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Cytoskeletal transition at the paranodes: the Achilles' heel of myelinated axons

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Abstract

Myelination organizes axons into distinct domains that allow nerve impulses to propagate in a saltatory manner. The edges of the myelin sheath are sealed at the paranodes by axon–glial junctions that have a crucial role in organizing the axonal cytoskeleton. Here we propose a model in which the myelinated axons depend on the axon–glial junctions to stabilize the cytoskeletal transition at the paranodes. Thus paranodal regions are likely to be particularly susceptible to damage induced by demyelinating diseases such as multiple sclerosis.

Keywords

Myelin; axon–glial junctions; paranode; node of Ranvier; axonal cytoskeleton; Caspr (contactin-associated protein); NCP1; paranodin; cytoskeletal transition

INTRODUCTION

In vertebrates, neurons project axons over long distances and, to ensure fast nerve-impulse conduction in an energetically economic fashion, some of these axons are targeted for myelination and become insulated by specialized glia (oligodendrocytes in the CNS and Schwann cells in the PNS). Myelination, ultimately, allows the impulse to propagate in a saltatory manner by preventing ionic leakage along insulated segments of the axons. There are differences between myelination in the CNS and PNS such as protein composition, number of axonal segments myelinated by each myelinating cell, the type of glia contacting the nodes of Ranvier, and the presence of a basal lamina. Most of the features that we discuss here are thought to be shared by both the CNS and the PNS. We focus our discussion on the changes in axonal cytoskeleton promoted by myelination and speculate on the contribution of these changes to the dependence of myelinated axons on proper contacts with the myelinating glial cells.

The landmarks for the myelination process include: (1) axonal outgrowth and glial migration to the region of future white matter; (2) targeting of axons for myelination; (3) bidirectional signaling between axons and glia that triggers glial differentiation, polarization of glia and axons into nodes of Ranvier and adjacent molecular domains, and local changes in both glial and axonal architectures; and (4) dependence on cell–cell interactions for survival of both

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myelinating glia and myelinated axons (Barres and Barde, 2000; Witt and Brady, 2000; Pedraza *et al.*, 2001; Corfas *et al.*, 2004; Edgar and Garbern, 2004).

Myelination: a selective process

Axons must have a minimum axonal diameter to become a target for myelination, and proper contact between axons and glia is required for glial differentiation and survival mediated by the axonal signals (Sherman and Brophy, 2005). In the PNS, the protein Neuregulin 1 (Nrg1) type III, which is expressed on the axolemma (axonal membrane), is proposed to serve as a measurement tool for this selection. Because a threshold level of Nrg1 type III is required for further development of the Schwann cells, only axons that display more than this threshold level (i.e. those with more than the minimum membrane surface) provide sufficient Nrg1 type III signaling to be myelinated by Schwann cells (Michailov *et al.*, 2004).

Myelination drives polarization of both the glia and the axons that is coordinated in time and space, resulting in the formation of distinct domains: nodes of Ranvier, paranodes, juxtaparanodes and internodes (Fig. 1A). Each segment of axon wrapped by a compact portion of the myelin sheath corresponds to an internode. Nodes of Ranvier are gaps in the myelin sheath where voltage-gated Na⁺ channels are clustered in the axonal membrane. The paranodes flank the nodes of Ranvier and form the axon–glial junctions (AGJs). Juxtaparanodes are a specialized region of the internodes that flank the paranodes. Here we provide a brief overview of these domains. Several recent reviews have focused on how the node of Ranvier and adjacent domains are established and maintained in myelinated axons (Arroyo and Scherer, 2000; Peles and Salzer, 2000; Pedraza *et al.*, 2001; Salzer, 2003; Bhat, 2003; Rasband, 2004).

At the internodes, the glial cytoplasm is extruded during myelination, which results in the formation of layers of a compact lipoprotein coat (Fig. 2A). Accumulation of glial cytoplasm at the lateral edges of each layer form the paranodal loops. Beneath the glial internodes, in some axons the axonal internodes also experience a myelination-induced regulation in their volume, and axonal diameter increases by up to five-fold in the internodal region compared with the nodes of Ranvier (Figs. 1B and 2A) (Windebank *et al.*, 1985). This expansion is a consequence of the trophic reorganization of the axonal cytoskeleton and correlates with the local phosphorylation of neurofilaments and a subsequent increase in neurofilament spacing (Fig. 1B) (de Waegh *et al.*, 1992). Phosphorylation has been proposed to increase the spacing between the filaments *in vitro* by generating electrostatic repulsion of negatively-charged side-arms at the C-terminal of neurofilament subunits M and H (Kumar and Hoh, 2004). However studies of neurofilament gene replacement in mice indicate that electrostatic repulsion is not the sole mechanism by which phosphorylation increases neurofilament spacing *in vivo* (Garcia *et al.*, 2003).

