

Molecular Pathogenesis of Genetic and Inherited Diseases

Induction of Inflammatory Mediators and Microglial Activation in Mice Transgenic for Mutant Human P301S Tau Protein

Arianna Bellucci,^{*,†} Andrew J. Westwood,^{*}
Esther Ingram,^{*} Fiorella Casamenti,[†]
Michel Goedert,[‡] and Maria Grazia Spillantini^{*}

From the Department of Clinical Neurosciences,^{*} Brain Repair Centre, University of Cambridge, Cambridge, United Kingdom; the Medical Research Council Laboratory of Molecular Biology,[†] Cambridge, United Kingdom; and the Department of Preclinical and Clinical Pharmacology,[‡] University of Florence, Florence, Italy

Mice transgenic for human P301S tau protein exhibit many characteristics of the human tauopathies, including the formation of abundant filaments made of hyperphosphorylated tau protein and neurodegeneration leading to nerve cell loss. At 5 months of age, the pathological changes are most marked in brainstem and spinal cord. Here we show that these changes are accompanied by marked neuroinflammation. Many tau-positive nerve cells in brainstem and spinal cord were strongly immunoreactive for interleukin-1 β and cyclooxygenase-2, indicating induction and overproduction of proinflammatory cytokines and enzymes. In parallel, numerous activated microglial cells were present throughout brain and spinal cord of transgenic mice, where they concentrated around tau-positive nerve cells. These findings suggest that inflammation may play a significant role in the events leading to neurodegeneration in the tauopathies and that anti-inflammatory compounds may have therapeutic potential. (*Am J Pathol* 2004, 165:1643–1652)

The most common neurodegenerative diseases are characterized by the presence of abnormal filamentous protein inclusions in nerve cells of the brain.¹ In Alzheimer's disease (AD), these inclusions are made of hyperphosphorylated tau protein. Together with the extracellular β -amyloid deposits, they constitute the defining neuropathological characteristics of AD. Tau inclusions, in the absence of extracellular deposits, are characteristic of progressive supranuclear palsy, corticobasal degenera-

tion, Pick's disease, argyrophilic grain disease, and frontotemporal dementia and parkinsonism linked to chromosome 17 (FTDP-17).² The identification of mutations in *Tau* in FTDP-17 has established that dysfunction of tau protein is central to the neurodegenerative process.^{3–5}

At an experimental level, the expression of mutant tau in nerve cells is leading to improved models of neurodegeneration.^{6–12} We have described a line of mice transgenic for human P301S tau,¹⁰ a FTDP-17 mutation with strong functional effects and an early age of disease onset.^{13,14} At 5 to 6 months of age, homozygous animals from this line develop a motor phenotype dominated by a severe paraparesis. Brains and spinal cords of these animals contain abundant neuronal inclusions made of hyperphosphorylated mutant human tau protein in a filamentous state. In the spinal cord, this is accompanied by a large reduction in the number of motor neurons and a marked reactive astrocytosis.

In recent years, much work has been devoted to the study of inflammatory processes in neurodegenerative diseases, especially AD.¹⁵ Neuropathological analysis has revealed many of the hallmarks of inflammation, including microglial cell activation, expression of cytokines and complement, and invasion of immune cells. There has been debate about the relevance of anti-inflammatory drugs for the treatment of AD.¹⁶ Comparatively little is known about inflammatory processes in diseases with only tau deposits; no information is available about inflammation in experimental animal models of the tauopathies. Here we report that brains and spinal cords from 5-month-old human P301S tau transgenic mice exhibit a strong inflammatory reaction, characterized by interleukin-1 β (IL-1 β) production and cyclooxygenase-2 (COX-2) induction, as well as by microglial cell activation.

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Address reprint requests to M.G. Spillantini, Brain Repair Centre, University of Cambridge, Robinson Way, Cambridge CB2 2PY, UK. E-mail: mgs11@cam.ac.uk.

Materials and Methods

Animals and Antibodies

Five-month-old homozygous mice transgenic for human P301S tau¹⁰ and age-matched C57BL/6J controls were used. In some experiments, 14-month-old mice transgenic for wild-type human tau (line ALZ17)¹⁷ and age-matched controls were used. Polyclonal antibodies specific for IL-1 β and COX-2 (Santa Cruz Biotechnology, Santa Cruz, CA) were used at 1:200. A monoclonal anti-actin antibody (Sigma-Aldrich, Poole, UK) was used at 1:1000. Two phosphorylation-dependent anti-tau antibodies were used. AT8 (1:750, Innogenetics, Ghent, Belgium) recognizes human and murine tau phosphorylated at S202 and T205 and AP422 (1:500) recognizes human and murine tau phosphorylated at S422.^{18,19} Rat monoclonal antibody OX-42 (1:250; Serotec, Oxford, UK), which binds CD11b receptors, was used to identify microglial cells. Activated microglial cells were detected using monoclonal antibodies OX-6 and M5/114.15.2 (1:200; PharMingen, San Diego, CA), which are directed against major histocompatibility complex class II antigens.

