

Intrinsic Resistance of Oligodendrocytes to Prion Infection

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Within the CNS, the normal form of cellular prion protein (PrP^C) is expressed on neurons, oligodendrocytes, and astrocytes. The contribution of these cell types to prion replication and pathogenesis is unclear. To assess the role of oligodendrocytes, we expressed PrP^C under the control of the myelin basic protein (MBP) promoter in mice lacking endogenous PrP^C. PrP^C was detected in oligodendrocytes and Schwann cells but not in neurons and astrocytes. MBP–PrP mice never developed scrapie after intracerebral, intraperitoneal, or intraocular challenge with scrapie prions. Transgenic brains did not contain protease-resistant prion protein and did not transmit scrapie when inoculated into PrP^C-overexpressing indicator mice. To investigate whether prion spread within the CNS depends on oligodendrocytic PrP^C, we implanted PrP^C-overexpressing neuroectodermal grafts into MBP–PrP brains. After intraocular prion inoculation, none of the grafts showed spongiform encephalopathy or prion infectivity. Hence oligodendrocytes do not support cell-autonomous prion replication, establishment of subclinical disease, and neural spread of prions. Prion resistance sets oligodendrocytes aside from both neurons and astrocytes.

Key words: prion; PrP^{Sc}; oligodendrocytes; myelin basic protein; transgenic mice; scrapie

Introduction

Prion protein (PrP) plays a central role in the pathogenesis of transmissible spongiform encephalopathies (TSEs) such as sheep scrapie, bovine spongiform encephalopathy, and Creutzfeldt–Jakob disease (Aguzzi and Weissmann, 1997; Aguzzi et al., 2001). The normal form of PrP, designated PrP^C, is encoded by the *Prnp* gene (Basler et al., 1986), and is almost ubiquitously expressed. In the CNS, PrP^C is predominantly neuronal, but expression on astrocytes and oligodendrocytes is also detectable (Moser et al., 1995). *Prnp*^{0/0} mice are devoid of PrP^C and develop and behave normally (Büeler et al., 1992) but are resistant to prion disease (Büeler et al., 1993) and do not support replication of prions (Sailer et al., 1994). Reintroduction of PrP transgenes into *Prnp*^{0/0} mice, even when carrying significant amino-proximal deletions, restores susceptibility to scrapie (Fischer et al., 1996; Shmerling et al., 1998; Flechsig et al., 2000).

Previous transgenic studies have revealed that expression of PrP^C on either neurons or astrocytes is sufficient for scrapie replication (Race et al., 1995; Raeber et al., 1997). Surprisingly, mice expressing PrP^C only within astrocytes or only within neurons develop indistinguishable pathologies after prion infection. Although oligodendrocytes express PrP^C and are histogenetically highly related to astrocytes, it is not known whether they play a role in prion replication and in development of brain damage. White-matter involvement in TSEs was proposed to be a result of direct modifications of oligodendrocyte physiology (El Hachimi et al., 1998). If this is the case, oligodendrocytes may contribute to the pathogenetic process of scrapie.

The mechanisms by which prions spread within the CNS are unknown. Although neuronal trans-synaptic spread appears plausible, expression of PrP^C in myelinating cells and the alleged potential of cultured Schwann cell lines to generate prion infectivity during infection (Follet et al., 2002; Archer et al., 2004) raise a possible role for oligodendrocyte-borne PrP^C in the cerebral spread of prions. The latter issue may be of practical interest, because neuroinvasion can occur, for example, via the heavily myelinated visual pathway after conjunctival instillation (Scott et al., 1993), corneal grafts (Duffy et al., 1974), and intraocular injection (Fraser, 1982).

These questions would be testable in animals that express PrP^C exclusively on myelinating cells. Expression would have to be sustained and highly selective, with no other cells expressing PrP^C. Here we describe transgenic mice expressing PrP^C under the control of the myelin basic protein (MBP) promoter. PrP^C was strongly and exclusively expressed in oligodendrocytes and Schwann cells, but not in neurons or astrocytes. Using this model, we have established that expression of PrP^C on myelinating cells does not support prion replication or spread of the agent along

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neural pathways. These results are in contrast to previous evidence that had been gathered with cultured cells and negate any cell-autonomous role for myelinating cell-borne PrP^C in the pathogenesis of prion diseases.

Materials and Methods

DNA constructions and generation and identification of transgenic mice. A blunt-ended 760 bp fragment containing the entire PrP open reading frame (ORF) region (corresponding to exon 3 of the *Prnp* gene) (Fischer et al., 1996) was introduced and ligated into the *EcoRI* restriction sites of the MBP construct (see Fig. 1A). The 1.3 kb promoter region of the mouse MBP gene contains the cap site of MBP mRNA and 5' noncoding sequences. Although the cDNA fragment used is only approximately two-thirds as long as the normal MBP cDNA and does not contain sequences corresponding to the 3' portion of MBP mRNA, it contains the full coding sequence of 14 kDa for the smallest mouse MBP. Poly(A) addition signals are provided from rabbit β -globin and simian virus 40 early genes, and the second intron of the rabbit β -globin gene is placed between the cap site and PrP ORF for splicing.

Plasmid DNA was digested with *HindIII* and *SaII* and microinjected into homozygous *Prnp*^{0/0} zygotes, as described previously (Fischer et al., 1996). Using PCR, founders were identified by the presence of *Prnp*⁰ alleles and *Prnp*⁺ transgenes. The primers P3 (specific for the disrupted *Prnp* allele), P10 (for the wild-type *Prnp* gene), and Mut217 (exon 3 primer) were used under standard PCR conditions (Fischer et al., 1996). Transgenic founders were mated to *Prnp*^{0/0} mice, and one transgenic line, designated tg640^{+/-}, was established from the F₁ progeny on a *Prnp*^{0/0} mixed background, C57BL/6 × 129Sv. Additional breeding yielded the homozygous line tg640^{+/+}, or MBP-PrP.