The juxtaparanodes form at the region where compaction of the myelin sheath relaxes the inner layer to allow the formation of the paranodal loops. The function of the juxtaparanode is still unclear, but this domain is recognized by clustering of specific proteins such as TAG-1 and connexin-29 on the glial side and K⁺ channels, Caspr-2, protein 4.1B and PSD-95 on the axonal side (Fig. 2B) (Rasband, 2004).

The paranodal loops provide the region of closest contact between the myelinating glia and the axons. Paranodes are the site of assembly of AGJs, which resemble the invertebrate septate junctions (Einheber *et al.*, 1997; Banerjee *et al.*, 2006). In mouse mutants with disrupted AGJs, the paranodal domain does not form transverse bands and juxtaparanodal proteins on the axonal side localize at the region coordinated with the glial paranodal loops (Dupree *et al.*, 1999; Bhat *et al.*, 2001; Boyle *et al.*, 2001). The consequence is a direct contact between proteins of juxtaparanodes and nodes of Ranvier, which indicates that AGJs normally serve as a ‘fence’ to segregate the axonal domains that flank the nodes of Ranvier (Rios *et al.*, 2003). The

molecular components of paranodes that are known are the 155 kDa isoform of neurofascin, NF155, on the glial side and Caspr (also known as Paranodin and NCP1), contactin (Cont), protein 4.1B, α II-spectrin, β II-spectrin and ankyrin-B on the axonal side (Fig. 2C) (Einheber *et al.*, 1997; Menegoz *et al.*, 1997; Bellen *et al.*, 1998; Bhat *et al.*, 2001; Charles *et al.*, 2002; Denisenko-Nehrbass *et al.*, 2003; Garcia-Fresco *et al.*, 2006; Ogawa *et al.*, 2006). Contactin is also synthesized by oligodendrocytes during maturation, but has not been detected on the glial side of the paranodes (Gennarini *et al.*, 1989; Faivre-Sarrailh *et al.*, 1992; Koch *et al.*, 1997).

The node of Ranvier provides the site for clustering of the voltage-gated Na⁺ and K⁺ channels for ion exchange (Devaux *et al.*, 2003; Poliak and Peles, 2003; Devaux *et al.*, 2004; Schwarz *et al.*, 2006). In addition to these channels, the main molecular components of the nodes are Nr-CAM and NF186 at the axolemma, and β IV-spectrin and ankyrin-G in the axonal cytoplasm (Fig. 2D) (Poliak and Peles, 2003).

Axon–glial communication

Proper communication between axons and glial cells is a feature of the myelination process. Axons communicate their diameter to glia, which use this information to adjust accordingly the number of myelin wraps, resulting in a highly regulated ratio between axonal diameter and myelin thickness, known as the g ratio. The diameter of a myelinated axon also correlates with the length of the paranode (which correlates with the number of paranodal loops that contact the axons) and the length of the internodes. The mechanisms that dictate the internodal length in myelinated axons are unclear (Friede and Samorajski, 1967; Friede and Bischhausen, 1980; Friede and Bischhausen, 1982).

Analysis of myelin thickness in mice that are deficient in phosphorylation of neurofilaments because of deletion of the phosphorylatable isoforms of neurofilament (NF-M, NF-H and NF-M/H) indicated that Schwann cells and oligodendrocytes read different axonal signals for establishment of the g ratio (Elder *et al.*, 2001). In these mutants, the expected g ratio was maintained in CNS axons, whereas it was decreased in PNS axons. This suggests that the CNS axons relay to oligodendrocytes about their diameter after myelination-induced axonal expansion, whereas PNS axons relay to Schwann cells about their future diameter during initial contact or shortly thereafter. Because PNS axons of the above mutants do not undergo the axonal expansion that is regulated normally via phosphorylation of neurofilaments, the myelination thickness is increased in relation to the final axonal diameter, and the g ratio is decreased (Elder *et al.*, 2001). Similar observations have been made in the sciatic nerve of periaxin-null mice that do not undergo significant myelination-induced axonal expansion and have a decreased g ratio (Williams and Brophy, 2002). Recent studies by Nave and colleagues have demonstrated that it is possible to alter the g ratio in the PNS through manipulating expression of Nrg1 type III in transgenic mice. This indicates that these axons communicate their diameter to glia through Nrg1 type III on the axolemma. At the glial membrane, ErbB2 binding to Nrg1 type III triggers signaling for glial extension into wrapping layers (Michailov *et al.*, 2004). These studies indicate that CNS and PNS glia might receive axonal signaling for regulation of the g ratio at different stages during myelination. However, it is generally thought that myelinating glia perceive axonal diameter at the initial stages of myelination, because a minimum initial diameter is required for axons to be myelinated.