Immunohistochemistry

Mice were perfused transcardially with 4% paraformaldehyde in 0.1 mol/L of phosphate buffer, pH 7.2. Brains and spinal cords were removed, postfixed for 2 hours, followed by cryoprotection in 30% sucrose in 0.1 mol/L of phosphate buffer, pH 7.4, for 24 hours. Sagittal brain sections and transverse spinal cord sections (30 μ m) were cut on a Leica SM2400 microtome (Leica Microsystems, Bucks, UK), placed in Tris-buffered saline (TBS) containing 0.1% sodium azide, and stored at 4°C. For single-labeling, sections were incubated for 20 minutes at room temperature in 0.1 mol/L of phosphate-buffered saline (PBS), pH 7.4, containing 0.3% Triton X-100, 20% methanol, and 1.5% H₂O₂, followed by a 30-minute incubation in blocking solution (0.1 mol/L PBS, containing 0.3% Triton X-100, 3% goat serum, and 2 g/L bovine serum albumin). This was followed by the overnight incubation at 4°C in primary antibody in blocking solution. After washing, the sections were incubated in biotinylated secondary antibody (1:1000 in blocking solution, Vectastain; Vector Laboratories, Burlingame, CA). The reaction product was visualized using avidin-biotin and 3',3'-diaminobenzidine as the chromogen (DAB kit, Vector Laboratories). For double labeling, sections underwent a further cycle of staining and the reaction product was visualized with the Vector NovaRED kit. In control experiments, 20 μ g/ml of recombinant IL-1 β (Upstate Biotechnology, Lake Placid, NY) was incubated overnight at 4°C with the IL-1 β antibody (1:200) before the addition to tissue sections. In additional control experiments, filamentous, sarkosyl-insoluble tau was extracted from human P301S tau transgenic mouse brain¹⁰ and the tau filaments solubilized as described.²⁰ The dialyzed material (50 μ l) was incubated overnight at 4°C with the IL-1 β antibody (1:200) or anti-tau antibody AT8 (1:1000), before the addition to tissue sections.

Immunofluorescence

For co-localization experiments, sections were incubated for 30 minutes at room temperature with blocking reagent and overnight at 4°C with the primary antibody in blocking solution. After washing, the sections were incubated for 1 hour in the dark with the appropriate fluorescent secondary antibody (Green Alexa, 1:200 in blocking solution; Molecular Probes, Leiden, The Netherlands). Sections were then incubated in the dark with the second primary antibody. After washing, they were incubated in the dark with the second fluorescent secondary antibody (Red Alexa). After washing, the sections were mounted onto slides using Vectashield (Vector Laboratories).

Immunoblot Analysis

Brainstem and spinal cord (10 mg each) from human P301S tau mice and age-matched controls were homogenized in 0.5 ml of cold RIPA buffer (0.1 mol/L PBS, 1% Nonidet P-40, 0.5% sodium deoxycholate, 0.1% sodium dodecyl sulfate, 1 mmol/L phenylmethyl sulfonyl fluoride) containing the complete protease inhibitor cocktail (Roche, Basel, Switzerland). The homogenates were spun for 10 minutes at 10,000 $\times g$ and the supernatants used for immunoblotting. Protein concentrations were determined using the Bio-Rad Protein Assay (Hercules, CA) and 25 μ g of protein run on 12% sodium dodecyl sulfate-polyacrylamide gel electrophoresis. Gels were blotted overnight at 4°C onto a polyvinylidene difluoride membrane (Pierce, Rockford, IL). Membranes were blocked for 2 hours at 25°C in TBS containing 5% milk and incubated overnight at 4°C with anti-IL-1 β or anti-COX-2 antibodies (1:200) in blocking buffer. Membranes were washed in TBS and incubated for 1 hour at 25°C with peroxidase-conjugated anti-rabbit antibody (DakoCytomation, Glostrup, Denmark) (1:2000) in blocking buffer. After washing in TBS, the membranes were incubated for 1 hour at 25°C with a monoclonal anti-actin antibody (1:1000) in TBS. This was followed by washing and a 1-hour incubation at 25°C with peroxidase-conjugated anti-mouse antibody (1:2000, DakoCytomation) in blocking buffer. Blots were developed using enhanced chemiluminescence (Amersham Biosciences, Arlington Heights, IL), followed by the scanning of the bands.

Results

Staining and Immunoblotting for IL-1 β and COX-2

Numerous IL-1 β -positive cells with a neuronal morphology were present throughout the central nervous system of 5-month-old human P301S tau transgenic mice. They were particularly abundant in brainstem and spinal cord (Figure 1, A and C). Nerve cell processes were also stained, particularly in spinal cord, where the majority of proximal dendrites was immunoreactive. In age-matched control mice, only a few nerve cells were weakly IL-1 β -immunoreactive (Figure 1, B and D). In transgenic mice,

