

Blocking effects of human tau on squid giant synapse transmission and its prevention by T-817 MA

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Filamentous tau inclusions are hallmarks of Alzheimer's disease and related neurodegenerative tauopathies, but the molecular mechanisms involved in tau-mediated changes in neuronal function and their possible effects on synaptic transmission are unknown. We have evaluated the effects of human tau protein injected directly into the presynaptic terminal axon of the squid giant synapse, which affords functional, structural, and biochemical analysis of its action on the synaptic release process. Indeed, we have found that at physiological concentration recombinant human tau (h-tau42) becomes phosphorylated, produces a rapid synaptic transmission block, and induces the formation of clusters of aggregated synaptic vesicles in the vicinity of the active zone. Presynaptic voltage clamp recordings demonstrate that h-tau42 does not modify the presynaptic calcium current amplitude or kinetics. Analysis of synaptic noise at the post-synaptic axon following presynaptic h-tau42 microinjection revealed an initial phase of increase spontaneous transmitter release followed by a marked reduction in noise. Finally, systemic administration of T-817MA, a proposed neuro-protective agent, rescued tau-induced synaptic abnormalities. Our results show novel mechanisms of h-tau42 mediated synaptic transmission failure and identify a potential therapeutic agent to treat tau-related neurotoxicity.

Keywords: tauopathies, synaptic transmission, hyperphosphorylation, Alzheimer's disease, neurodegenerative, human tau, synaptic vesicles, neuro-protective agent

INTRODUCTION

Classical neuropathological studies characterized the intracellular accumulation and aggregation of abnormal filaments composed primarily of the microtubule associated protein tau as a hallmark for a variety of neurodegenerative disorders known as tauopathies, as exemplified by progressive supranuclear palsy, Pick's disease, corticobasal degeneration, and frontotemporal dementia with Parkinsonism linked to chromosome 17 (FTDP-17; Lee et al., 2001). The most common tauopathy, Alzheimer's disease (AD), is also characterized by additional filamentous structures paired helical filaments (PHFs) and straight filaments (SFs). These filaments eventually form large aggregations, known as neurofibrillary tangles (NFTs). In addition, diffuse and mature senile plaques, which are predominantly composed of amyloid beta (A β) peptides, are present in AD brains too (Selkoe, 2001).

A number of studies have investigated the role of different forms of A β peptides and aggregates in synaptic function (Arancio et al., 2004; Moreno et al., 2009), but so far the role of tau or tau aggregates in neurotransmission has not been elucidated.

The association between tau filaments, neuron loss, and brain dysfunction in vertebrates and invertebrates originally led to the hypothesis that NFTs invariably cause brain dysfunction and neurodegeneration. However, mouse tauopathy studies indicate that

severe abnormalities in synaptic function can precede neuronal loss and even NFTs formation (LaFerla et al., 2007; Yoshiyama et al., 2007). While the molecular mechanisms responsible for this early malfunction (as well as those responsible for tau polymerization dependent pathogenesis) remain unknown (Marx, 2007). Neurofibrillary degeneration is accompanied by lysosomal hypertrophy (Nixon et al., 1992), beading and degeneration of distal dendrites (Braak et al., 1994; Marx, 2007) and axonal damage (Kowall and Kosik, 1987).

In the present study physiological concentrations of recombinant human tau isoform (full length h-tau42; Perez et al., 2001) was directly injected into the presynaptic terminal of the squid giant synapse, to examine possible acute effects of h-tau on the synaptic release mechanism. The results showed that heavy exogenous h-tau42 accumulation induces a rapid and short-lasting increase in spontaneous transmitter release followed by a drastic decrease and failure of synaptic transmission. This synaptic block does not affect presynaptic calcium current flow or spike generation at the presynaptic terminal. Immunohistochemistry, performed in h-tau42 injected synapses, demonstrated that h-tau42 becomes phosphorylated rapidly in good temporal agreement with the time course of the transmitter failure. Electron microscopy and electrophysiological experiments unambiguously indicate that h-tau42-mediated synaptic transmission block is due to exocytosis failure.

Finally, systemic administration of the novel compound T-817MA prevented the structural, biochemical, and functional deleterious effects h-tau42 microinjection.

The present set of studies identifies several mechanisms of tau-mediated toxicity at the presynaptic terminal, and introduces a potential disease modifier for AD and other tauopathies for which there is no specific treatment presently.

MATERIALS AND METHODS

TAU PROTEINS

Recombinant human tau, h-tau42 (isoform with four tubulin binding motifs and two extra exons in the N-terminal domain) was isolated as previously described (Perez et al., 2001).

