

NIH Public Access

Author Manuscript

Exp Neurol. Author manuscript; available in PMC 2007 April 17.

Published in final edited form as: *Exp Neurol.* 2005 September ; 195(1): 208–217.

Myelin-associated glycoprotein and complementary axonal ligands, gangliosides, mediate axon stability in the CNS and PNS: Neuropathology and behavioral deficits in single- and double-null mice

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Abstract

Complementary interacting molecules on myelin and axons are required for long-term axon-myelin stability. Their disruption results in axon degeneration, contributing to the pathogenesis of demyelinating diseases. Myelin-associated glycoprotein (MAG), a minor constituent of central and peripheral nervous system myelin, is a member of the Siglec family of sialic acid-binding lectins, and binds to gangliosides GD1a and GT1b, prominent molecules on the axon surface. Mice lacking the ganglioside biosynthetic gene Galgt1 fail to express complex gangliosides, including GD1a and GT1b. In the current studies, CNS and PNS histopathology and behavior of Mag-null, Galgt1-null and double-null mice were compared on the same mouse strain background. When backcrossed to >99% C57BL/6 strain purity, Mag-null mice demonstrated marked CNS, as well as PNS, axon degeneration, in contrast to prior findings using mice of mixed strain background. On the same background, Mag- and Galgt1-null mice exhibited quantitatively and qualitatively similar CNS and PNS axon degeneration, and nearly identical decreases in axon diameter and neurofilament spacing. Double-null mice had qualitatively similar changes. Consistent with these findings, Mag-, Galgt1null mice had similar motor behavioral deficits, with double-null mice only modestly more impaired. Despite their motor deficits, Mag- and Galgt1-null mice demonstrated significant hyperactivity, with spontaneous locomotor activity significantly above that of wild type mice. These data demonstrate that MAG and complex gangliosides contribute to axon stability in both the CNS and PNS. Similar, neuropathological and behavioral deficits in Galgt1-, Mag-, and double-null mice support the hypothesis that MAG binding to gangliosides contributes to long-term axon-myelin stability.

Keywords

MAG; axon degeneration; axon diameter; neurofilament spacing; motor deficit; hyperactivity

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Introduction

Axon degeneration secondary to dysmyelination is responsible for long-term pathogenesis in demyelinating disorders, such as multiple sclerosis and Charcot-Marie-Tooth disease (Bjartmar et al., 1999; Bjartmar et al., 2003; Coleman and Perry, 2002; Wujek et al., 2002; Bjartmar et al., 2000; Bruck and Stadelmann, 2003; Krajewski et al., 2000; Frei et al., 1999). The relationship between myelination and axon survival is supported by studies of mice lacking certain myelin proteins, in which progressive axonal degeneration is a major phenotype (Bjartmar et al., 1999; Frei et al., 1999). Recognition molecules on the innermost myelin sheath bind to complementary ligands on the apposing axon surface to initiate signaling cascades that control axon physiology (Edgar and Garbern, 2004).

Knowledge of the molecules underlying axon-myelin interactions may reveal insights into axonal degeneration in demyelinating diseases. Among these molecules is myelin-associated glycoprotein (MAG), a minor constituent found on periaxonal myelin in the CNS and PNS (Trapp et al., 1989; Trapp, 1990). Although Mag-null mice myelinate axons, older mice exhibit perturbations in axon integrity (Fruttiger et al., 1995; Weiss et al., 2001). In addition to mild deficits in myelination, Mag-null mice exhibit late-onset progressive PNS axonal atrophy and increased Wallerian degeneration (Bjartmar et al., 1999). MAG also modulates the axonal cytoskeleton. Myelinated axons of Mag-null mice have smaller diameters and fail to exhibit myelin-regulated neurofilament phosphorylation and increased neurofilament spacing, as do wild type mice (Yin et al., 1998). Additionally, MAG, on residual myelin, inhibits axon outgrowth from adult nerve cells after injury (Mukhopadhyay et al., 1994; McKerracher et al., 1994; Filbin, 2003; Yiu and He, 2003; Sandvig et al., 2004). Thus, MAG is required for longterm axon stability, controls axon cytoarchitecture, and regulates axon outgrowth. These effects may be mediated by multiple complementary MAG ligands on the axon, including the Nogo receptor, NgR (Liu et al., 2002; Domeniconi et al., 2002), and gangliosides, the major glycans on nerve cells and axons (Schnaar, 2000) that are the focus of the current studies.

