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The BMP co-receptor *RGMb* promotes while the endogenous BMP antagonist Noggin reduces neurite outgrowth and peripheral nerve regeneration by modulating BMP signaling

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Abstract

Repulsive guidance molecule b (*RGMb*) is a bone morphogenetic protein (BMP) co-receptor and sensitizer of BMP signaling, highly expressed in adult dorsal root ganglion (DRG) sensory neurons. We used a murine *RGMb* knockout to gain insight into the physiological role of *RGMb* in the DRG, and address if *RGMb*-mediated modulation of BMP signaling influences sensory axon regeneration. No evidence for altered development of the peripheral and central nervous systems was detected in *RGMb* ^{-/-} mice. However, both cultured neonatal whole DRG explants and dissociated DRG neurons from *RGMb* ^{-/-} mice exhibited significantly fewer and shorter neurites than those from wildtype littermates, a phenomenon that could be fully rescued by BMP-2. Moreover, Noggin, an endogenous BMP signaling antagonist, inhibited neurite outgrowth in wild type DRG explants from naïve as well as nerve injury-preconditioned mice. Noggin is downregulated in the DRG after nerve injury, and its expression is highly correlated and inversely associated with the known regeneration-associated genes, which are induced in the DRG by peripheral axonal injury. We show that diminished BMP signaling *in vivo*, achieved either through *RGMb* deletion or BMP inhibition with Noggin, retarded early axonal regeneration after sciatic nerve crush injury. Our data suggest a positive modulatory contribution of *RGMb* and BMP signaling to neurite extension *in vitro* and early axonal re-growth after nerve injury *in vivo* and a negative effect of Noggin.

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Introduction

Cell specification, differentiation, proliferation, patterning and migration during vertebrate development are profoundly influenced by transforming growth factor beta (TGF β) superfamily ligands, particularly bone morphogenetic proteins (BMP) (Balemans and Van Hul, 2002; Wu and Hill, 2009). These diverse functions require both tight spatiotemporal control of ligand production and activation of specific intracellular signaling pathways by type I and II serine-threonine kinase receptors. BMPs direct skeletal patterning, chondrogenesis and bone formation in the embryo (Babitt et al., 2006; Rosen, 2006), and in the nervous system both act as instructive signals for neuronal lineage commitment and promote neuronal differentiation (Liu and Niswander, 2005).

BMP signaling is greatly enhanced in those cells that express the glycosyl-phosphatidylinositol (GPI) anchored repulsive guidance molecule a (RGMa), RGMb, and RGMcBMP co-receptors, sensitizing these cells to low levels of ligand (Severyn et al., 2009). RGMb (also termed “Dragon”) is a 436 amino acid membrane-bound protein expressed in the adult in a wide range of neural sites including the DRG, spinal cord, optic nerve, retina, and several regions of the brain (Niederkofler et al., 2004; Oldekamp et al., 2004; Samad et al., 2004; Severyn et al., 2009), as well as, the reproductive tract (Xia et al., 2005) and kidney (Xia et al., 2010b). RGMb potentiates BMP signaling via direct binding to BMP-2 and BMP-4, and subsequent association with type I and II BMP receptors (Samad et al., 2004). RGMb is expressed in developing mouse bone and nervous system (Samad et al., 2004; Samad et al., 2005) and in the branchial arches, somites and tail bud of developing mouse and *Xenopus* (Samad et al., 2005). While the role of RGMc (also known as hemojuvelin) in regulating iron homeostasis is well established (Babitt et al., 2006; Niederkofler et al., 2005), less is known about the roles of RGMa and RGMb in the nervous system. RGMa is suggested to contribute to neuronal differentiation and axonal guidance (Matsunaga et al., 2006; Liu et al., 2009; Severyn et al., 2009; Conrad et al., 2010) however, RGMa deficient mice do not display alterations in neuronal patterning and retinal projection topography (Niederkofler et al., 2004).

Several lines of evidence suggest that BMP-signaling may be involved in neuronal regeneration. Specifically, BMP receptors, ligands, and downstream effectors are expressed in the PNS, and there is an injury-induced regulation of these BMP-signaling pathway components (Ai et al., 1999; Tsujii et al., 2009; Zou et al., 2009). Interestingly, BMP signaling appears to have differential effects on neurite extension and axonal growth in the central (Matsuura et al., 2007; Matsuura et al., 2008; Liu et al., 2009) and peripheral (Hodge et al., 2007; Moon and Birren, 2008; Tsujii et al., 2009) nervous systems, BMP pathway activation appears to be predominantly inhibitory in the CNS and permissive in the PNS.

Here we show that deletion of the *RGMb* gene in mice or the use of the endogenous BMP signaling antagonist Noggin, decrease BMP-mediated signaling in primary sensory neurons and alter their axonal regenerative response to injury both *in vitro* and *in vivo*. These findings provide further support for BMP-signaling as a growth promoting pathway in the PNS following injury, and insight into the contribution of positive and negative BMP-signaling modifiers in this process.

Materials and Methods

Animals

All surgical procedures were performed in accordance with Massachusetts General Hospital and Children’s Hospital Animal Care guidelines. *RGMb* KO and WT littermate mice of either sex at postnatal day 8–9 were used in nerve crush and regeneration studies. In the

Noggin experiment on adult mice, C57BL/6 adult male mice were obtained from Charles River Laboratories (Wilmington, MA).

Generation and phenotypes of *RGMB* knockout (KO) mice

RGMB KO mice (C57BL/6/129) lacking exon II of the *RGMB* gene were generated and were genotyped as described (Xia et al., 2010a). Heterozygous animals were bred to obtain homozygous *RGMB* KO mice; this was necessary as *RGMB*-null animals die at approximately 21 days of age. Histological analysis and magnetic resonance imaging of an *RGMB* KO mouse and its WT littermate (LM) failed to reveal any gross differences in brain or ventricular structure (data not shown). The KO mice display less pulmonary alveolarization, as well as, delayed eruption and poor root formation of the teeth (data not shown).

In situ hybridization and immunohistochemistry

Tissue was rapidly removed, embedded in Tissue-Tek OCT and frozen. Sections were cut serially at 12 μ m, and *in situ* hybridization was performed using digoxigenin-labeled antisense and sense riboprobes according to published methods (Blackshaw and Snyder, 1997). *RGMB* primers were as follows: 5'-TGCCAACAGCCTACTCAATG-3' (*RGMB* exon 1-3) and 5'-GACCAACAGGCAAATCCCTA-3' (*RGMB* exon 1-3). We detected the hybridized riboprobes using Fab fragments from an anti-digoxigenin antibody conjugated to horseradish peroxidase (Roche). Visualization was conducted using 5-bromo-4-chloro-3-indolyl-phosphate (BCIP; Roche) and 4-nitro blue tetrazolium chloride (NBT; Roche) in a solution containing 0.1 M Tris pH 9.5, 0.1 M NaCl and 50 mM MgCl₂. The probes for Ret, CGRP, Trk A, Trk B, Trk C, Ngn1, Math1 and Mash1 have been described previously (Ma et al., 1997; Abdel Samad et al., 2010). Trk A or neurofilament (NF-200) immunostaining was performed on the same section after *in situ* hybridization according to published methods (Brenner et al., 2004). Sections were blocked with 4% donkey serum/0.1% Triton in PBS, and incubated with anti-Trk A (rabbit polyclonal, 1:1000, Chemicon) or anti-NF-200 (mouse monoclonal, 1:2500, Abcam) antibody. Following primary antibody incubation, sections were incubated with Cy3-conjugated secondary antibody (1:300, Jackson Lab).