IMMUNOHISTOCHEMISTRY

A variation of the array tomography method by Micheva and Smith (2007) was followed. The ganglia were fixed by immersion in 4% paraformaldehyde (EM grade EM Sciences) plus 7.0% sucrose in calcium-free sea water for 3 h; rinsed with 7% sucrose and 50 mM glycine in 0.1 M PBS; dehydrated with graded ethanol dilutions (50%, 70%, 90%, and 3× 100%), embedded in LR White resin (medium grade, SPI), and polymerized in gelatin capsules at 49°C for 48 h. Semithin sections (500 nm) were mounted on subbed slides and encircled on the slides with a PAP pen (EM Sciences, USA). The immunocytochemistry was done as follows: (a) blocked in 50 mM glycine in tris buffer pH 7.6, 5 min; (b) primary antibody incubation, anti-tau PHF (AT8, Thermochemical, USA) diluted 1:50 in 1% BSA in tris (tris-BSA), 4 h; (c) rinses in tris-BSA 2× 5 min; (d) secondary antibody incubation, goat anti-mouse Alexa Fluor 594 diluted 1:150 in tris-BSA, 30 min; (e) tris and distilled water rinses, 4× 5 min each; (f) mounting of slides with coverslips and anti-fading mounting media; (g) image under fluorescent microscopy (Zeiss Axioimager, Germany). Controls were performed with the same protocol omitting the primary antibody. Primary antibody: Anti-tau PHF (AT8; Thermochemical, USA), secondary antibody: Alexa Fluor 594 goat anti-mouse (Invitrogen).

T-817MA ADMINISTRATION

Squid received an oral dose of T-817MA (0.25 ml of 3.68 mmol l⁻¹ T-817MA) every 12 h using a methodology recently developed by our group (Berk et al., 2009). At 24 h after the first dose a second dose was given, and the electrophysiological experiments were performed 1 h after the second feeding. The optimal dose, absorption, and concentration reached in the optic lobes of T-817MA in squid had been obtained previously (Berk et al., 2009). Indeed, it was determined that after 24 h of the double administration, a T817-MA concentration of 26.7 μmol l⁻¹ was measured in the CNS of the gastrically intubated squid ($n = 10$ squid). Synapses from T-817MA-treated squid were microinjected at the presynaptic axon with h-tau42 (an approximate 80 nM final concentration h-tau42 ($n = 19$), considering a 100× dilution factor. Pre- and post-synaptic potentials were recorded for 90 min.

ELECTROPHYSIOLOGY AND MICROINJECTIONS

The squid (*Loligo paelli*) stellate ganglia isolation from the mantle and the electrophysiological techniques used have been described previously (Llinas et al., 1985). Two glass micropipette electrodes

impaled the largest (most distal) presynaptic terminal digit at the synaptic junction site while the post-synaptic axon was impaled by one microelectrode at the junctional site. One of the pre-electrodes was used for pressure microinjection of h-Tau 42 and also supported voltage clamp current feedback, while the second monitored membrane potential. The total volume injected fluctuated between 0.1 and 1 pl. (Llinas et al., 1985). The exact location of injection and the diffusion and steady-state distribution of the protein/fluorescent dye mix (0.001% dextran fluorescein) were monitored using a fluorescence microscope attached to a Hamamatsu camera system (Middlesex, NJ). In all experiments a good correlation was observed between the localization of the fluorescence and the electrophysiological findings.

ELECTRON MICROSCOPY

Immediately following the electrophysiological study the ganglia were removed from the recording chamber, fixed by immersion in glutaraldehyde, post-fixed in osmium tetroxide, stained in block with uranium acetate, dehydrated and embedded in resin (Embed 812, EM Sciences). Ultrathin sections were collected on Pioloform (Ted Pella, Redding, CA) and carbon-coated single slot grids, and contrasted with uranyl acetate and lead citrate. Morphometry and quantitative analysis of the synaptic vesicles were performed with the in house program developed with LabVIEW (National Instruments, Austin, TX, USA). Electron micrographs were taken at an initial magnification of ×16,000 and ×31,500 and photographically enlarged to a magnification of ×40,000 and ×79,000 for synaptic vesicles and clathrin-coated vesicle (CCV) counting, respectively. Vesicle density at the synaptic active zones was determined as the number of vesicles per μm², on an average area of 0.8 μm² per active zone. CCV density was determined within the limits of the presynaptic terminal on an average terminal area of 3.3 μm².

