

Factors Related to Alzheimer's Disease, Tau Pathology in Alzheimer's Disease: Possible Treatments for Tau Pathology

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Abstract

The Tau protein is a microtubule-associated protein that acts as a three-dimensional "railroad tie" for the microtubule. Accumulation and Aggregation of the Tau is the pathogenesis of Alzheimer's disease. Risk factors like ApoE risk alleles, changes in the endoplasmic reticulum, and Kinases and phosphatases dysregulation have identified as the most critical factors. In Tau pathology, the abnormal hyperphosphorylation of tau appears as its accumulation in the affected neurons in Alzheimer's disease.

Neurofibrillary Tangles has shown truncated tau in both Glu-391 and Asp-421. Truncated tau associated with apoptosis in cultured cells. All six molecules of tau are the hyperphosphorylated state in PHF. In AD, hyperphosphorylated tau is present as a cytosolic protein and PHF. Treatments related to tau pathology are under research. Tau phosphorylation inhibitors and Tau aggregation inhibitors tested in people with AD. In tau phosphorylation inhibitors, Lithium has multiple targets and inhibits GSK-3b, and in tau aggregation inhibitors, many drugs block aggregation of tau in cell-free conditions. Methylene blue has multiple targets; it slows disease progression. Tau pathology appears to be a primary cause of neurodegeneration in AD. Risk factors showed a relation between AD and Tau pathology clearly. Abnormal hyperphosphorylation of tau leads to AD, and truncated tau is the main finding in tau pathology. Tau phosphorylation inhibitors and Tau aggregation inhibitors are emerging treatments.

Keywords: Alzheimer's disease, Paired Helical Filaments, Neurofibrillary tangles, Tau pathology, Abnormal Hyperphosphorylation, Truncation, Conformation and Isoforms, Toxicity, Tau Phosphorylation Inhibitors and Tau Aggregation Inhibitors.

Introduction

Tau is the essential Microtubule-Associated proteins (MAP) in the neurons and acts as a three-dimensional "railroad tie" for the microtubule. Accumulation and Aggregation of the Tau is the pathogenesis of Alzheimer's disease. Phosphorylation of tau binds it to the microtubules and helps in maintaining the structure,

stability of neurons¹. Accumulation of phosphate (Hyperphosphorylation) on the tau proteins cause "paired helical filaments" (PHFs) that accumulate and lead to the neurofibrillary tangles (NFTs)². PHFs are the main component in NFTs. Abnormally hyperphosphorylated, insoluble, and filamentous tau was the main component of Neurofibrillary Tangles^{3,4,5}. NFTs are neurological hallmark of AD⁶. It expressed in the central and peripheral nervous system and less amount observed in the kidney, lungs, and testis⁷. Abundantly seen in neuronal axons⁸. The human tau gene is located over 100 kb on the long arm of chromosome 17 at band position 17q21 and has 16 exons⁹. Tau divided into four regions: N-terminal Projection Region, a Proline-Rich Domain, a Microtubule-Binding Domain (MBD), a C-terminal region¹⁰. Tau can bind to outside and inside of microtubules by N- and C- terminal regions projecting outwards^{11,12}.

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Factors related to Alzheimer's disease

ApoE Risk alleles

ApoE is a risk factor related to AD in relation to tau pathology. ApoE, as a vital component of chylomicrons and IDLs involved in the catabolism of lipoproteins¹³. It has a crucial role in the metabolism of fats in the body. ApoE is a cholesterol carrier in the brain. ApoE transfers lipoproteins, fat-soluble vitamins, cholesterol to the lymph vessels, and into the blood. ApoE is abundant in CNS but synthesized in the liver¹⁴.

Accumulated cholesterol in the neurons converted into 24OHC (24S-hydroxycholesterol) in neurodegeneration¹⁵. The CSF tau levels related to neurodegeneration in AD¹⁵. The association between ApoE, 24S-hydroxycholesterol, and tau shows its direct involvement in the generation of NFTs¹⁵.

