

Commentary

The Tauopathies

Toward an Experimental Animal Model

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Abundant intraneuronal neurofibrillary lesions within certain brain regions constitute a defining neuropathological characteristic of Alzheimer's disease.¹ Ultrastructurally, the neurofibrillary lesions consist of abnormal filamentous deposits in the form of paired helical filaments (PHFs) and the related straight filaments (SFs). These filaments are made of the microtubule-associated protein tau in a hyperphosphorylated state. In normal brain, tau protein is soluble and nonfilamentous. Its ordered assembly into filaments is therefore a pathological event. Tau pathology is not limited to Alzheimer's disease but is also present in a number of other dementing disorders, such as Pick's disease, progressive supranuclear palsy, and corticobasal degeneration.^{2,3} In these disorders, as in Alzheimer's disease, the hyperphosphorylated tau protein is filamentous. However, the filament morphologies and tau isoform compositions differ from those of Alzheimer's disease. The good correlation between the presence of tau pathology and the degree of cognitive impairment has suggested that the events leading to the formation of tau filaments or the mere presence of these filaments are sufficient to produce nerve cell degeneration. Recently, this view has been significantly reinforced by the discovery of mutations in the tau gene in familial frontotemporal dementia and parkinsonism linked to chromosome 17 (FTDP-17).⁴⁻⁸ The new work will no doubt lead to increased efforts aimed at producing experimental animal models of the tau pathology of Alzheimer's disease and other tauopathies.

Tau Protein and its Assembly into Filaments

Tau protein promotes microtubule assembly and binds to microtubules, which are thus stabilized. In adult human brain six tau isoforms are expressed; they are produced by alternative mRNA splicing from a single gene located on the long arm of chromosome 17 (Figure 1). They differ by the presence of three or four tandem repeats of 31 or

32 amino acids each located in the carboxyl-terminal region in conjunction with 0, 29, or 58 amino acid inserts located in the amino-terminal region.^{9,10} There is also a larger tau isoform, with an additional insert in the amino-terminal region, which is mainly expressed in the peripheral nervous system.^{11,12} Eleven exons contribute to the longest human brain tau isoform, with exons 2, 3, and 10 being subject to alternative mRNA splicing.^{9,10,13} Tau expression is developmentally regulated in that only the tau isoform with three repeats and no amino-terminal inserts is present in fetal brain. There exist true species differences in the expression of tau isoforms in adult brain. Thus, only four-repeat tau isoforms are expressed in rodent brain. By contrast, all six tau isoforms are expressed in adult human brain, where tau isoforms with three repeats are slightly more abundant than tau isoforms with four repeats. The repeat regions of tau and sequences flanking the repeats constitute microtubule-binding domains.^{14,15} Tau is expressed predominantly in nerve cells, with lower levels in some glial cells. Within nerve cells, it is found mainly in axons.¹⁶ Inactivation of the tau gene by homologous recombination leads to no overt phenotype, indicating that tau is not an essential protein.¹⁷

Tau is a phosphoprotein and phosphorylation is also developmentally regulated. Thus, tau from developing brain is phosphorylated more than tau from adult brain. Tau from the PHFs and SFs of Alzheimer's disease brain is hyperphosphorylated and abnormally phosphorylated on all six isoforms compared to tau from normal adult human brain.^{18,19} This contrasts with progressive supranuclear palsy and corticobasal degeneration, where only four-repeat tau isoforms are found in the abnormal filaments.²⁰⁻²² In Pick's disease, the tau filaments consist only of three-repeat isoforms.²³ Hyperphosphorylation and abnormal phosphorylation are major biochemical abnormalities of filamentous tau. They are early events in the development of tau filaments and as a result tau is unable

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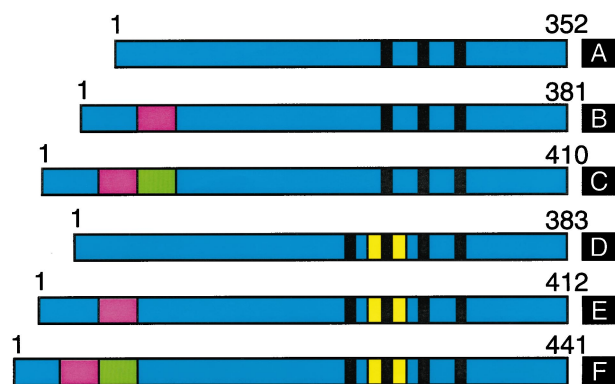


Figure 1. Isoforms of human brain tau. The region common to all isoforms is shown in blue, with the amino-terminal inserts encoded by exons 2 and 3 shown in red and green, respectively. The alternatively spliced repeat encoded by exon 10 is in yellow. The three or four tandem repeats are indicated by black bars. The tau isoforms range from 352 to 441 amino acids in length. Isoform A is expressed in fetal brain, whereas all six isoforms (A-F) are expressed in adult human brain. Transgenic mouse models based on the expression of isoform F⁴⁸ or isoform A⁴⁹ have been described.

to bind to microtubules.^{24–26} However, it is unclear whether hyperphosphorylation and abnormal phosphorylation are sufficient for the assembly of tau into filaments.