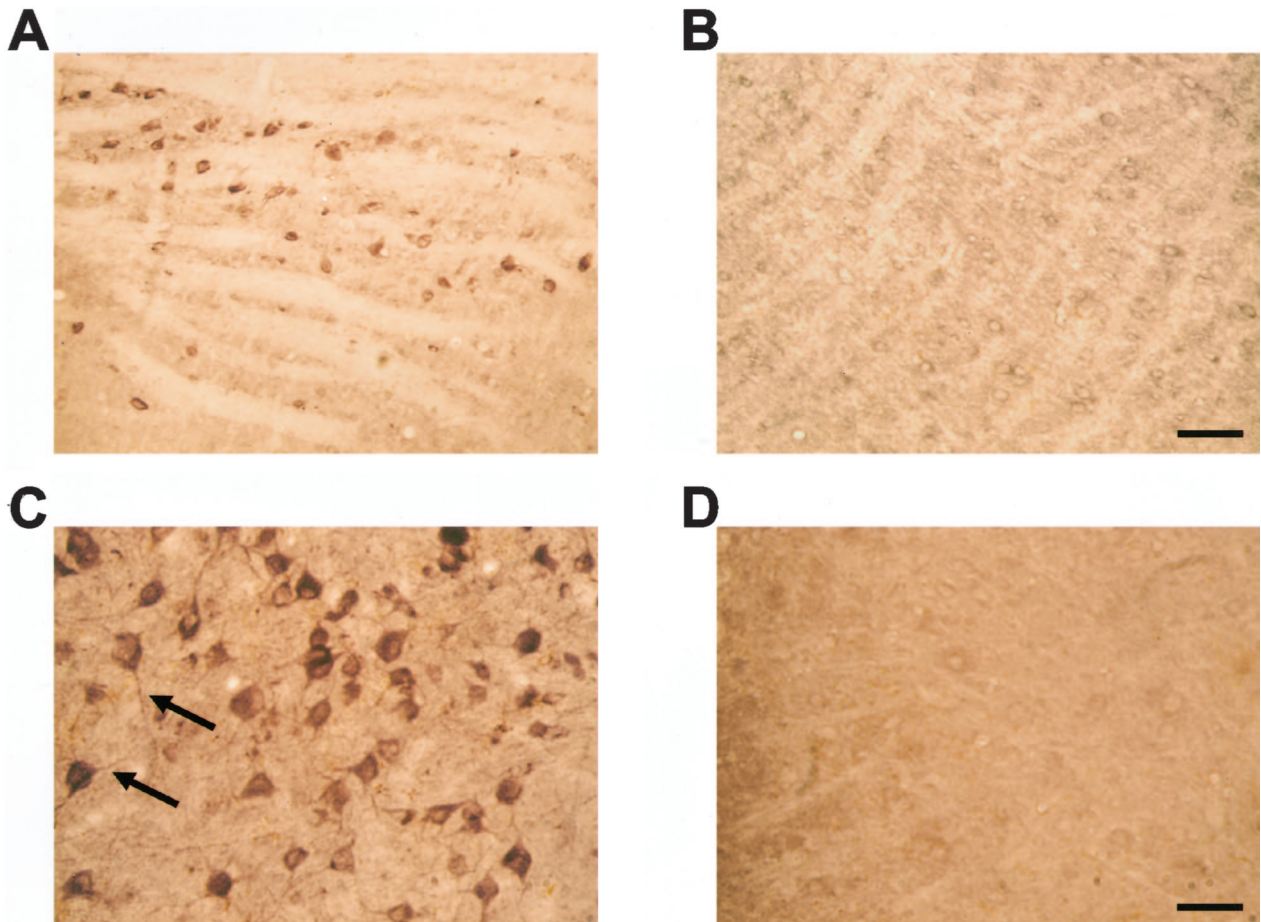


Figure 1. IL-1 β immunoreactivity in brainstem and spinal cord of human P301S tau transgenic mice and age-matched controls. **A** and **B:** Brainstem of 5-month-old transgenic (**A**) and control (**B**) mice. **C** and **D:** Spinal cords of 5-month-old transgenic (**C**) and control (**D**) mice. Note the strong staining of some nerve cell bodies and processes (arrows in **C**). Scale bars: 150 μ m (**B**, also representative for **A**); 71 μ m (**D**, also representative for **C**).

numerous COX-2-immunoreactive nerve cell bodies and processes were present in cerebral cortex, hippocampus, brainstem, cerebellum, and spinal cord (Figure 2; A, C to F). In control mice, a proportion of nerve cells was positive for COX-2 (Figure 2B). Double-labeling immunofluorescence showed co-localization of IL-1 β and COX-2 with staining for the phosphorylation-dependent anti-tau antibody AT8 (Figures 3 and 4). IL-1 β staining was completely abolished after preincubation of the primary antibody with recombinant IL-1 β . It was not changed after preincubation of the IL-1 β antibody with hyperphosphorylated tau protein extracted from the brains of human P301S tau transgenic mice. Preincubation with the latter totally abolished staining with anti-tau antibody AT8. Microglial cells were not immunoreactive for IL-1 β or COX-2. By immunoblotting, the levels of proIL-1 β , IL-1 β and COX-2 were markedly increased in brainstem and spinal cord from human P301S tau transgenic mice, when compared with age-matched controls (Figure 5). Immunoblotting with an anti-actin antibody was used to ensure equal loading.

Staining for Microglial Cells

In transgenic human P301S tau mice, numerous strongly OX-42-immunoreactive microglial cells were present,

particularly in close proximity to AP422-positive nerve cells in brainstem, cerebral cortex, hippocampus, and spinal cord (Figure 6; A, C, E). Some had the morphology of activated cells, in that they were characterized by a bushy appearance with swollen cell bodies and intensely stained, thickened, and branched processes. In the remainder of the brain, OX-42-immunoreactive cells were characterized by a more ramified morphology, even when showing features of activation. In brain and spinal cord from control mice, microglial cells were only weakly OX-42-immunoreactive and displayed a sessile morphology, with thin, finely branching processes extending radially from small, oblong cell bodies (Figure 6, B and D). In transgenic mice, double-labeling immunohistochemistry with OX-42 and IL-1 β or COX-2 antibodies gave the same staining pattern as that obtained using OX-42 and AP422 antibodies, with OX-42-positive, activated microglial cells concentrated around anti-IL-1 β and COX-2-positive cells (not shown).

Major histocompatibility complex class II-positive microglial cells were numerous in the spinal cord of human P301S tau mice, as evidenced by staining with antibodies OX-6 and M5/114.15.2 (Figures 6F and 7A). Double-labeling immunohistochemistry showed the presence of OX-6-positive microglial cells around AP422-positive nerve cells (Figure 6F). The microglia had the morphol-