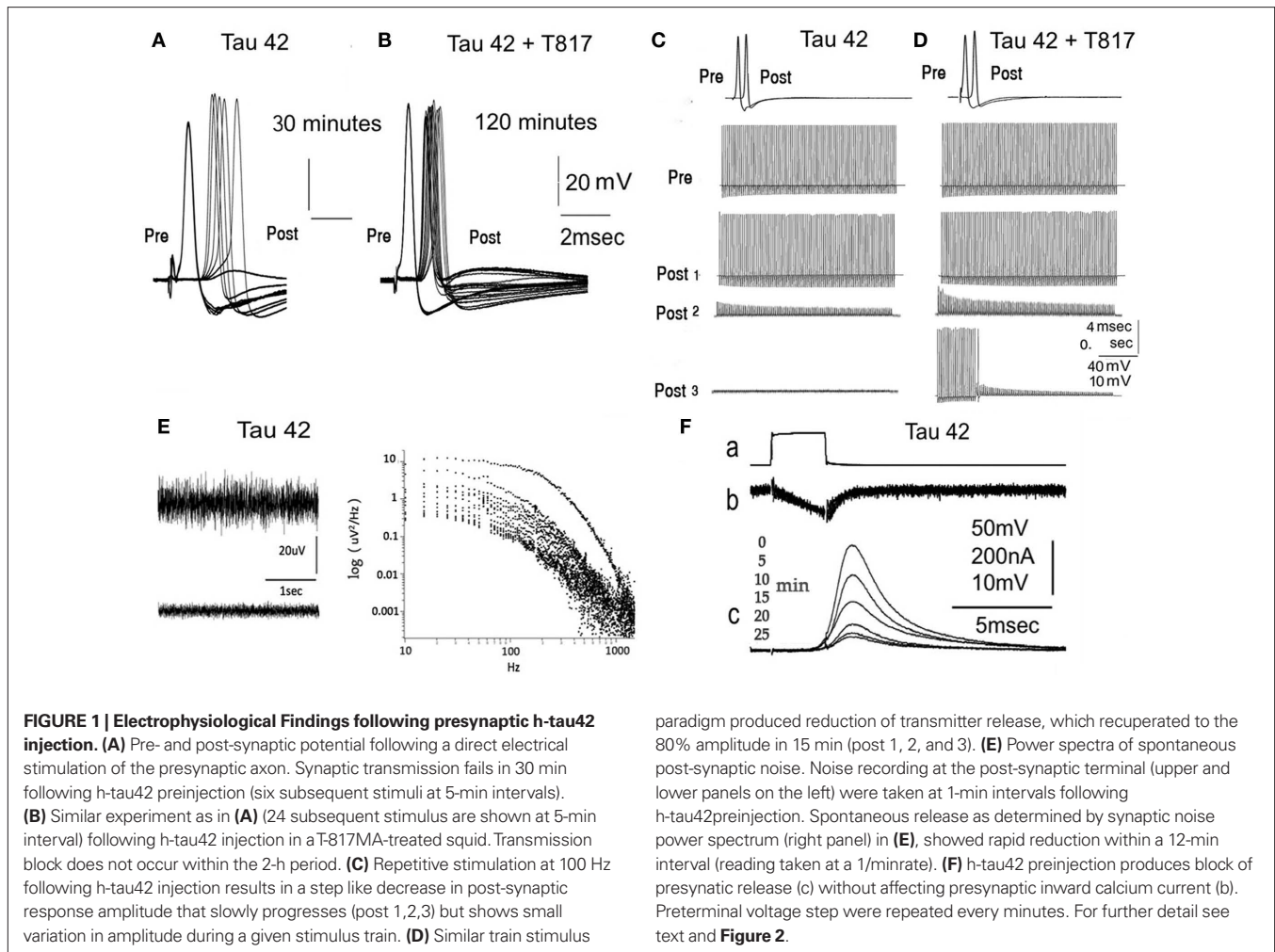
PHARMACOLOGICAL TOOLS

T-817MA (1-{3-[2-(1-benzothiophen-5-yl) ethoxy] propyl}azetidin-3-ol maleate), was synthesized at Toyama Chemical Co. Ltd., Japan.

RESULTS

INTRA-AXONAL H-TAU42 ACUTELY BLOCKS SYNAPTIC TRANSMISSION

Following presynaptic and post-synaptic axon impalements and the determination of normal synaptic transmission (Llinas et al., 1985, 1994; Lin et al., 1990), the effect of human tau on synaptic release was evaluated by presynaptic microinjections administered under direct visualization using a fluorescent dye/protein mix, reaching a final concentration after diffusion of approximately 80 nM (see Materials and Methods). Presynaptic and post-synaptic potentials were recorded simultaneously under current-clamp configuration. Presynaptic spikes were activated every 5 min (low-frequency protocol). With this paradigm, it was determined that 5–10 min after an injection of h-tau42, a reduction of transmitter release could be observed. With further time, a total block of transmission resulted, within 30–40 min depending on the length of the release zone in the preterminal axon (Figures 1A and 2 blue dots, $n = 8$). No modification of presynaptic spike amplitude or duration ensued (Figure 1A).