Damage in Endoplasmic reticulum

Damage in Endoplasmic reticulum is a significant risk factor related to AD in relation to tau pathology. The Endoplasmic reticulum functions are energy production, folding, and trafficking of proteins and apoptosis¹⁶. Defects of ER results in cell stress response and triggers apoptosis. Role of ER in the development and progression of neurodegenerative diseases. Inositol-Requiring kinase 1 (IRE1) starts the ER stress pathway by triggering Apoptosis signal-regulating Kinase 1 (ASK 1), activates c-Jun N-terminal Kinase (JNK) signaling route¹⁷. This cascade has the strength to trigger AD pathology through hyperphosphorylation of tau and aggregation of neurofibrillary tangles¹⁸. ER stress causes deposition of misfolded like tau¹⁹, and long-time existence of these toxic misfolded proteins triggers intrinsic apoptosis pathways²⁰. Overexpression increase the intracellular concentration of tau may inhibit the plus-end-directed transport of vesicles along microtubules by kinesin, and the minus-end-directed transport by dynein becomes dominant²¹.

Tau degradation decreased by 20% in ER stress due to a decrease in the tau binding to CHIP (carboxyl-terminus of Hsc70-interacting protein), which delayed the tau degradation through the ubiquitin-proteasome pathway²².

Kinases and phosphatases dysregulation

Dysregulation of kinases and phosphatases is a critical risk factor related to AD in relation to tau

pathology. Several protein kinases like cyclic AMP-dependent Protein Kinase A (PKA), Glycogen Synthase Kinase-3 (GSK-3), cyclin-dependent protein kinase-5 (cdk5), Calcium/Calmodulin-dependent Protein Kinase-II (CaMKII), Mitogen-Activated Protein (MAP) kinase, Extracellular signal-regulated kinase (ERK 1/2), Stress-activated protein kinases are involved in abnormal hyperphosphorylation of tau²³. Some kinases like Protein Kinase A (PKA), Protein Phosphatases 2A (PP2A), Glycogen Synthase Kinase-3 (GSK-3), Calcium/Calmodulin-dependent Protein Kinase-II (CaMKII) plays a role in Tau phosphorylation and dephosphorylation²⁴. Abnormally hyperphosphorylated sites in tau are proline-directed, serine, and threonine, followed by proline are official sites of proline-directed protein kinases (PDPKs). GSK-3, cdk5, PDPKs, and ERK 1/2 are subjected to phosphorylate tau the sites in AD²⁵.

The preparation of tau by PKA or CAMKII is enough to initiate the abnormal hyperphosphorylation of tau. Dysregulation of Tau kinase CAMKII effects AD progression²⁶, and CAMKII inhibits tau-microtubule interaction by tau phosphorylation²⁷. When GSK-3 overexpressed and it results in hyperphosphorylation of tau²⁸. GSK-3 activation results in brain aging and AD initiates detrimental events like NFT formation and neuronal death pathways²⁹. PP-2A and PP-1 do 90% of serine/threonine protein phosphatase activity³⁰. Phosphorylation of ERK inhibits activity in response to neuronal stimuli³¹.

Tau pathology

In AD, the regular role of tau impaired because this protein loses its capacity to bind to microtubules. Tau pathology starts from the entorhinal cortex and continues to the hippocampus, neocortex³², frontal, temporal cortices, and all isocortex areas. Tau pathology observed in the form of abnormal Hyperphosphorylated protein. There is a very robust correlation between tau pathology and clinical measures of dementia. In AD, tau pathology also plays a significant role because this has the potential to trigger AD in humans.