MAG is a member of the Siglec family of carbohydrate binding proteins (Crocker, 2002). Siglecs bind to cell surface glycans that carry a terminal sialic acid residue. We previously identified potential MAG ligands as gangliosides GD1a and GT1b (Fig. 1). These two gangliosides, major sialic acid-containing glycans on nerve cells and axons, bind to MAG *in vitro* (Yang et al., 1996; Collins et al., 1997b; Collins et al., 1997a). Mice engineered to lack the MAG binding glycan determinant on GD1a and GT1b (*Galgt1*-null mice, see Fig. 1) exhibit progressive axon degeneration and other sequelae similar to *Mag*-null mice (Sheikh et al., 1999; Chiavegatto et al., 2000).

We now report results from a genetic approach that compares the functional relationship between MAG and gangliosides *in vivo*. We engineered *Mag*-null, *Galgt1*-null and double-null mice on the same mouse strain background (C57BL/6). Quantitative analysis of neuropathic and behavioral changes reveal remarkable similarities in the phenotypes of the three mouse strains.

Materials and Methods

Animals

Mag-null founder mice, kindly provided by Dr. Bruce Trapp, The Cleveland Clinic Foundation, Cleveland, OH, were constructed by disruption of exon 5 of the *Mag* gene as previously reported (Li et al., 1994). The strain provided (identical to that available from The Jackson Laboratory, Bar Harbor, ME) is a mixture of C57BL/6 and 129 inbred strains and the CD1 random bred strain. Mice with a disrupted *Galgt1* gene, which lack the "NeuAc α 2-3 Gal β 1-3

GalNAc" terminus on gangliosides (Fig. 1), were constructed as described previously (Liu et al., 1999). Founder mice were kindly provided by Dr. Richard Proia, National Institutes of Health, Bethesda, MD. The strain provided was a mixture of C57BL/6 and 129 inbred strains.

To enhance comparisons between mutant strains, mutant mice were repeatedly back-crossed onto a C57BL/6 background. A PCR-based 80-loci full genome screen revealed 99.5% strain purity of the *Mag*-null mice and 94% strain purity of the *Galgt1*-null mice (personal communication, Peter Sobieszczuk, Ph.D., Director, Transgenics Core, Consortium for Functional Glycomics, The Scripps Research Institute, La Jolla, CA). The two strains were crossed to generate double-null offspring. Both back-crossed single-null strains were deposited with the Consortium for Functional Glycomics (www.functionalglycomics.org).

Mice were genotyped at the *Galgt1* locus using PCR with tail-tissue DNA as template as described previously (Sun et al., 2004). The *Mag* locus was similarly genotyped using TGCCGCTGTTTTGGATAA and CGCCTCGGAAATAGTATTTG as the forward and reverse primers, resulting in products of 0.6 and 2 kb for wild type and disrupted alleles respectively.

At the time of tissue harvest for histopathology or behavioral studies, mice averaged approximately 6 months old (ages of each genotype are provided in the figure legends). Most mice used were males; no significant differences between males and females were observed.

Histopathology

Animals from each of *Mag*-null, *Galgt1*-null, double null and wild type C57BL/6 mice were perfused with 4% paraformaldehyde and 1.5% glutaraldehyde in Sorenson's buffer (Morris et al., 1980). Sciatic nerves and lumbosacral spinal cord were harvested from all animals. Tissues were trimmed and postfixed in osmium and embedded in Epon. Thick sections (1 µm) were stained with toluidine blue for light microscopy, and thin sections (80 nm) were examined by electron microscopy.

Axon degeneration—For evaluation of Wallerian degeneration in the PNS, the number of degenerating myelinated nerve fibers per total nerve transverse section was determined using 1- μ m-thick Epon cross sections of sciatic nerve. For evaluation of nerve degeneration in the CNS, 1- μ m-thick Epon transverse sections of the cervical spinal cord were used. Degenerating fibers were determined in the dorsal and ventral funiculus and the pathological features were confirmed on thin sections by electron microscopy.

Axon caliber—Sciatic nerves were analyzed at light and ultrastructural levels, respectively. Thick sections $(1 \ \mu m)$ of sciatic nerve were sampled and nerve fiber counts were obtained at light level by standard stereological methods as described previously (Sheikh et al., 1999; Mayhew and Sharma, 1984; Mayhew, 1988). The mean axonal caliber of myelinated fibers in sciatic nerves was calculated from the diameter of a circle with an area equivalent to that of each axon.