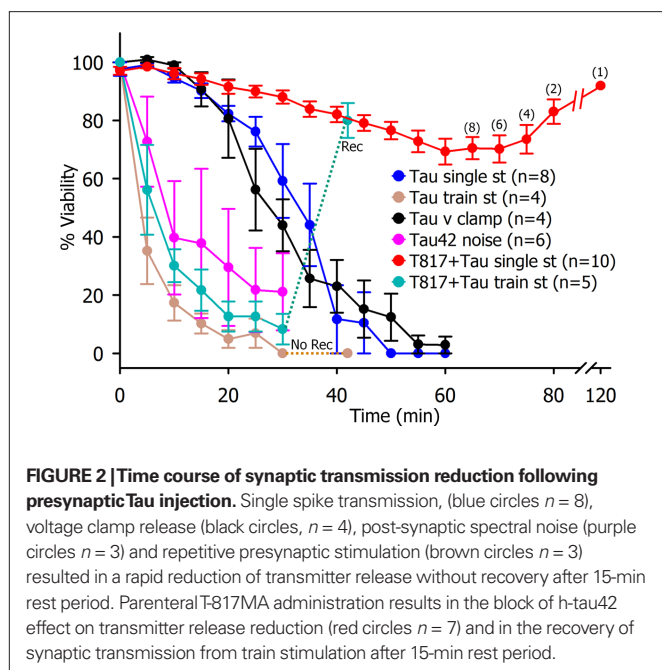


By contrast, following T-817MA administration to the squid (see below) h-tau42-dependent transmitter block was prevented (**Figures 1B and 2** red dots, $n = 7$).

To determine whether the transmission block described above was produced by a reduction of transmitter availability (as would be expected by inhibition of any step in the synaptic vesicle recycling pathway, e.g., endocytosis, refilling of vesicles with transmitter, or docking,) or by a defect in synaptic vesicle fusion, the effect of trains of presynaptic high frequency stimuli (100 Hz) was tested. This form of activation rapidly depleted the transmitter, as evidence by the rapid decrease in post-synaptic potential amplitude during a stimulus train (**Figure 1D** Post 2). The rapid time course of this decay has been shown to give an estimate of transmitter availability (Llinás et al., 1994), and is a reflection of a decrease in either synaptic vesicle mobilization or docking. On the other hand, synaptic block accompanied by a slow, progressive reduction of post-synaptic amplitude, without amplitude reduction during the tetanic stimulus, is a direct indication that the block is due to a defect in transmitter release (Llinás et al., 1994), i.e., vesicular fusion.

The test paradigm implemented to address this query consisted of synaptic high frequency (100 Hz) spike activation of the presynaptic axon of synapses preinjected with h-tau42 (**Figure 1C**, 1 Pre

Figure 2 light brown dots, $n = 4$) which initially generated post-synaptic repetitive activation (**Figure 1C**, Post 1). This train stimulation was repeated once a minute until post-synaptic spike failure occurred from the first stimulus in the train. Note that the amplitude of the subthreshold synaptic potentials showed little reduction during the stimulus train itself but steadily reduced in amplitude as h-tau42 mobilizes into the preterminal. The fact that the amplitude of the evoked post-synaptic potentials remained unchanged during the duration of a given repetitive stimulation barrage indicates that the limiting factor was the vesicular release process (**Figure 1C** Post 2). This synaptic failure did not recuperate after 15 min of rest (as normally occur in this preparation; Llinás et al., 1985, 1994; Lin et al., 1990), but rather diminished to just noticeable amplitude (Post 3). In synapses from squid pretreated with T-817MA, microinjection with h-tau42 (**Figures 1D and 2**, green dots $n = 6$), show the normal reduction of the EPSP amplitude during the repetitive stimulus barrage, however, the treated animals showed normal recovery after a 15-min rest period. These findings indicate that T-817MA administration results in the normal recovery of synaptic transmission after high frequency stimulation, suggesting that the h-tau 42 effects could relate to vesicular availability and not to the actual vesicular release process (**Figure 1D** Post 3 and **Figure 2**, green dot, $n = 5$).



TAU MODIFIES SPONTANEOUS NEUROTRANSMITTER RELEASE AND PRODUCES AN EARLY TRANSIENT INCREASE IN INTRACELLULAR CALCIUM

Beyond spike-initiated release, the possibility that h-tau42 may affect spontaneous transmitter release was directly evaluated using post-synaptic noise analysis. A detailed description of the technique has been published by our team (Lin et al., 1990). In all preparations tested ($n = 6$) the membrane noise recorded from the post-synaptic axon increased during the initial 5 ± 1 min after presynaptic microinjection of h-tau42, this was followed by rapid noise level reduction in parallel with decreased amplitude of the evoked transmitted release. An example of the time course of noise level changes is illustrated in Figure 1E (left panel and Figure 2 purple dots $n = 3$). Note the biphasic changes in noise levels. A frequency spectrum of the membrane noise at different times after the h-tau42 injection is shown in Figure 1E (right panel). This pattern indicates that block by h-tau42 interferes with both spontaneous and evoked transmitter release.

MECHANISMS UNDERLYING TAU-INDUCED SYNAPTIC BLOCK

Following the initial finding that transmission is rapidly blocked by h-tau42; the possibility that this block was associated with changes in presynaptic calcium currents (ICa^{2+}) was tested. The amplitude and time course of ICa^{2+} (Figure 1F) were directly determined by presynaptic voltage clamp steps, after blocking voltage dependent K^+ and Na^+ currents ($n = 6$), as previously described (Llinas et al., 1985). ICa^{2+} amplitude and time course were determined at 5-min intervals over a period of 25 min following presynaptic injection of h-tau42, 80 nM.

