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Demyelination and axonal preservation in a transgenic mouse model of Pelizaeus-Merzbacher disease

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It is widely thought that demyelination contributes to the degeneration of axons and, in combination with acute inflammatory injury, is responsible for progressive axonal loss and persistent clinical disability in inflammatory demyelinating disease. In this study we sought to characterize the relationship between demyelination, inflammation and axonal transport changes using a *Plp1*-transgenic mouse model of Pelizaeus-Merzbacher disease. In the optic pathway of this non-immune mediated model of demyelination, myelin loss progresses from the optic nerve head towards the brain, over a period of months. Axonal transport is functionally perturbed at sites associated with local inflammation and 'damaged' myelin. Surprisingly, where demyelination is complete, naked axons appear well preserved despite a significant reduction of axonal transport. Our results suggest that neuroinflammation and/or oligodendrocyte dysfunction are more deleterious for axonal health than demyelination *per se*, at least in the short term.

INTRODUCTION

Multiple sclerosis (MS) is an inflammatory demyelinating disorder in which the previously formed myelin sheath breaks down. MS is considered the archetypal myelin disorder and much current research focuses on the associated axonal loss; the major cause of persistent disability (De Stefano et al, 1998; Scolding & Franklin, 1998; Trapp & Nave, 2008). Axonal damage in MS occurs in both the acute-inflammatory and late chronic phases of the disease. The causes are not known, but inflammation and loss of trophic support due to chronic demyelination, respectively, could contribute (Bjartmar et al, 2003; Scolding & Franklin, 1998).

Axonal transport impairment has been implicated in a number of neurodegenerative disorders such as MS (Ferguson et al, 1997) or spastic paraplegia type 2 (Edgar et al, 2004) in which the primary defect is in the oligodendroglial proteolipid protein (*PLP1*) gene. The aim of the present study was to determine to what extent demyelination impairs axonal transport and whether perturbed transport correlates with the chronic absence of myelin or the acute inflammatory phase of demyelination, or both. We selected as our model the optic pathway of a *Plp1*-transgenic mouse (line #72) defined by about twofold overexpression of PLP/DM20 (Anderson et al, 1999; Readhead et al, 1994). In the optic nerve, active demyelination with associated inflammation is succeeded by complete demyelination in the absence of inflammation. For comparison, we also examined the dysmyelinating *Plp1*-mutant *rumpshaker* (Schneider et al, 1992). In *rumpshaker*, formation of the myelin sheath is perturbed leading to a reduced amount of myelin, but there is no evidence of demyelination and the inflammatory response is moderate. Both the *Plp1*-transgenic mouse and *rumpshaker* are models of Pelizaeus-Merzbacher disease (PMD), an X-linked leukodystrophy most commonly caused by duplication of the *PLP1* gene (model: *Plp1*-transgenic mouse), but also by point mutations (model: *rumpshaker*).

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Here we show that in the *Plp1*-transgenic mouse, axonal transport impairment occurs at sites where active demyelination is taking place and also at sites where axons remain ensheathed by mutant oligodendrocytes. In contrast, axons in the completely demyelinated region of the nerve appear structurally and functionally preserved. These data suggest that inflammation and/or oligodendrocyte dysfunction in the *Plp1*-transgenic mouse cause axonal pathology while complete demyelination facilitates axonal survival under stress, at least in the short term.

RESULTS

The optic pathway of *Plp1*-transgenic mice undergoes temporo-spatially determined demyelination

The optic pathway of the *Plp1*-transgenic mouse exhibits moderate dysmyelination followed by progressive demyelination that shows both temporal and spatial evolution. By

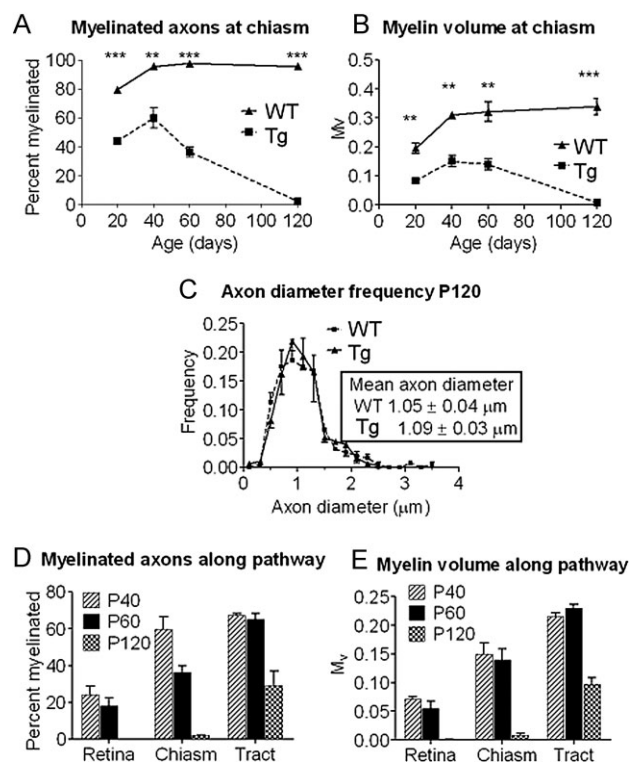


Figure 1. Demyelination progresses in a rostral to caudal direction with increasing age in optic pathway of *Plp1*-transgenic mice.