By contrast, axons rearrange their cytoskeleton in response to glial signaling. Myelination, thus, triggers a second expansion of axonal diameter (Windebank *et al.*, 1985). Whereas the total number of neurofilaments grossly dictates the diameter of unmyelinated axons, the myelination-induced axonal expansion maximizes the initial diameter of axons by modulating the spacing between neurofilaments through phosphorylation (Mata *et al.*, 1992; Hsieh *et al.*, 1994b). Griffin and colleagues have developed a methodology based on electron micrographic

data to quantify the levels of neurofilament packing in PNS axons through a parameter that the authors named nearest neighbour neurofilament distance (NNND). The NNND value indicates the spacing between orderly distributed neurofilaments and correlates nicely with the phosphorylation levels of neurofilaments and the diameter of normal myelinated axons (Hsieh *et al.*, 1994a; Hsieh *et al.*, 1994b; Lunn *et al.*, 2002).

A key glial component in the signaling pathway of myelination-induced axonal expansion is myelin associated glycoprotein (MAG) (Yin *et al.*, 1998). MAG localizes to the periaxonal membranes of glial internodes in both CNS and PNS. In the PNS, MAG is enriched at the paranodal loops of mature Schwann cells (Trapp *et al.*, 1989). Cleveland and colleagues have analyzed mouse mutants that lack phosphorylation sites at the C-terminal of neurofilaments and proposed a model in which axonal expansion triggered by myelination starts with MAG-dependent signaling and, ultimately, targets phosphorylation of neurofilament-M via p75NTR-dependent activation of cyclin-dependent kinase 5 (cdk-5) and/or ERK1/2 (Garcia *et al.*, 2003). Moreover, cdk5 mediates phosphorylation of important cytoskeletal proteins in axons such as tau and MAP1B, and, consequently, regulates the attachment of these proteins to microtubules, which modulates rates of axonal transport (Matsushita *et al.*, 1996; Wada *et al.*, 1998; Bu *et al.*, 2002; Dashiell *et al.*, 2002; Kawauchi *et al.*, 2005).

Alternating axonal diameter: an Achilles' heel of myelinated axons

Axonal expansion caused by phosphorylation of neurofilaments is reversible, thus, axons decrease their internodal diameter upon demyelination and expand again after remyelination. Consistent with this observation, axons do not undergo myelination-induced expansion in unmyelinated segments like at the nodes of Ranvier, where neurofilaments remain unphosphorylated and, hence, tightly packed (Hsieh *et al.*, 1994b; Sanchez *et al.*, 1996). Therefore, because of the axon–glial communication, some myelinated axons present alternating diameter along their length. Here we argue that this alternating pattern is a key contributor to the axonal dependence on the integrity of the axon–glial interactions after myelination (Fig. 1B).

While still unmyelinated, the diameter of the axons tends to be uniform along their length, which provides a linear track for axonal transport. After myelination, some axons display remarkably uneven diameter, with long segments of spaced phosphoneurofilaments interrupted by short segments of packed, non-phosphorylated neurofilaments (Fig. 1B). Alternation of diameter along these axons generates points of retention for axonal transport (i.e. nodal and paranodal regions present reduced rates of axonal transport and increased accumulation of organelles when compared with internodes) (reviewed in Salzer, 2003). Finally, one might picture the paranode as an Achilles' heel in these myelinated axons, where the creation of an alternating pattern for axonal diameter might generate regions of potential complications in axonal transport. Does myelination provide an adaptation mechanism to ensure smooth transport at each change in axonal diameter?

Cytoskeletal transition at the paranodes

Although perhaps unconventional, we refer to paranodes as the region of transition between alternations in diameter of myelinated axons. In axons that undergo alternating pattern, the paranodal cytoskeleton tapers as it approaches the nodes of Ranvier; and we suggest that this transition between expanded and nonexpanded cytoskeleton is maintained actively by paranodal modulation of local cytoskeleton (Figs. 1B and 2). The finding that AGJs are linked to axonal cytoskeleton through the Caspr-dependent recruitment of a paranodal complex with protein 4.1B, α II-spectrin, β II-spectrin, ankyrin B and actin supports this idea (Gollan *et al.*, 2002; Denisenko-Nehrbass *et al.*, 2003; Garcia-Fresco *et al.*, 2006; Ogawa *et al.*, 2006). *CGT* and *Caspr* mutants, which are deficient in proteins that are synthesized in glia and axons,

respectively, exhibit disrupted AGJs stemming from very different mechanisms (Coetzee *et al.*, 1996; Bhat *et al.*, 2001). The study of these mutants reveals a previously unappreciated function for the Caspr-mediated link between AGJs and the axonal cytoskeleton in the maintenance of the cytoskeletal organization as well as axonal transport at the paranodal region. Disruption of this link leads to the development of axonal swellings at the paranodal region of the Purkinje axons (Fig. 3B). *CGT*- and *Caspr*-mutants exhibit normal myelination-induced expansion of the axonal cytoskeleton, but do not form the paranodal transverse bands. These mice, thus, provide a model to selectively study the role of AGJs in the intrinsic transition between expanded and nonexpanded cytoskeleton that normally takes place in the paranodal region (Fig. 3A). Ultrastructural analysis revealed that the paranodal regions in *CGT* and *Caspr* mutants are disorganized severely in their local cytoskeleton, as depicted in Fig. 2B. It is, thus, appealing to propose that the AGJ link with the cytoskeleton stabilizes the transition between expanded and nonexpanded cytoskeletons along the myelinated axons.