Tau Mutations in FTDP-17

Over the past few years, familial frontotemporal dementias, some with parkinsonism, have been recognized as FTDP-17, a previously unknown group of dementia disorders.²⁷ Their unifying pathological characteristic is the presence of abundant filamentous hyperphosphorylated tau deposits in the absence of A β amyloid plaques. In some of these families tau deposits are found in both nerve cells and glial cells, whereas in others only nerve cells are affected.²⁸

Besides having a filamentous tau pathology in common, the familial frontotemporal dementias also share linkage to chromosome 17q21-22, the same region that contains the tau gene.²⁹ Recently, the first mutations in the tau gene have been identified in several of these families.^{4–8} They are either missense mutations in the microtubule-binding repeat region and the carboxy-terminal region or intronic mutations that change the ratio of three-repeat to four-repeat tau isoforms. Missense mutations have been found in exons 9, 10, 12, and 13 of the tau gene; they change glycine residue 272 to valine (G272V), asparagine residue 279 to lysine (N279K), proline residue 301 to leucine (P301L), valine residue 337 to methionine (V337M), and arginine residue 406 to tryptophan (R406W) (numbering accords with the 441-amino acid isoform of human tau). The N279K and P301L mutations lie in the extra repeat of tau, thus affecting only four-repeat tau isoforms. By contrast, the other three missense mutations are found in all six brain tau isoforms. Four different intronic mutations are found in the region of the exon 10 splice-donor site, where they disrupt a predicted stem-loop. This disruption leads to increased splicing of exon 10, resulting in the overproduction of

four-repeat tau isoforms and reduced levels of tau isoforms with three repeats.^{5,6}

The functional consequences of missense mutations in tau have been studied in microtubule assembly experiments.³⁰ All the mutations investigated showed a markedly reduced ability to promote microtubule assembly. The P301L mutation produced the largest effect, the R406W mutation the smallest effect, and the G272V and V337M mutations intermediate reductions. This partial loss of function may be the primary effect of these missense mutations in tau. It may be followed by the hyperphosphorylation of tau and, through interaction with other cellular factors, by assembly into filaments. Similarly, overproduction of four-repeat tau isoforms in cases with intronic mutations may result in the inability of some of the excess tau to bind to microtubules, leading to its hyperphosphorylation and assembly into filaments. Most missense mutations are likely to lead to a reduced ability of tau to interact with microtubules. The N279K mutation may be an exception, since it creates an exon splice enhancer sequence, which may lead to increased splicing of exon 10.⁸ There may be mutations in tau that produce effects on both microtubule assembly and on mRNA splicing of exon 10.

In Seattle family A (with the V337M mutation), in familial multiple-system tauopathy with presenile dementia (with the +3 mutation in the intron following exon 10), in the Iowa family (with the R406W mutation), in pallido-pontonigral degeneration (with the N279K mutation), and in Dutch family 1 (with the P301L mutation), tau is hyperphosphorylated at the same sites as in Alzheimer's disease.^{31–35} Pick-like bodies have been described in Dutch family 2 (with the G272V mutation) that show tau staining characteristics similar to those of classical Pick bodies.^{35,36}

The balance between tau protein levels and available binding sites on microtubules appears to be critical for determining whether or not tau assembles into filaments. Thus, a reduced ability to interact with microtubules appears to be the shared primary abnormality in tau protein resulting from the different exonic and intronic mutations described thus far. A partial loss of function may be necessary for the assembly of tau into filaments.

The locations of the tau mutations appear to determine the nature of the pathology. Mutations in exon 10 or in the intron following exon 10 lead to a filamentous neuronal and glial cell tau pathology.^{32,34,35} For exon 10 mutations, the filaments are narrow twisted ribbons consisting predominantly of tau isoforms with four microtubule-binding repeats.³⁵ In the case of the intronic mutations, the filaments are wide twisted ribbons consisting exclusively of four-repeat tau isoforms.³² This is reminiscent of progressive supranuclear palsy and corticobasal degeneration, suggesting that these largely sporadic diseases may also result from abnormalities in the splicing of exon 10 of the tau gene. Missense mutations located outside exon 10 lead to a predominantly neuronal pathology.^{31,33} The tau filaments are PHFs and SFs and consist of all six tau isoforms. In the case of the V337M mutation in exon 12, the tau filaments have been shown to be indistinguishable from those of Alzheimer's disease.³¹