Voltage clamp experiments were implemented with presynaptic voltage steps that generated presynaptic inward calcium current and post-synaptic EPSPs. Following h-tau42 injection presynaptic voltage steps were repeated at 5 min intervals (low-frequency stimuli), which resulted in a progressive reduction

of post-synaptic response amplitude, to total failure, without a change in the amplitude or time course of the presynaptic ICa^{2+} (Figure 1F). Under normal conditions this paradigm results in transmitter release that last, unaltered, for up to 2 h, the maximum period utilized (Llinas et al., 1985). The results show, therefore, that neither the time course nor the amplitudes of the presynaptic calcium currents were altered concomitantly with the transmitter release block induced by h-tau42 (Figures 1F and 2). The time course and number of electrophysiological experiments are summarized in Figure 2.

INTRA-AXONAL H-TAU42 BECOMES PHOSPHORYLATED AND PRODUCES SYNAPTIC VESICLE AGGREGATION

Since the aggregation of typical tau filaments is accompanied by the development of tau hyperphosphorylation, we investigated whether h-tau42 residues serine 202, threonine 205 and/or 231 were phosphorylated in the squid synapse. We used AT8 antibodies, as commonly used in neuropathological studies (Goedert et al., 1995). Here, we used immunohistochemistry in a variance of the array tomography technique (Micheva and Smith, 2007); we found that single sections (500 nm) allowed a clear view of the pre- and post-synaptic compartments (Figure 3A). Anti-phospho-tau immunohistochemistry was detected as dot-like profiles in the presynaptic compartment in h-tau42 injected synapses but drastically reduced in T-817 treated squid (Figure 3B lower panel). These were absent in synapses injected with vehicle (Figure 3D lower panel). This finding demonstrates that h-tau42 becomes phosphorylated in the squid axon.

ULTRASTRUCTURAL PRESYNAPTIC CHANGES SECONDARY TO H-TAU42 INJECTION

The structural changes that follow h-tau42 injection were addressed by rapidly fixing stellate ganglia (see Materials and Methods) after high or low-frequency stimulation protocols. The material consisted of injected synapses (62 synaptic active zones from 10 different squid) and vehicle-injected synapses (control, 27 active zones in five synapses). The synapses were fixed ~75–90 min after h-tau42 injection and processed for ultrastructural microscopy (see Materials and Methods). There was a statistically significant reduction in the number of “docked vesicles” in h-tau42-injected synapses compared to axons injected with vehicle (Table 1). This reduction was not seen in T817MA-treated squid following h-tau42 injection (Table 1).

As shown in representative control synapses (Figure 4A), vesicles are normally present at the active zone, some in contact with the presynaptic terminal membrane (docked). By contrast, in h-tau42-injected synapses (Figure 4B) vesicles were often closely aggregated with electron dense material serving as a bonding matrix (red dot). Similar electron dense material was also observed around vesicles in contact with the active zone (red arrows). At a lower magnification (Figure 4C) a large number of aggregated vesicular profiles are evident in the vicinity of the active zone (red dots). In T-817MA administered squid, the synaptic morphology was quite similar to the vehicle-injected synapses (compare Figures 4A,D). Quantification of number of undocked, docked, and clathrin-coated vesicles (CCV) among the three groups (control, h-tau42 injected, and h-tau42 injected + T-817MA) is tabulated in Table 1.