Abnormal hyperphosphorylation of tau

The main characteristic feature of tau pathology is the abnormal phosphorylation of tau. The abnormal hyperphosphorylation of tau and significant protein subunit of PHF, which results in neurodegeneration by sequestration of MAPs, self assembles bundles of

PHF and forms NFTs. In Tau pathology, accumulation of abnormal Hyperphosphorylated tau associated with neurofibrillary degeneration and dementia. The abnormal hyperphosphorylation of tau observed in both NFTs³³ and the cytosol of AD brains³⁴. Mutations in the tau gene and their cosegregation linked to chromosome-17 (FTDP-17) have abnormalities in tau as a leading, and the first event leads to neurodegeneration and dementia³⁵. The study of mAb Tau1 revealed deposits of abnormally hyperphosphorylated tau in neurons without tangles (stage "0" tangles) in AD³⁶. In tau, Ghost tangles are ubiquitinated^{37,38}, and in abnormally hyperphosphorylated tau, cytosol has no ubiquitin reactivity. Abnormal hyperphosphorylation of tau might be due to a conformational change in tau, and tau conformationally altered in AD^{39,40}. On hyperphosphorylation, murine tau self assembles into tangles of filaments (PHF/SF)⁴¹. Abnormal hyperphosphorylation of tau comes before its accumulation into NFTs⁴². In our research, tau hyperphosphorylation seen at many sites and even HMW-tau hyperphosphorylated at many locations. An essential finding of the study is overexpressing p25 initiates and promotes hyperphosphorylation. This abnormal hyperphosphorylation leads to filaments self-assembly.

Truncated tau

Presence of truncation connected to neurofibrillary pathology in the brain of AD patients⁴³. Neurofibrillary Tangles has shown truncated tau in both Glu-391 and Asp-421. Truncated Tau associated with apoptosis in cultured cells⁴⁴. There is no report on what percentage of tau truncated at affected sites in different stages of AD. Neurodegeneration in AD seen for a long time from months to years, and there is a fair chance to view truncation in tau. Truncation is seen in both affected neurons and ghost tangles (extracellular space) when NFTs exposed to hydrolases⁴⁵. Truncation of tau was observed by immunolabeling with the monoclonal antibody MN423⁴⁶. MN423 recognizes Glu-391. Glu-391 was associated with NFTs and progression related to neurofibrillary pathology described by Braak's stages⁴⁷. Braak stages I and II – NFT involvement confined to the transentorhinal region of the brain, Stages III and IV – Also involvement of limbic regions (hippocampus) and Stages V and VI – Extensive neocortical involvement⁴⁷.

In vitro studies described that truncated tau at Glu-391 showed elevated rates of polymerization over full-length tau, promoted by arachidonic acid⁴⁸. Truncation

of tau was observed by immunolabeling with the monoclonal antibody Tau-C3⁴⁹. Tau-C3 recognizes Asp-421. Asp-421 was associated with the neurofibrillary pathology in the AD brain⁵⁰. *In vitro* studies described that truncated tau at ASP-421 showed elevated rates of polymerization over full-length tau⁴⁹. Cells transfected with truncated tau protein, it has demonstrated that both soluble and insoluble forms can induce toxicity^{51,52}. In COS7 cells that Asp-421 is highly phosphorylated and abnormally redistributed. In HEK-293T cells that Glu-391 has a problem in binding with microtubule⁵¹, and there is toxicity resulted in cell death. In the progression of the disease, Early event truncation of Asp-421 and preceded by truncation of C-terminal of Glu-391, further occurring from intermediate to advanced stages of NFTs evolution⁵³. In truncation, this research specially confined to truncated tau in Glu-391 and Asp-421. In our findings, COS7 cells highly phosphorylated in Asp-421 and HEK cells have a binding problem with microtubule.