Synthetic Tau Filaments

Phosphorylated full-length recombinant tau has consistently failed to assemble into PHF-like filaments in *in vitro* experiments. By contrast, incubation of recombinant tau with sulphated glycosaminoglycans, such as heparin and heparan sulphate, results in the bulk assembly of tau into Alzheimer-like filaments.³⁷⁻⁴² Tau isoforms with three repeats assemble into twisted paired helical-like filaments, whereas tau isoforms with four repeats assemble into straight filaments.³⁷

Immunoelectron microscopy shows that the paired helical-like filaments are decorated by antibodies directed against the amino- and carboxy-termini of tau, but not by an antibody directed against the microtubule-binding repeat region.³⁷ These results, which indicate that in the filaments the repeat region of tau is inaccessible to the antibody, are identical to those previously obtained with PHFs from the brains of Alzheimer's disease patients.¹⁸ They establish that the microtubule-binding repeat region of tau is essential for sulphated glycosaminoglycan-induced filament formation. Three microtubule-binding repeats of tau are also believed to form the core of the PHFs found in the brains of Alzheimer's disease patients, supporting the evidence for a similar organization of the two types of filament. Previous experiments had shown that three recombinant microtubule-binding repeats of tau assemble into twisted filaments *in vitro*.^{43,44} This assembly is phosphorylation-independent and occurs in the absence of sulphated glycosaminoglycans. It confirms that three repeats are required to give the morphology of the PHF. However, these experiments do not shed light on the mechanisms that lead to tau filament formation in the brains of Alzheimer's disease patients, because PHF-tau is made of full-length tau. The dimensions of tau filaments formed in the presence of sulphated glycosaminoglycans are similar to those of filaments extracted from brains of Alzheimer's disease patients, with diameters of approximately 20 nm for twisted and 15 nm for straight filaments and a crossing-over spacing of approximately 80 nm for paired helical-like filaments, although their twist is in general less regular than that found in Alzheimer's disease filaments.

Sulphated glycosaminoglycans also stimulate phosphorylation of tau by a number of protein kinases, prevent the binding of tau to taxol-stabilized microtubules, and disassemble microtubules assembled from tau and tubulin.^{37,40} Moreover, heparan sulphate has been detected in nerve cells in the early stages of neurofibrillary degeneration.^{37,45} Sulphated glycosaminoglycans stimulate tau phosphorylation at lower concentrations than those required for filament formation. The pathological presence of heparan sulphate within the cytoplasm of some nerve cells, perhaps as a result of leakage from membrane-bound compartments, would first lead to hyperphosphorylation of tau, resulting in its inability to bind to microtubules. At higher concentrations of heparan sulphate, tau would then assemble into PHFs and SFs.

Formation of tau filaments is also observed after incubation of recombinant tau with RNA, which has been shown to be sequestered in the neurofibrillary lesions of

Alzheimer's disease.^{40,46,47} Whether the presence of RNA is an early event remains to be determined. Sulphated glycosaminoglycans and RNA share a repeat sugar backbone and negative charges in the form of sulphates or phosphates. Tau protein is thought to be an extended molecule with little secondary structure that becomes partially structured upon binding to microtubules. Binding of sulphated glycosaminoglycans or RNA to tau may induce or stabilize a conformation of tau that brings the microtubule-binding repeats of individual tau molecules into close proximity, creating sites which favor polymerization into filaments.

Transgenic Mice

The work on synthetic tau filaments has provided the first robust method for producing Alzheimer-like filaments from full-length tau. The same cannot yet be said of tau filaments in nerve cells. To date, there has been no demonstration of Alzheimer-like filaments in transgenic mice. Two studies have directly addressed this issue by expressing wild-type human tau in the brains of transgenic mice.^{48,49} It has been indirectly addressed in the transgenic mouse models of A β amyloid deposition, which are based on the expression of mutated amyloid precursor protein (APP).⁵⁰⁻⁵² Although some staining for hyperphosphorylated tau has been described in nerve cell processes around A β deposits in transgenic mice expressing mutated APP,^{52,53} no somatodendritic staining of hyperphosphorylated tau was observed in these mice. Two of these mouse lines did not exhibit nerve cell loss,^{54,55} whereas a third showed a 17% reduction in the number of nerve cells in layer CA1 of the hippocampus.⁵⁶ However, it remains to be seen whether this cell loss is mechanistically related to the nerve cell loss observed in Alzheimer's disease hippocampus. Mutated APP is expressed at high levels in these mice and this could in itself result in degeneration of some nerve cells. It is well established that in Alzheimer's disease brain there exists an inverse correlation between the number of extracellular tangles and the number of surviving nerve cells in the hippocampus,⁵⁷⁻⁵⁹ suggesting that nerve cell loss is due to the formation of neurofibrillary lesions.