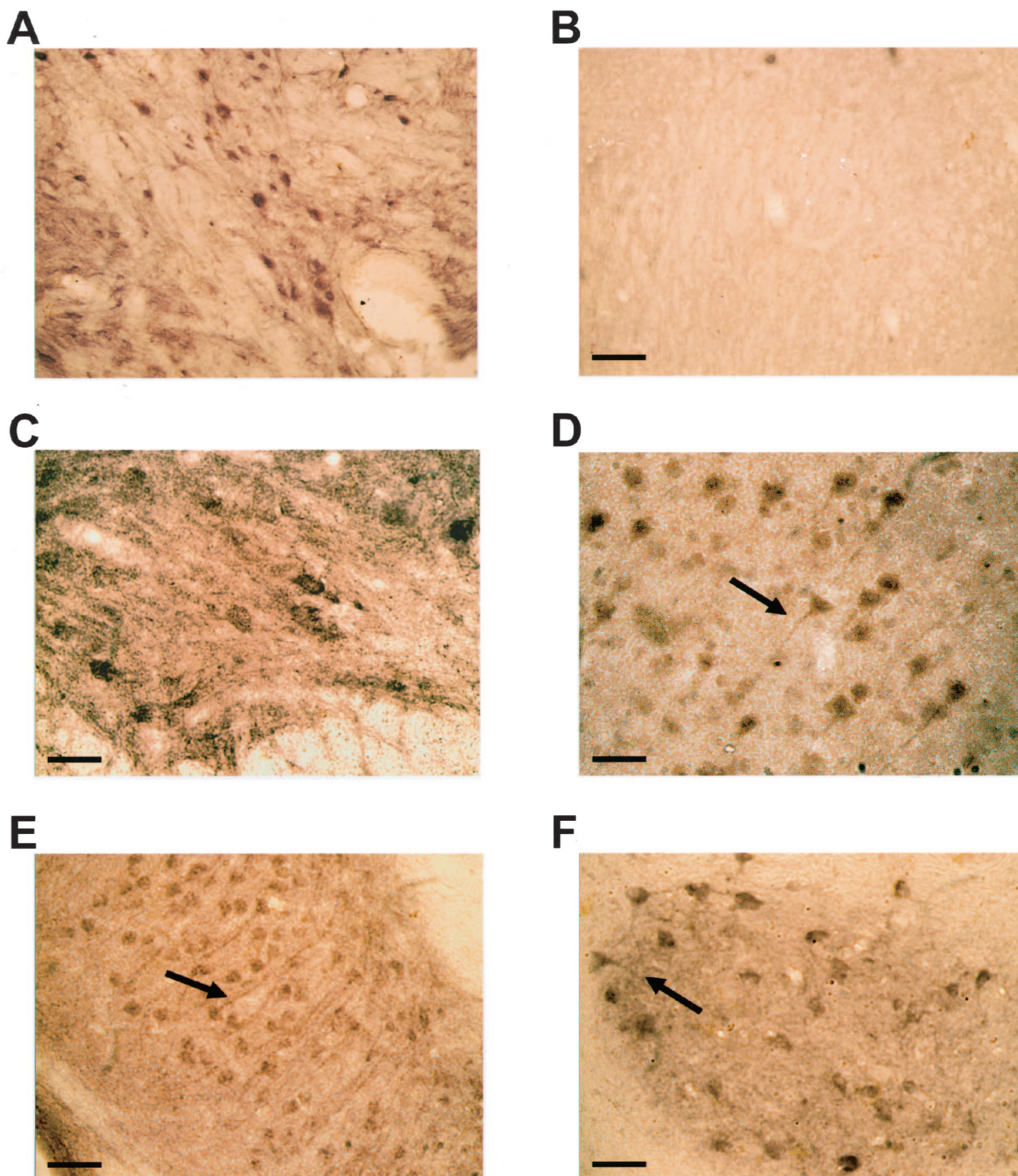


Figure 2. COX-2 immunoreactivity in brain and spinal cord of human P301S tau transgenic mice. Strongly stained nerve cells and processes (**arrows**) in brainstem (**A**), spinal cord (**C**), cerebral cortex (**D**), hippocampus (**E**), and cerebellum (**F**) of 5-month-old human P301S tau transgenic mice. **B:** COX-2 immunoreactivity in brainstem of an age-matched control mouse. Note the strong staining of some nerve cell bodies and processes in **A**, **C**, **D–F**. Scale bars: 150 μ m (**B**, also representative for **A**); 85 μ m (**C**); 80 μ m (**D**); 170 μ m (**E**); 70 μ m (**F**).

ogy of activated cells, as evidenced by an amoeboid, bushy appearance with large, round cell bodies and short, thick processes. In some instances, several microglial cells clustered around a single tau-positive cell. Mice transgenic for wild-type human tau from the ALZ17 line and control mice showed no specific immunoreactivity for OX-6 (not shown) or M5/114.15.2 (Figure 7; B to D).

Discussion

The present findings demonstrate that the intraneuronal accumulation of human P301S tau protein is associated with the activation of inflammatory processes. The latter were characterized by increased IL-1 β and COX-2 staining of nerve cells and the activation of microglial cells. In