Northern blot analyses. Mouse-brain total RNA was isolated with RNeasy Midi kits (Qiagen, Hilden, Germany), and samples (15 μ g) were electrophoresed through a 1% agarose gel in 0.02 M sodium borate, pH 8.3, 0.5 mM EDTA, and 5% formaldehyde. RNA was blotted onto Hybond-N⁺ membranes (DuPont, Billerica, MA) using 20× SSC and cross-linked to membranes by ultraviolet radiation. Prehybridization occurred for 4 hr at 42°C in 50% formamide, 1 M NaCl, 1% SDS, and 10% dextran sulfate. Hybridization occurred overnight at 42°C in the same buffer using ³²P-labeled probe A (10⁶ cpm/ml), prepared by random priming, which is known to hybridize to transcripts produced by both wild-type and disrupted *Prnp* genes (Büeler et al., 1992), and 0.1 mg/ml denatured salmon-sperm DNA. Membranes were washed for 10 min at room temperature (RT) in 2× SSC, washed twice for 30 min at 60°C in 2× SSC and 1% SDS, and washed twice for 30 min at RT in 1× SSC. Autoradiography was performed for 7–24 hr at RT using Eastman Kodak (Rochester, NY) XAR-5 film and an intensifier screen.

Western blot analysis. Brain homogenates were adjusted to 5 mg/ml, and 50 μ g of total protein from each sample was electrophoresed through a 12% SDS-PAGE gel. Proteins were transferred to nitrocellulose by semidry blotting. Membranes were blocked with TBS-Tween 20 and 5% nonfat milk, incubated with antibody 6H4 or POM1 (monoclonal antibody to PrP; M. Polymenidou and A. Aguzzi, unpublished data), and developed by enhanced chemiluminescence (Amersham Biosciences, Braunschweig, Germany), as described recently (Prinz et al., 2002). Quantification was performed using NIH Image software (<http://rsb.info.nih.gov/nih-image/>).

In situ hybridization. A 290 bp *Asp718-BstEII* fragment of the mouse PrP cDNA was cloned into pBluescript KS and SK vectors (Stratagene, La Jolla, CA). Digoxigenin-labeled sense and antisense probes were synthesized (Boehringer Mannheim, Mannheim, Germany) from *Asp718*-cleaved KS and SK constructs, respectively, using T3 RNA polymerase (Raeber et al., 1997). Sections were deparaffinized and postfixed in 4% paraformaldehyde in PBS, RNA was denatured for 10 min in 0.1 M HCl, and sections were digested with 10 μ g/ml proteinase K (PK) at 37°C for 10 min. After a second fixation step in 4% paraformaldehyde for 10 min, acetylation was performed in 0.1 M triethanolamine and 0.25% acetic anhydride, and prehybridization was performed in 5× SSC, 50% formamide, 5× Denhardt's solution, and 50 μ g/ml yeast tRNA for 90 min at room temperature. Hybridization was performed in the same mixture, containing the digoxigenin-labeled probe (50 ng/ml; 12 hr; 58°C). Sam-

ples were washed for 1 hr at 65°C in 0.2 and 0.1× SSC, respectively, labeled with anti-digoxigenin Fab fragments (1:5000), and developed with 5-bromo-4-chloro-3-indolyl phosphate and 4-nitroblue tetrazolium chloride. Sections were mounted in glycerol gelatin.

Transmission electron microscopy. White matter was freshly prepared from mouse brains and fixed for 4 hr at 4°C in PBS containing 4% formaldehyde and 0.1% glutaraldehyde. These small tissue fragments were immersed in 2 M sucrose containing 15% polyvinyl pyrrolidone (10 kDa; Sigma, St. Louis, MO), mounted on aluminum pins, frozen, and stored in liquid nitrogen. Grids with ultrathin sections [prepared according to Tokuyasu (1976)] were conditioned with PBS containing 0.5% milk powder and 0.02% Tween 20 for 10 min, incubated with PrP-specific mouse IgG (10 μ g/ml in conditioning buffer; clone 6H4; Prionics AG, Schlieren, Switzerland) for 2 hr at room temperature, rinsed with PBS, and incubated with 8 nm gold-labeled goat anti-mouse IgG (diluted to an absorbance of 0.1 at 525 nm with conditioning buffer) for 1 hr. After rinses with PBS and distilled water, grids were embedded and stained with methylcellulose and uranyl acetate. Micrographs were taken on a Zeiss (Oberkochen, Germany) EM 910 at an original magnification of 20,000×.

Preparation of oligodendrocytes and immunofluorescence. Optic nerve oligodendrocytes were isolated from postnatal day 7–8 wild-type, transgenic, and *Prnp*^{0/0} mice, as described previously for the rat (Schwab and Caroni, 1988). Four days after isolation, cells were fixed with 4% paraformaldehyde and 5% sucrose in PBS for 15 min at RT. The cultures were permeabilized and blocked in PBS supplemented with 0.1% Triton X-100 and 10% FCS and then incubated with anti-myelin-associated glycoprotein (MAG; 1:100; Roche Diagnostics, Rotkreuz, Switzerland) and anti-PrP rabbit antiserum XN (1:800; raised against full-length murine PrP) for 1 hr at RT. After washing, the cells were incubated with goat anti-mouse secondary antibodies conjugated with Alexa 488 (1:150; Molecular Probes, Eugene, OR) and goat anti-rabbit Alexa 546 (1:500). After additional washing, stained cultures were embedded in Mowiol (Calbiochem, La Jolla, CA) and mounted on slides for fluorescence microscopy.

Scrapie infection and determination of infectivity. Mice were inoculated intraperitoneally with 100 μ l of brain homogenate containing 6 logLD₅₀, intracerebrally with 30 μ l (3 × 10⁵ LD₅₀) of the Rocky Mountain Laboratory (RML) scrapie strain (passage 5), or intraocularly with 10 μ l (1 × 10⁵ LD₅₀) of the same RML scrapie strain, prepared as described previously (Büeler et al., 1993). Mice were monitored every second day, and scrapie was diagnosed according to standard clinical criteria. Mice were killed on the day of onset of terminal clinical signs of scrapie. Infectivity of tissues was determined on 1% spleen or brain homogenates. Tissues were homogenized (10% w/v) in 320 mM sucrose with a microhomogenizer, passed several times through 18 and 22 gauge needles, and diluted in PBS with 5% BSA. When the solution appeared homogenous, it was spun for 5 min at 500 × g. Supernatants (30 μ l) were inoculated intracerebrally into groups of four tg20 mice (Fischer et al., 1996). Indicator mice were killed after development of terminal scrapie and the relationship $y = 11.45 - 0.088x$, where y is logLD₅₀/ml homogenate and x is incubation time in days to terminal disease, was used to calculate infectivity titers (Prusiner et al., 1982). The presence of a protease-resistant isoform of PrP (PrP^{Sc}) in the infected brains was investigated on proteinase K-treated (50 μ g/ml; 30 min; 37°C) homogenates by Western blot analysis, as described above.