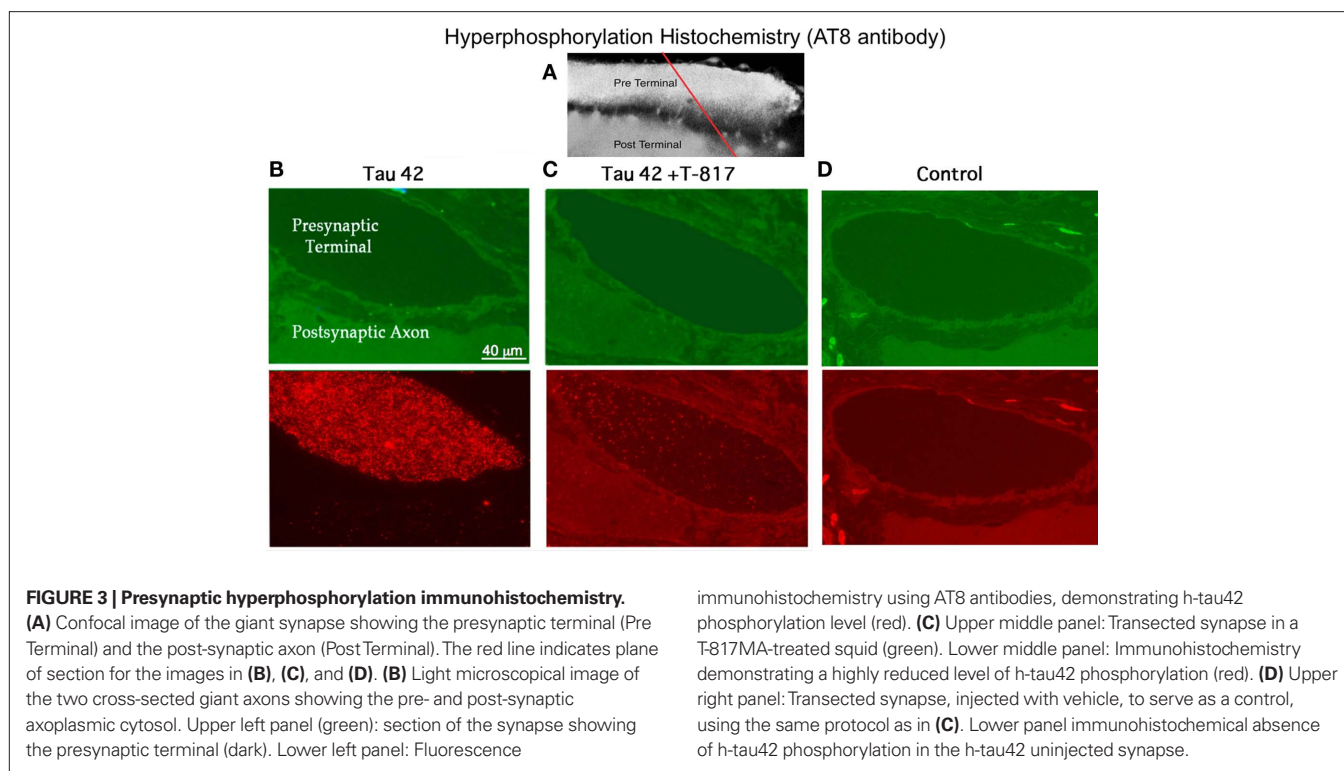


Table 1 | Quantification of preterminal docked vesicles in: control, h-tau42 injected and h-tau42 injected in T-817MA pretreated squid.

	Undocked	Clathrin-coated	Docked
Control	67.3 ± 5.8	3.2 ± 0.4	10.1 ± 0.7
h-tau42	61.2 ± 3.3	4.3 ± 0.3	5.8 ± 0.4*
T817-MA + h-tau42	56.7 ± 3.5	3.9 ± 0.4	10.0 ± 0.6*

Number of vesicles in 1 μm^2 of active zone, Average \pm SEM, * p 0.001.

SYSTEMIC ADMINISTRATION OF T-817MA PREVENTED TAU-MEDIATED SYNAPTIC BLOCK, SYNAPTIC VESICLE AGGREGATION, AND DECREASED H-TAU42 PHOSPHORYLATION

T-817MA (1-{3-[2-(1-benzothiophen-5-yl)ethoxy]propyl}azetidin-3-ol maleate) is a newly developed neuro-protective agent that prevents oxidative stress, nitric oxide-induced neurotoxicity, and acts as a neurotrophic factor (Hirata et al., 2005). Although it has been shown that T-817MA administration may ameliorate amyloid beta related neuronal dysfunction (Nguyen et al., 2007; Kimura et al., 2009), its effect on tau related pathology is unknown. To study the potential effects of T-817MA on tau neuropathology, squids received an oral dose of T-817MA as described in the Section “Materials and Methods”. Electrophysiologically, as shown in **Figure 1B** no significant changes in the amplitude or time course of the pre- or post-synaptic potentials were observed. Further, ultrastructural studies in synapses used for the electrophysiological experiments demonstrated the number of docked vesicles recovered to the normal range in h-tau42/T-817MA squid (31 active zones in six synapses; 10.0 ± 0.6) compared to control synapses (10.1 ± 0.7), with the presence of normal CCV profiles (3.9 ± 0.4). Also clear was a significant reduction, of electron dense vesicles clusters and electron dense active zones (see **Table 1**; **Figure 4**). It is thus concluded that T-817MA

prevented the h-tau42 dependent synaptic vesicle clustering, indicating a close relation between such morphology and the synaptic transmitter release block observed electrophysiologically. Finally squid pretreated with T-817MA showed a significantly reduced signal of intra-axonal h-tau42 phosphorylation, as detected by AT8 immunohistochemistry (**Figure 3**).

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DISCUSSION

While the underlying pathogenic mechanisms of tau related neuronal abnormalities remain obscure, the acute effects of preterminal injection of h-tau42 presented here suggest that synaptic dysfunction is an early mechanism in AD and other tauopathies. We used the squid giant synapse in this study because it provides unique advantages in addressing the cellular and molecular mechanisms involved in chemical synaptic transmission. In this set of experiments we determined that h-tau42 produces a rapid failure in exocytosis. Our results also indicate that h-tau42 has previously unknown physiological properties that may be relevant in tau related neurodegenerative process.