Primary DRG culture

P1 DRG cultures were prepared (Benn et al., 2002) and cultured in Neurobasal (Full NB) media supplemented with B27 (Invitrogen), 50ng/ml NGF(Invitrogen), 2ng/ml GDNF(Sigma), 10 μ M Ara-C, and 200mM L-glutamine. Dissociated DRG neurons were plated onto poly-D-lysine-laminin coated 8-well chamber slide (Lab-Tek) for 24 hr, washed once and replaced with fresh NB medium with indicated concentrations of BMP-2. After an additional 24 hr, neuronal cultures were fixed with 4% paraformaldehyde and immunostained with both anti- β -tubulin III (rabbit monoclonal, 1:1000, Sigma) and anti-pSmad1/5/8 (rabbit polyclonal, 1:1000, Cell Signaling) for pSmad induction studies. pSmad expression was analyzed by using Image J (NIH) to assess signal intensity. For neurite outgrowth assay, neurons were cultured with BMP for 17 hr, immunostained with anti- β -tubulin III, and the longest neurites were measured using Image J (NIH). Data were obtained from at least three separate experiments.

Western blot

Cultured DRG neurons were treated with 100ng/ml BMP-2 for 1 hr and were sonicated in a lysis buffer, and Western blot was performed as described (Samad et al., 2004). Briefly, protein samples were separated on a 4-15%SDS-PAGE gel (Bio-Rad) and transferred to a polyvinylidenedifluoride filter (Immobilon-P; Millipore). The blots were blocked with 6% non-fat milk and incubated with anti-*RGMB* antibody (rabbit polyclonal, 1:6000, produced in

our laboratory as previously described (Samad et al., 2004)). The blots were incubated with HRP-conjugated anti-rabbit secondary antibody (1:2000; Amersham Biosciences), developed in ECL solution (PerkinElmer Life Sciences) and exposed to film.

DRG explant culture

DRG from P9 *RGMB* KO and WT (Fig 3B) or 8 weeks old C57Bl/6 (Fig 5A) mice were cleaned of spinal and peripheral roots and plated onto 8-well chamber slide dish (Lab-Tek) coated with thin layer of Matrigel (BD Biosciences). DRG were cultured in Full NB media and 50ng/ml BMP (Fig 3B) or Noggin 2 μ g/ml (Fig 5A) as indicated. After 48hr, DRG were fixed with 4% paraformaldehyde, washed and then blocked with 2% goat serum/0.3% Triton, and immunostained with anti- β -tubulinIII antibody (rabbit monoclonal, 1:500, Sigma). Non-overlapping quadrant images were taken at low magnification (4 \times) for analysis. The total length of neurite attached to the DRG explants was measured using WIS-NeuroMath (Weizmann Institute of Science).

Sciatic nerve crush

Nerve injury and the *in vivo* measurement of regeneration were performed under isoflurane anesthesia. Crush injury was performed with smooth forceps (5/45, Fine Scientific Tools) which were used to clamp the sciatic nerve for 30s at the level of the proximal thigh. The crush site was marked with a 10-0 nylon suture knotted to the epineurium.

Measurement of axonal regeneration

The sciatic nerve pinch test was used to determine the rate nerve regeneration *in vivo* (Sjoberg and Kanje, 1990; Mills et al., 2005). At 24 or 48 hr post sciatic nerve crush injury animals were anesthetized with 2.5% isoflurane anesthesia and the left sciatic nerve was exposed. Animals were then brought to a lighter plane of anesthesia (0.8% isoflurane) for measurement of regeneration. Starting distally, a series of pinches using fine smooth forceps (5/45) were delivered to the sciatic nerve, moving proximally toward the injury site. We determined the rate of regeneration by measuring the distance from the injury site, marked with an epineural suture, to the most distal point on the nerve that produced a reflex withdrawal response when pinched. The entire procedure was conducted using a dissecting microscope. For experiments comparing *RGMB* KO mice to LM controls, an individual blind to the pinching assessed reflex withdrawal. GAP-43 fluorescent immunostaining was utilized for histological characterization of regeneration following crush injury. GAP-43 expression is highly induced in regenerating fibers in the sciatic nerve following injury (Seijffers et al., 2007). Two days after crush injury sciatic nerves were removed and fixed. 10- μ m longitudinal sections were immunostained for anti-GAP-43 (rabbit polyclonal, 1:1000, Chemicon), images captured, and the maximal regeneration (as indicated by GAP-43 fiber staining distal to the crush site) was determined.

Peripheral nerve injections

Male C57BL/6 mice were anesthetized with 2.5% isoflurane and the left sciatic nerve was exposed at the mid-thigh level. Noggin or vehicle was injected directly into the sciatic nerve through a finely-pulled glass pipette using a Narishige IM300 Micro-injector. A total volume of 1 μ l was used for noggin (0.5 or 1 μ g, Fig 5B; 2 μ g, Fig 5C) and vehicle injections, all of which were performed blind to substance injected. 60s following injection, sciatic nerve crush was performed as described above.

Bioinformatic Analysis

Weighted gene co-expression network analysis was performed using methods detailed in (Oldham et al., 2006; Oldham et al., 2008) with expression profiling data detailed in (Lewis

et al., 1999; Ma et al., 2011). Primary expression array data is deposited in Gene Expression Omnibus (NCBI) GSE30691. Briefly, after selecting genes present in at least five samples, the absolute Pearson correlation coefficients between one gene and every other screened gene were computed, weighted and used to determine the topological overlap, a measure of connection strength, or ‘neighborhood sharing’ in the network. A pair of nodes in a network is said to have high topological overlap if they are both strongly connected to the same group of nodes. In gene networks, genes with high topological overlap have been found to have an increased chance of being part of the same tissue, cell type or biological pathway. Network Neighborhood Analysis provides a set neighborhood for an initial seed or node. Using Noggin (Affymetrix probe U31203) as the chosen seed, the top 64 nearest neighbors (65 genes total) were selected using topological overlap as a measure of connection strength. Visual C++ implementation of the multinodeTOM software can be found at <http://www.genetics.ucla.edu/labs/horvath/MTOM/>.

Statistics

Analysis was conducted either by Student’s t-test or by one-way ANOVA, the latter with *post-hoc* analysis via Newman-Keuls test as appropriate.

Results

***RGMb* deletion alters BMP-mediated Smad1/5/8 phosphorylation in DRG neurons**

RGMb expression starts early in development (< E12.5) and is maintained in adult mouse DRG, spinal cord, and brain (Samad et al., 2004; Niederkofler et al., 2004). In the DRG, *RGMb* is highly expressed in a subset of small (C-fiber) and large diameter (A-beta and A-delta) neurons as suggested by its partial colocalization with the DRG sensory neuron molecular markers TrkA, Peripherin and Neurofilament 200 – expressed in 33%, 62%, and 20% of *RGMb* positive DRG neurons, respectively (Fig 1A).

To investigate the role of *RGMb* in the nervous system, we first determined whether *RGMb* mutation in mice affects neuronal specification in the DRG. We examined the expression of several well-characterized sensory neuron subpopulation markers (Ret, CGRP, TrkA, TrkB, and TrkC) and found no detectable postnatal difference in the phenotype of sensory neurons between *RGMb* mutant and wild-type mice (Fig. 1B). Since *RGMb* is also expressed in the spinal cord, we looked and found no difference in expression of the neurogenic transcription factors Mash1, Math1 and Ngn1 (Ma et al., 1997; Abdel Samad et al., 2010) in the developing spinal cord (E12.5) between *RGMb* mutant and wild-type (WT) mice (data not shown). These data indicate that *RGMb* deletion does not appear to affect early neuronal specification of DRG sensory neuron subpopulations.