Neurofilament Spacing—Electron micrographic images of myelinated axons of sciatic nerves at a final magnification of 100,000X were used to determine neurofilament spacing. Nearest-neighbor distance was used as a measure of neurofilament spacing in myelinated axons in sciatic nerve as described previously (Yin et al., 1998; Xu et al., 1996). The mean neurofilament nearest-neighbor distance for 15 or more axons from three animals in each group was compared.

Behavior

Mice were individually housed for one week prior to behavioral experiments and were maintained in a 12 h light-dark cycle in a temperature- and humidity-controlled room. The mice were allowed free access to food and water. Mice of each genotype (*Mag*-null, *Galgt1*-null, double-null and wild type C57BL/6) were used for behavioral testing. Spontaneous locomotor activity was tested for a 24-h period starting at 1700 h. All other behavioral tests were conducted between 1300–1800 h. Mice were allowed to acclimate in the testing room for 5 h prior to initiating experiments. All mice were tested on the same day for each behavioral parameter. Care and handling procedures were in accordance with institutional and government guidelines.

Motor coordination and balance—A rotarod apparatus (Economex, Columbus Instruments, Columbus, OH) was used in fixed speed mode. Each mouse was placed in a separate lane of the apparatus on a rotating cylinder (5 cm diameter) at 10 rpm (for up to 10 min) or 5 rpm (for up to 5 min). Their performance was recorded as the mean latency to fall from the cylinder in 3 consecutive trials with a resting period between each trial.

Hindlimb reflex extension—Each mouse was suspended by the tail for 10 s in three consecutive trials. The position of the hindlimbs was scored as follows: 0, one or both hindlimbs paralyzed; 1, loss of reflex and hindlimbs and paws held close to the body with clasping toes; 2, loss of reflex with flexion of hindlimbs; 3, hindlimbs extended to form $<90^{\circ}$ angle; 4, hindlimbs extended to form $\geq 90^{\circ}$ angle.

Beam balance and platform—Each mouse was placed at the center and perpendicular to a rounded wooden beam (60-cm length, 2-cm diameter) suspended 60 cm above a foam pillow. The mean latency to fall (up to 120 s) was registered in three consecutive trials. Subsequently, each mouse was placed at the center of the same wooden beam set as a bridge, and facing one of two platforms (15 x 15 cm) set at either end of the bridge. The mean latency to reach either platform with four paws (up to 60 s) was registered in three consecutive trials. In each trial the mouse faced alternate platforms and a score of 60 s was given to the animal that failed to reach the platform or fell during the test.

Forelimb grip strength—Each mouse was suspended by its forepaws on a rounded metal bar (diameter 3 mm) positioned 60 cm above a foam pillow. The forelimb grip strength was assessed by measuring the mean time (up to 30 s) the mice hung from the bar in three consecutive trials.

Tremor and Catalepsy—Each animal was scored for the presence or absence of whole body tremor (resting or during movement) in an observational plastic cage with acrylic cover in three consecutive observations of 10 s interspersed with an interval of 10 s. The scoring began 1 min after being placed in the test cage. To test catalepsy, each mouse was placed with its hindlimbs on a flat surface and its forelimbs on a horizontal metal bar suspended 5 cm above the surface. The immobility state (time that the animal spent motionless hanging from the bar) was recorded in three consecutive trials. If an animal fell or climbed the bar, it was placed immediately on the bar again; the test was terminated after a maximum of three such escapes.

Spontaneous locomotor activity—Locomotor activity was quantified by placing mice into individual automated Plexiglas boxes (29 x 50 cm), each with 12 2-cm high infrared beam detectors arranged in a 4 x 8 grid (San Diego Instruments; San Diego, CA). Computer-recorded beam breaks were accumulated every 20 min for the duration of a 24-h test period starting at 1700 h, with changes in beam status assessed 18 times/s. Animals were habituated to test cages

for at least 5 h prior to data collection, and had access to food and water ad libitum throughout the experiments.