Histoblots. Histoblots were performed as described previously (Prusiner et al., 1982). Frozen brains that were cut into 12- μ m-thick slices were mounted on nitrocellulose membranes. Total PrP, as well as PrP^{Sc} after digestion with 50 μ g/ml proteinase K for 4 hr at 37°C, were detected with PrP antibody 6H4 (1:2000 in 1% nonfat milk, overnight at 4°C) and alkaline phosphatase immunconjugates. Visualization was performed using 5-bromo-4-chloro-3-indolyl phosphate and 4-nitroblue tetrazolium chloride (Boehringer Mannheim).

Establishment and analysis of intracerebral neuronal grafts. Embryos were obtained at day 12.5 after conception from homozygous matings of tg20 transgenic mice and transferred into modified HBSS that was supplemented with 10% FCS and 2% glucose at 4°C. The neuroectodermal anlage was prepared, homogenized in a volume of 30 μ l, and grafted into the caudoputamen of adult wild-type mice, as described previously

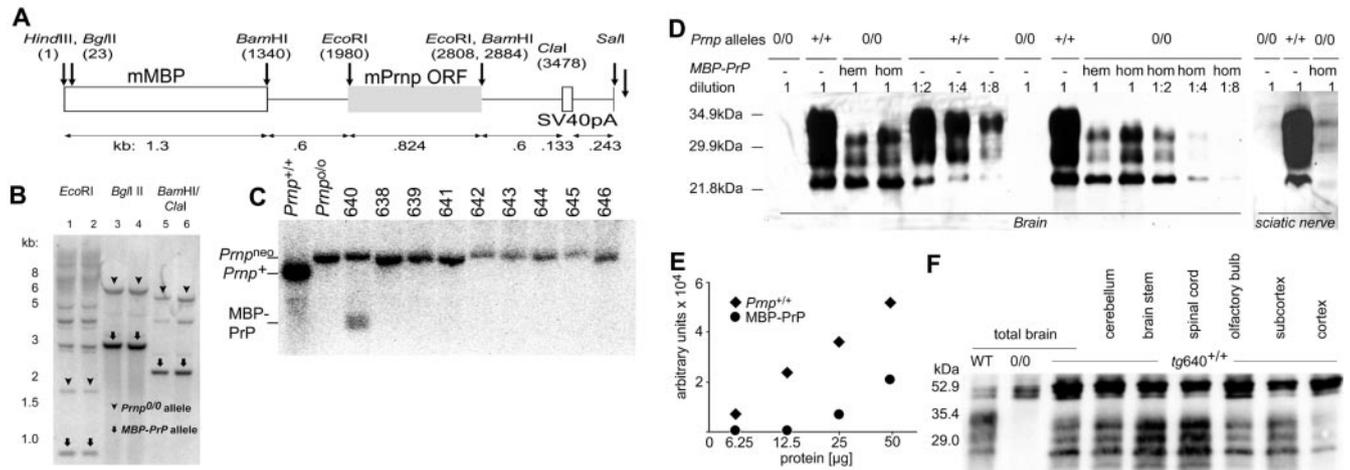


Figure 1. Generation and identification of MBP–PrP transgenic mice. *A*, Schematic drawing of the MBP–PrP transgene used for generation of transgenic mice. Arrows and parenthetical numbers indicate the respective restriction sites. The second intron of rabbit β -globin was placed between the cap site and MBP cDNA to improve transgene expression. mMBP, Mouse MBP; mPrnp, mouse *Prnp*; SV40, simian virus 40. *B*, Genomic Southern blot analysis of MBP–PrP transgenic mice. *EcoRI* exactly excises the PrP ORF, confirming the presence of the transgene. (All tested mice showed the same banding pattern.) The *BglII* and the *BamHI*–*Clal* digests verify the integrity of the 5' and 3' ends, respectively. *C*, Northern blot analysis of brain RNA. The numbers above each lane indicate the various transgenic lines analyzed. RNA from wild-type (*Prnp*^{+/+}) and *Prnp*^{0/0} brains was used as a control. *D*, PrP^C expression in brains and sciatic nerves of hemizygous (hem) or homozygous (hom) *tg640* transgenic mice. Expression levels were estimated by comparison with serial twofold dilutions of wild-type brain homogenate. *E*, Brain PrP^C quantification by chemiluminescence analysis. The abscissa displays the amount of protein in micrograms; the ordinate shows chemiluminescence intensity (arbitrary units). *F*, Anatomical dissection of the CNS in the cerebellum, brainstem, spinal cord, olfactory bulb, and gray or white matter (cortex, subcortex) revealed strongest expression of PrP^C in white-matter or white-matter-rich regions of the CNS. MBP–PrP mice display a dominant unglycosylated PrP^C band, whereas wild-type mice show a dominant diglycosylated band. To facilitate sample comparisons, only 25% of total protein was loaded in wild-type samples, resulting in PrP^C bands of similar intensity. The uppermost bands represent actin. WT, Wild type.

(Brandner et al., 1996a; Isenmann et al., 1996a,b). For histological analysis, mice were killed while under deep anesthesia. For paraffin histology, whole mouse brains were fixed for at least 72 hr in 4% paraformaldehyde and PBS and processed for paraffin embedding. Hematoxylin and eosin staining and immunohistochemistry for GFAP (anti-GFAP antiserum; 1:300; Dako, High Wycombe, UK) and synaptophysin (synaptophysin antiserum; 1:40; Dako) were all performed on sections of paraffin-embedded tissues. Biotinylated secondary antibodies (goat anti-rabbit and rabbit-anti mouse; Dako) were used at a 1:200–1:300 dilution. Visualization was achieved using biotin–avidin–peroxidase (Dako) and diaminobenzidine as a chromogen, according to the protocols suggested by the manufacturer.

Results

MBP–PrP transgenic mice express PrP^C exclusively on myelinating cells

To drive expression of PrP^C in myelinating cells, we prepared a DNA construct in which the entire coding region of the murine *Prnp* gene was placed under the transcriptional control of a murine MBP promoter fragment. A similar strategy has been used for expressing myelin basic protein antisense transcripts to produce a *shiverer*-like hypomyelinating mouse model (Katsuki et al., 1988). The construct, consisting of the 1.3 kb promoter region of the mouse MBP gene and two 0.6 kb rabbit β -globin introns that flank the 0.76 kb PrP open reading frame region (Fig. 1*A*), was excised from its prokaryotic backbone and injected into the pronuclei of *Prnp*^{0/0} zygotes.