- A, B.** The percentage of myelinated axons (A) and the relative volume of myelin (B) at the chiasm end of the optic nerve in wild type (WT) and *Plp1*-transgenic (T_g) mice between P20 and P120.
- C.** Axon diameter frequency distribution at the chiasm end of the optic nerve of WT and *Plp1*-transgenic mice aged P120. Axonal diameter distributions are not significantly different.
- D, E.** The percentage of myelinated axons (D) and the relative volume of myelin (E) along the optic pathway of *Plp1*-transgenic mice at P40, P60 and P120 showing the temporal and spatial progression of demyelination.

examining the mid point of the pathway, we showed that myelination commenced on schedule and 44 and 60% of axons were myelinated at P20 and P40, respectively; compared to 80 and 96% in the wild type (WT). However, demyelination led to a complete absence of myelin by P120 (Fig 1A). These profiles were mirrored by myelin volume (M_v) measurements (Fig 1B). Myelin sheaths in the transgenic mice were of normal structure and periodicity, but were thinner than in WT (g-ratio of 0.766 ± 0.007 in mutant vs. 0.732 ± 0.003 in controls at P60; $p < 0.05$). Importantly, there was no significant reduction in absolute axonal number at any age studied. At P120, axon number in the *Plp1*-transgenic mice was $49,683 (\pm 964 \text{ SEM})$ versus $48,768 (\pm 910 \text{ SEM})$ in WT, and axon diameter frequency distributions were not different from WT (Fig 1C). P120 is the oldest age examined, because the *Plp1*-transgenic mice rarely live beyond this age.

To determine the spatial pattern of demyelination we quantified the proportion of myelinated fibres and M_v at the retinal and chiasm end of the optic nerve and at the mid optic tract of transgenic mice. The data (Fig 1D, E) showed a distinct pattern of demyelination progressing from the retinal end towards the brain. At P30, the retinal end of the nerve had 40% of axons myelinated (data not shown). At P120 the pathway shows two distinct zones; a proximal (retinal) completely demyelinated region and a distal actively demyelinating portion. Electron microscopy of the proximal nerve revealed numerous naked axons that appeared morphologically normal, with interspersed astrocytic processes (Fig 2A and Supporting Information Fig S1A). Microglia/macrophages were infrequent. In contrast, the distal (chiasm) end of the nerve, although

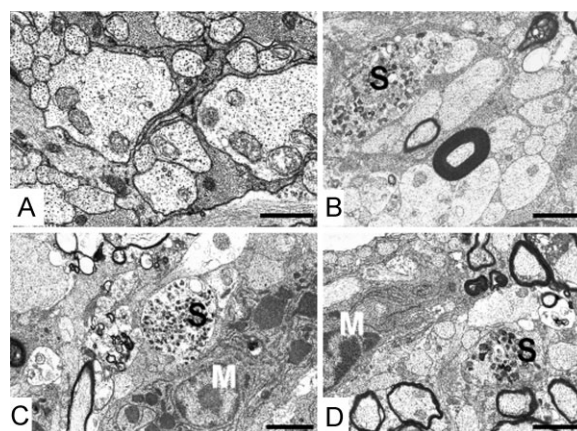


Figure 2. Distribution of myelin and axonal pathology along the optic pathway of *Plp1*-transgenic mice aged P120.

- A.** Retinal end of optic nerve: all myelin has been removed leaving naked axons, which appear morphologically normal and are interspersed with astrocyte processes. Bar: $1 \mu\text{m}$.
- B, C.** Chiasm end of optic nerve: demyelination is on-going although most axons have lost their sheaths. Microglia (M) and axonal spheroids (S) are present. Bar: $2 \mu\text{m}$.
- D.** Optic tract: demyelination is present but not as advanced as at the chiasm. Microglia and axonal spheroids are evident. Bar: $3 \mu\text{m}$.

largely demyelinated, contained numerous active microglia/macrophages (Fig 2B, C). The optic tract was similar, but with a greater proportion of myelinated axons (Fig 2D and Supporting Information Fig S1B). In both the chiasmatal end of the nerve and the tract, some axons showed disruption of the cytoskeleton and/or accumulations of membranous organelles (Fig 2B–D).

Electrical conduction in the dysmyelinated and demyelinated optic nerve

To assess conduction in dysmyelinated and in demyelinated nerves, stimulus-evoked compound action potentials (CAP) were recorded from P40 and P120 WT and *Plp1*-transgenic mice (Fig 3A, B). In WT mice, the CAP has a stereotyped