The mechanisms and signaling cascades that direct the cytoskeleton transition at the paranodes are largely unknown. Does the AGJ/microfilament complex link the microtubules and/or neurofilaments to the axolemma? Are the paranodes sites for modifications of neurofilaments that promote a decrease in neurofilament spacing, such as dephosphorylation of C-terminal neurofilament sidearms or phosphorylation of the neurofilament N-terminal? Are paranodes sites for regulation of proteins that crosslink microtubules and neurofilaments? Although we do not have answers for these questions, the model proposed here suggests that they are relevant and need to be addressed. We propose that the AGJs actively modulate the cytoskeletal organization beneath the paranodes. This would ensure the spatial segregation and a smooth transition between expanded internodal and non-expanded nodal cytoskeletons.

Although the molecular link between AGJs and the cytoskeleton and its implications on axonal transport have been reported only recently, earlier studies by Griffin and others using the neurotoxin β,β' -iminodipropionitrile (IDPN) pioneered the correlation between paranodal integrity and disruption in axonal transport (Griffin *et al.*, 1984; Griffin *et al.*, 1987; Parhad *et al.*, 1987; Griffin and Sheikh, 1999). It was demonstrated that impairments of axonal transport of neurofilaments induced by IDPN leads to the formation of neurofilament-enriched swellings and to a passive paranodal demyelination, with displacement of the paranodal loops towards the internodes (Griffin *et al.*, 1987). These results are consistent with our model and indicate that the link between AGJs and the paranodal cytoskeleton is sufficiently strong to create interdependence. In other words, disorganization of the paranodal cytoskeleton leads to mechanical displacement of myelin loops through their link with the AGJs.

IDPN-induced paranodal swellings indicate that paranodes provide a cytoskeletal track that is highly susceptible to complications in axonal transport, which is consistent with the idea of a cytoskeletal transition at the paranodes. However, these experiments did not rule out the possibility that mechanical displacement of AGJs insults the axolemma, which might be the major cause of the formation of paranodal swellings. Trauma induced in myelinated axons is reported to lead to rapid formation of swellings at the nodes of Ranvier (nodal blebs), the formation of which has been attributed to Ca^{2+} influx into the axon through the axolemma (Maxwell *et al.*, 1991; Maxwell and Graham, 1997). Other studies support the hypothesis that mechanical disruptions of the axolemma lead to formation of axonal swellings through influx of free Ca^{2+} and local activation of Ca^{2+} -dependent enzymes, such as proteases (Fitzpatrick *et al.*, 1998). The development of paranodal swellings in *CGT*- and *Caspr*-mutants, where AGJs do not form and no detectable paranodal demyelination is observed, indicate a causal link between the formation of AGJs and the organization of paranodal cytoskeleton (Garcia-Fresco *et al.*, 2006).

Paranodal swellings in *CGT* and *Caspr* mice are observed only in a limited range of myelinated axon (i.e. axons of Purkinje neurons). This indicates that other compensatory mechanisms act to stabilize the cytoskeletal transition at the paranodes in the absence of AGJs in other neurons. AGJ disruption also leads to accumulation of mitochondria at the nodal region in older *Caspr* mutants (Einheber *et al.*, in press). The presence of a basal lamina in the PNS nerves might physically protect the paranodes from developing swellings and, thus, might cause the accumulation of axonal transport cargo instead at the nodes. Because mitochondria are peculiar in the way they are transported through the axons with high rates of releases from the molecular motors, they might be more likely to be trapped on the disorganized paranodal cytoskeleton than other organelles (Einheber *et al.*, in press).

Other possible mediators of cytoskeletal transition at the paranodes

AGJs are unlikely to be the sole mediators of cytoskeletal transition at the paranodes. *PLP/DM20*-null and mosaic mice develop paranodal swellings and myelination-induced axonal expansion but AGJs form normally in these mice (Griffiths *et al.*, 1998). This observation indicates that isoforms of PLP might also affect paranodal cytoskeleton through unknown mechanisms.

Paranodal demyelination with the formation of swellings is also a feature of giant axonal neuropathy (GAN), an autosomal recessive disorder caused by mutations in the protein gigaxonin that interferes with axonal transport. *In vitro*, gigaxonin regulates microtubule stability by binding and inducing degradation of the microtubule-associated protein 1B (MAP1B) (Allen *et al.*, 2005). Mutations in gigaxonin disrupt interaction between gigaxonin and MAP1B (Ding *et al.*, 2002). Interestingly, MAP1B mediates cross talk between microtubules and actin filaments in axons. Cdk5 phosphorylates MAP1B and the phosphorylated form of MAP1B was reported recently to be enriched at the paranodes (Soares *et al.*, 2002).