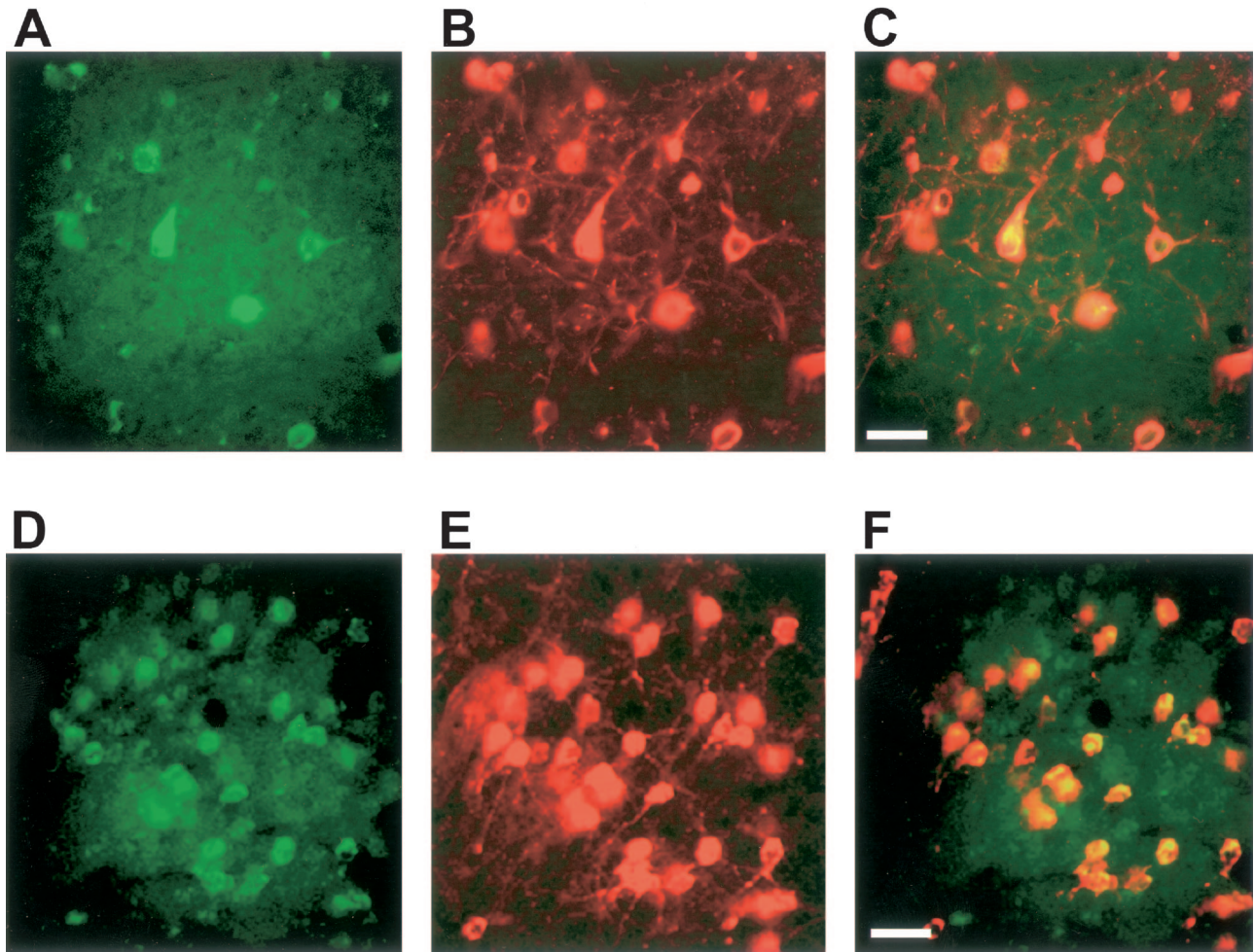


Figure 3. Double-labeling immunofluorescence staining for IL-1 β (green) and tau phosphorylated at S202 and T205 (antibody AT8) (red) in brainstem and spinal cord of human P301S tau transgenic mice. **A** and **B**: Brainstem stained for IL-1 β (**A**) and phosphorylated tau (**B**). **C**: Merged images shown in **A** and **B**. **D** and **E**: Spinal cord stained for IL-1 β (**D**) and phosphorylated tau (**E**). **F**: Merged images shown in **D** and **E**. Co-localization is indicated by the yellow color. Scale bars: 60 μ m (**C**, also representative for **A**, **B**); 125 μ m (**F**, also representative for **D**, **E**).

age-matched control brains, only a small number of nerve cells was weakly immunoreactive for IL-1 β and COX-2, in agreement with previous findings.^{21,22} By immunoblotting, markedly increased levels of IL-1 β and COX-2 were present in brainstem and spinal cord from human P301S tau transgenic mice. Cells strongly immunoreactive for IL-1 β and COX-2 were also stained by the phosphorylation-dependent anti-tau antibodies AT8 and AP422, which recognize both soluble and filamentous tau proteins. This indicates that the accumulation of abnormal mutant human tau protein triggers the production of proinflammatory cytokines and enzymes in nerve cells.

These findings are reminiscent of work on transgenic animal models of amyotrophic lateral sclerosis (ALS) resulting from the overexpression of mutant superoxide dismutase, in which IL-1 β and COX-2 were expressed at abnormally high levels in motor neurons.^{23,24} It suggests that the induction of proinflammatory cytokines and enzymes may be a general response of neurons to the intracellular accumulation of aggregation-prone proteins.²⁵ IL-1 β levels are elevated in AD brain²⁶ and COX-2 expression is increased in tangle-bearing nerve

cells.^{27,28} In addition, COX-2 protein expression is induced in nerve cells after ischemia, with COX-2 selective inhibitors preventing ischemia-induced delayed nerve cell death.²⁹ Cytokines, such as IL-1 β , are known to induce COX-2 expression in nerve cells,^{30,31} suggesting that IL-1 β expression may have triggered COX-2 induction in the human P301S tau transgenic mice. It remains to be determined whether IL-1 β was produced by neuronal or glial cells.

COX-2-derived prostanoids may play a role in the cascade of events that leads to the death of motor neurons and other nerve cells in the human P301S tau transgenic mice. Neuronal overexpression of COX-2 has been shown to lead to nerve cell death and signs of cognitive impairment.³² Conversely, inhibition of COX-2 in mice transgenic for mutant human superoxide dismutase delayed neurodegeneration and the appearance of clinical symptoms,^{33,34} possibly through a reduction of glutamate release.³⁵ In the light of the present findings, inhibition of COX-2 may be a potential therapeutic target for the human tauopathies.