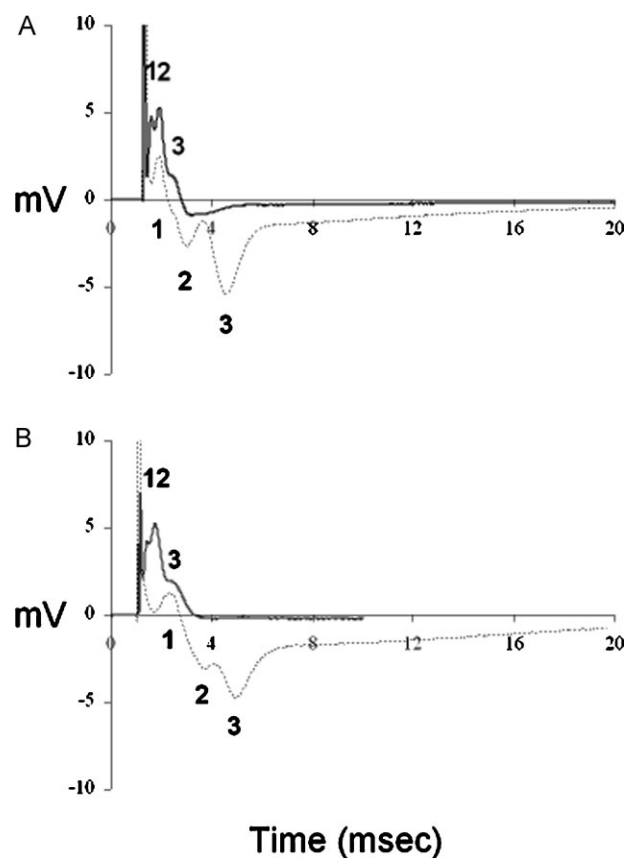


Figure 3. Electrophysiological recordings from WT and *Plp1*-transgenic optic nerves.

- A.** Representative recordings from P40 WT (—) and *Plp1*-transgenic (----) mice. In WT mice, the CAP has the stereotyped triple-peaked profile that reflects populations of axons with different conduction velocities. The CAP recorded from *Plp1*-transgenic mice resolved into three peaks, but the duration of the CAP was longer than that of the control (6 ms vs. 3 ms) signifying a decrease in the conduction velocity.
- B.** Representative recordings from P120 WT (—) and *Plp1*-transgenic (----) mice. The CAP recorded from P120 *Plp1*-transgenic mice was similar to that of the P40 transgenic animals, but the peaks were less well defined and their amplitudes were smaller. A total of 10 WT and 10 *Plp1*-transgenic nerves were examined.

triple-peaked profile (Brown et al, 2003; Stys et al, 1998). In transgenic mice the duration of the CAP was longer (6 ms in mutants vs. 3 ms in controls) signifying a decrease in conduction velocity, consistent with hypomyelination. However, the CAP could still be resolved into three peaks. The CAP recorded from P120 transgenic mice differed from P40 mice in that the amplitudes of the peaks were smaller. In the absence of axonal loss (see above), reduced amplitudes were most likely caused by conduction blocks along demyelinated axons.

The inflammatory response is associated with active demyelination

Since active demyelination is associated with inflammation (Ip et al, 2006), we used the pan-leukocyte marker CD45 to identify inflammatory cells in the *Plp*-transgenic mouse. Whereas in the more distal portions of the P120 optic nerves the numbers of CD45+ cells per optic nerve section were elevated >10-fold compared with WT; the completely demyelinated retinal portion showed no significant increase (Fig 4A). At P40, prior to overt demyelination the cell numbers were strikingly lower than at P120, although elevated compared with WT, probably reflecting the underlying dysmyelination (Fig 4B). Direct correlation of active demyelination and inflammation was achieved by immunostaining for myelin basic protein (MBP) and CD45 (Fig 4C, E).

Using the T-cell marker CD3 we identified approximately 10–20 cells per 10 μ m thick longitudinal section (data not shown). B-cells were rarely observed. We then focused on the microglial/macrophage population using CD169 (sialoadhesin), since CD169 expressing cells contribute to pathogenesis in another *Plp*-transgenic line (line #66; Ip et al, 2006; Readhead et al, 1994). Very few CD169+ cells were present in the demyelinated retinal end of the P120 optic nerve, in contrast to the abundant population at the chiasmatal end (Fig 4C, D). CD169+ cells were absent in WT (Fig 4C) and infrequent in P40 optic nerve, prior to overt demyelination (Fig 4E, F). In fact, the pattern of expression of CD169 closely mirrored the progression of active demyelination at all ages examined (Supporting Information Fig S2). Note that in Fig 4E and Supporting Information Fig S2, the variation in the intensity of MBP staining probably reflects improved antibody penetration into the demyelinating tissue.

Impaired retrograde axonal transport correlates with active inflammation

The occurrence of axonal spheroids (Fig 2B–D) suggests impaired axonal transport, as reported previously in the *Plp1* gene knockout mouse (Edgar et al, 2004). To examine retrograde transport, we injected fluorescein isothiocyanate (FITC)-labelled cholera toxin B subunit (CtB) into the right rostral colliculus of P40 or P120 mice and measured fluorescence signal in the contralateral retinal ganglion cells (RGC) after 18 h. At P40 there was no significant difference in either CtB-labelled area or mean signal intensity between transgenic and WT mice (Fig 5A). At P120, mean signal intensity in RGCs was significantly lower in transgenic mice than in WT, whereas