Transgenic founders were identified by PCR analysis and Southern blotting (Fig. 1*B*). Nine of 11 PCR-positive founder (F_0) mice transmitted the transgene to their offspring. Transgene-carrying animals were bred to *Prnp*^{0/0} mice. Transgene expression in the resulting mouse colony (mixed C57BL/6 \times 129Sv) was analyzed in organs of 12-week-old F_1 offspring. Unless otherwise specified, all subsequent analyses were performed with homozygous offspring of one line (designated *tg640*) that showed sustained transcription of the transgene (Fig.

1*C*). Northern blot analysis failed to detect expression of the transgene in spleen, kidney, thymus, liver, testes, lung, heart, and muscle (data not shown).

We amplified and sequenced genomic DNA of six randomly chosen mice, as well as RNA from brain samples of four randomly chosen mice, after retrotranscription, with primers encompassing the *Prnp* open reading frame, which is contained in one single exon. We only detected sequences corresponding to the wild-type *Prnp* open reading frame, thereby excluding mutations in the transgene or RNA editing leading to a mutated protein (data not shown).

We then assessed expression of PrP^C by Western blot analysis. Homogenates of whole *tg640* brains, as well as of subregions thereof, yielded a characteristic three-band pattern of unglycosylated, monoglycosylated, and diglycosylated PrP (Fig. 1*D,F*). Western blots of spleen, heart, lung, thymus, muscle, liver, and kidney did not reveal any PrP^C expression (data not shown). *Tg640* mice developed normally and remained healthy, without any abnormal clinical signs, for >670 d.

To quantitate total PrP levels in transgenic brains (Fig. 1*D*), serial twofold dilutions of *tg640* and wild-type brain homogenates in *Prnp*^{0/0} homogenates were blotted, and the relative expression levels were calculated from calibration curves obtained by direct acquisition of chemiluminescence (Fig. 1*E*), as described previously (Heppner et al., 2001). The PrP^C content in *tg640* brains was ~35% of that found in wild-type mice. Immunoblotting of sciatic nerves of MBP–PrP, *Prnp*^{0/0}, and *Prnp*^{+/+} mice (Fig. 1*D*) revealed expression of PrP^C in sciatic nerves of MBP–PrP mice. However, the amount of PrP^C was only 7% of that found in wild-type mice.

We then analyzed the details of intracerebral PrP^C distribution in *tg640* mice (Fig. 1*F*). Western blot analysis of anatomically dissected cortical and subcortical forebrain compartments indicated strong PrP^C signals in the white matter (subcortex), yet

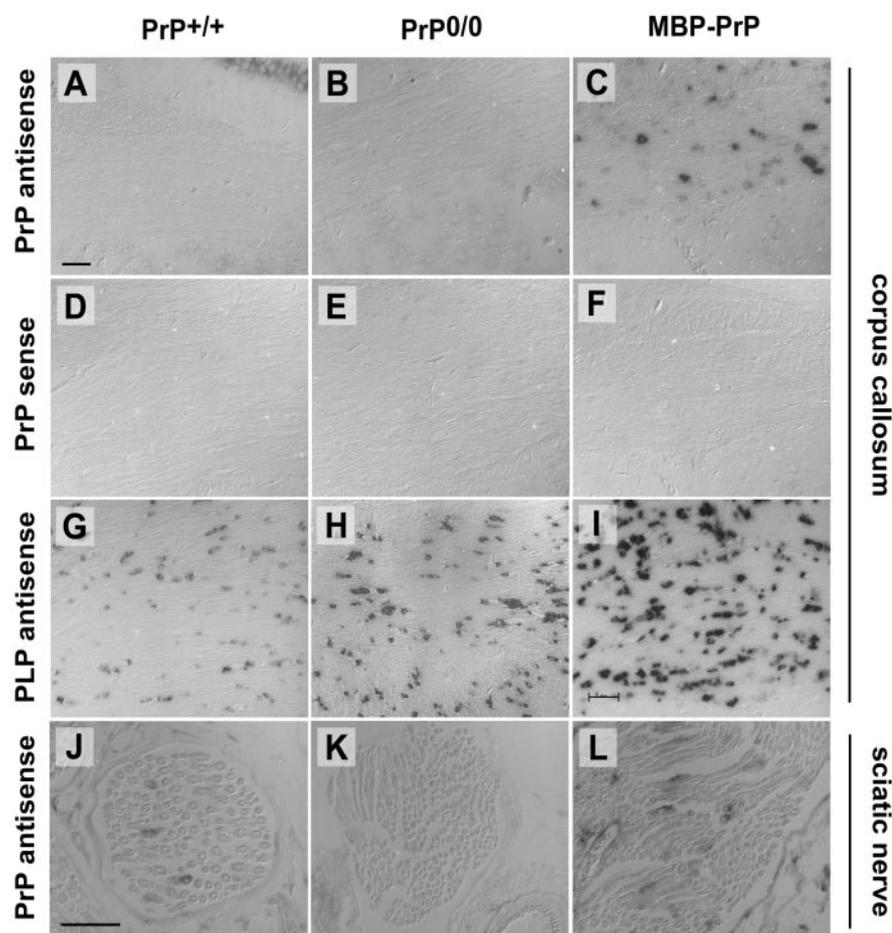


Figure 2. Localization of PrP mRNA in MBP-PrP transgenic mice. *In situ* hybridization of forebrain sections (A–I) with corpus callosum and of sciatic nerve sections (J–L) of adult *Prnp*^{+/+}, *Prnp*^{0/0}, and MBP-PrP transgenic mice is shown. Expression of PrP mRNA occurs in neurons of wild-type (*Prnp*^{+/+}) forebrains (A) and in *Prnp*^{+/+} Schwann cells (J), whereas *Prnp*^{0/0} brains (B) and nerves (K) lack any signal. In contrast, strong PrP signals are visible in callosal oligodendrocytes (C) and sciatic Schwann cells (I–L) of transgenic mice. D–F, Sense probes did not reveal any signals in brains and sciatic nerves. G–I, The oligodendrocyte-specific PLP antisense probe revealed strong mRNA signals in the forebrains of all genotypes investigated. Scale bars, 50 μ m.

no PrP^C was detected in the gray matter (cortex). Instead, in wild-type (*Prnp*^{+/+}) mice, both compartments contained sizeable levels of PrP^C (data not shown). MBP-PrP mice displayed a predominant unglycosylated PrP^C band, whereas wild-type mice show a dominant diglycosylated band, suggesting that post-translational processing of PrP^C differs between oligodendrocytes and other CNS cells.