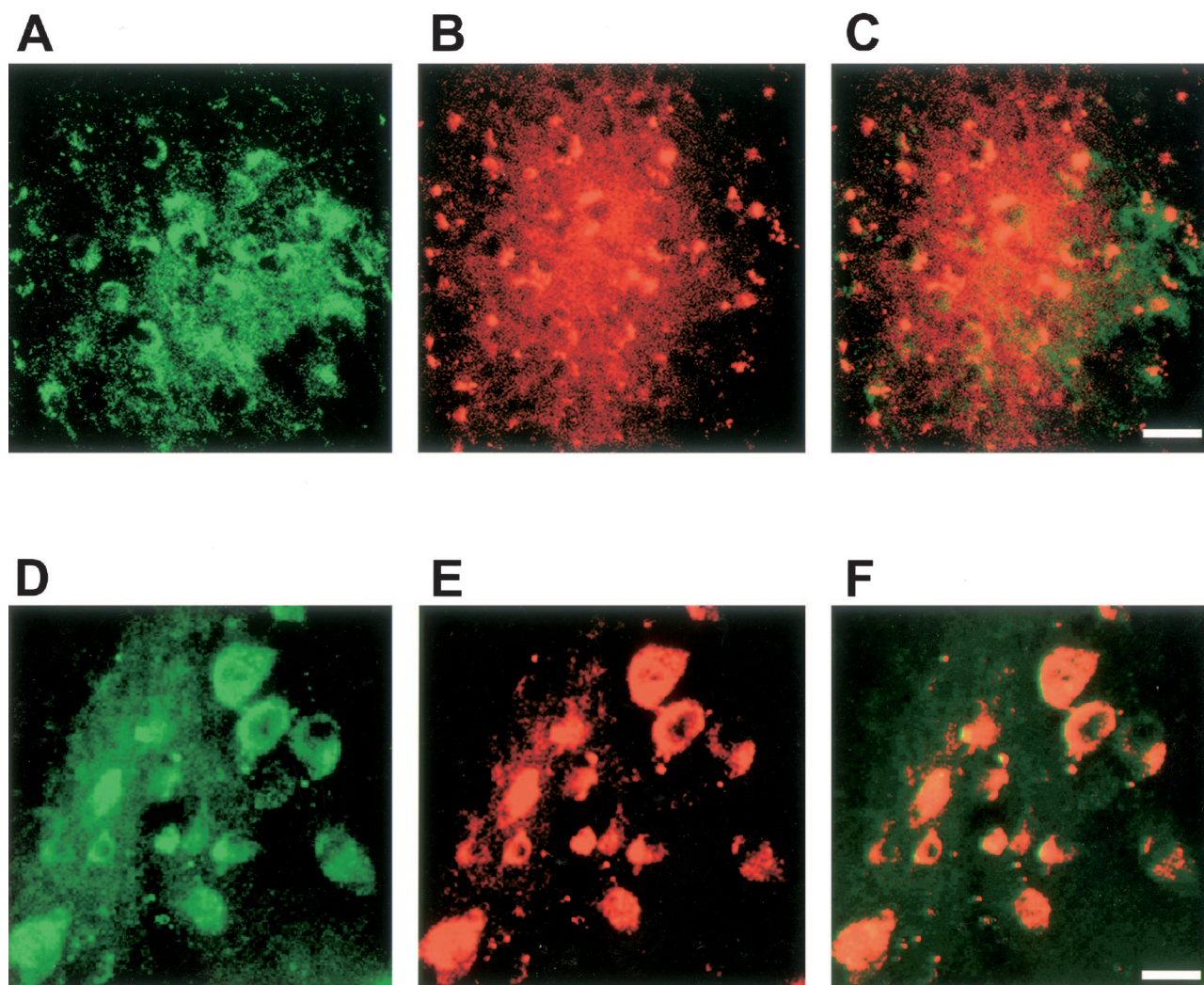


Figure 4. Double-labeling immunofluorescence staining for COX-2 (green) and tau phosphorylated at S202 and T205 (antibody AT8) (red) in brainstem and spinal cord of human P301S tau transgenic mice. **A** and **B**: Brainstem stained for COX-2 (**A**) and phosphorylated tau (**B**). **C**: Merged images shown in **A** and **B**. **D** and **E**: Spinal cord stained for COX-2 (**D**) and phosphorylated tau (**E**). **F**: Merged images shown in **D** and **E**. Co-localization is indicated by the yellow color. Scale bars: 45 μ m (**C**, also representative for **A**, **B**); 40 μ m (**F**, also representative for **D**, **E**).

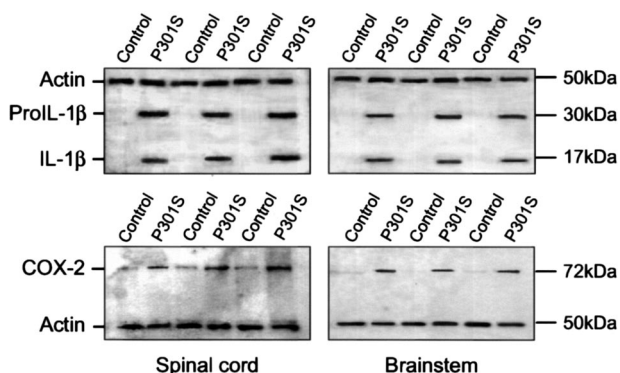


Figure 5. Immunoblotting for IL-1 β and COX-2 of spinal cord (**left**) and brainstem (**right**) from three human P301S tau transgenic mice and three age-matched controls. Immunoblotting for actin was used to ensure equal loading. Note the increased levels of proIL-1 β , IL-1 β and COX-2 in tissues from the transgenic mice.