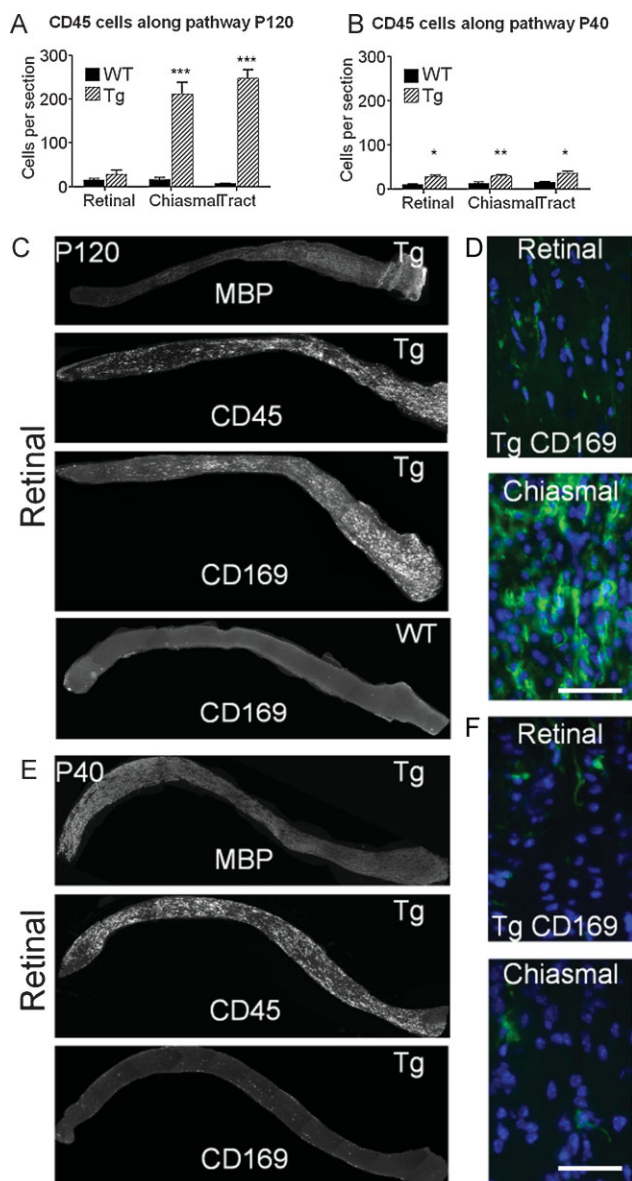


Figure 4. Inflammatory cell numbers change in relation to chronic or active demyelination.

- A.** CD45⁺ cell numbers along the optic pathway of WT and *Plp1*-transgenic (*Tg*) mice aged P120. Whereas the demyelinated retinal region has no increase in these cells there is a massive increase in the actively demyelinating distal regions of the nerve.
- B.** At P40, prior to overt demyelination, the CD45⁺ cell numbers are similar along the entire pathway. The increased number, relative to WT, reflects the developmental dysmyelination.
- C.** Longitudinal sections of optic nerve from the retinal end (left) to the chiasm from a *Plp1*-transgenic (*Tg*) mouse aged P120 immunostained for MBP, CD45 and CD169, showing the gradient in myelin and macrophages between the two ends of the optic nerve. Only CD169 is shown for the WT mouse.
- D.** Enlarged images from the retinal and chiasmatal ends of the optic nerve from the *Plp1*-transgenic mouse illustrate the marked difference in the densities of CD169⁺ cells between the two sites.
- E.** Longitudinal sections of optic nerve, as for C, from a *Plp1*-transgenic (#72) aged P40. MBP immunoreactivity shows that the entire length of the nerve (except the unmyelinated lamina cribrosa) is myelinated at this age. Although CD45⁺ cells are present along the nerve, very few CD169⁺ cells are evident.
- F.** Enlarged images from the retinal and chiasmatal ends of the optic nerve from the *Plp1*-transgenic (#72) mouse illustrating the scarcity of CD169⁺ cells prior to overt demyelination. Bar = 50 μ m.

Within axons, CtB accumulations often showed a punctate appearance (Fig 5D), suggesting an association with endosomes or multivesicular bodies (MVBs). Electron microscopy following 3,3',5,5'-tetramethylbenzidine (TMB) labelling of peroxidase-conjugated CtB showed that accumulations occurred at sites in which axons appeared intact, and in some cases, myelinated (Fig 5E, F).

Simultaneous injection of FITC-labelled CtB into the right colliculus and tetramethyl rhodamine iso-thiocyanate (TRITC)-labelled CtB into the left eye, led to the co-localization of the tracers in some axonal swellings (Fig 5G–I and Supporting Information Fig S3), demonstrating that these foci result from the impairment of both anterograde and retrograde transport and indicating that the associated axons are still intact.

Inflammation correlates with transport defects

The microglial/macrophage population in the transgenic mouse optic nerve revealed a biphasic temporal course. There was a low level of inherent microglia activation associated with dysmyelination, followed by a massive activation in areas of overt demyelination. To compare transport defects in solely dysmyelinated optic nerves, we turned to the *Plp1* mutant *rumpshaker* mouse (Schneider et al, 1992) that exhibits no demyelination. The level of hypomyelination in P40 *rumpshaker* mice was similar to that of age-matched *Plp1*-transgenic mice. However, CD169⁺ cells appeared to be slightly more frequent in *rumpshaker* (Fig 6A). Only a small number of CtB foci, indicating local impairment of retrograde axonal transport (Fig 6B), were present in *rumpshaker* and in contrast to *Plp1*-transgenic mice at age P120, these foci occurred in similar numbers in both retinal and chiasmatal portions.

the CtB-labelled area (an indication of the number of labelled RGC) was similar (Fig 5A). Since axonal counts are not reduced in the mutants, this suggests that the average amount of material transported along the length of an individual axon within 18 h is reduced in the P120 mutant nerves. To visualize perturbations of retrograde transport directly, we examined longitudinal sections of the left optic nerve of *Plp1*-transgenic animals. At P40, occasional focal accumulations of CtB in axons occurred more often in the retinal portion of the nerve, where demyelination is commencing, than in the chiasmatal portion. However, the difference between the two regions was not significant (Fig 5B). Interestingly, at P120, numbers of focal accumulations of tracer were significantly greater at the inflamed chiasmatal end of the nerve compared to the completely demyelinated retinal end (Fig 5B, C).