We performed *in situ* hybridizations on brain and sciatic nerve sections with digoxigenin-labeled riboprobes for PrP mRNA (Fig. 2). In *Prnp*^{+/+} mice, PrP mRNA was found mainly in neuronal cell bodies but was not detectable in glial cell bodies within fiber tracts, such as in the corpus callosum (Fig. 2A). No *Prnp* transcripts were detected in the corpus callosum of *Prnp*^{0/0} mice (Fig. 2B). In contrast, strong mRNA signals were observed in the corpus callosum and, to a lesser extent, in the cerebral cortex (data not shown) of adult *tg640* brains, indicative of strong PrP expression in oligodendrocytes (Fig. 2C). Control hybridization with digoxigenin-labeled sense transcripts did not yield any signal (Fig. 2D–F). *In situ* hybridizations of sciatic nerves yielded PrP signals in both *Prnp*^{+/+} (*tga20*) and MBP-PrP mice but not in *Prnp*^{0/0} mice (Fig. 2J–L). Hybridization signal intensity was less prominent in MBP-PrP mice than in wild-type mice, which is in line with the weaker protein expression. Again, sense probes

did not hybridize (data not shown). Oligodendrocyte-restricted expression of the PrP transgene in brains of *tg640* mice was demonstrated by colocalization of mRNA for *Prnp* and the oligodendrocyte-specific proteolipid protein (PLP) (Fig. 2G–I).

Transmission electron microscopy was used to trace the subcellular distribution of PrP^C within the white matter of adult mice (Fig. 3A–C). Punctuate distribution of colloidal gold-labeled antibodies to PrP^C (arrows) was found in axons as well as in surrounding myelin sheaths of wild-type mice (Fig. 3A), whereas only occasional randomly distributed gold particles were found in *Prnp*^{0/0} mice. In contrast, *tg640* mice displayed the largest number of gold particles within myelinating oligodendrocyte processes (Fig. 3C).

Expression of PrP was also studied by fluorescence microscopy in optic nerve explants from 7- to 8-d-old mice. Strong PrP^C labeling was observed only in oligodendrocytes derived from *tg640* mice, but not from *Prnp*^{+/+} or *Prnp*^{0/0} mice (Fig. 3D–F, top row). Costaining with the early oligodendrocyte marker MAG (Fig. 3D–F, bottom row) confirmed the assignment of signals to oligodendrocytes, as well as the purity of the cultured cells. The majority (but not all) of MAG⁺-cultured cells also expressed PrP^C. Thus, only more mature cells that had formed membrane sheets between their processes were PrP positive, which correlates with late MBP expression. Ectopic PrP expression did not affect oligodendrocyte density, morphology, or MAG expression.

Resistance of *tg640* mice to scrapie

To test whether oligodendrocyte-restricted expression of PrP^C suffices to render *Prnp*^{0/0} mice susceptible to scrapie, *tg640* mice were challenged with scrapie prions (RML strain, passage 5) intracerebrally, intraperitoneally, or intraocularly. However, challenge of homozygous (intraperitoneally, *n* = 12; intracerebrally, *n* = 10; intraocularly, *n* = 7) or hemizygous (intraperitoneally, *n* = 11; intracerebrally, *n* = 10; intraocularly, *n* = 6) *tg640* mice and, as a control, *Prnp*^{0/0} mice (intraperitoneally, *n* = 4; intracerebrally, *n* = 5) did not result in clinical signs of scrapie in any of the mice as late as 641 d postinoculation (dpi). This period of time approaches the natural life span of laboratory mice. Instead, wild-type mice that had been challenged with an inoculum of the same size died of scrapie at 195 \pm 3 d after intraperitoneal (*n* = 6), 163 \pm 4 d after intracerebral (*n* = 6), and 194 \pm 11 d after intraocular (*n* = 7) inoculation (Table 1). All of these mice developed typical clinical symptoms of scrapie, such as ataxia, paralysis, kyphosis, foot-clasp reflex, and mincing gait. Thus, expression of full-length PrP^C by oligodendrocytes of *Prnp*^{0/0} mice does not restore clinical susceptibility to scrapie.

A major hallmark of TSE diseases is brain deposition of PrP^{Sc} (McKinley et al., 1983). We assessed formation of PrP^{Sc} in scrapie-infected MBP-PrP mice by Western blot analysis of brain

extracts (Fig. 4A). Terminally scrapie-sick, intracerebrally scrapie-inoculated wild-type mice (*Prnp*^{+/+}) showed high levels of PrP^{Sc}. However, clinically asymptomatic hemizygous *tg640* mice killed 210 d after intracerebral inoculation did not exhibit any detectable PrP^{Sc} in their brains. All brains of scrapie-inoculated mice surviving until termination of the experiments (641 d postinoculation) were analyzed by histology. This analysis failed to indicate any morphological signs of scrapie, such as spongiosis or gliosis (data not shown).

Next, we investigated whether myelin-derived PrP^C is required for the accumulation of PrP^{Sc} *in situ* by histoblotting. Mice were killed at the indicated time points and analyzed for the presence of disease-associated PrP^{Sc} (Fig. 4B). Large amounts of proteinase K-resistant prion protein were found in infected *Prnp*^{+/+} mice, whereas MBP–PrP and *Prnp*^{0/0} mice lacked any detectable PK-resistant PrP^{Sc}. As reported previously (Brandner

et al., 1996a), myelinated structures show faint homogeneous background staining in inoculated *Prnp*^{0/0} mice. However, these myelinated structures were much less intensive than the PrP^C-containing white-matter tracts in MBP–PrP mice.

To determine whether scrapie infectivity might subclinically propagate in brains or spleens of asymptomatic *tg640* mice, tissue homogenates from animals killed at various time points were injected intracerebrally into indicator *tga20* mice (Fischer et al., 1996), which overexpress *Prnp* under transcriptional control of orthologous regulatory elements (Table 2). Brain homogenates of *Prnp*^{+/+} mice 140 d after intraperitoneal inoculation or 167 d after intracerebral inoculation elicited disease in *tga20* mice after 58–71 d. This result was compared with a calibration curve and found to correspond to a prion titer of ~4.9–6.3 logLD₅₀. In contrast, brains of *tg640* transgenic mice never induced scrapie after transmission to *tga20* mice. We conclude that *tg640* brains do not replicate prion infectivity despite PrP^C expression in oligodendrocytes. The detection threshold of the mouse bioassay with *tga20* mice under the conditions used here is ~1.5 logLD₅₀ per gram of tissue.