To test whether lack of *RGMb* expression in DRG alters BMP-mediated signaling in sensory neurons, we exposed cultured neonatal DRG neurons from P14 *RGMb* KO mice and WT (littermates) LMs to varying concentrations of BMP-2 and assayed for induction of phospho-Smad1/5/8. Smad1/5/8 proteins, which are activated by phosphorylation, are the primary downstream effectors of BMP receptor activation (Liu and Niswander, 2005). We observed a dose-dependent effect on BMP-2 mediated Smad phosphorylation in cultured DRG neurons from *RGMb* KO mice compared with WT LMs (Fig 2A, B). pSmad induction in *RGMb* KO DRG neurons was significantly reduced at 5 ng/ml (6-fold) and 50 ng/ml (3-fold) of BMP-2, but was not different at higher concentrations of BMP-2 (200 ng/ml), when compared to BMP-2 action on neurons from *RGMb* WT littermates. Consistent with these immunohistochemical data, a marked induction of pSmad by BMP2 was also observed by Western blot in cultured WT DRG neurons (Fig 2C). Our data indicate that *RGMb* expression in DRG neurons enhances signaling of BMP ligands, and support a role for

RGMb as a BMP sensitizer in sensory neurons, similar to the role of hemojuvelin (RGMc) – another member of the RGM family - in hepatocytes (Babitt et al., 2006).

RGMb deletion and inhibition of BMP signaling by Noggin reduce neurite formation in DRG neurons *in vitro*

Smad1, a key intracellular effector required for BMP-2 and -4 signaling, is required for the initiation of sensory neuron axonal regeneration following injury in adult mice (Zou et al., 2009). To test whether RGMb, which can potentiate BMP signaling and hence Smad activation, enhances axonal growth in DRG neurons, we prepared both dissociated DRG neuronal cultures and DRG explants from *RGMb* mutant mice and their WT LMs. The dissociated neuron cultures allow assessment of neurite formation in individual cells, while the cultured whole explanted DRG maintain important interactions between resident neurons and supporting cells. Dissociated DRG neurons from *RGMb* KO mice developed shorter neurites, as compared with LM controls ($p < 0.001$) (Fig 3A). The addition of BMP-2 to cultured *RGMb*-deficient DRG neurons fully restored average neurite length; there was no significant difference of average length between *RGMb* KO+BMP-2 and WT LM (Fig 3A). Analogous results were obtained using cultured whole DRG explants. *RGMb* KO mice exhibited 50% less neurite outgrowth than explants from WT LMs, and more importantly, this loss of function could be fully rescued by addition of exogenous BMP-2 (Fig 3B, KO vs WT $p < 0.001$; KO vs KO-BMP $p = 0.001$). There was no significant effect of the addition of BMP-2 to dissociated WT DRG neurons or to WT DRG explants (Fig 3). The apparent lack of effect of BMP-2 on cultured WT DRG neurons suggests that there may be a low ceiling tonic BMP signaling effect on neurite outgrowth *in vitro* such that additional BMP ligand has no further effect. To further test whether the reduced regenerative capacity in DRG explants from *RGMb* deficient mice is due to decreased BMP signaling, we cultured wild type naïve and preconditioned (i.e., 48 hrs post sciatic nerve crush) adult mouse DRG explants with and without the endogenous BMP-inhibitor, Noggin (Fig 5a). Exposure to Noggin caused a 60 % decrease in neurite formation in both naïve and preconditioned DRG explants ($p < 0.01$, $n = 8$ per group) (Fig 5a, b). Overall, these data show that *RGMb* expression enhances BMP-dependent neurite outgrowth in DRG neurons *in vitro* and that Noggin reduces neurite outgrowth.

RGMb deficiency and inhibition of BMP signaling by Noggin lead to slower axonal regeneration following sciatic nerve injury

We next assessed whether *RGMb* deficiency alters axonal regeneration *in vivo* after sciatic nerve injury. We used the sciatic nerve crush model for assessment of axonal regeneration in sensory neurons (Sjoberg and Kanje, 1990). Axonal regrowth in the sciatic nerve following crush injury was assayed both via immunohistochemical evaluation of GAP-43 expressing fibers (Seiffers et al., 2006) and by a functional measure, the pinch test (Seiffers et al., 2007). As a control, we assessed axonal density in the sciatic nerves of *RGMb* WT and KO mice and found no significant difference (KO/WT ratio of neurofilament 200 positive fibers: 0.93 ± 0.072 , $P = 0.568$, $n = 9$ sections per group, 3 animals per genotype), thus suggesting that there are no major defects in axonal development in the PNS. GAP-43 expression revealed an average of 1.3 ± 0.1 mm regrowth in *RGMb* KO mice compared to 2.5 ± 0.2 mm in WT LM controls 48 hours following crush injury (Fig 4A shows representative images; $p < 0.01$, $n = 3$ per group). The pinch test similarly demonstrated delayed axonal regeneration in *RGMb* KO animals compared with their WT LMs; Two-way ANOVA of these data indicated a significant one-way effect of *RGMb* genotype (KO vs WT, $F(1,13) = 35.8$, $p < 0.001$) and of time post crush (24 vs 48 hours, $F(1,13) = 47.9$, $p < 0.001$), but no significant interaction between the two (Fig 4B, $n = 8$ and 9 for KO and WT, respectively). Of note, there was no significant difference in regeneration between *RGMb* heterozygous (mean $3.3 \text{ mm} \pm 0.3 \text{ mm}$, $n = 5$) and WT ($2.7 \text{ mm} \pm 0.1 \text{ mm}$, $n = 5$) mice at 48 hours ($p = 0.25$).

The diminished axonal regrowth in *RGMb* KO mice was not due to DRG cell death as there is no detectable apoptosis in DRG from *RGMb* KO mice 2 and 7 days after sciatic nerve crush *in vivo* as assessed by TUNEL staining (data not shown). Our *in vitro* data suggest that the reduced regenerative capacity of DRG neurons and explants from *RGMb* deficient mice is due to decreased BMP signaling. To test this hypothesis *in vivo*, we injected the sciatic nerve of WT adult mice with 0.5 or 1 μ g of Noggin immediately prior to a sciatic nerve crush. Assessment of regeneration using the pinch test 48 hours after crush injury revealed a significant inhibition ($p < 0.05$, $n = 6$) associated with Noggin treatment (at both doses) compared with vehicle-injected control mice (Fig 5B). A similar effect of Noggin injection (2 μ g) on *in vivo* axonal regeneration was found at 4 days post crush (Vehicle: $6.55\text{mm} \pm 0.3$; Noggin: $3.08\text{mm} \pm 0.72$; $p < 0.01$, $n = 6$). These data suggest that BMP signaling is required for axonal regrowth following sciatic nerve crush, and that *RGMb* as a BMP signaling enhancer, plays a pivotal role in the BMP effect on early axonal regeneration.