Results

Histopathology in Mag-null, Galgt1-null, and double-null mice

Axon degeneration—In both the CNS and the PNS, axon degeneration was the predominant pathological feature in mice lacking expression of MAG or complex gangliosides (Figs. 2–4). In comparison to wild type mice, in which essentially no degenerating fibers were seen, singleand double-null mice had a highly significant occurrence of degeneration of myelinated axons, including myelin figures and ovoids representing various stages of degeneration (Figs. 2 & 3). Quantification revealed highly significant axon degeneration in the PNS (sciatic nerve, Figs. 2 & 4) of all mutant strains compared to wild type mice. *Galgt1*-null and *Mag*-null mice displayed significantly greater PNS axon degeneration, whereas double-null mice displayed significantly greater PNS axon degeneration than did either single-null strain. In the CNS (spinal dorsal and ventral columns, Figs. 3 & 4), all mutant strains displayed highly significant axon degeneration compared to wild type mice, with no significant differences among mutant strains. The discovery of axon degeneration in the CNS is in contrast to prior studies of *Mag*-null mice when studied on a mixed-strain background (see Discussion).

Axon caliber and neurofilament spacing—Axon calibers were determined from cross sections of sciatic nerves of wild type and mutant mice (Fig. 5). Histograms reveal a distinct shift to smaller axon calibers in all mutant mice compared to wild type, with *Mag*-null, *Galgt1*-null and double null histograms nearly overlapping. The number of axons \geq 5 µm diameter was significantly reduced (to about half that in wild type mice) in all of the mutant strains, whereas axons \leq 2.5 µm diameter were about double those in wild type (Fig. 5). There were no significant differences among the three mutant strains.

Axon caliber is dependent, in part, on the spacing of axon neurofilaments. Electron micrographs of sciatic nerve cross sections revealed closer neurofilament spacing in *Mag*-null, *Galgt1*-null and double-null mice compared to wild type mice (Fig. 6). Quantitative analysis revealed a significant 25% reduction in neurofilament spacing in all mutant strains compared to wild type mice, with no significant difference among the mutant strains.

Dysmyelination—Dysmyelination was seen in both PNS and CNS of all mutant mouse strains (Figs. 2 & 3), similar to the changes documented previously in *Galgt1*-null and *Mag*-null single-null mice (Sheikh et al., 1999;Montag et al., 1994;Li et al., 1994). These changes appeared to be more prominent in the double null animals.

Behavior of Mag-null, Galgt1-null, and double-null mice

Motor behavior, reflexes, and tremor—Consistent with their histopathology, mutant mice displayed motor and behavioral deficits compared to their wild type counterparts. Rotarod testing, in which mice are timed for their ability to remain on a rotating cylinder, is a combined measure of motor balance, coordination, and muscle control (Kuhn et al., 1995). Compared to wild type mice, all mutant strains were less capable of remaining on a cylinder rotating at 10 rpm, with double-null mice significantly more impaired than *Mag*-null or *Galgt1*-null mice (Fig. 7). This differential in motor coordination was even more apparent when the cylinder was rotating at 5 rpm, a speed at which only double-null mice were significantly impaired compared to wild type mice.

Hindlimb reflex extension was also clearly impaired in all mutant mouse strains (Fig. 8). When a wild type mouse is gently lifted by the tail, it typically extends its hindlimbs outward in a

A marked whole body tremor was recorded in double-null mice (Fig. 9). Wild type mice showed no signs of tremor, whereas double-null mice showed tremors in nearly all trials. Although tremor was detected in a proportion of *Mag*-null and *Galgt1*-null mice, only double-null mice were uniformly affected.

In the beam balance, platform, forelimb grip strength, and catalepsy tests, the mutant mouse strains were not significantly different from wild type mice (data not shown).

Locomotor hyperactivity—Despite motor behavioral deficits, all of the mutant mouse strains were hyperactive, with total spontaneous locomotor activity over a 24-h period increased 60–130% above that of wild type mice (Fig. 10). During the latter half of the dark period, when spontaneous locomotor activity of wild type mice decreased compared to the first half of the dark period (open bar, Fig. 10, top), the activity of mutant strains remained high, resulting in significant hyperactivity of each mutant strain compared to wild type mice during that period (p < 0.03, Student's t-test). When summed over the entire 24-h test period, *Galgt1*-null were 2.3-fold more active than wild type mice, whereas *Mag*-null and double-null mice displayed ~1.6-fold higher activity than wild type mice (Fig. 10, bottom).