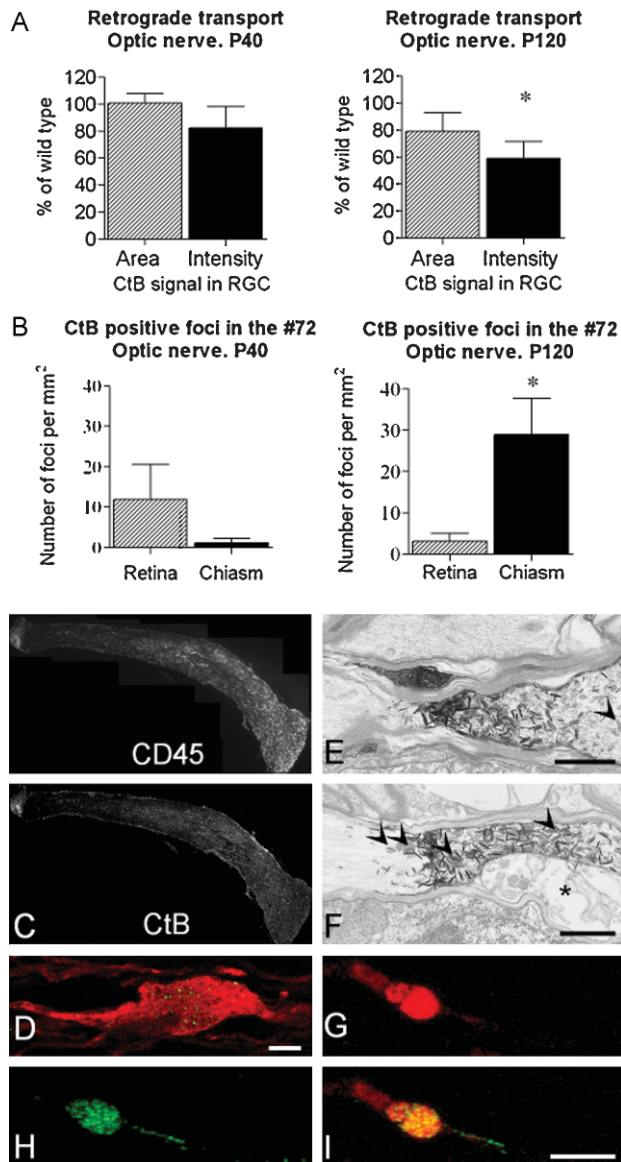


Figure 5. Impaired retrograde axonal transport is associated with active inflammatory demyelination.

- A.** Following injection of CtB into the rostral colliculus, the mean signal intensity in the retinal ganglion cells (RGC) and the area of retina occupied by labelled cells was determined in WT and transgenic littermates aged P40 and P120. At P40, there is no significant difference between transgenic and WT. However, the signal intensity is significantly lower in the P120 transgenic group, when there is active, inflammatory demyelination. As the axon counts in the optic nerves of the two genotypes are also similar the data suggest that the reduced signal is due to impaired transport rather than axonal loss.
- B.** Quantification of CtB foci in retinal or chiasmal portions of the P40 or P120 nerve. At P40, only a few foci are observed and they tend to be more prevalent in the retinal half of the nerve. However, the difference between retinal and chiasmal halves is not significant. At P120, sites of transport stasis are located predominantly in the distal portion of the nerve.
- C.** Longitudinal sections of nerve (retinal end left) from a *Plp1*-transgenic mouse aged P120 immunostained for CD45 and showing CtB positive foci. The sections show the gradient in macrophages and CtB between the two ends of the optic nerve. Lipofuscin is present in these nerves, particularly in the chronically demyelinated region, and is indistinguishable from CtB in these images.
- D.** Single z step through an SMI32-labelled (red) axon in which FITC-CtB (green) appears punctuate, suggesting that it may associate with endosomes and multivesicular bodies. Bar: D = 10 μ m.
- E, F.** Electronmicrographs of horseradish peroxidase (HRP)-conjugated CtB, labelled with TMB, in axons in the optic tract of a P120 mouse. TMB appears as electron dense 'spikes'. The axons remain patent. Axons in these images remain myelinated, however in (F), the inner tongue (asterisk) of the oligodendrocyte is swollen. Arrowheads indicate axonal organelles. Bar: E,F = 1 μ m.
- G, H, I.** Eighteen hours after TRITC-CtB (red) and FITC-CtB (green) were simultaneously injected into the eye and the rostral colliculus, respectively, confocal microscopy showed that many axonal swellings contained both tracers. This suggests that the swellings are associated with defects in both retrograde and anterograde axonal transport, and also indicates that the associated axon remains intact. Bar: I = 5 μ m.

tion. The C3H/101 *rumpshaker* optic nerve exhibits mild dysmyelination associated with modest cellular activation. Sites of axonal transport stasis were most prevalent at regions of active demyelination and inflammation, suggesting that the presence of inflammatory cells and/or damaged myelin are key factors.