HUMAN TAU-42 ACUTELY BLOCKS CHEMICAL SYNAPTIC TRANSMISSION WITHOUT AFFECTING THE PRESYNAPTIC CALCIUM CURRENTS OR THE ENDOCYTIC PATHWAY

The data presented here indicate that an excess of h-tau42 protein produces synaptic transmission block by interfering with a mechanism of synaptic vesicle exocytosis. Our conclusion, that h-tau42 induces a failure in neurotransmitter availability due to reduced synaptic vesicle release, is derived from morphological (**Figures 4B,C**), high frequency stimulation (**Figure 1C**), and spontaneous neurotransmitter release data (**Figure 1E**). Moreover, all the h-tau-42 injected synapses demonstrated a drastic block of

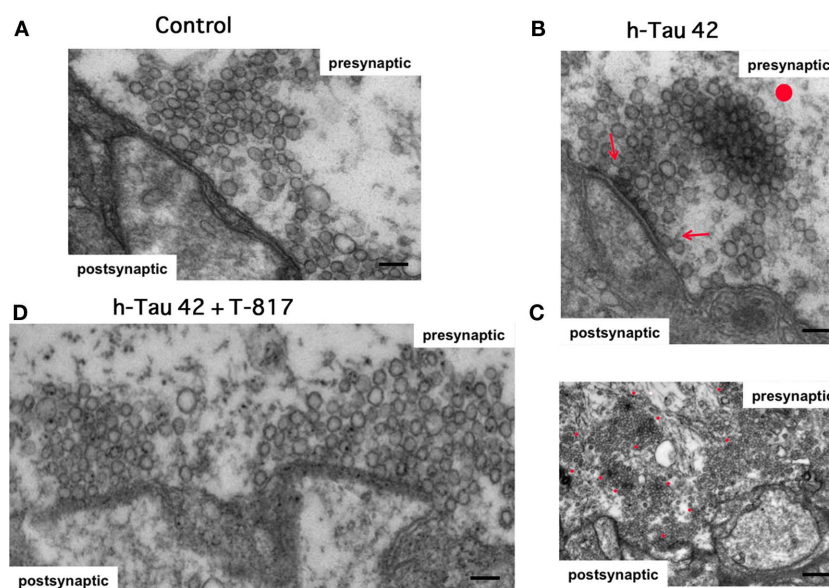


FIGURE 4 | Ultrastructure of the synaptic junction demonstrating differences in normal and h-tau42 injected synapses in untreated and T-817MA-treated squid. (A) Control synapse showing the typical presynaptic ultrastructure with synaptic vesicles grouped at the synaptic active zone (scale bar: 100 nm). **(B)** Ultrastructure of ah-tau42 injected synapse showing a synaptic vesicle cluster congealed around electron dense

material (red dot, scale bar: 100 nm). **(C)** low magnification image of a h-tau42 injected synapse showing a large number of vesicular clusters as in the upper panel also marked with red dots (scale bar: 500 nm). **(D)**: Ultrastructural image of a T-817MA-treated squid demonstrating normal synaptic morphology and properly docked vesicles at the presynaptic active zone (scale bar: 100 nm).

both spontaneous and evoked transmitted release, without affecting presynaptic spike generation (Figure 1A) or the associated calcium current (Figure 1F). We interpret these findings as reflecting the reduced vesicle count at the active zone, the vesicles being instead concentrated in groups away from the active zone. It was also noted that these electron dense vesicular congregations were characterized by profiles resembling vesicular adhesions to microfilaments as would be expected if synapsin 1 were to be dephosphorylated affording a strong adhesion to such microfilaments (Llinas et al., 1985). As a result h-tau42 would lead to the failure in exocytosis due to both a defect in the release mechanism and a reduction in vesicular availability (Llinas et al., 1991).

HUMAN TAU IS PHOSPHORYLATED IN THE ISOLATED PRESYNAPTIC TERMINAL AND INDUCES ABNORMAL VESICULAR CLUSTERING

Previous experiments in drosophila have shown that misexpression of human – tau (h-tau), the same isoform as the one used in our experiments, produced significant neurodegeneration (Jackson et al., 2002; Avila et al., 2004; Steinhilb et al., 2007). In the drosophila model tau co-expressed with Shaggy, which generated a single fly homolog of GSK-3 β , the phenotype was aggravated (Jackson et al., 2002). Dysfunctional phenotypes were also found in the central neurons of lamprey, where long-term expression (2–38 days) of several h-tau isoforms produced neurodegenerative changes as a result of accumulation of h-hyperphosphorylated tau which correlated with the appearance of structures that resemble AD characteristic – “straight like filaments” (Hall et al., 1997). In the latter experiments the isoform hyperphosphorylated moiety was, to a larger extent, the long form of h-tau (h-tau42; Hall et al., 1997; Lee et al., 2009). It has been proposed that the physiological function of

tau is adversely affected by excess phosphorylation resulting in tau being displaced from microtubules and aggregating, which in turn leads to microtubule disassembly, disruption of axonal transport, and finally synaptic failure (Stamer et al., 2002).