The first study expressing human tau protein in transgenic mice was published in 1995 and described the expression of the longest human brain tau isoform (four repeats and the 58-amino acid amino-terminal insert) under the control of the human Thy1 promoter⁴⁸ (Figure 1). The new study, which describes expression of the shortest human brain tau isoform (three repeats and no amino-terminal inserts) under the control of the mouse 3-hydroxy-methyl-glutaryl CoA reductase promoter, is published in this issue of the *Journal*⁴⁹ (Figure 1).

Both studies describe broadly similar results with some minor differences. They show strong somatodendritic and axonal staining for hyperphosphorylated tau of subpopulations of nerve cells. The somatodendritic staining is pathological, because in control mouse brain tau staining is largely axonal. Götz et al described only nerve cell staining,⁴⁸ whereas Brion et al also describe some astro-

cytic staining,⁴⁹ presumably reflecting the use of a different promoter. The presence of hyperphosphorylated human tau in mouse brain astrocytes is interesting in view of the extensive glial tau pathology seen in some FTDP-17 pedigrees, as well as in progressive supranuclear palsy and corticobasal degeneration. Both studies show somatodendritic staining of nerve cells with a number of phosphorylation-dependent anti-tau antibodies that also stain the neurofibrillary pathology of Alzheimer's disease and other tauopathies. These antibodies also recognize tau from normal adult human brain, albeit more weakly than PHF-tau. Brion et al show that antibodies which are entirely specific for PHF-tau, such as AP422 and AT100,^{60–62} do not stain transgenic mouse brain, a finding in agreement with the lack of tau filaments. By electron microscopy, they show that transgenic human tau is associated with microtubules in axons and dendrites, but not in nerve cell bodies, where it is associated with ribosomes or distributed more diffusely.⁴⁹ Overexpression of human tau in lamprey neurons has also been shown to lead to the presence of hyperphosphorylated human tau in the somatodendritic compartment.⁶³ It thus appears that an excess of tau over available binding sites on microtubules results in the accumulation of tau in nerve cell bodies. The same may be true of the FTDP-17 cases with intronic mutations in the tau gene.

Somatodendritic staining for hyperphosphorylated tau has been described as an early pathological change in human brain, where it is characteristic of the so-called pre-tangle stage of Alzheimer's disease.²⁴ In human brain, the pre-tangle pathology progresses to the tangle stage, which is followed by nerve cell degeneration and death. In the case of the classical neurofibrillary tangle, thick bundles of tau filaments survive the death of affected nerve cells and are found in the extracellular space in the form of ghost tangles.¹ The presence of neurofibrillary tangles does not appear to be a necessary prerequisite for nerve cell degeneration, because they are absent from a number of FTDP-17 brains.²⁸ The invariant feature of the various tauopathies is the presence of filaments made of hyperphosphorylated tau protein. So far, such filaments have not been observed in the brains of mice transgenic for tau protein. There is no evidence to suggest nerve cell loss in the mice, indicating that the prolonged presence of hyperphosphorylated tau in the somatodendritic compartment of nerve cells is not sufficient to lead to nerve cell degeneration. The current transgenic mouse models thus go only part of the way towards establishing a filamentous tau pathology.

The levels of expression of human tau protein in the transgenic mouse lines were relatively modest, ranging between 10–20% of total mouse brain tau. Adult mouse brain tau consists of three four-repeat isoforms, whereas only one human tau isoform was expressed in each of the transgenic mouse studies. However, human tau appeared to be concentrated in a relatively small number of nerve cells, suggesting that the levels of human tau per cell may be much higher. Nevertheless, the failure to form tau filaments in mouse brain may be due to insufficient levels of human tau. From the experiments on synthetic tau filaments, it is clear that the assembly of recombinant

tau in presence of sulphated glycosaminoglycans is strongly concentration-dependent, as befits a nucleation-dependent process.^{37–40,42} Other differences between mice and humans may also play a role. Mice express only three four-repeat tau isoforms in adult brain, whereas humans express an additional three isoforms with three repeats. If cellular factors are needed to induce tau filament formation, they may not be present in sufficient concentrations in mouse brain. Finally, differences may be a function of the very different life spans of mouse and human.

Outlook

The discovery of missense mutations in tau in FTDP-17 has demonstrated that tau dysfunction produces neurodegeneration. The existence of mutations in the intron following exon 10 of the tau gene has shown that the simple overproduction of four-repeat tau is sufficient to lead to a filamentous pathology and to produce a dementia disorder. This knowledge will be invaluable for future efforts aimed at producing mouse lines transgenic for tau. Higher expression levels of human transgenic tau than have been achieved so far may be the key to success. Animal models of FTDP-17, Pick's disease, progressive supranuclear palsy, and corticobasal degeneration are eagerly awaited. Perhaps most importantly, there will be no true transgenic mouse model of Alzheimer's disease without a filamentous tau pathology.

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