Discussion

Qualitative and quantitative similarities in the histopathological and behavioral deficits of *Mag*-null and *Galgt1*-null mice, along with prior data indicating that MAG selectively and efficiently binds to the terminal saccharide sequence on gangliosides GD1a and GT1b (Yang et al., 1996; Collins et al., 1997b; Collins et al., 1997a), support the conclusion that MAG (on myelin) and gangliosides GD1a/GT1b (on axons) function as complementary receptor and ligands enhancing axon-myelin stability. The distribution of MAG on myelin (Trapp et al., 1989) and gangliosides GD1a and/or GT1b on axolemma (DeVries and Zmachinski, 1980) is consistent with this conclusion. The potential relevance of MAG-ganglioside binding to human axon-myelin stability is supported by the observations that human and mouse MAG share >94% amino acid identity, and that the structures of major brain gangliosides (including GD1a and GT1b) are essentially identical among mammals (Tettamanti et al., 1973).

In addition to mild dysmyelination (Li et al., 1994; Montag et al., 1994; Bartsch et al., 1997; Fujita et al., 1998), prior studies demonstrated axon degeneration in the PNS of aging *Mag*-null mice (Fruttiger et al., 1995; Weiss et al., 2001), but notably reported an absence of axon degeneration in the CNS (Lassmann et al., 1997; Uschkureit et al., 2000; Quarles, 2002). In contrast, the current study found clear evidence of CNS, as well as PNS, axon degeneration in *Mag* single-null mice (Figs. 3 & 4). We infer that the genetic background impacts the CNS phenotype. A widely used *Mag*-null mouse strain (Li et al., 1994) (distributed by The Jackson Laboratory), is maintained on a mixture of C57BL/6 and 129 inbred strains and the CD1 random bred strain. A second widely used strain (Montag et al., 1994) was derived by injecting 129 strain ES cells into B6CBAF1 hybrid blastocysts. In the current study, extensive backcrossing of the former strain to C57BL/6 mice resulted in 99.5% strain purity, resulting in clear indications of CNS (as well as PNS) axon degeneration in aging *Mag*-null mice.

The functional relationship between MAG and gangliosides is supported by comparisons of axon degeneration and axon architecture in *Mag*-null and *Galgt1*-null mice. In addition to supporting axon survival, MAG signals increased neurofilament phosphorylation in

myelinated axons (Yin et al., 1998), resulting in increased neurofilament spacing and larger axon caliber. The findings that, compared to *Mag*-null mice, *Galgt1*-null mice have similar occurrence of axon degeneration (Fig. 4) and essentially identical decreases in axon caliber and neurofilament spacing (Figs. 5 & 6), strongly implicate gangliosides in this intercellular signaling pathway.

The observation of close similarities in the behavioral phenotypes of *Mag*- and *Galgt1*-null mice further support a functional relationship between MAG and complex gangliosides. In nearly every behavioral measure where a deficit was detected, *Mag*- and *Galgt1*-null mice had qualitatively and quantitatively similar responses.

MAG is a member of the Siglec family of sialic acid binding lectins, and binds preferentially to the terminal glycan sequence "NeuAc α 2-3 Gal β 1-3 GalNAc" (Crocker et al., 1996). In the brain and peripheral nerves, unlike in other tissues, gangliosides (rather than glycoproteins) are the predominant sialoglycoconjugates, representing >75% of the bound sialic acid in both rodents and humans (Tettamanti et al., 1973; Schnaar, 2000). Furthermore, an abundant terminal sequence on brain gangliosides is the preferred MAG-binding determinant, NeuAc α 2-3 Gal β 1-3 GalNAc (Schnaar, 2000). Based on these observations, it is reasonable to propose that MAG and gangliosides GD1a and GT1b are functional receptor and ligand. The *Galgt1* gene product, UDP-N-acetylgalactosamine:GM2/GD2 β -Nacetylgalactosaminyltransferase, is specific for biosynthesis of glycolipids and does not

transfer GalNAc to glycoproteins (Furukawa et al., 2002). Deleting this gene results in loss of all complex gangliosides and a concomitant increase in the simple gangliosides, GM3 and GD3 ((Liu et al., 1999), see Fig. 1). *Galgt1*-null mice express similar concentrations of brain gangliosides and ganglioside sialic acid as their wild type counterparts, only differing in complexity of the structures expressed (Sun et al., 2004; Wu et al., 2001). We conclude that the phenotypes reported in *Galgt1*-null mice are due to the specific depletion of complex gangliosides.