It is tempting to speculate that MAP1B has an additional role in the paranodal cytoskeletal transition. Most importantly, is the phosphorylation of MAP1B at the paranodes, a downstream event of MAG signaling triggered during myelination? Ablation of MAP1B leads to delayed myelination and myelination-induced axonal expansion (Takei *et al.*, 1997). MAP1B localizes at neuronal membranes and immuno-precipitates with MAG; moreover, MAP1B phosphorylation increases when neurons are cocultured with cells that express MAG (Tanner *et al.*, 2000; Franzen *et al.*, 2001; Dashiell *et al.*, 2002). Is there a correlation between disruption of gigaxonin binding to MAP1B in GAN and the formation of paranodal swellings? Identification of additional players in the cytoskeletal transition at the paranodes should help to elucidate the mechanisms by which myelination modifies the paranodal cytoskeleton.

Limitations of the cytoskeletal transition model

One of the premises of this model is that the feature of myelination-induced axonal expansion is, in some measure, shared by both the CNS and the PNS axons. At least for the CNS optic nerves, cytoskeletal expansion of the myelinated part of the axons is dependent on oligodendrocyte signaling, and the unmyelinated portion of the same axons does not expand (Sanchez *et al.*, 1996). However, it must be pointed out that the key studies characterizing the differences between internodal and nodal diameter have been performed in the PNS. It is likely that the larger diameter of PNS axons together with the existence of a structurally protective basal lamina facilitates the preservation of local diameter in axonal segments during sample preparation and ultrastructural detection of the alternating pattern in the PNS rather than CNS.

The existence of nodes of Ranvier in CNS myelinated axons was received initially with much skepticism, because morphological features such as the 'nodal constriction' were not detected

in these CNS myelinated axons. This was later attributed to difficulties in obtaining proper cytological fixation in the CNS because under optimal conditions the 'nodal constrictions' are detected in the CNS axons (Bodian, 1951; Nakai, 1954; Pease, 1955). However, CNS axons seem to be morphologically more diverse than PNS axons. Even in optimal tissue preparations, CNS nodes of Ranvier are more permeable than the internodes, which probably led to variability in the morphology and stain of these domains (Bodian, 1951). Another possible explanation for the variability of morphology in the CNS nodes of Ranvier is the fact that they are exposed to different environmental cues in a less patterned way compared to PNS fibers from, for example, the sciatic nerve. External cues modulate the dynamics of actin-based cytoskeleton and lead to different forms of structural plasticity such as synapse formation and axonal branching that influence the shape of the nodes of Ranvier (Rosenstein and Leure-DuPree, 1976; Yiu and He, 2006). This local, actin-based activity might lead to nodal protrusions and contractions and, thus, mask the morphological detection of neurofilament-based alternating pattern in the CNS.

Palay and Chan-Palay have described an anatomical point just before the onset of the myelin sheath as the narrowest portion of the initial segment of the Purkinje axons, where the diameter drops from about 1 μm to 0.5 μm , and expands again beneath the myelinated sheath (Palay and Chan-Palay, 1974), however, we are unaware of studies that show precisely that Purkinje axons indeed undergo alternating pattern. Together these circumstantial observations indicate that some CNS axons, including Purkinje axons, undergo alternating pattern after myelination, but further studies are needed to confirm this assumption.

Predictions of the cytoskeletal transition model

Understanding the mechanisms of degeneration of myelinated axons in demyelinating diseases is relevant to the development of treatments for MS and CMT because axonal degeneration is the major cause of neurological disabilities associated with these diseases (Bjartmar *et al.*, 1999; Bjartmar *et al.*, 2003). Here we present a view of the paranodes as a region of axonal susceptibility to injuries caused by demyelination. Consistent with this view, an early sign of myelin loss in MS is the mislocalization of Caspr and NF155 (Wolswijk and Balesar, 2003; Coman *et al.*, 2006; Howell *et al.*, 2006).

The model of cytoskeletal transition at the paranodes makes several predictions that can be tested and correlated with the available data. Probably the most important prediction is that partial myelination would insult some axons more than complete demyelination. This would be the case for myelinated axons that display alternating pattern in their diameter but have lost the adaptation mechanisms to modulate transition at each alternation. This prediction is consistent with the phenotype of mutants that have myelination-induced axonal expansion but lack proper AGJs, such as Caspr and *CGT* mutants, which develop paranodal swellings. By contrast, the model predicts that complete abolition of this alternating pattern on axonal diameter would not lead to the development of axonal swellings. This prediction is also consistent with the phenotype of mutants that are deficient in the signaling for phosphorylation of neurofilaments, such as MAG and mice deficient in phosphorylation of neurofilament by truncation or ablation of neurofilament subunits M or H/ M (Yin *et al.*, 1998; Garcia *et al.*, 2003).

A simplified summary of relevant observations is presented in Fig. 4, to conceptualize the predictions of our model. Interpretation of axonal degeneration in most demyelinating diseases and animal models should not be narrowed down to a single general mechanism. We have selected mutants for discussion that display changes in axonal diameter because of changes in neurofilament spacing, and we have analyzed a very specific parameter of axonal damage, the development of paranodal swellings. Studies using NNND as a parameter should be

instrumental in better characterizing the intrinsic deficiencies in the alternating pattern generated by partial myelination and to test the model proposed here.