Rostral to caudal progression of demyelination

The rostral to caudal progression of demyelination in the *Plp1*-transgenic mouse mirrors that of developmental myelination which commences at the optic nerve head approximately 7 days before it does so at the chiasm (Foran & Peterson, 1992). However, it seems unlikely that the small age difference between retinal and chiasmal oligodendrocytes explains the pattern of demyelination, which progresses over weeks in the *Plp1*-transgenic mouse. One possibility is that remyelination occurs concurrently with demyelination in the chiasmal portion of the nerve, but is less efficient in the retinal portion, manifesting as an apparent gradient of demyelination.

DISCUSSION

Axonal loss is widely accepted as the main determinant of neurological disability in MS (De Stefano et al, 1998). Axonal degeneration correlates with inflammatory demyelination in MS (Ferguson et al, 1997; Trapp et al, 1998), in EAE (Papadopoulos et al, 2006) and in aged mice in which demyelination is induced with cuprizone (Irvine & Blakemore, 2006). Whether axonal injury persists in MS patients once inflammation has resolved has been debated (Frischer et al, 2009; Trapp et al, 1998). To determine the relationship between axonal changes and myelin-related pathology we studied two different models of myelin disease. The *Plp1*-transgenic mouse optic nerve shows a tempo-spatially separated active demyelination with robust microglial/macrophage activation and chronic hypomyelination (due to dys- and demyelination) with low-grade inflamma-

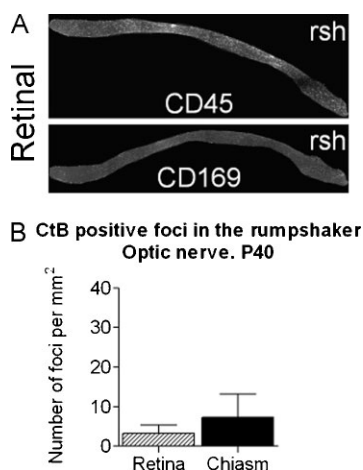


Figure 6. Relatively few sites of axonal transport stasis are associated with diffuse, low level inflammation.

- A.** Longitudinal sections of optic nerve (retinal end left) from the dysmyelinating *rumpshaker* mouse aged P40, immunostained for CD45 and CD169 (sialoadhesin). There is no gradient in the distribution of microglia/macrophages along the optic nerve and activated, sialoadhesin positive cells are present in low numbers compared to the P120 *Plp1*-transgenic nerve.
- B.** Eighteen hours after injection of CtB into the rostral colliculus of P40 *rumpshaker* mouse, a relatively small number of CtB positive foci are present at similar densities in retinal and chiasm portions of the nerve.

Surprisingly, axonal diameter frequency distributions at the chiasm portion of the P120 nerve were not altered compared to WT. In the retinal portion of the nerve, where complete demyelination occurred earlier, a greater proportion of axons were less than 0.6 μm in diameter in the *Plp*-transgenic mouse compared to the WT (average 34.0% vs. 17.2%, respectively); however the overall frequency distributions were not significantly different. Whereas peripheral nervous system (PNS) axons and large diameter spinal cord axons are profoundly influenced by myelination (De Waegh et al, 1992; Tanaka et al, 2009), the subtle differences we observed may be due to the fact that the size of small diameter optic nerve axons is only minimally increased by myelination (Kirkpatrick et al, 2001; Perge et al, 2009).

Sites of transport impairment and swelling do not necessarily represent axonal transection

Some markers of axonal damage, such as focal accumulations of APP, represent sites of axonal transport stasis but may not necessarily indicate that the associated axon is transected (Ferguson et al, 1997). Thus, the damage may be reversible. For example, in EAE models, transient accumulations of APP have been correlated with iNOS expression by perivascular macrophages (Aboul-Enein et al, 2006) or with co-transfer of antibodies to neurofascin (Mathey et al, 2007). In these models, the extent of APP immunoreactivity correlates with clinical symptoms and subsides during remission, providing indirect evidence that the affected axons do not degenerate. Here, we took advantage of

the anatomy of the optic pathway to show directly, using simultaneous injection of CtB into the eye and the colliculus that spheroids belong to intact axons in the demyelinating *Plp1*-transgenic mouse.

Similarly, in a mouse model of Alzheimer's disease, axons remain continuous along their entire length despite containing large focal swellings, indicative of a partial block of axonal transport (Adalbert et al, 2008). Thus, it appears that axonal transport has an in-built overcapacity, such that most axons can survive limited transport defects for some time.

Effectors of axonal transport stasis

Sites of transport impairment were noted in the chiasm and optic tract portions of the visual pathway at P120, which at this age represent sites of active, inflammatory demyelination. Spheroids were noted in both myelinated internodes and in demyelinated axonal segments. In contrast, focal sites of transport stasis were much rarer in completely demyelinated regions where axons appeared structurally intact and were able, in the case of some axons, to conduct electrical impulses.

Activated microglia/macrophages probably contribute to axonal transport stasis at sites of active demyelination. For example, microglia/macrophage 'activation' has been associated with low levels of acute axonal injury in pre-demyelinating lesions in rats receiving focal injection of lipopolysaccharide (Marik et al, 2007), and inflammatory factors (e.g. nitric oxide and tumour-necrosis factor (TNF- α)) produced by this population, can possibly contribute directly to the dysregulation of molecular motors (De Vos et al, 2000; Stagi et al, 2006).