Low levels of scrapie infectivity were detected in one spleen of a *tg640* mouse killed 35 d after intraperitoneal inoculation, whereas three of four recipient *tga20* mice died. This finding is most likely a result of the presence of residual inoculum in this investigated *tg640* mouse. Similar findings have long been known to occur, even after inoculation of PrP^C-deficient mice (Büeler et al., 1993; Sailer et al., 1994). Neuropathological investigations of brains from clinically sick wild-type mice revealed typical signs of scrapie (e.g., spongiform changes and pronounced astrogliosis predominantly in the neocortex and hippocampus; data not shown). In contrast, no typical histopathological signs of scrapie were visible in any of the examined clinically healthy *tg640* mice (*n* = 9; killed at 641 dpi; data not shown). These observations virtually exclude the possibility that scrapie prions replicate subclinically in MBP–PrP mice.

Oligodendrocyte-borne PrP^C does not support oculocerebral prion spread

Intraocular prion inoculation leads to progressive scrapie pathology along the optic nerve and optic tract to the contralateral superior colliculus and lateral geniculate nucleus, followed by generalized encephalopathy (Fraser, 1982; Brandner et al., 1996b). These results suggest that the infectious agent travels along fiber tracts of the CNS, such as the retinotectal projection.

To determine whether oligodendrocyte-restricted PrP^C is suf-

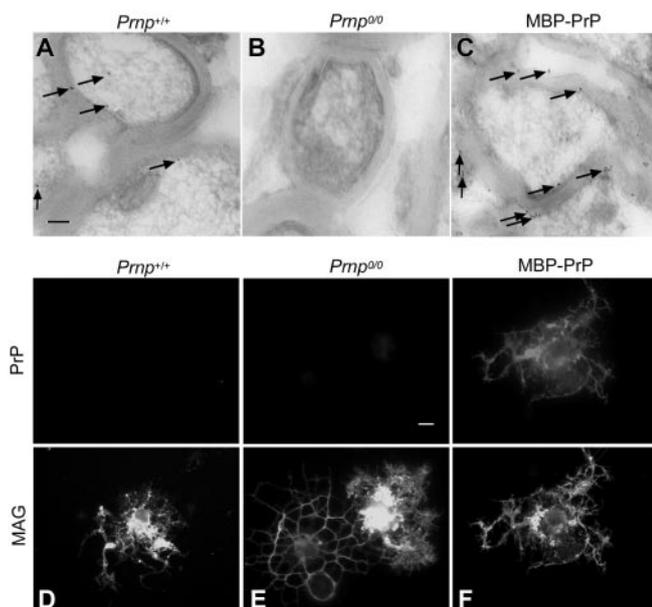


Figure 3. Localization of PrP^C in MBP–PrP mice. *A–C*, Cellular distribution of PrP^C within the white matter of adult mice revealed by immunoelectron microscopy of gold-labeled anti-PrP antibody. Arrows indicate the location of the black gold particles. MBP–PrP mice display abundant decoration of myelin with gold particles. Scale bar, 5 nm. *D–F*, Immunofluorescence of primary optic nerve oligodendrocytes cultured from 7- to 8-d-old mice. Top row, expression of PrP is detectable only in transgenic oligodendrocytes (*F*), whereas the early oligodendrocyte marker MAG is expressed on all oligodendrocytes (bottom row). Scale bar, 5 μ m.

Table 1. Resistance to scrapie of MBP–PrP mice

Type of prion challenge (dose)	Mouse genotypes							
	MBP–PrP ^{+/-}		MBP–PrP ^{+/+}		<i>Prnp</i> ^{+/+}		<i>Prnp</i> ^{0/0}	
	Attack rate	Disease latency ^a	Attack rate	Disease latency ^a	Attack rate	Disease latency ^a	Attack rate	Disease latency ^a
Intraperitoneal (6 logLD ₅₀)	0/11	>641	0/12	>467	6/6	195 ± 3	0/4	>641
Intracerebral (3 × 10 ⁵ LD ₅₀)	0/10	>641, 2 × >210 ^b	0/10	>467, 1 × >197 ^b	6/6	163 ± 4	0/5	>467, 1 × >243 ^b
Intraocular (1 × 10 ⁵ LD ₅₀)	0/6	>531 ^c	0/7	>410, 1 × >376 ^d	7/7	194 ± 11	ND	

Only control wild-type mice (*Prnp*^{+/+}) developed scrapie after intraperitoneal, intracerebral, and intraocular prion challenge, whereas mice carrying the MBP–PrP transgene hemizygously or homozygously and *Prnp*^{0/0} mice never developed clinical signs of scrapie. Average incubation times and SDs were calculated for the groups of mice that developed scrapie. For mice that remained free of disease, the total observation time is reported. All mice were bred to a similar mixed genetic background (C57BL/6 × 129Sv). ND, Not determined.

^aAverage ± SD.

^bClinically healthy mice were killed at the time points indicated, and organs were used for infectivity analysis.

^cOne mouse died 24 hr after inoculation.

^dDeath of one mouse 376 d after inoculation resulting from abdominal tumor growth.

ficient for centripetal scrapie spread, prions were inoculated intraocularly, intraperitoneally, and intracerebrally into either *tg640* or *Prnp^{0/0}* mice containing a PrP-overexpressing *tga20* neurograft (Fig. 5) (Brandner et al., 1996a). Prion injections were performed ≥ 62 d after transplantation to ensure full differentiation of embryonic tissue into normal neuronal and glial components and its integration into the host brain (Isenmann et al., 1996a).

Tg640 or *Prnp^{0/0}* mice grafted with *tga20* neuroectoderm were killed at ≥ 247 dpi. By this time, all intracerebrally infected grafts in both *tg640* ($n = 8$) and *Prnp^{0/0}* ($n = 7$) hosts had developed severe scrapie encephalopathy, as diagnosed by the presence of typical histopathological features, including spongiosis. Pronounced gliosis typical of scrapie was visualized by immunocytochemistry for GFAP, and synaptic loss was illustrated by coarse granular deposits and patchy staining for synaptophysin. Predominantly synaptic deposits of PrP^{Sc} were apparent with the anti-PrP antibody SAF84. In contrast, none of nine *tg640* and four *Prnp^{0/0}* mice that were inoculated intraocularly showed typical histopathological features of scrapie encephalopathy. Identical

results were obtained with five intraperitoneally inoculated *tg640* and five *Prnp^{0/0}* hosts. In two instances, brain grafts of intraocularly inoculated *tg640* mice were assayed by transmission of tissue homogenates to groups of four *tga20* mice. However, none of the recipient mice developed disease, indicating that the grafts were devoid of prion infectivity. We therefore exclude the possibility that PrP^C, when expressed solely on myelin components, supports prion spread within the CNS.