Isoforms of tau

Six isoforms of tau differ by contents of three (3R) or four (4R) tubulin-binding domains of 31 or 32 amino acids in C-terminal and one (1R), two (2R) or inserts of 29 amino acids each in N-terminal of tau. All six molecules of tau are the hyperphosphorylated state in PHF 3,4,54. 441-residue tau is the longest tau isoform seen in the human CNS. In AD, hyperphosphorylated tau is present as a cytosolic protein⁴² and PHF^{3,4,54}. Cytosolic hyperphosphorylated tau (P tau) has 5-9 mol of phosphate per mole of the protein (contains 2-3 phosphate groups)⁴². Tau self-assembles by intermolecular hydrophobic interaction and microtubule-binding repeat R3 (3R taus), and R2 and R3 (4R taus), when the rest of the molecule neutralized⁴¹. Inhibitory amino-terminal and carboxyl-terminal regions neutralized by abnormal hyperphosphorylation in AD⁴¹. Cytosolic and PHF are abnormally hyperphosphorylated taus are readily dephosphorylated by phosphatases *in vitro*^{33,54,55,56}. Tau molecule changing its conformation to form a paperclip-like conformation by folding the N- and C-terminal portions back on the microtubule-binding repeats⁵⁷. FTDP-17 mutations alter the conformation of the protein, and it becomes a more favorable substrate to brain protein kinases⁵⁸. Mutated taus are soon hyperphosphorylated and self-assemble at a lower level of hyperphosphorylation⁵⁸. In tau, Dephosphorylation inhibits self-assembly, and hyperphosphorylation promotes self-assembly, but Deglycosylation of AD promotes self-assembly. This study found that all six tau

isoforms self-assembled into PHF/SF. This result clearly explains that hyperphosphorylation is enough to initiate self-assembly of tau into filaments. The 3R isoforms are more toxic than 4R.

Possible treatments for tau pathology

Treatments targeting different aspects of tau pathology are under research. Tau phosphorylation Inhibitors and Tau Aggregation Inhibitors are in clinical trials for people with AD, while tau reduction strategies are still in preclinical trials.

Tau Phosphorylation Inhibitors

In tau phosphorylation inhibitors, Lithium has multiple targets and inhibits GSK-3b⁵⁹. Lithium has a narrow safety margin⁶⁰. Studies in geriatric patients taking chronic Lithium for BPAD showed a reduced risk of developing AD. Lithium inhibits chemical changes in tau that lead to the formation of NFT's. Studies on Lithium shown benefit with low doses in mild cognitive impairment, Reduction of GSK-3b impairs NMDAR-mediated long-term depression⁶¹ and memory consolidation⁶², raising concerns about potential side effects of GSK-3b inhibitors. CDK5 is essential for multiple cell signaling pathways and adult neurogenesis, limiting its appeal as a tau-targeting approach in AD. However, CDK5 and p25, a truncated form of the CDK5 subunit p35, promote neurodegeneration through mechanisms that are independent of tau phosphorylation⁶³.

Tau Aggregation Inhibitors

Filamentous tau aggregates are damaging forms of tau and Filamentous tau is toxic. Tau enters dendritic spines, and adverse effects of tau aggregates are seen only in intracellular compartments⁶⁴. Many of the drugs that block the aggregation of tau also prevent the pathological aggregation of other proteins under cell-free conditions⁶⁵. Some tau aggregation inhibitors are effective in Neuro2A cell lines overexpressing a 4R tau microtubule repeats domain fragment with a K280 deletion and promotes its aggregation⁶⁶. In AD patients, the Methylthioninium Chloride (methylene blue) showed slowing disease progression⁶⁷. Methylene blue is the first drug targeting tau. The drug derived from the dye used to stain NFT's in neuropathological studies. Primarily inhibits tau aggregation and showed cognitive benefits. Methylene blue inhibits tau-tau interactions⁶⁸ and reduces soluble tau by mechanisms⁶⁹. Inhibition

of tau aggregation mediated by direct binding of tau to the FK506 binding protein 52⁷⁰. It is unknown that tau assembly is responsible for tau-dependent neuronal dysfunction and degeneration⁷¹.