Alternatively, these effects could be indirect, for example triggered by oligodendrocytes that appear physically stressed in an inflammatory milieu (see Fig 2 Butt et al, (2004)). In this context, we note that axonal transport stasis can precede microglial activation and can occur in the absence of demyelination when axons are ensheathed by dysfunctional oligodendrocytes, such as in the myelinated *Plp1* and *Cnp1* knockout mice (Edgar et al, 2004, 2009; Lappe-Siefke et al, 2003). Thus, transport impairment in the *Plp*-transgenic mouse (which was sometimes observed within myelinated internodes) most likely indicates that axons are ensheathed by dysfunctional oligodendrocytes (Edgar & Nave, 2009; Nave & Trapp, 2008).

Good evidence for a functional link between demyelination, inflammation and axonal changes is provided by a comparison with *shiverer* mice, which lack compact myelin in the optic nerve. Indeed, *shiverer* mice show no signs of axonal loss/swelling, unless confronted additionally with the lack of PLP/DM20 from mutant oligodendrocytes (see Fig 1E (Griffiths et al, 1998)). This confirms that *shiverer* axons are in principle at risk for a similar axonal breakdown, but that it requires a 'second hit' on top of the lack of myelin. This could be either inflammation or deficiency of a critical oligodendroglial molecule. Taking these data together, we propose a model in which both neuroinflammation and oligodendrocyte dysfunction provide the highest risk for axonal loss, with demyelination surprisingly improving the chance of axon survival, at least in the short term.

The paper explained

PROBLEM:

Axonal loss is widely accepted as the major cause of permanent neurological disability in multiple sclerosis (MS), an inflammatory demyelinating disorder of the central nervous system. However, the relationship between demyelination, inflammation and axonal damage in neuroinflammatory disorders is not yet fully understood.

RESULT:

Using a mouse model of the leukodystrophy, Pelizaeus-Merzbacher disease, we show that fast transport within optic nerve axons is focally impaired at sites of damage, where active,

inflammatory demyelination takes place and also at sites where axons remain ensheathed by mutant oligodendrocytes. Most interestingly, where demyelination is complete, axons appear preserved.

IMPACT:

Our results challenge the field to re-think the mechanisms of glial support to axons as they suggest that, at least in the short-term, neuroinflammation and associated changes are more deleterious for axonal health than overt demyelination *per se*.

MATERIALS AND METHODS

Mice

The mice used in this study (line #72) carry a cassette harbouring three copies of a *Plp1* genomic transgene (Readhead et al, 1994). These were backcrossed for more than six generations to the C57BL/6N (Charles River) background. WT and homozygous mice were examined at P20, P30, P40, P60 and P120. Only males were used for quantitative analysis, except for the electrophysiological studies, when only females were used. See Supporting Information for genotyping methods. For axonal transport assays (see below), homozygous transgenic mice were compared to age and sex-matched littermates. For all other analyses, comparisons were between age and sex-matched littermates or closely related individuals from the same inbred colony.

Rumpshaker (*rsh/y*) and WT (+/y) male offspring were also examined. These were distinguished on the basis of the neurological phenotype of *rumpshaker*. All animal studies were approved by the Ethical Committee of the University of Glasgow and licensed by the UK Home Office.

Tissue collection and preparation

For electron microscopy, tissues were prepared as described previously (Griffiths et al, 1981). For immunohistochemistry, frozen tissue sections were used. Details of the fixatives employed can be found in Supporting Information.

Antibodies and immunohistochemistry

Antibodies to CD3 (1:100), CD45 (1:600), CD169 (sialoadhesin; 1:100), B-cells (CD45R/B220; 1:100) and MBP (1:500) were obtained from Serotec Ltd (Oxford, UK) SMI-32, which recognizes non-phosphorylated neurofilaments (1:1000) was obtained from Sternberger Monoclonals, Inc. (Lutherville, MD). Secondary antibodies were conjugated with FITC or Texas Red (Cambridge Biosciences). In all cases (except for B cells) sections were permeabilized with methanol (−20 °C), blocked in 10% goat serum in PBS and incubated

overnight in primary antibody in blocking solution. Secondary antibodies in blocking solution were applied for 1 h at room temperature (RT). For the B-cell marker, frozen sections were blocked and the primary antibody was applied for 1 h at RT. Sections were then fixed in 4% paraformaldehyde and the secondary antibody applied, as above.

Digital images

Fluorescence images were taken using an Olympus IX70 microscope, QIcam Fast digital camera (QImaging Surrey, BC, Canada) and Image-Pro Plus 6 software (Media Cybernetics, Silver Spring, MD). Images were processed using Adobe Photoshop. No significant electronic manipulations were made except to images of longitudinal optic nerve sections in which tissue debris around the nerve was digitally removed.

Collicular injections for retrograde transport

To study retrograde axonal transport, WT and *Plp1*-transgenic homozygous littermates at P40 and P120, and WT and *rumpshaker* littermates at P40, were used. Fluorescein-conjugated CtB subunit was injected into the right rostral colliculus and its accumulation in the retina was measured 18 h later (Edgar et al, 2004). The total area occupied by fluorescent signal and the mean density of the signal were determined using Image-Pro Plus 6 software; data are expressed as area of fluorescence per mm² of retina and average mean density (intensity) of signal.