Noggin expression after peripheral nerve injury negatively correlates over time with the expression of a cluster of regeneration associated genes

We explored bioinformatically, using a network neighborhood analysis (NNA) (Oldham et al., 2006; Oldham et al., 2008), how the pattern of Noggin expression after peripheral nerve injury in the DRG correlates over time with the expression of ~1200 genes whose expression alters significantly due to nerve injury (Lewis et al., 1999). Of the 65 most closely related genes identified by this analysis (Fig 6), including Noggins seed, only one other gene was down regulated (*Lin7b*) and 97% were upregulated (63 genes). Remarkably, 29 of 30 genes whose expression across multiple data sets most closely resembled the expression pattern of *Hspb1*, a newly defined regeneration associated gene (RAG) (Ma et al., 2011) were present in this list. Of the *Hspb1* related genes present 18 have been shown in the literature to increase neurite formation (Ma et al., 2011). These RAGs are *Arg1*, *Atf3*, *C1qb*, *Camk2d*, *Crabp2*, *Fabp5*, *Gal*, *Gap43*, *Gfra1*, *Gpc1*, *Hspb1*, *Jun*, *Ptpn5*, *Sox11*, *Spr1a*, *Spr1al*, *Stmn4*, and *Tgm1*. The inverse correlation between the expression of Noggin, a negative modulator of BMP signaling, and that of a group of RAGs further supports our findings with respect to the contribution of BMP signaling to axonal regeneration after peripheral nerve injury.

Discussion

RGMb was initially identified in embryonic DRG, and was the first member of the RGM family shown to potentiate BMP signaling by acting as a BMP co-receptor (Samad et al., 2005). *RGMb* binds directly BMP ligands, associates with BMP type I (*ALK2*, *ALK3*, and *ALK6*) and type II (*ActRII* and *ActRIIB*) receptors, and its signaling is reduced by dominant negative *Smad1* and *ALK3* or -6 receptors (Samad et al., 2005). The physiological role of *RGMb in vivo* and its functional implications as a BMP sensitizer in DRG neurons have not, however, been elucidated, until now.

Several lines of evidence suggest that BMP signaling plays a pivotal role in neuronal development; genetic mouse models with null mutations of genes encoding BMPs, BMP receptors, and their downstream signal transducers indicate such a role at several developmental stages, especially for neural patterning (Zhao, 2003). *RGMb* is widely expressed in the nervous system, mostly in regions complementary to and non-overlapping with *RGMa* (Oldekamp et al., 2004; Samad et al., 2004; Niederkofler et al., 2004). Both, *RGMa* and *RGMb*, act as BMP co-receptors, and their expression levels are upregulated after nerve injury (Schwab et al., 2005). These common patterns suggest potential involvement in comparable physiological roles. Interestingly, while *RGMa* inhibition has been shown to promote axonal growth and recovery after spinal cord injury (Hata et al., 2006), *RGMa* expression was shown to promote retinal ganglion cell survival (Koeberle et

al., 2010). In addition to *RGMs*, BMP-2, -4 and -7 as well as their downstream effector pSmad are upregulated after spinal cord injury (Xiao et al., 2010). Increased levels of pSmad1/5/8 in neurons, astrocytes and oligodendrocytes following injury, indicates that BMP signaling pathways are likely involved in the cellular response to CNS injury. The selective neural expression pattern of *RGMB* and its high levels in developing and adult DRG neurons led us to assess the role of this BMP co-receptor in the development and growth of sensory neurons.

We utilized an *RGMB* deficient mouse line to study the role of *RGMB* in DRG neurons and its contribution to axonal growth following injury. We found no evidence for a major developmental defect in sensory neurons in *RGMB* KO mice, and our data show that deletion of *RGMB* leads to impaired DRG neurite outgrowth *in vitro* and axonal regeneration *in vivo*. These data are in keeping with previous studies showing BMP involvement in these phenomena (Tsuji et al., 2009; Zou et al., 2009). Injury-induced axonal growth was shown to require the function of the BMP signaling effector Smad1. After peripheral axotomy, neuronal Smad1 is upregulated, and phosphorylated Smad1 accumulates in the nucleus. Both events precede the onset of axonal extension (Zou et al., 2009). We show here that *RGMB*, as a BMP coreceptor, contributes to PNS regeneration after nerve injury via the BMP signaling pathway. Since the nerve injury experiments were conducted at P8 in the mutant mice, an age at which the PNS is still developing (Court et al., 2004), we looked at the effect of Noggin in the adult to address a possible contribution of a developmental disorder in the postnatal *RGMB* mutant mice. We show that Noggin functions as a negative regulator of axonal regeneration in the PNS; it inhibits neurite outgrowth in embryonic and adult DRG neurons and explants *in vitro*, and reduces adult peripheral nerve regrowth after crush *in vivo*.

Although *in vivo* intra-DRG injection of BMP2 was shown to promote axonal growth (Zou et al., 2009), we found that increasing BMP signaling above basal levels did not promote increased neurite outgrowth in WT DRG neurons *in vitro*, suggesting that the endogenous levels of BMP provided under our experimental conditions *in vitro* are sufficient for enabling normal axonal growth and no further growth is produced beyond this ceiling effect.

Lack of *RGMB* in KO mice diminished basal and BMP-induced levels of Smad1/5/8 phosphorylation in DRG neurons, reaffirming the role of this co-receptor as a BMP ligand sensitizer. Our data suggest that the reduction of BMP signaling resulting from lack of *RGMB* expression reduced neurite outgrowth *in vitro* in naïve and in injury-preconditioned DRG neurons and explants, by decreasing the responsiveness of the neurons to endogenous BMP. *In vivo*, *RGMB* mutation led to reduced axonal regrowth after a sciatic nerve crush injury. That this is likely due to a deficit in BMP signaling is supported by our finding that exogenous administration of the BMP antagonist, Noggin (Smith, 1999) also impaired axonal regeneration *in vitro* and *in vivo*. Further, Smad1 phosphorylation and subsequent activation has been implicated in mediating BMP signaling from axon terminals (Hodge et al., 2007), and Smad1 knockdown in adult DRG neurons reduces axonal growth (Zou et al., 2009).

Interestingly, Noggin expression is down-regulated in the DRG after nerve injury, as opposed to the upregulation observed for many RAGs. This paradox, due to Noggin's pattern of regulation following nerve injury, closely resembling an inverse plot of the upregulated transcripts that it is grouped with in the NNA (Oldham et al., 2006; Oldham et al., 2008), corresponds well with our finding that Noggin as a BMP antagonist negatively regulates neurite formation while induction of the other RAGs increase regeneration. It is entirely possible that other genes whose expression correlates with Noggin may also be therefore RAG genes. The action of *RGMB* as a co-receptor sensitizing BMP-2 signaling

through type I and II BMP receptors is analogous to the role of GFRa1 (a co-receptor for RET) which sensitizes GDNF signaling and therefore axonal growth within the injured PNS (Mills et al., 2007), interestingly GFRa1 is one of the RAGs inversely associated with Noggin expression. Collectively, these data point to a possible co-regulation of the many positive and few negative regulators of axonal growth by common master transcription factors, and that Noggin is the first documented example of the latter and suppression of its expression after injury might contribute to regenerative capacity.

The specific molecular mechanisms by which BMP receptor activation and downstream Smad signaling regulate axonal growth remain poorly understood. Furthermore, a number of BMP-mediated Smad independent mechanisms have recently emerged as a potential modulators of axonal growth and elongation (Hocking et al., 2009). The actin cytoskeleton in extending axonal and dendritic processes undergoes continuous assembly and disassembly. Extrinsic factors modulate actin turnover through controlling the activity of LIM kinase 1 (LIMK1), which phosphorylates and inactivates the actin depolymerizing factor cofilin. BMPRII is a modulator of LIMK1, and via such an interaction BMPs were shown to induce positive and negative responses in growth cones. BMP ligand binding to the receptor complex modulates this inhibitory interaction and activates Rho GTPase, Cdc42 and LIMK1 (Sanchez-Camacho and Bovolenta, 2009). Further supporting a Smad independent pathway, expression of a BMPRII mutant lacking the LIMK1 binding region was shown to alter axonal growth (Hocking et al., 2009). Whether the effect of BMP signaling on axonal growth that we observe in *RGMb* deficient mice is mediated by Smad1 signaling or LIMK1 activation will require additional analysis.

