxP_{5/6} treatment demonstrating reduction of endogenous Tau-Fyn PLA in neurons by Tau-PxxP_{5/6}. Tau-PxxP_{5/6} reduced endogenous Tau-Fyn interaction in neurons by 62% (Student's *t*-test, $t(13) = 6.124$, $p < .0001$, average of 5–7 images per coverslip from $n = 7$ –8 coverslips). Scale bar = 25 μ m.

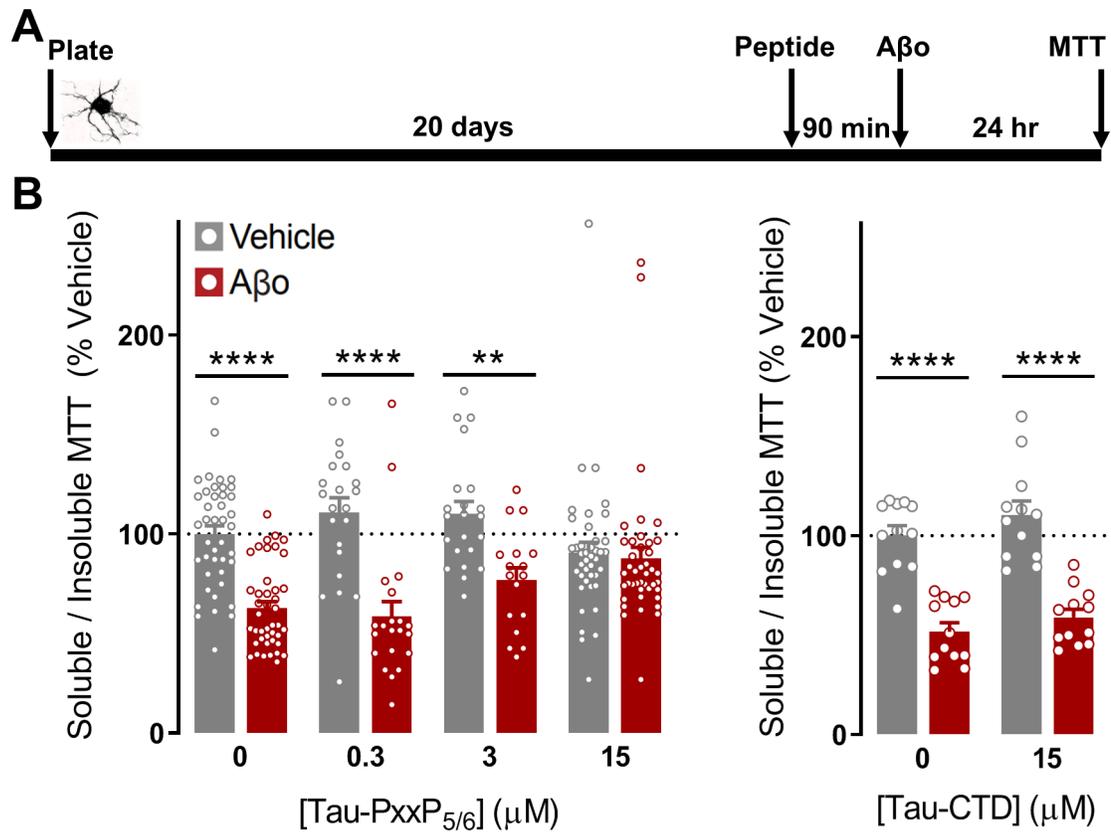


Fig. 5.

Inhibiting Tau-SH3 interactions ameliorates amyloid- β -induced membrane trafficking abnormalities. **A:** Timeline of Tau-PxxP_{5/6} modified MTT assay. Neurons were pretreated with increasing doses of Tau-PxxP_{5/6} for 90 min before 2.5 μ M A β o application. After 24 h, the modified MTT assay detailed earlier was performed. **B:** Tau-PxxP_{5/6} prevented A β o-induced membrane trafficking abnormalities in a dose-dependent fashion (Two-way ANOVA interaction for factors A β o and Tau-PxxP_{5/6} $F(3,241) = 7.74$, $p < .0001$, $n = 17-42$ wells per group from $N = 3$ embryo harvests, 3-6 wells per group per plate with 2 plates per harvest for 2 harvests and 9-12 wells per group per plate with 2 plates per harvest for 1 harvest) while the negative control peptide Tau-CTD failed to ameliorate A β o-induced membrane trafficking abnormalities (Two-way ANOVA interaction for factors A β o and Tau-CTD $F(1, 44) = 0.092$, $p = .76$, $n = 12$ wells per group from $N = 2$ embryo harvests, 6 wells per group per plate with 1 plate per harvest). Differences indicated: ** $p < .01$, and **** $p < .0001$, by Dunnett's post hoc test with correction for multiple comparisons relative to vehicle control at that dose of peptide. Bars indicate mean \pm SEM.