Conclusions and remarks

Here, we propose a model in which alternations in axonal diameter induced by myelination generate axonal dependence on the proper cytoskeletal transition at each paranode. An important prediction from this model is that disruptions in AGJs will promote less damage if axons do not present the alternating pattern in their diameter. One might wonder whether the development of paranodal swellings in mice with disrupted AGJs (CGT and Caspr) would be attenuated in a mouse background in which phosphorylation of neurofilaments induced by myelination is abolished (neurofilament-M mutants). In addition, this raises the hypothesis that drugs that disrupt phosphorylation of neurofilament-M specifically might have the potential to either attenuate or partially prevent paranodal swellings that are promoted by disruption in the AGJs. Finally, the model for cytoskeletal transition at the paranodes provides a novel perspective to speculate on the role of axon–glial interactions beyond the membrane specializations, down into the core of the specialized axonal cytoskeleton.

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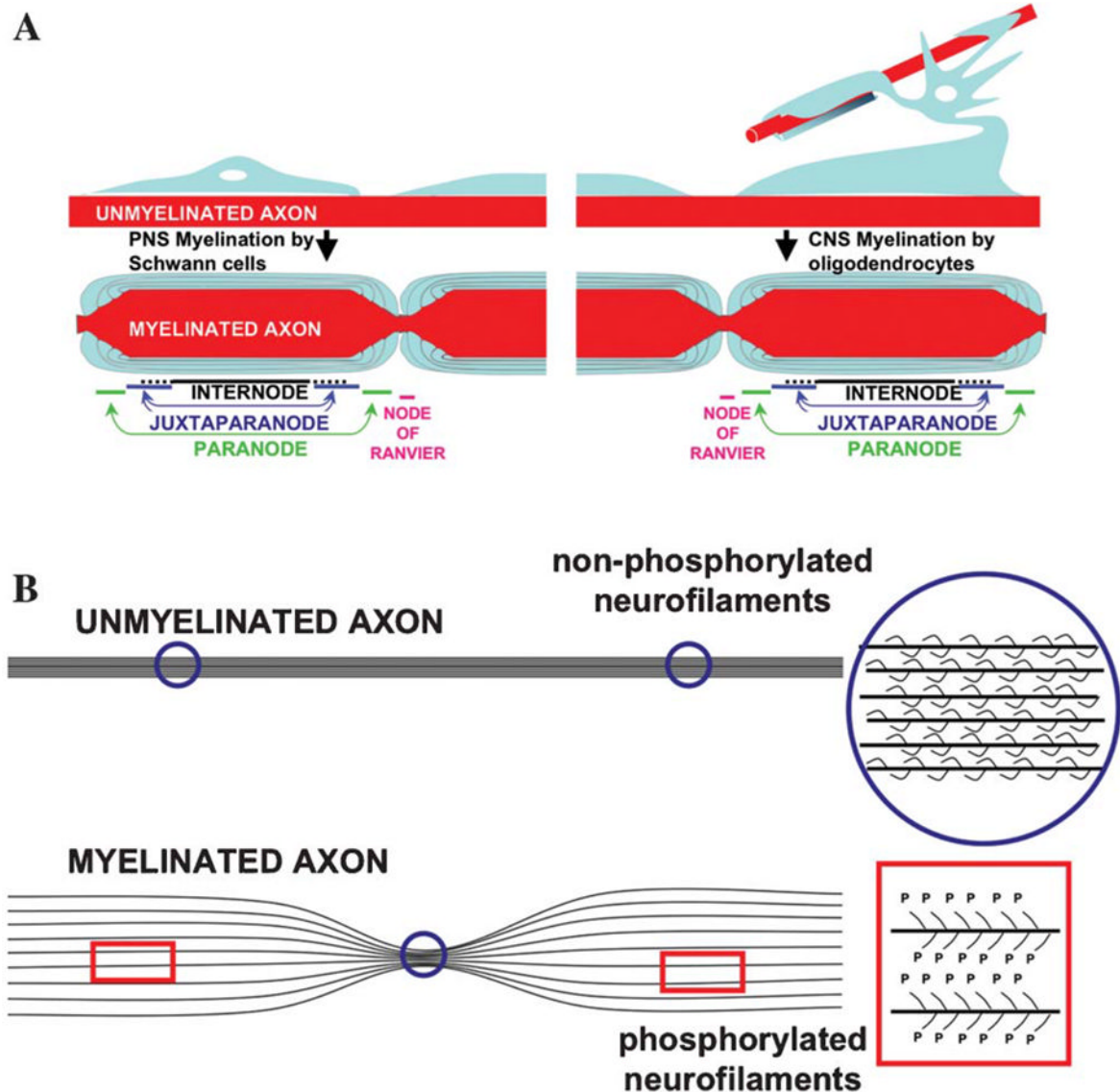


Fig. 1. Landmarks for changes in axons and glia after myelination

(A) Polarization of both axons and glia in the node of Ranvier and adjacent domains. The schematic is not to scale but the difference in axonal diameter between nodes of Ranvier and internodes is reported to be up to five fold. Similarities between myelination by oligodendrocytes (CNS) and Schwann cells (PNS) are presented. Nodes of Ranvier contact processes of perinodal astrocytes in the CNS and microvilli of Schwann cells in the PNS. We have omitted this information as well as the basal lamina in the PNS from the schematic, aiming to emphasize the similarities between the two myelination processes. (B) Phosphorylation of neurofilaments has a major role in increasing axonal diameter through myelination-dependent signaling.