Our data suggest that *RGMb* plays a pivotal role in contributing to a tonic level of BMP signaling in DRG neuron that is necessary for normal axonal regrowth. While the neurotrophic properties for BMPs are well established, the tissue distribution and cellular source of these ligands in the peripheral nervous system *in vivo* have not yet been fully established. In addition to neurons, BMPs (e.g. BMP-2 and -7) and their receptors are expressed in Schwann cells in the sciatic nerve (Tsuji et al., 2009). Schwann cells represent a major source of trophic factors supporting intact and injured neurons. Their responsiveness to BMP ligands represents, therefore, a pathway by which BMPs may modulate neuronal survival and growth indirectly. Interestingly, Schwann cells also express *RGMb* (data not shown); their contribution to the axonal growth phenotype in *RGMb* mutant mice cannot be excluded. However, because in dissociated DRG primary cultures lack of neuronal *RGMb* results in reduced neurite outgrowth, we conclude that neuronal *RGMb* signaling directly contributes to axonal growth.

Although our data strongly suggest a role for *RGMb* in promoting BMP signaling and mediating axonal growth, these data do not preclude additional roles for *RGMb* via BMP-independent mechanisms. It is important to note that *RGMa*-induced growth cone collapse in DRG neurons is BMP-independent and acts via the GTP-RhoA signaling pathway (Conrad et al., 2007). Noggin also has no effect on *RGMa*-induced collapse response, further supporting the BMP signaling-independent nature of this phenomena (Conrad et al., 2007).

Taken together, the data presented here support a role for *RGMb*, as well as, for BMP signaling in enhancing peripheral nerve regeneration after axonal injury, and suggests that Noggin expression in naïve animals may suppress intrinsic growth, an effect relieved by Noggin down regulation after nerve injury.

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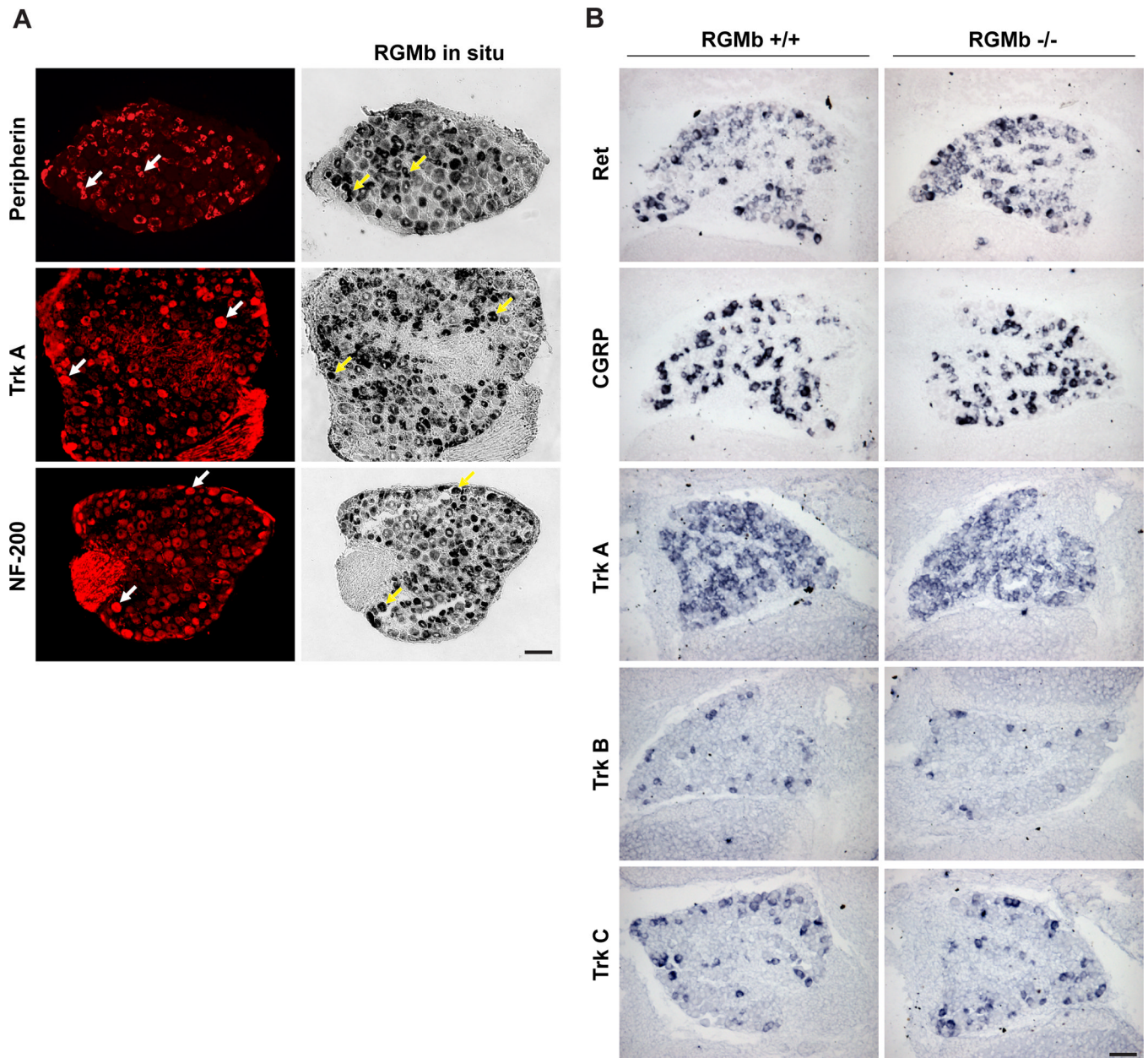


Figure 1. *RGMb* deletion does not alter development of neurons as revealed by *in situ* hybridization

(A) *RGMb* is widely expressed in small and large DRG neurons of adult C57BL/6 mice as demonstrated by co-expression with the sensory neuron markers peripherin, TrkA and neurofilament 200 (NF-200), visualized by fluorescent immunohistochemistry (right panel) and *in situ* hybridization (left panel). Arrows indicate cells that express peripherin, TrkA, or NF-200 (white arrows) and *RGMb* (yellow arrows). Scale bar, 100 μ m. (B) *In situ* hybridization revealed no difference between *RGMb* KO and WT LM in the expression of a variety of key markers of sensory neurons in P0 DRG.