It should be noted that the *Galgt1* mutation results in loss of all complex gangliosides, not just GD1a and GT1b ((Sun et al., 2004), and see Fig. 1). Although a role for GD1a and GT1b in axon-myelin stability is inferred based on their selective interaction with MAG, one cannot rule out a role for other complex gangliosides in the phenotypes detected. A directed mutation in the sialyltransferase(s) responsible for adding the key terminal sialic acid residue on GD1a and GT1b responsible for MAG binding would provide stronger support for the role of these particular gangliosides in axon-myelin stability. However, there are at least six sialyltransferases capable of transferring sialic acid to the 3-position of galactose (Takashima et al., 2000), and the one(s) responsible for transfer of the terminal sialic acid on GD1a and GT1b has(have) not yet been identified by gene deletion. Until a more directed mutation is achieved, the conclusion that the loss of GD1a and GT1b, in particular, are responsible for the pathologies *Galgt1*-null mice remains a working hypothesis.

If MAG and GD1a/GT1b act via complementary binding to one another, *Mag-Galgt1* doublenull mice would be expected to have a similar phenotype to the single null mice. To a significant extent, this appears to be the case, at least qualitatively. Double-null mice had similar axon degeneration, similarly decreased axon caliber, and similarly decreased neurofilament spacing compared to the single null strains. When behavior was tested, the double-null mice showed similar qualitative deficits in motor coordination, reflexes, and tremor, although to a greater extent than in single-null mice. These observations, combined with prior MAG binding specificity studies (Collins et al., 1997a), are consistent with the conclusion that MAG and complex gangliosides function, at least in part, via complementary interaction. The somewhat more marked behavioral deficits in the double-null mice compared to *Galgt*-null mice may be due, in part, to the presence of GM3 in the *Galgt1*-null mice. GM3 retains some MAG-binding capability, although significantly less than GD1a or GT1b (Collins et al., 1997a). Any residual MAG-interaction capability in *Galgt1*-null mice provided by GM3 would be lost in the double-null mice. However, the more marked behavioral deficits of double-null (and to some extent *Galgt1*-null) mice compared to *Mag*-null mice (Figs. 7–9) indicate that complex gangliosides also function via mechanisms independent of MAG. This is a reasonable hypothesis since all complex gangliosides are ablated in *Galgt1* mice, whereas among the major brain gangliosides only GD1a and GT1b interact with MAG. Therefore, functions of complex gangliosides other than those that interact with MAG (e.g. the major brain gangliosides GM1, GD1b) would be seen only in *Galgt1*-null and double-null mice.

A previously unanticipated shared phenotype of the mutant mice was marked hyperactivity (Fig. 10). Recently, a null mutation in the ganglioside biosynthetic gene responsible for synthesis of GM3 (*SIAT9*, see Fig. 1) in humans was found to result in seizures and progressive neurological degeneration (Simpson et al., 2004). Similarly, *Siat8a/Galgt1* double-null mutant mice, which express only GM3, also display a seizure disorder (Kawai et al., 2001). Prior studies noted that children with central dysmyelination, specifically periventricular leukomalacia (PVL) resulting from neonatal hypoxia in very low birth weight babies, are prone to hyperactivity (Galli et al., 2004). In a recently reported mouse model of PVL, neonatal hypoxia resulted in decreased myelin markers, including MAG, myelin basic protein, and Nogo (among others), presumably due to selective loss of periventricular white matter (Curristin et al., 2002;Weiss et al., 2004). Adult mice that had been subjected to neonatal hypoxia were hyperactive compared to normoxic controls (Weiss et al., 2004). This was interpreted, at least in part, to be due to inappropriate sprouting of axons due to reduction in myelin-derived axon outgrowth inhibitors. Whether similar inappropriate axon sprouting occurs in *Mag-* or *Galgt1*-null mice remains to be tested.

The underlying molecular defects that lead to the documented neuropathological changes in *Mag*- and *Galgt1*-null mice have yet to be fully elucidated. MAG participates in defining the distribution of axon molecules at nodes of Ranvier; the distribution of juxtaparanodal K^+ channels and the cell recognition marker Caspr2 on axons is altered in MAG mutants (Marcus et al., 2002). Additional studies on the roles of MAG and axonal gangliosides in maintaining nodal molecular expression and architecture may be particularly revealing.