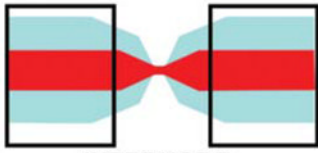
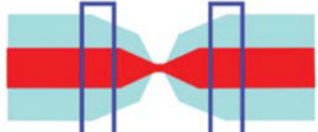


	GLIA	AXON
A  INTERNODE	Cytoplasmic extrusion Localization of MBP, P0, PMP22, PLP, MAG at the inner layer	Cytoskeletal expansion
B  JUXTAPARANODE	Clustering of TAG-1 and connexin 29	Clustering of K ⁺ channels, Caspr2, 4.1B, and PSD-95
AXO-GLIAL JUNCTIONS		
C  PARANODE	Swollen at paranodal loops Clustering of NF155 Enrichment of MAG (PNS)	Transition between expanded and non-expanded cytoskeleton Clustering of Caspr, Cont, 4.1B, α II spectrin, β II spectrin and AnkB
D  NODE	No compact myelin, contacted by microvilli of Schwann cells (PNS) or processes of perinodal astrocytes (CNS)	No cytoskeletal expansion Clustering Na ⁺ channels, NF186, AnkG and β IV spectrin

Fig. 2. Schematic of the main changes in axons and glia that are promoted by myelination
 (A) In the internodal region the glial cytoplasm is extruded, which leads to compaction of the myelin sheath and remarkable expansion of the cytoskeleton in some axons. (B) In the juxtaparanodal region, the specific glial and axonal components cluster during myelination and establish an axon–glial scaffold with no specialized ultrastructure. (C) In the paranodal region, local clustering of glial NF155 and axonal Caspr, Cont, 4.1B, α II and β II spectrin, and ankyrin B results in the establishment of axon–glial junctions. (D) In the nodal region, either Schwann cell microvilli (PNS) or perinodal astrocytes (CNS) interact with nodal axolemmal components. Nodal cytoskeletal proteins NF186, ankyrinG and β IV spectrin coordinate the clustering of the voltage-gated Na⁺ channels.