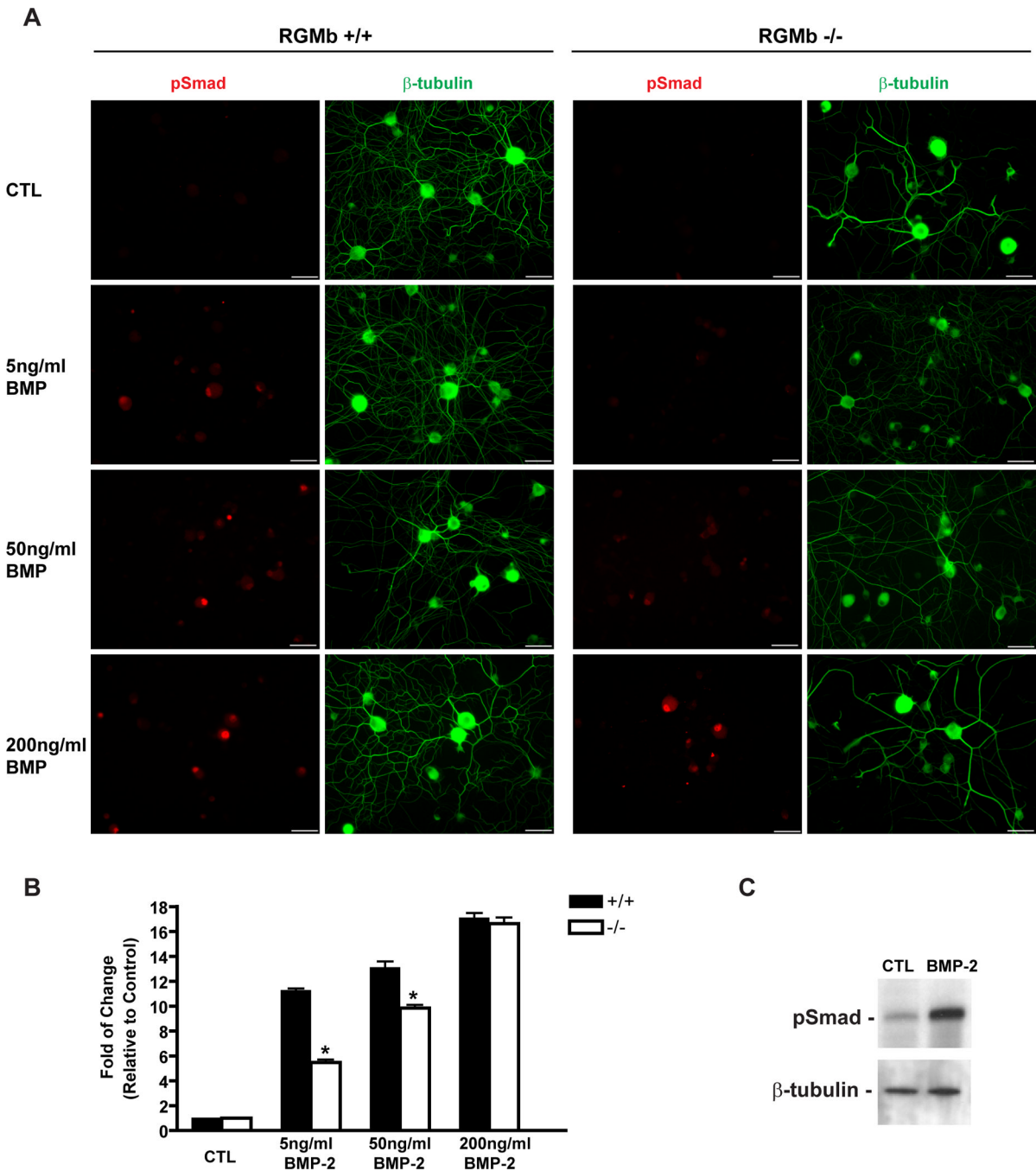


Figure 2. Cultured DRG neurons from *RGMb* deficient mice have decreased BMP-2 mediated signaling

DRG neurons from *RGMb* deficient and WT LMs were cultured in varying concentrations of BMP-2 and then assayed for phospho-Smad (pSMAD) expression. (A) Dissociated DRG neurons were immunostained for anti-pSmad (red) and anti- β -tubulin (green). Scale bars, 10 μ m. (B) Quantification of pSmad intensity. Values represent mean \pm SEM. * indicates statistically significant difference from WT LM, $p < 0.001$. (C) Western blot demonstrating BMP-2 (100ng/ml for 1 hour) increased pSmad1/5/8 induction in cultured DRG neurons from adult WT mice. β -tubulin was used as a loading control.

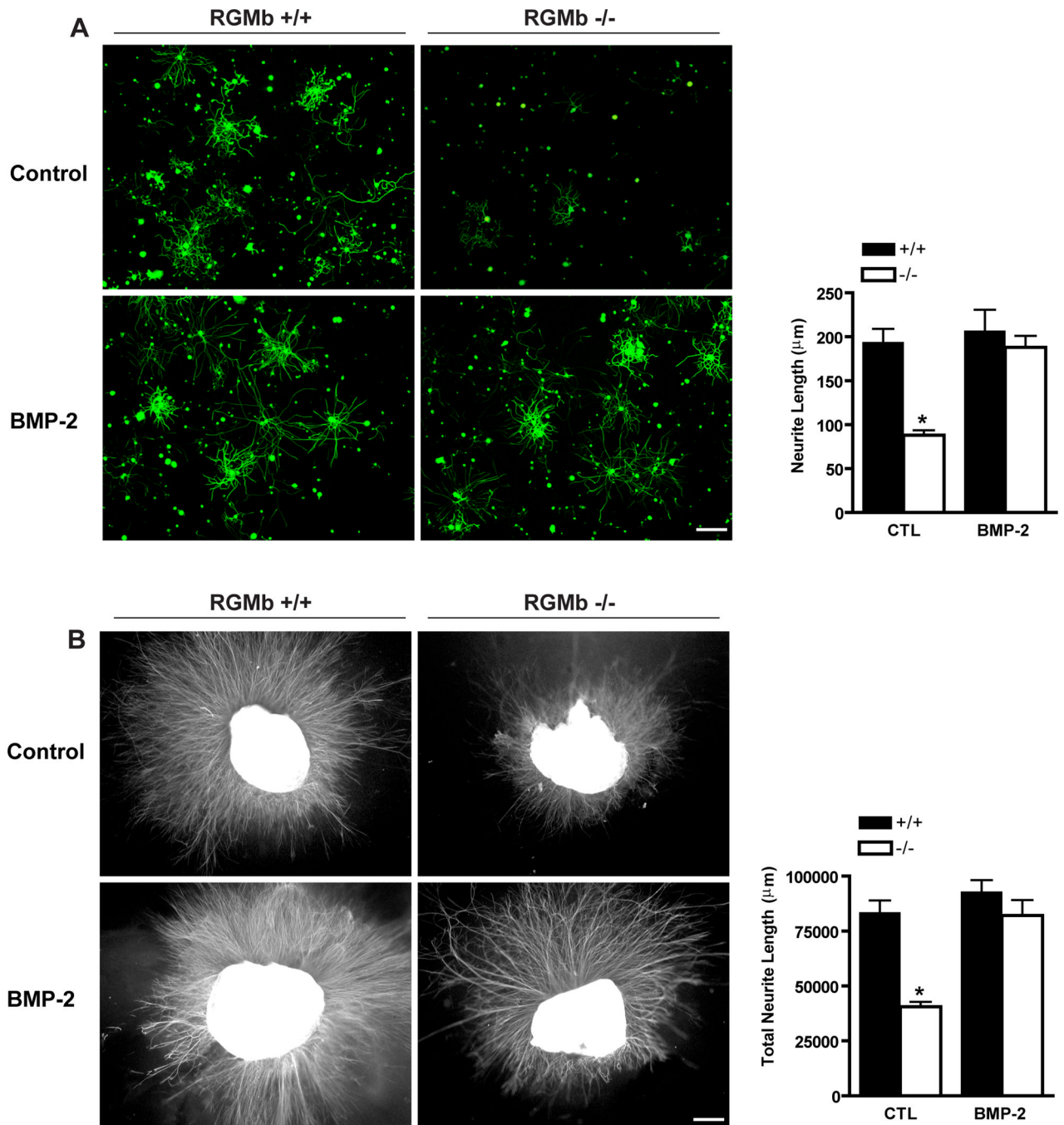


Figure 3. Neurite outgrowth in axotomized *RGMb* KO DRG neurons is reduced and can be rescued by BMP-2