Cell surfaces, including those of axons, are dominated by glycan structures. In the nervous system and elsewhere, evidence is mounting that glycan structures serve as recognition molecules that may interact with complementary carbohydrate binding proteins (lectins) on apposing cell surfaces (Taylor and Drickamer, 2003). The data presented here support the hypothesis that axonal gangliosides serve as functional ligands for myelin-associated glycoprotein, providing recognition required for appropriate axon cytoarchitecture and long-term axon-myelin stability. A more complete understanding of the biosynthesis of the key MAG-binding terminal sialic acid determinant on GD1a and GT1b, and the signaling pathways by which MAG-ganglioside binding is translated into enhanced axon architecture and long-term stability, promise to improve our understanding of axon-myelin interactions.

Acknowledgements

The authors thank Carol Rubright and Alene Carteret, Peripheral Nerve Laboratories, for their excellent assistance in tissue preparation and David Fromholt for his gracious help with behavioral studies. This work was supported by National Institutes of Health Grants NS37096 and NS42888, and a research grant from the National Multiple Sclerosis Society.

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Fig. 1.

Biosynthesis of major brain gangliosides. The relationships between major brain gangliosides and their precursors are shown schematically, along with the ganglioside nomenclature of Svennerholm (Svennerholm, 1994). The block in ganglioside biosynthesis due to disruption of the *Galgt1* gene (Liu et al., 1999) is indicated by a vertical double line (the genes responsible for synthesis of GM3 and GD3, *Siat9* and *Siat8a*, respectively, are also indicated). MAG ligands GD1a and GT1b appear at the right, with the key MAG-binding determinant (NeuAc α 2–3 Gal β 1–3 GalNAc) indicated with a dotted line.



Fig. 2.

Neuropathology in sciatic nerve axons of *Mag-*, *Galgt1-* and double-null mice. Light microscopic toluidine blue-stained 1- μ m Epon cross sections of sciatic nerve. Myelinated fibers undergoing axonal degeneration (arrowheads) and tomacula (arrows) in *Galgt1-*null (**B**), *Mag-*null (**C**) and double-null (**D**) nerves compared with wild type nerves (**A**). Bar = 10 μ m.



Fig. 3.

Neuropathology in cervical spinal cords of *Mag-*, *Galgt1-* and double-null mice. Low magnification electron microscopic images of transverse sections of the dorsal funiculus of wild type (**A**), *Galgt1-*null (**B**), *Mag-*null (**C**), and double-null (**D**) mice. Wallerian degeneration (arrowheads), dysmyelinated axons (arrows) and axon swelling (asterisk) are noted. Bar = $2 \mu m$.



Fig. 4.

Increased axon degeneration in *Mag-*, *Galgt1-* and double-null mice. Sciatic nerve (PNS, *top panel*) and spinal dorsal column (CNS, *bottom panel*) cross sections were viewed for evidence of degenerating axons in the following mice (number, average age): *Mag-*null (n=6, 6.3 months), *Galgt1-*null (n=5, 7.4 months), double-null (n=6, 7.9 months), and wild type C57BL/ 6 (n=6, 6 months). Data are presented as means \pm SE. One-way ANOVA for the number of degenerating axons revealed highly significant differences based on genotype (p < 0.01) for both PNS and CNS. Post-hoc analysis (Fisher's PLSD) revealed significant increases (*, p < 0.0001) in the number of degenerating PNS axons in *Galgt1-*, *Mag-* and double-null mice compared to wild type mice, and significantly more degenerating axons in the double-null mice

compared to the single-null mice (\dagger , p < 0.01). Post-hoc analysis of the CNS data indicated highly significant increases in degenerating axons (\$, p < 0.01) in all mutant strains compared to wild type, but no difference among the mutant strains.



Fig. 5.

Decreased axon caliber in sciatic nerves of *Mag-*, *Galgt1-* and double-null mice. Micrographs of sciatic nerve cross sections were subjected to image analysis to determine the diameters of myelinated axons from the following mice (average age): *Mag-*null (6.2 months) *Galgt1-*null (7.4 months), double-null (6.3 months), and wild type (6 months). Diameters of ~500 axons each from 3 mice of each genotype were determined. *Top panel:* Histograms of pooled axon diameters for each genotype. *Bottom panel:* Combined data representing the percent of axons $\geq 5 \mu m$ diameter (black bars) and $\leq 2.5 \mu m$ diameter (gray bars). The data are presented as mean $\pm SE$ (n = 3). One-way ANOVA for the number of axons >5 μm diameter revealed a significant difference based on genotype (p < 0.05). Post-hoc analysis (Fisher's PLSD) revealed significant

decreases (*) in axon caliber for *Galgt1*-null mice (p < 0.05), *Mag*-null mice (p < 0.05) and double-null mice (p < 0.01) compared to wild type mice. Although *Galgt1*-null and *Mag*-null mice demonstrated a trend toward greater numbers of axons $\leq 2.5 \mu$ m diameter (p = 0.08), only double-null mice expressed significantly more of these smaller myelinated axons (†, p < 0.05).