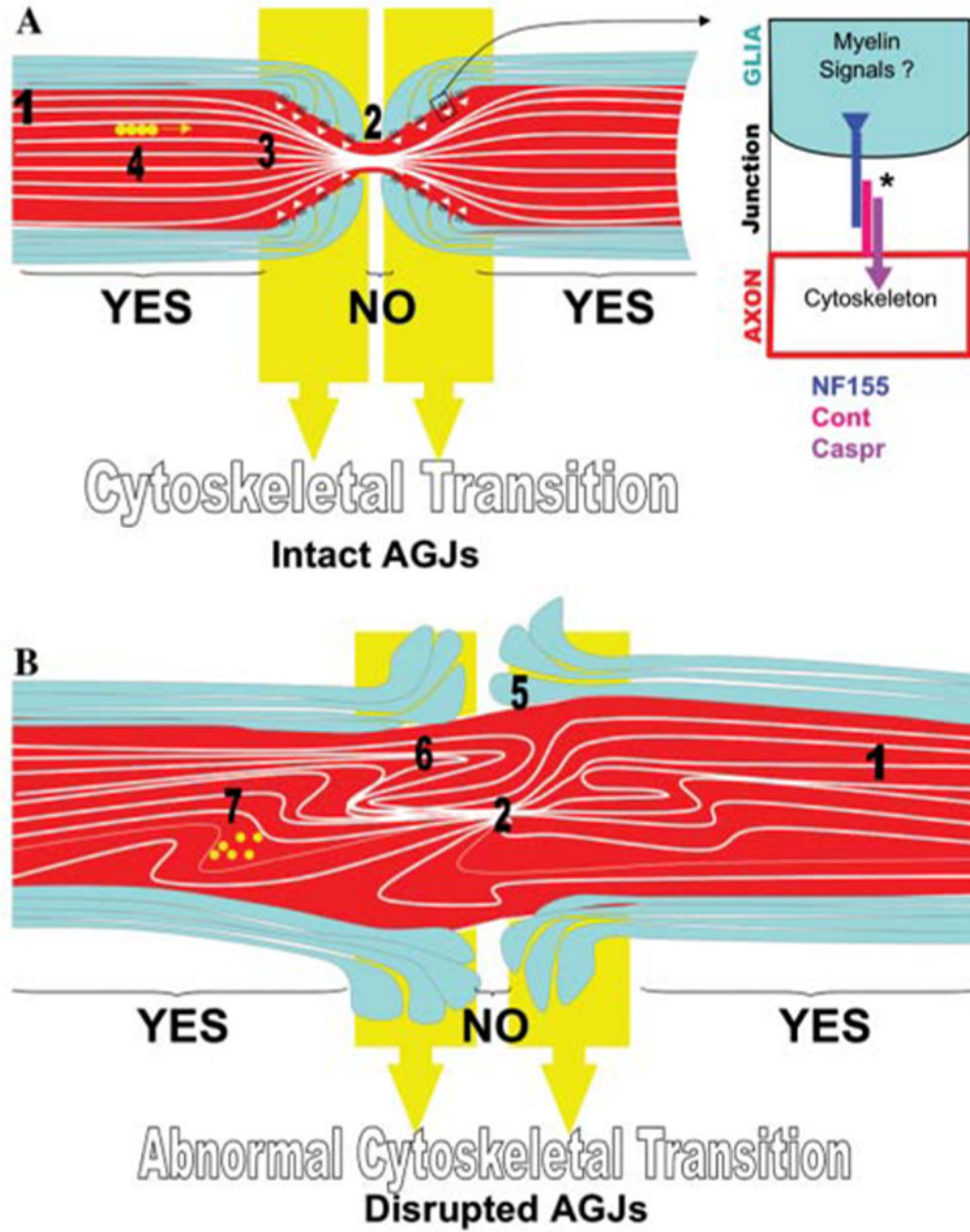


Fig. 3. Model for cytoskeletal transition at the paranodes

(A) Myelination induces phosphorylation of neurofilaments and increases the spacing between neurofilaments at the internodes (1), but not at the nodes of Ranvier where neurofilaments are more tightly packed (2). This periodic signaling along myelinated axons creates what we call the alternating pattern for axonal diameter. The transition between expanded and nonexpanded cytoskeleton happens at the paranodal region (3), where AGJs (white arrowheads) have a role in maintaining this smooth transition, ensuring organization of tracks for axonal transport (4). The inset shows the known molecular components of AGJs that link to the axonal cytoskeleton. Although Caspr is delivered to the paranodal axolemma only when associated with Cont, this association has been shown to inhibit Cont binding to NF155 (Charles *et al.*, 2002; Gollan *et*

al., 2003). * indicates that the association of Caspr to NF155/Cont might be mediated by other proteins. Yellow circles represent organelles that are transported along microtubules. (B) If AGJs are disrupted and the signaling for cytoskeletal expansion at the internodes is still intact, the axonal diameter undergoes alternating pattern (1 and 2) but the transition of diameters is not modulated by AGJs (5). In the absence of proper transition, expanded cytoskeleton invades regions of nonexpanded cytoskeleton, and vice-versa, causing cytoskeletal disorganization at the paranodal region (6) and generation of a site for impairment in transport (7) and potential formation of swellings.

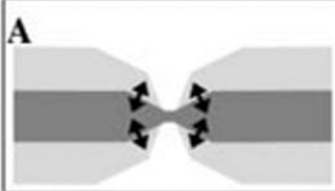
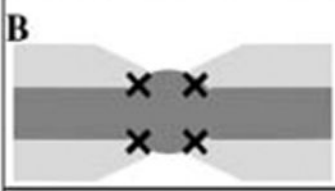
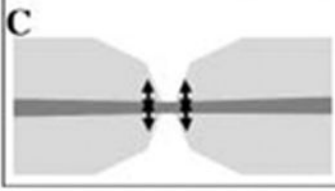
	Mouse Models	AGJ	Axonal Expansion	Swellings
A 	Wild type	Yes	Yes	No
B 	<i>CGT</i> <i>CASPR</i>	No	Yes	Yes
C 	<i>MAG</i> <i>NFL</i>	Yes	No	No

Fig. 4. Correlation between predictions of the cytoskeletal transition model and data from mouse mutants

(A) Normal myelinated axons present myelination-induced axonal expansion at the internodes, intact AGJs and proper transition of cytoskeleton at the paranodes. (B) Disruption of AGJs in axons under proper myelination-induced cytoskeletal expansion leads to disorganized transition of cytoskeleton, impairments in axonal transport and the formation of paranodal swellings, which is consistent with the phenotypes of *CGT* and *Caspr* mutants (Garcia-Fresco *et al.*, 2006). (C) Disruption of signaling in myelination-induced axonal expansion *per se* would not lead to formation of paranodal swellings as in the case of *MAG* and neurofilament mutants (Yin *et al.*, 1998; Elder *et al.*, 2001; Garcia-Fresco *et al.*, 2006). Note that, as discussed for the PLP mutants under the heading ‘Other possible mediators of cytoskeletal transition at the paranodes’, not all the reported paranodal swellings correlate with disruptions of the axon–glial junctions.