(A) Fluorescent microscopy (anti- β -tubulin immunostaining) of dissociated DRG neurons from *RGMb* KO and WT LM mice. Fewer cultured DRG neurons from *RGMb* $-/-$ mice form neurites, and neurites that do form are shorter than those from neurons derived from WT LMs (* $p < 0.001$). Addition of BMP-2 (50 ng/ml) to the *RGMb* $-/-$ neurons reverses this effect (no significant difference compared with *RGMb* $+/+$). Scale bar, 250 μm . (B) Fluorescent micrographs demonstrating that cultured whole DRG explants from *RGMb* $-/-$ mice generate significantly fewer neurites than explants from WT LM mice and that this loss of function is reversed by the addition of BMP-2 (50 ng/ml). * indicates statistically

significant difference from all other groups, $p < 0.001$. Values represent mean \pm SEM as shown in all graphs. Scale bar, 250 μ m.

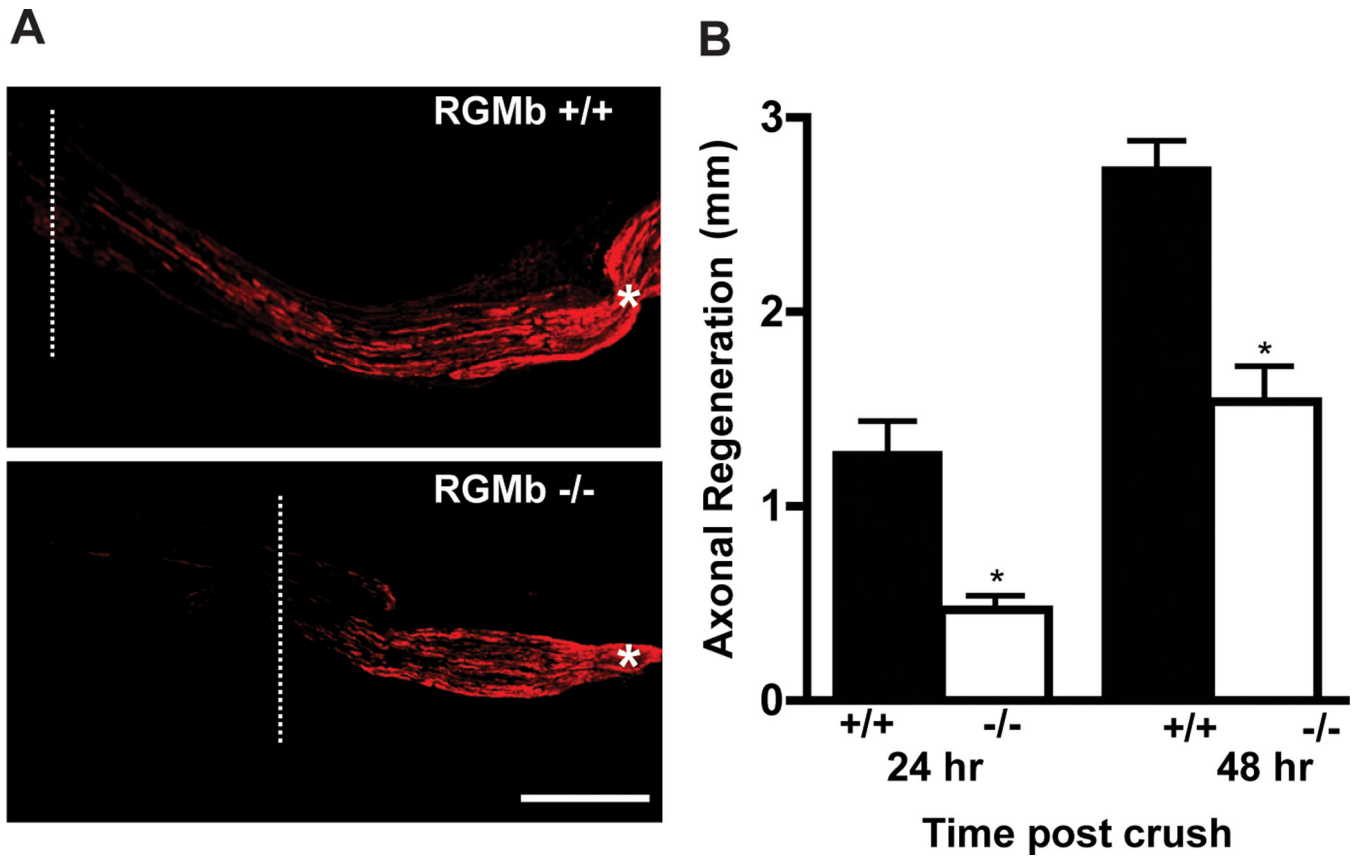


Figure 4. *RGMb* deficient mice show retarded axonal regeneration following sciatic nerve crush (A) GAP-43 fluorescent immunohistochemistry demonstrates decreased axonal growth 48 hours following sciatic nerve crush in *RGMb* KO mice compared with WT LMs; sciatic nerve crush injury was performed at post-natal day 8; Scale bar, 500 μ m. (B) Functional regeneration assessed using the pinch test shows decreased regeneration at 24 and 48 hours post-injury in *RGMb* KO mice compared with WT LMs; sciatic nerve crush injury was performed at post-natal day 8. * $p < 0.001$.