Brain and spinal cord of 5-month-old human P301S tau transgenic mice exhibited large numbers of OX42-immunoreactive microglial cells. They were much larger in number and more strongly immunoreactive for CD11b receptors than microglia from age-matched controls. In tissues from transgenic mice, a proportion of OX42-positive cells had the morphological appearance of activated microglia. These cells, which were concentrated around nerve cells immunoreactive for phosphorylated tau, IL-1 β , and COX-2, were immunoreactive for major histocompatibility complex class II antigens. Activated microglial cells were not present in the spinal cord of mice from line ALZ17, which accumulate wild-type human tau in nerve cells, but do not develop filamentous tau deposits or nerve cell death.¹⁷ They express a four-repeat human brain tau isoform (441 amino acids) under the control of the murine Thy1 promoter. Expression levels of human tau in brainstem and spinal cord are very similar between line ALZ17 and the line transgenic for human P301S tau

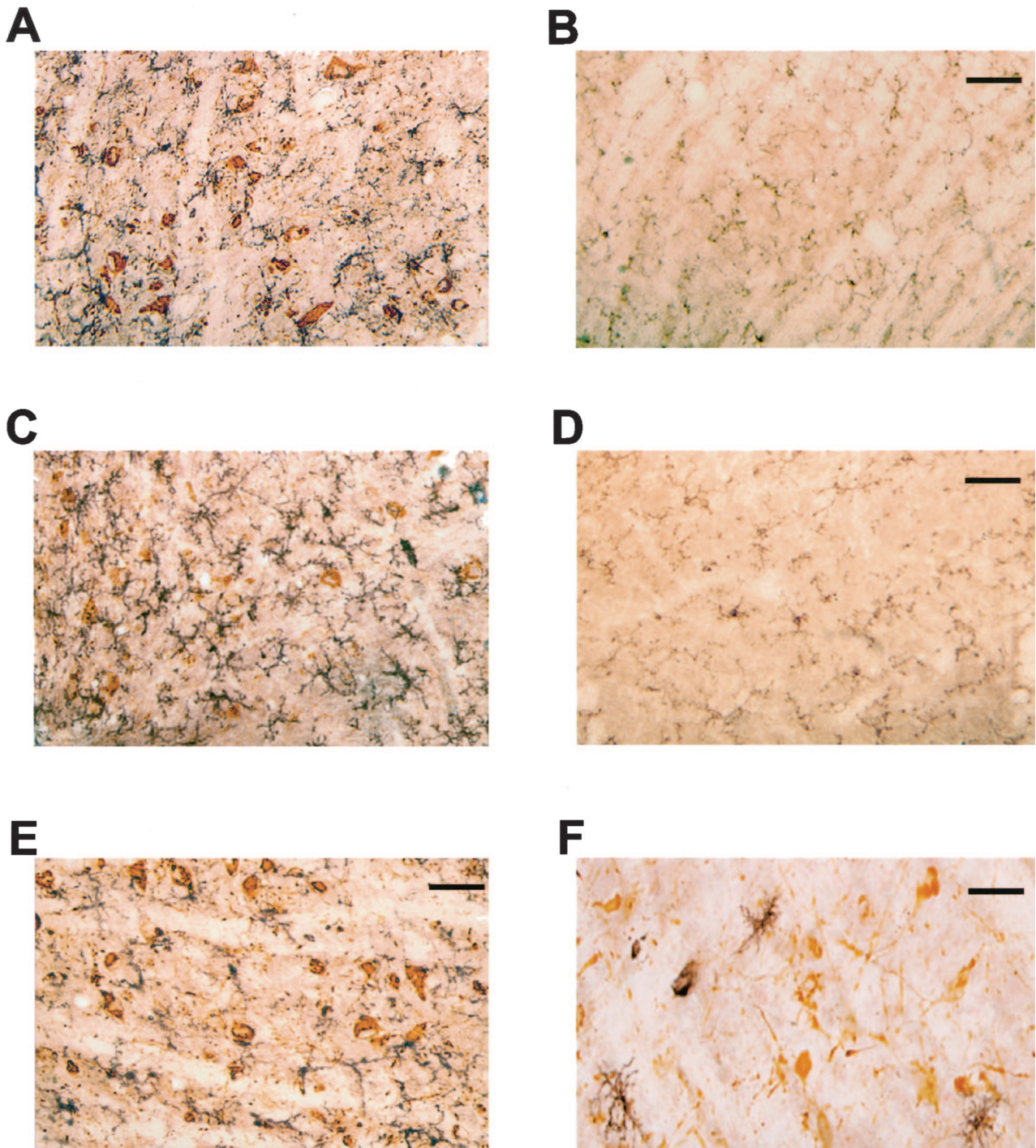


Figure 6. Double-labeling immunostaining for OX-42 or OX-6 (brown) and tau phosphorylated at S422 (antibody AP422) (red) in brainstem and spinal cord of human P301S tau transgenic mice and age-matched controls. **A, B, and E:** Brainstem of transgenic (**A, E**) and control (**B**) mice. **C, D, and F:** Spinal cord of transgenic (**C, F**) and control mice (**D**). Sections **A** to **E** were stained for OX-42 and phosphorylated tau. Section **F** was stained for OX-6 and phosphorylated tau. Scale bars: 70 μm (**B**, also representative for **A**); 90 μm (**D**, also representative for **C**); 60 μm (**E, F**).

protein.^{10,17} It follows that activation of microglial cells in the human P301S tau mice was linked to tau filament formation and neurodegeneration and was not the mere consequence of the overexpression of human tau.