Discussion

Spongiosis, neuronal loss, astroglial and microglial activation, and accumulation of PrP^{Sc} are the most prominent neuropathological features of TSEs. Although neurons are clearly affected in the disease process, it is still unclear whether other brain cells also play a role. Although it is undisputed that during the course of prion diseases neurons can undergo pathological changes culminating in cell death, it is by no means clear whether neurons are the sole or even the primary target of prions. Previous studies have shown that astrocyte-specific expression of PrP driven by the GFAP promoter is sufficient to mediate susceptibility to scrapie in *Prnp^{0/0}* mice (Raeber et al., 1997) and, most surprisingly, to produce a pathology similar to that of wild-type mice. Furthermore, it was speculated that oligodendrocytes may also participate in scrapie pathogenesis, because PrP^{Sc} accumulated most strongly in the white-matter areas, such as the corpus callosum or the fiber tracts of the striatum, during scrapie disease (Bendheim et al., 1992; Taraboulos et al., 1992). In addition, substantial PrP mRNA expression in the white-matter tracts, as shown previously (Moser et al., 1995), implied that oligodendrocytes might be the primary producers of PrP^{Sc} and may be the primary target for prions as well.

Here we describe transgenic mice expressing full-length PrP^C under the control of the MBP promoter. The MBP expression vector was shown to drive ex-

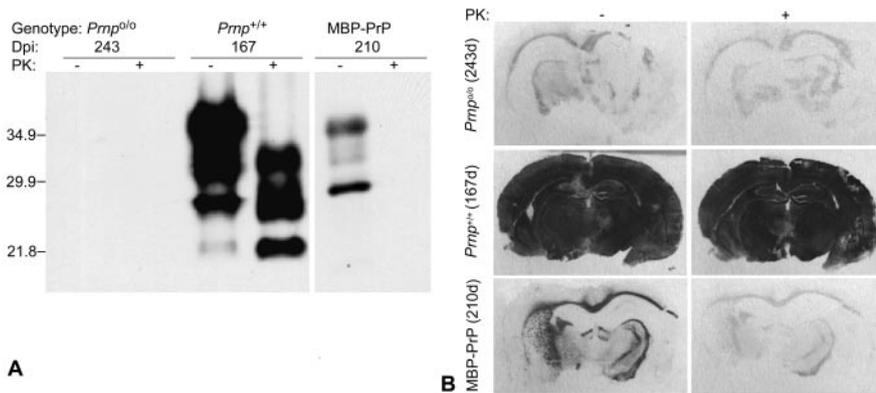


Figure 4. Absence of conversion of PrP^C into PrP^{Sc} after scrapie challenge of MBP-PrP mice. *A*, Western blots of homogenized brain material electrophoresed natively (–) or after digestion with proteinase K (+). Large amounts of PK-resistant prion protein (PrP^{Sc}) are detectable in brains of wild-type mice (*Prnp^{+/+}*) that developed terminal scrapie 167 d postinoculation. No PrP^{Sc} is visible in *Prnp^{0/0}* or MBP-PrP mice that had been challenged intracerebrally with a high dose of the RML strain. Molecular weight markers are indicated on the left. *B*, Histoblot analysis of PrP^{Sc} expression. Protease-resistant PrP was detectable only in intracerebrally infected *Prnp^{+/+}* mice. MBP-PrP and *Prnp^{0/0}* mice did not accumulate PrP^{Sc}. As reported previously (Brandner et al., 1996b), myelinated structures show faint homogeneous background staining in inoculated *Prnp^{0/0}* mice.

Table 2. Prion load in brains and spleens of individual *tg640* mice

Mouse genotypes	Days after inoculation; route of infection	Transmission of brain to indicator mice		Transmission of spleen to indicator mice	
		Brain infectivity (logLD ₅₀ per gram)	Attack rate (mean ± SD days)	Spleen infectivity (logLD ₅₀ per gram)	Attack rate (mean ± SD days)
MBP-PrP	35 d; intraperitoneal	<1.5	0/4	<1.5	0/4
		<1.5	0/4	<1.5	3/4 (99,148,150) ^a
	140 d; intraperitoneal	<1.5	0/4	<1.5	0/4
	210 d; intracerebral	<1.5	0/4	<1.5	0/4
		<1.5	0/3	<1.5	0/4
<i>Prnp^{+/+}</i>	35 d; intraperitoneal	<1.5	0/4	5.0	4/4 (73 ± 5)
		<1.5	0/4	5.1	4/4 (72 ± 4)
	140 d; intraperitoneal	5.3	4/4 (68 ± 6)	4.0	3/3 (84 ± 4)
		4.9	4/4 (74 ± 7)	5.1	4/4 (72 ± 8)
		6.3	4/4 (58 ± 6)	5.1	4/4 (72 ± 5)
167 d; intracerebral	6.1	4/4 (60 ± 7)	5.3	4/4 (70 ± 2)	
<i>Prnp^{0/0}</i>	42 d; intraperitoneal	<1.5	0/4	<1.5	0/4
		<1.5	0/4	<1.5	1/4 (99)

Wild-type (*Prnp^{+/+}*) and *tg640* mice homozygous for the MBP-PrP transgenic cluster were inoculated intraperitoneally and intracerebrally and killed for analysis at the time points indicated.

^aPrion disease in three of four *tga20* indicator mice that had received spleen extracts from a 35 dpi-challenged *tg640* mouse is most likely a result of prions persisting from the inoculum.

pression of the smallest isotype of myelin basic protein and cure the hypomyelinating mouse model *shiverer* (Kimura et al., 1989). Only one of nine transgenic lines expressed sustained levels of PrP^C. Inherent toxicity of the transgene is unlikely, because the incidence of transgene-positive founder mice was normal and line *tg640* exhibited undistorted Mendelian transmission of the MBP–PrP transgene. The relative inefficiency of expression may be specific to the MBP–PrP minigene with a coding exon of 1.3 kb flanked by two small artificial introns. In *tg640* mice, full-length PrP^C was found exclusively on oligodendrocytes and on Schwann cells but was absent from other tissues and cell types. Although the constitutive level of PrP^C in the total brains of transgenic mice was only ~30–35% of that observed in the normal mouse brain, PrP^C protein was clearly overexpressed on transgenic oligodendrocytes compared with wild-type oligodendrocytes.