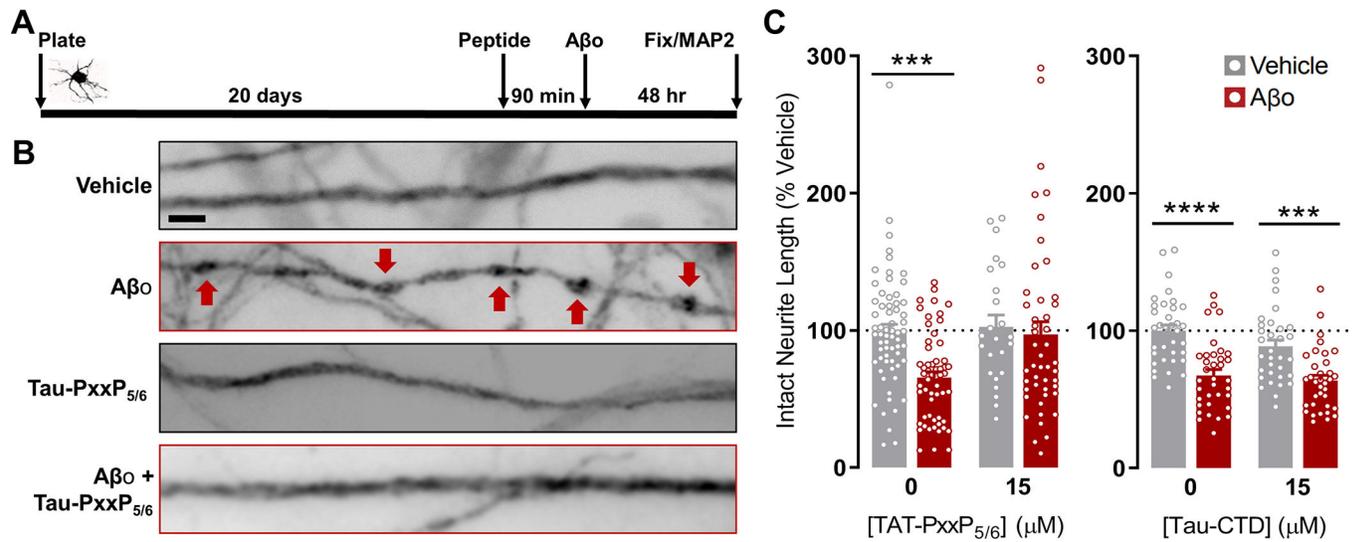


Fig. 6. Inhibiting Tau-SH3 interactions ameliorates amyloid- β -induced neurite degeneration. **A:** Timeline of Tau-PxxP_{5/6} MAP2. **B:** Representative images of neurons show signs of neurite blebbing and degeneration by immunofluorescent labeling of MAP2 (grayscale, blebbing indicated by arrows) after 48 h exposure to A β o. Neurons pretreated with 15 μ M Tau-PxxP_{5/6} showed no blebbing or signs of neurite degeneration. Scale bar = 5 μ m. **C:** Unbiased quantification of intact neurite length from MAP2 staining (see Supplementary Fig. 4) demonstrate that Tau-PxxP_{5/6} ameliorated A β o-induced neurite degeneration (Two-way ANOVA interaction for factors A β o and Tau-PxxP_{5/6} $F(1,115) = 3.98$, $p = .048$, $n = 24-64$ wells per group from $N = 5$ neuron harvests, 3-6 wells per group per plate with 2 plates per harvest for 4 harvests and 6-12 wells per group per plate with 2 plates per harvest for 1 harvest) while Tau-CTD did not (Two-way ANOVA interaction for factors A β o and Tau-CTD $F(1,135) = 0.857$, $p = .35$, $n = 33-36$ wells per group from $N = 3$ embryo harvests, 11-12 wells per group per plate with 1 plate per harvest). Differences indicated: *** $p < .001$, and **** $p < .0001$, by Dunnett's post hoc test with correction for multiple comparisons relative to vehicle control at that dose of peptide. Bars indicate mean \pm SEM.