These findings mirror what has been described in AD,^{15,36,37} in other tauopathies,^{38,39} and in diseases with intracellular filamentous deposits, such as Parkinson's

disease,⁴⁰ dementia with Lewy bodies,⁴¹ multiple system atrophy,⁴² and ALS.⁴³ The lesions of AD display evidence of localized inflammation, with reactive microglia being present in neuritic plaques and in and around neurofibrillary lesions. Many of the latter are extracellular tangles,^{44,45} suggesting that microglia may be activated by extracellular tau deposits and involved in their attempted

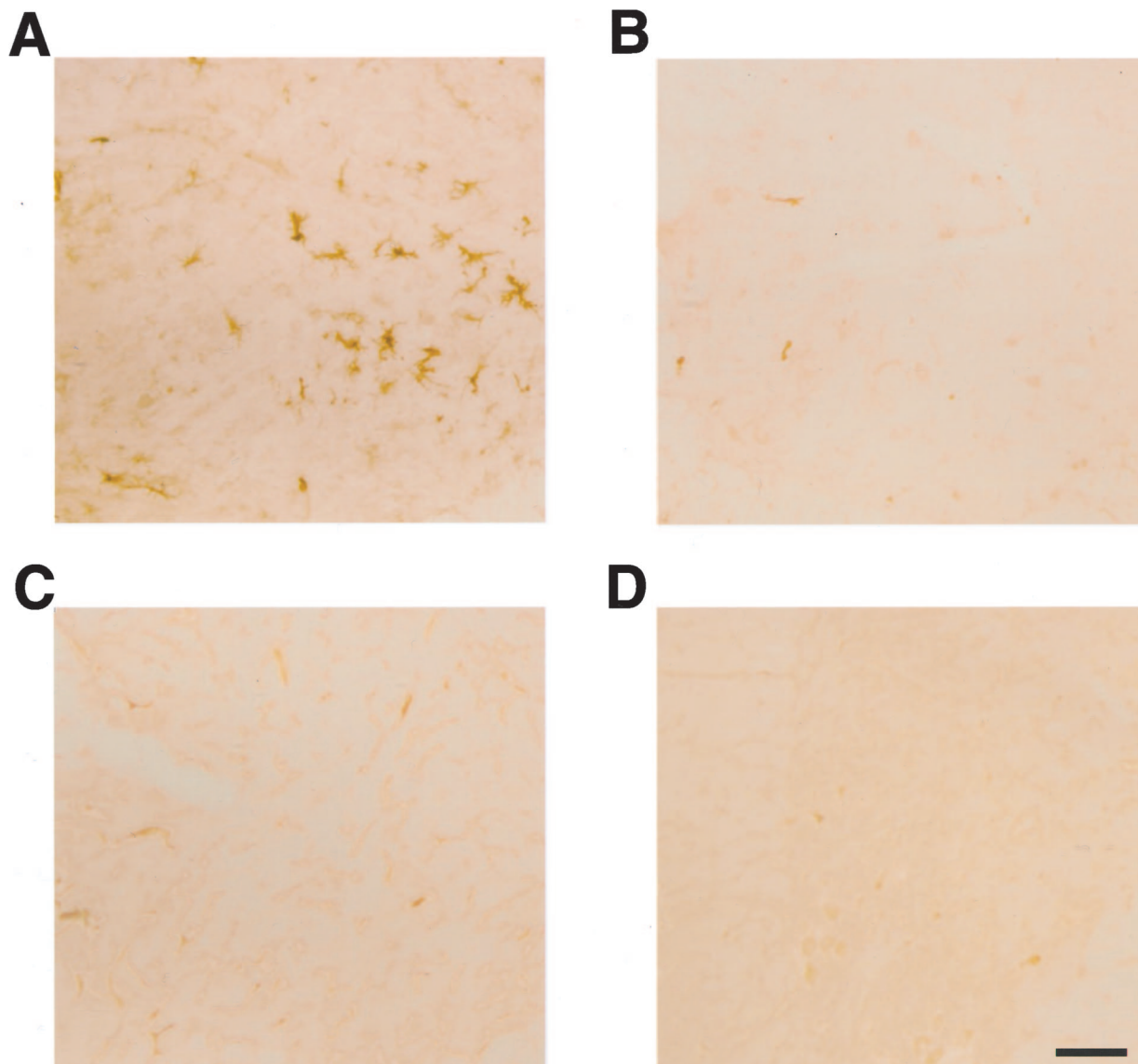


Figure 7. Major histocompatibility class II-immunoreactivity (antibody M5/114.15.2) in spinal cord of mice transgenic for human wild-type tau, human P301S tau, and age-matched controls. **A** and **B**: Five-month-old mouse transgenic for human P301S tau (**A**) and age-matched control (**B**). **C** and **D**: Fourteen-month-old mouse transgenic for human wild-type tau (**C**) and age-matched control (**D**). Scale bars: 50 μ m (**D**, also representative for **A–C**). Note the staining of cells with the morphology of activated microglial cells in **A**.

removal. However, activated microglial cells are also present in human tauopathies that lack extensive extracellular tangles, such as progressive supranuclear palsy and corticobasal degeneration.³⁹ The same is also true of the human P301S tau transgenic mice, in which nerve cells degenerate without tau deposits lingering on in the extracellular space.¹⁰

Microglia have long been known to migrate to regions of nerve cell damage, where they contribute to the elimination of dead cells through their marked phagocytic activity.⁴⁶ More recent work has shown that they may also cause the death of nerve cells. Thus, in the embryonic chicken retina, death of nerve cells expressing the p75 nerve growth factor receptor results from exposure to nerve growth factor released by microglia.⁴⁷ Furthermore, the generation of superoxide ions and tumor necrosis factor by microglial cells has been shown to result

in the death of Purkinje cells and motor neurons, respectively.^{48,49} In the nematode *Caenorhabditis elegans*, genetic analysis has established that signals released by phagocytic cells actively promote programmed nerve cell death.^{50,51}

Inflammation is not the root cause of disease in the human P301S tau transgenic mice. However, it may well play an important role in the series of events leading to neurodegeneration. It will be interesting to determine whether anti-inflammatory compounds can have a beneficial effect on the neurodegenerative process.

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