Intraocular injections for anterograde transport

Some mice were simultaneously injected with FITC-CtB and with 1 μl of 1% TRITC-CtB in physiological saline (Invitrogen Ltd., Paisley, UK) into the left eye.

TMB labelling and visualization of CtB-HRP by electronmicroscopy

Twenty-four hours after injection of peroxidase-conjugated CtB (0.5% CtB/1% peroxidase in sterile PBS; Sigma-Aldrich Company

Ltd.) into the colliculus of P120 homozygous #72 mice, the animals were perfused with 1% paraformaldehyde/1% glutaraldehyde/0.1 M PB. Nerves were treated with 0.07% solution of the chromogen TMB and stabilized with 0.7% 3,3'-diaminobenzidine (DAB) tetrahydrochloride/0.33% Cobalt chloride as described by van der Want et al, (1997). Resin embedded tissue was mounted on blank Araldite stubbs and 70 nm section cut and stained with Reynold's lead citrate for EM viewing.

Axonal morphometry and quantification of axonal changes

Axon diameter, density and the myelination status were assessed from electron micrographs (initial magnification $\times 6700$) of randomly selected fields at the chiasmal end of optic nerve from P20, P40, P60 and P120 mice. EM negatives were scanned for analysis using Image-Pro Plus 6 software. Optic nerve areas were measured from light micrographs. Axon densities and nerve area measurements were used to derive total numbers. Axon diameter frequency distributions and the percentage of myelinated axons were also determined. The myelin volume (Mv) and the axonal volume (Av) were determined by a point counting method (Williams, 1977). Myelination status was also ascertained for the retinal end of the optic nerve and the optic tract from *Plp*-transgenic mice.

Cell counts

Cell counts were made in the retinal and chiasmal portions of the optic nerve and in the optic tract. Ten micrometer thick transverse sections, 90 μm apart, were immunostained for CD45, counterstained with DAPI then photographed at $\times 40$ magnification. At least six sections per region were quantified. The densities of nucleated CD45+ cells were counted within an area of interest using Image-Pro Plus 6 software (Al-Saktawi et al, 2003). Optic nerve and tract transverse areas were measured and the average number of cells per 10 μm thick section was calculated.

Quantification of CtB in retinal and chiasmal portions of the optic nerve

Longitudinal sections (10 μm thick) of the optic nerve from animals injected with CtB were imaged using a confocal microscope (Leica SP2 AOBs, Leica Microsystems Semiconductor GmbH, Wetzlar, Germany) at $\times 63$ magnification. Images were taken through the entire depth of the tissue, in retinal and chiasmal portions, at 0.5 μm steps. The image stack was flattened and the numbers of CtB+ foci, greater than 10 μm^2 were counted within an AOI, using Image-Pro Plus 6 software. Foci $< 10 \mu\text{m}^2$ could be observed in WT nerves and therefore were excluded from analysis. At least nine images per nerve were taken from each animal. Lipofuscin granules, which were visible in both red and green channels, were excluded from the analysis.

Electrophysiology

WT and homozygous *Plp1*-transgenic female mice were killed by cervical dislocation. The optic nerves were dissected and placed in an interface perfusion chamber (Medical Systems Corp., Greenvale, NY), maintained at 37°C and superfused with artificial cerebrospinal fluid (aCSF; see Supporting Information) and aerated by humidified gas mixture of 95% O₂/5% CO₂. The CAP was recorded

following supramaximal stimulation, as described previously (Allen et al, 2006).

Statistical analyses

A two-sample Kolmogorov–Smirnov test was used to compare axonal diameter frequency distributions. Calculations were made using Microsoft Office Excel and significance was assessed by reference to published tables (Siegal & Castellan, 1988). All other statistical analyses were made using GraphPad Prism 5 (GraphPad Software, Inc., San Diego, CA). Comparisons between the morphometry of WT and homozygous mice were made using a grouped *t*-test; comparisons along the length of the optic pathway used ANOVA. The comparisons between CtB transport in pairs of WT and transgenic littermates were performed with a paired *t*-test, as were comparisons between CtB foci at retinal and chiasmal portions of the optic nerve. Data are expressed as Mean \pm SEM (*N* was at least 3) and significance in bar charts indicated as < 0.05 (*), < 0.01 (**) and < 0.001 (***).

Authors contributions

J.E., I.R.G., K.-A.N. conceived and designed the experiments and wrote the manuscript. J.E., I.R.G., M.M., P.M., A.M.B., S.T., L.P. and F.G. performed experiments. J.E., I.R.G. and A.M.B. analysed the data.

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Supporting information is available at EMBO Molecular Medicine online.

The authors declare that they have no conflict of interest.

For more information

NINDS Pelizaeus-Merzbacher Disease Information Page
http://www.ninds.nih.gov/disorders/pelizaeus_merzbacher/pelizaeus_merzbacher.htm

Multiple Sclerosis Society UK
<http://www.mssociety.org.uk/index.html>

Proteolipid protein gene information
http://www.ncbi.nlm.nih.gov/gene/5354?ordinalpos=2&itool=EntrezSystem2.PEntrez.Gene.Gene_ResultsPanel.Gene_RVDocSum

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