There was no evidence for PrP^C expression in either neurons or astrocytes. If any expression occurred, it would be under the detection limit of the methods used. Perhaps the strongest argument for the strict specificity of expression in oligodendrocytes is that we could not detect replication of prions in transgenic mice, whereas we know from studies published previously that even very low levels of expression in astrocytes or neurons would restore cerebral prion replication (Büeler et al., 1994; Race et al., 1995; Raeber et al., 1997). Inoculation of *tg640* mice with scrapie prions by different routes did not induce clinical disease or cerebral PrP^{Sc} deposition. This was unexpected in view of the proposed role of oligodendrocytes during disease, as described above.

The fact that PrP^C expression was genuinely restricted to oligodendrocytes allowed us to ask questions related to the spread of prions within the CNS. Intraocular inoculation into *Prnp*^{0/0} mice carrying PrP^C-overexpressing grafts did not provoke scrapie in grafts, supporting the conjecture that lack of PrP^C on cells belonging to the visual pathway prevents prion spread within the CNS (Brandner et al., 1996b). However, it is unclear which mechanisms, be they axonal or nonaxonal, may be involved. Non-neuronal cells, including oligodendrocytes in the CNS and Schwann cells, may play a supportive or obligate role. Within the framework of the prion-only hypothesis, one might hypothesize a “domino” mechanism, by which incoming PrP^{Sc} converts resident PrP^C on the axolemmal or myelin surface, thereby spatially propagating the infection (Aguzzi and Weissmann, 1997). Indeed, one study showed that the velocity of neuronal prion spread is extremely slow (Kimberlin et al., 1983) and may not follow the canonical mechanisms of fast axonal transport. Additional findings support nonaxonal transport mechanisms that result in periaxonal deposition of PrP^{Sc} (Hainfellner and Budka, 1999; Glatzel and Aguzzi, 2000).

In a neurograft paradigm, we find that oligodendrocyte-borne PrP^C does not support transfer of prion infectivity from the eye to intracerebral compartments. Could the negative outcome of the grafting experiment be caused by technical problems of grafting

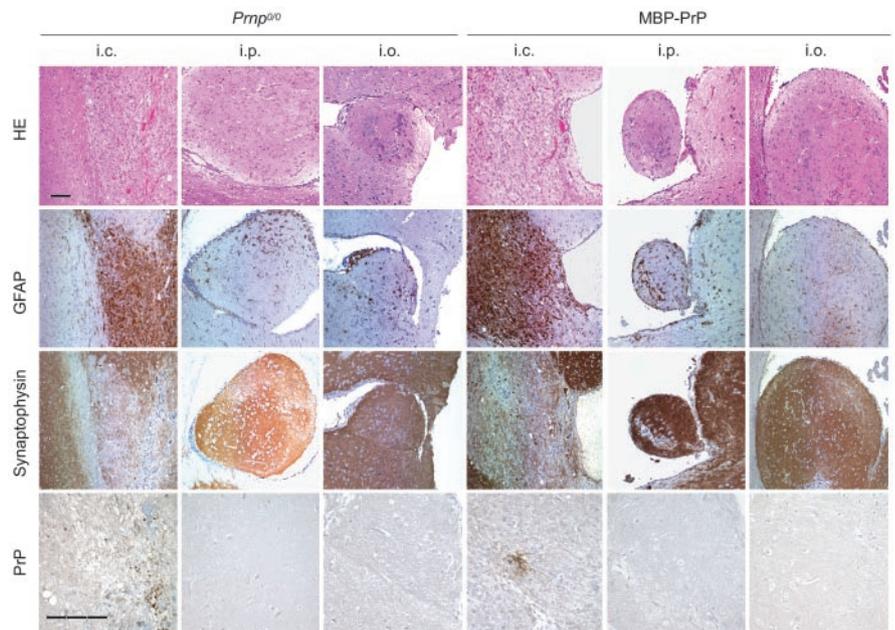


Figure 5. Oligodendrocytic PrP^C does not support intraneural prion transport. Embryonal PrP-overexpressing *tga20* neural tissue was transplanted into *Prnp*^{0/0} (left) or MBP–PrP (right) mice. Hosts were inoculated with scrapie prions intraocularly (i.o.), intraperitoneally (i.p.), or intracerebrally (i.c.). Neuropathological signs of scrapie (gliosis visualized by overexpression of glial fibrillary acid protein, synaptic loss shown by loss of synaptophysin, and deposition of PrP) were only visible in grafts with hosts that had been inoculated intracerebrally. In contrast, neural grafts remained free of disease after intraocular or intraperitoneal inoculation. The trauma deriving from transplantation procedure induced a slight gliosis in all grafts. HE, Hematoxylin and eosin. Scale bars, 100 μ m.

or of intraocular inoculation? This is unlikely, because *tga20* neurografts are easily infectible via the optic route when implanted into *Prnp*^{+/-} hosts, but not when grafted into *Prnp*^{0/0} hosts (Brandner et al., 1996b).

Our findings on the ineffective spread of prions along the visual pathway of *tg640* mice do not favor the nonaxonal theory. The hypothesis that spread of prions occurs axonally rests mainly on the demonstration of progressive spongiform changes along the retinal pathway after intraocular infection (Fraser, 1982). Within these experiments, the spread of scrapie pathology was observed first in the contralateral superior colliculus followed by the lateral geniculate nucleus and visual cortex.

The negative outcome of all prion-related tests prompted us to verify that MBP–PrP mice express intact PrP^C on myelinating cells. Northern and Western blot analyses, as well as *in situ* hybridization, confirmed correct transcription of the transgene and translation into PrP-immunoreactive protein with the expected molecular characteristics. Because prion replication may be compromised even by immunohistochemically silent point mutations, we sequenced the integrated transgene from PCR-amplified genomic DNA. Again, no deviations from the expected sequence were identified. Finally, we verified MBP–PrP transcripts by sequencing reverse transcription-PCR amplification products from brain homogenates. The results of the latter assay rule out RNA editing, which might lead to expression of mutated proteins despite correct genomic sequences.

We still do not know how prions propagate within the brain. The current study excludes any oligodendrocytic contribution to prion pathogenesis by a PrP^C-dependent cell-autonomous mechanism. It is still possible, however, that oligodendrocytic PrP^C modulates the speed of prion neuroinvasion in a non-cell-autonomous manner, perhaps by transcellular “painting” of glycosylphosphatidylinositol-linked proteins to adjacent axolem-

mal surfaces. Conversely, other experiments suggest that PrP^{Sc} of neuronal origin is unlikely to disturb oligodendrocyte physiology to a clinically significant degree (Mallucci et al., 2003). The mouse model described here will make it possible to address this and related questions in a direct way.

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