Other treatments

Hyperacetylation increases half-life⁷² and due to this microtubule-binding impairs and initiates aggregation⁷³. Lysosomal pathway degrades proteasome, removes aggregated tau⁷⁴, and inhibition of this pathway produces NFT-like tau deposition⁷⁵. Lysosomal pathway of tau degradation involved in Niemann-Pick type C disease (NPC), seen with neurological symptoms and NFT formation⁷⁶. NPC caused by loss of function of NPC1 (Lysosomal trafficking protein)⁷⁷. We found that that degradation of tau occurs in Lysosomal and ubiquitin-proteasome pathways. Destabilization of microtubules and problem binding to microtubules are two main problems related to overexpression of tau. Microtubule stabilizers have shown excellent results in preclinical and clinical trials of AD. Epothilone D has better BBB permeability, improved microtubule density, and cognition⁷⁸. Peptide NAP stabilizes microtubules⁷⁹ and reduces tau phosphorylation⁸⁰. In our study, Microtubule stabilizers have shown more than one mechanism of action.

Conclusion

Tau pathology is the leading and primary cause of neurodegeneration in AD. In the brain of AD, the level tau expression has no change concerning healthy aging, and toxicity depends on the contribution of many factors, like loss of normal function, the impermanent aggregation, and state of tau processing. Abnormal posttranslational modifications, like hyperphosphorylation and truncation, are responsible for altered tau structure in AD. Abnormal hyperphosphorylation and truncation are supported by *in vitro* experiments explain modifications elevate fibrillization of tau and induce cell toxicity⁸¹. Conformational changes of tau promoted by posttranslational modifications and the role of fibrillization important for checking the potential of tau-directed therapies⁸¹. Inhibition of the abnormal Hyperphosphorylation of tau and sequestration of MAPs by the hyperphosphorylated tau⁴¹ is the most critical therapeutic targets for AD. The pathogenic tau slowed or prevented by treatments that reduce levels of extracellular tau, so neurons cannot internalize it⁸². Identification and characterization of mechanisms of tau release and uptake

are essential for therapeutic interventions that may slow or prevent neurodegeneration in AD. Drugs are needed to develop more effectively to slow or prevent tau pathology. Methods to be prepared to restrict or inhibit the hyperphosphorylation of tau because inhibiting abnormal hyperphosphorylation does not self-assemble tau, and there will be no formation of PHF/SF.

Abbreviations

AD: Alzheimer's disease, MAP: Microtubule Associated Protein, PHF: Paired Helical Filaments, SF: Straight Filaments, NFT'S: Neurofibrillary tangles, MBD: Microtubule-Binding Domain, IDLs: Intermediate-Density Lipoproteins, CNS: Central Nervous System, CSF: Cerebrospinal fluid, 24OHC: 24S-hydroxycholesterol, ER: Endoplasmic reticulum, IRE1: Inositol-Requiring kinase 1, ASK1: Apoptosis signal-regulating Kinase 1, JNK: c-Jun N-terminal Kinase, CHIP: Carboxyl-terminus of Hsc70-interacting protein, Camp: cyclic AMP-dependent, PKA: Protein Kinase A, GSK3: Glycogen Synthase Kinase-3, cdk5: cyclin-dependent protein kinase-5, CaMKII: Calcium/Calmodulin-dependent Protein Kinase-II, MAP kinase: Mitogen-Activated Protein kinase, ERK 1/2: Extracellular signal-regulated kinase, PP2A: Protein Phosphatases 2A, PP1: Protein Phosphatases 1, FTDP-17: Frontotemporal dementia with parkinsonism-17, MN423 – Recombinant Mouse Antibody, COS-7: CV-1 in Origin with SV40 genes, HEK293T cells: Human embryonic kidney 293T cells, NMDAR: N-methyl-D-aspartate receptor, NPC: Niemann-Pick type C, BBB: Blood-Brain barrier.

Conflict of Interest: The authors declare no conflict of interest

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Ethical Considerations: Compliance with ethical guidelines

There is no ethical principle to be considered during this research.

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