Fig. 6.

Decreased neurofilament spacing in sciatic nerve axons of *Mag-*, *Galgt1-* and double-null mice. *Top panel*: Electron micrographic images of cross sections of sciatic nerves, showing individual neurofilaments and their spacing, in wild type (**A**), *Galgt1-*null (**B**), *Mag-*null (**C**), and double-null (**D**) mice. Bar = 0.2 µm. *Bottom panel*: Nearest neighbor spacing was determined for \geq 15 axons from each of three mice of each genotype (age): *Mag-*null (6.2 months), *Galgt1-*null (7.4 months), double-null (6.3 months), and wild type C57BL/6 (6 months). Values are presented as mean \pm SE. One-way ANOVA for neurofilament spacing was highly significant (p < 0.01). Post-hoc analysis (Fisher's PLSD) revealed that

neurofilaments of *Galgt1-*, *Mag-* and double-null mice had significantly reduced spacing compared to wild type mice (*, $p \le 0.001$).



Fig. 7.





Fig. 8.

Impaired hindlimb reflexes in *Mag*-null, *Galgt1*-null and double-null mice. Wild type mice typically hold their hindlimbs extended steadily at an angle of $\geq 90^{\circ}$ when lifted by the tail. The position of the hindlimbs of wild type and mutant mice were scored on a scale from 0 (most impaired, no reflex) to 4 (normal). The best performance in three trials was averaged for 3–5 mice of each genotype. Data were analyzed using the Kruskal-Wallis nonparametric test and are shown as means for clarity. There was a significant effect of genotype (*, p < 0.01) with all mutant strains impaired compared to wild type mice. Ages and numbers of mice used were as detailed in Fig. 7, legend. [Note: non-parametric statistical analysis was used, precluding the use of error bars.]



Fig. 9.

Tremor in *Mag/Galgt1* double-null mice. All double-null mice displayed severe episodes of body tremor while resting and during movement, whereas no wild type mice displayed tremor under any circumstance. To quantify this observation, whole body tremor was scored (as present or absent) in wild type, *Mag*-null, *Galgt1*-null and double-null mice. Data were analyzed using logistic regression and are shown as a percentage of mice exhibiting tremor for each genotype (n = 3–5). There was a significant effect of genotype with*Galgt1*-null and double-null mice exhibiting significantly more tremor than wild type mice (*, p < 0.05 and p < 0.001 respectively). Although tremor was recorded in a portion of the single-null mice, tremor was a consistent behavior in double-nulls. Ages and numbers of mice used were as detailed in Fig. 7, legend. [Note: non-parametric statistical analysis was used, precluding the use of error bars].



Fig. 10.

Hyperactivity in *Mag*-null, *Galgt1*-null and double-null mice. Spontaneous locomotor activity was determined for 3–6 individual mice of each genotype in two 24-h sessions in 29 x 50 cm Plexiglas boxes, each fitted with a grid of infrared beams. *Top panel:* The number of beam breaks per interval was averaged for mice of each genotype as a function of time. Data were smoothed over five data points (100 min). The solid bar designates the 12-h dark cycle. Over the time period constituting the latter half of the dark cycle (open bar), each mutant mouse strain was significantly more active than wild type mice (*Mag*-null, p < 0.01; *Galgt1*-null, p < 0.001; double-null, p < 0.03, pairwise Student's t-tests). *Bottom panel:* The average total beam breaks over the entire 24-h test period was determined for mice of each genotype. Data are

presented as mean \pm SE. One-way ANOVA for total activity was highly significant (p < 0.01). Post-hoc analysis (Fisher's PLSD) revealed that both *Galgt1*-null mice (p < 0.001) and*Mag*-null mice (p < 0.05) were significantly more active (*) than wild type mice over the entire test period, with double-null mice trending in the same direction (p = 0.06). The following mice were used (number, average age): *Mag*-null (n=6, 5.4 months); *Galgt1*-null (n=3, 6.9 months); double-null (n=5, 4.9 months); and wild type C57BL/6 (n=4, 6.5 months).