Concerning cephalopods, it has been demonstrated that h-tau binds to the squid axonal microtubules, but monomeric h-tau did not affect fast axonal transport (FAT), while filamentous h-tau42 did block anterograde FAT (LaPointe et al., 2009). However, other studies have found that flies misexpressing tau show defects in neuronal traffic without evidence of tau aggregation (Jackson et al., 2002; Mudher et al., 2004). Finally, extracellular applied h-tau42 to cell cultures produced aberrant signaling through muscarinic receptor activation (Gomez-Ramos et al., 2008; Diaz-Hernandez et al., 2010), suggesting that even “normal” tau may be detrimental when its expression becomes elevated or when it accumulates extracellularly. From these observations, it appears that an optimal level of tau phosphorylation is required to achieve the balance in the level of “free” and “microtubule bound” tau that is essential in maintaining microtubule dynamics and subsequent axonal transport.

As in the experiments mentioned above, in our experiments h-tau42 also became phosphorylated in the isolated axon (separated from the cell body) as demonstrated by immunohistochemistry using AT8 antibodies (Figure 3B). AT8 recognizes epitopes phosphorylated by GSK3 and cdk5 kinases both of which are found in squid axoplasm (Takahashi et al., 1995; Morfini et al., 2002; Hanger et al., 2009), suggesting that either one or both kinases may be involved in the effects of h-tau42 in the presynaptic terminal. Whichever the specific kinase, the results demonstrate that isolated axons have the complete machinery to produce local post

translational modifications and that these changes may explain, in part, the detrimental effects of excessive “normal tau” on the function of the presynaptic terminal.

Moreover, the vesicle clustering observed in h-tau42 injected synapses, resembled the effect of unphosphorylated synapsin 1 on synaptic vesicle (Jackson et al., 2002). The fact that h-tau42 is phosphorylated intra-axonally and that unphosphorylated synapsin 1 restrains the vesicle pool to the cytoskeleton – producing a decreased number of vesicles available for exocytosis was actually demonstrated some time ago (Llinas et al., 1985).

A possible explanation of our findings is that h-tau42 induces changes in the balance of kinases and phosphatases, perhaps influenced by the concentration of h-tau aggregates. We hypothesize that this could decrease the phosphorylated/dephosphorylated ratio in proteins involved in synaptic vesicle function, such as synapsin 1, which would result in a reduction in the available vesicles and ultimately synaptic transmission failure. Tau is phosphorylated by several protein kinases and this is balanced by protein phosphatases dephosphorylation. The potential kinases and phosphatases involved are so numerous that biochemical experiments dedicated to solve this issue are necessary. If this process were to involve also constitutive vesicular dynamics a secondary dying-back event (Moreno et al., 2009) would result in the synaptic disconnection encountered in AD pathomorphology.

POTENTIAL PHARMACOLOGICAL TARGETS OF TAU-MEDIATED NEUROPATHOGENESIS

The present results identify the protective effect of systemic administration of T-817MA on h-tau42 mediated axonal/synaptic dysfunction as shown in **Figures 1B,D, 2, and 3C**. This finding is highly significant, as it has been demonstrated that reduction of endogenous tau in an AD mouse model, ameliorates amyloid beta induced neurodegeneration at several levels (Roberson et al., 2007). Therefore, T-817MA has a potential to treat tau pathology. Addressing the mechanisms of T-817MA neuro-protection in relation to tau pathology remains to be elucidated. But the fact that both

functional and biochemical h-tau42 induced abnormalities in the presynaptic axon are prevented/ameliorated by this compound is a fundamental initial step toward the identification of therapeutic candidates to treat tauopathies.

CONCLUSION

Our results indicate that h-tau42 affects synaptic release by modifying intracellular phosphorylation homeostasis as a resultant of the h-tau42 hyperphosphorylation. This dynamic change, we propose, leads to a marked reduction of synaptic vesicle availability, most probably due to the reduction of synapsin 1 phosphorylation, known to be a powerful modulator of synaptic release (Llinas et al., 1985). Beyond affecting synaptic release the reduction of such vesicular fusion on constitutive vesicular dynamics would also result in a disconnection event ultimately generating a “dying-back” phenomenon (Stamer et al., 2002; Pigino et al., 2007; Serulle et al., 2007). Systemic administration of T-817MA, a proposed neuro-protective agent, rescued tau-induced synaptic abnormalities, markedly reduced h-tau42 hyperphosphorylation and prevented synaptic vesicle clustering, as determined by ultrastructural analysis.

AUTHOR CONTRIBUTIONS

Herman Moreno, Soonwook Choi, Eunah Yu, Mutsuyuki Sugimori, and Rodolfo R. Llinás designed research; Herman Moreno, Soonwook Choi, Eunah Yu, Jorge E. Moreira, Mutsuyuki Sugimori, and Rodolfo R. Llinás performed research; Herman Moreno, Soonwook Choi, Eunah Yu, Janaina Brusco, Jorge E. Moreira, Mutsuyuki Sugimori, and Rodolfo R. Llinás analyzed data; Herman Moreno and Jesus Avila implemented molecular biology, and Herman Moreno, Soonwook Choi, G.P., Jorge E. Moreira, Jesus Avila, Mutsuyuki Sugimori, and Rodolfo R. Llinás wrote the paper.

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