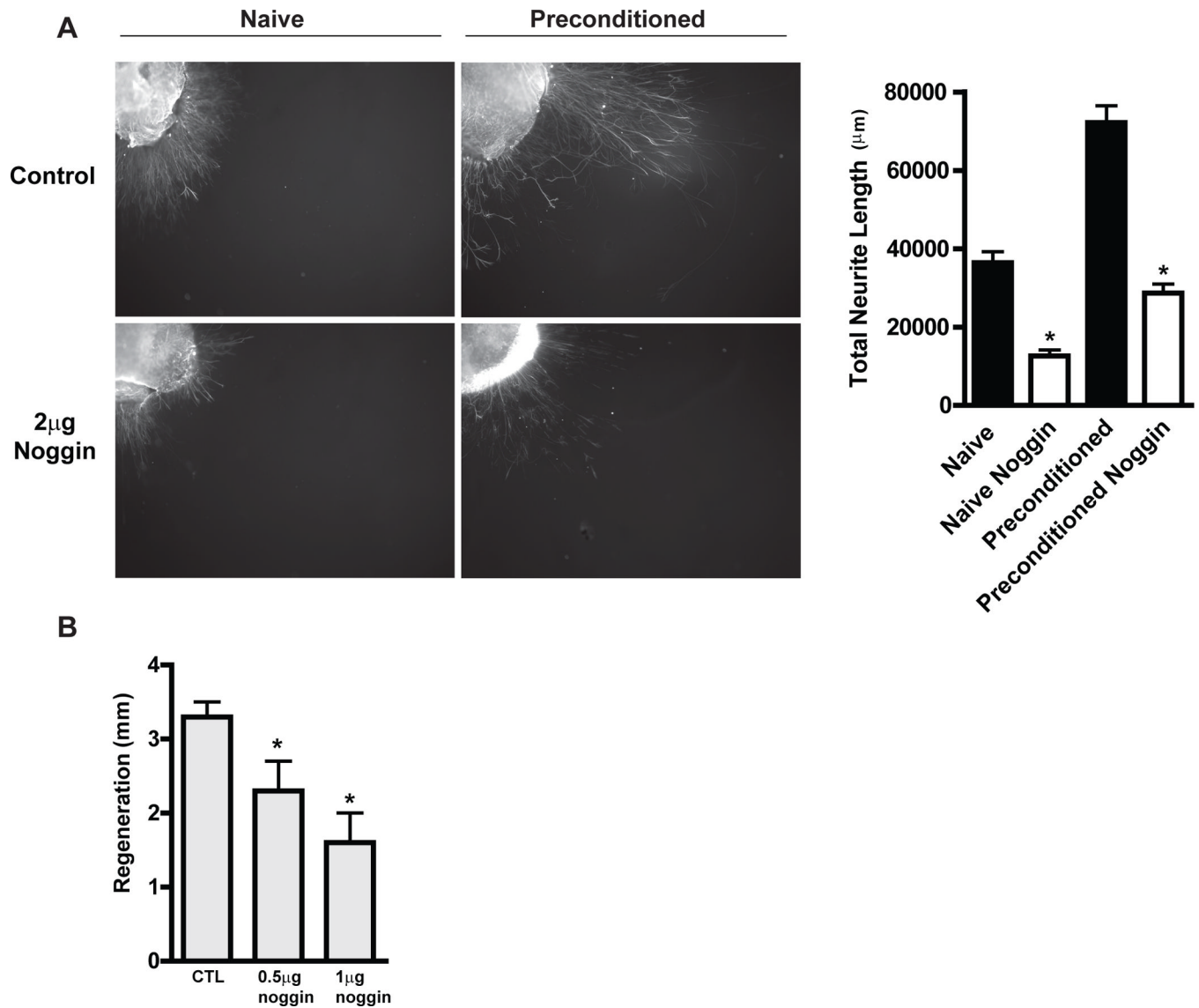


Figure 5. Inhibition of BMP signaling using the endogenous inhibitor, noggin, reduces neurite formation *in vitro* and axonal regeneration *in vivo*

(A) Addition of noggin to culture media causes decreased neurite formation in both naïve and preconditioned DRG explants from adult mice (* $p < 0.001$, $n = 8$ per group). (B) Intrasciatic injection of the BMP inhibitor, noggin, at both 0.5 µg and 1 µg decreased axonal regeneration 48 hours post-injury as assayed by the pinch test. * indicates statistically significant difference from control (CTL), $p < 0.05$, one-way ANOVA followed with *post-hoc* Newman-Keuls test. $n = 6$ per group. Values represent mean \pm SEM as shown in all graphs.

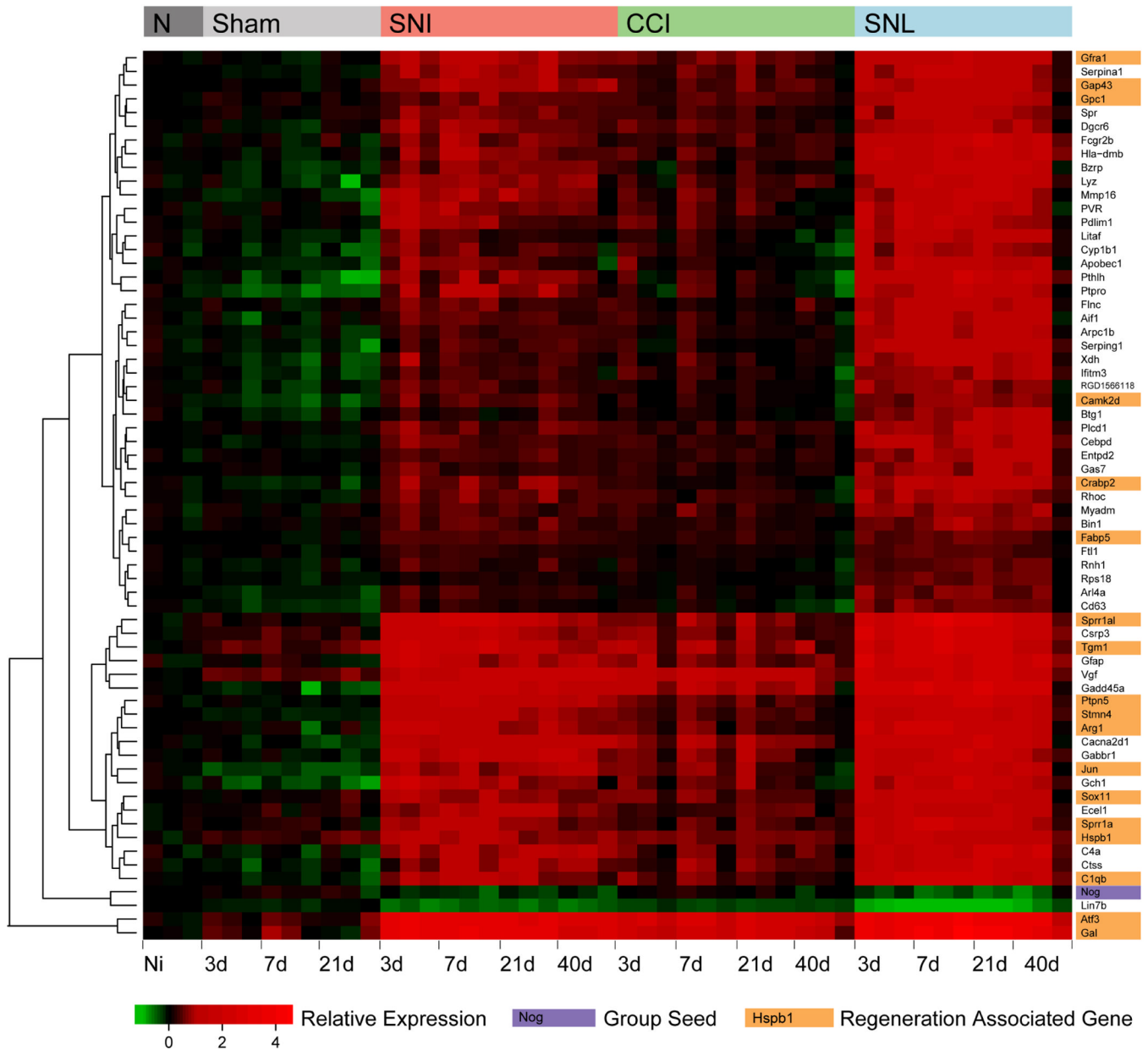


Figure 6. *Noggin* mRNA expression after peripheral nerve injury negatively correlates over time with the expression of a cluster of regeneration associated genes

Weighted gene co-expression network analysis/neighborhood network analysis was used to determine a group of 65 genes whose expression across an extensive dorsal root ganglion (DRG) nerve injury dataset most closely relate to *Noggin* regulation within that group. Heat map showing regulation of these genes in the DRG before (Naïve, N) and after sciatic nerve injury (Spared nerve injury (SNI), Chronic constriction injury (CCI) and Spinal nerve ligation (SNL) –top colored bar) as well as after sham injury (Sham) over time (3, 7, 21 and 40 days) - given below. Each colored block within the heat map represents relative expression of that gene in one array in relation to average naïve expression. Red on this plot represents upregulated, with green representing downregulated relative to average naïve levels. *Noggin* marked in purple is one of only two downregulated genes present, despite acting as the seed gene whose relative expression across this axonal injury time course was

used to choose the most closely related group of 65 genes. The other genes are upregulated after nerve injury but have highly correlated inverse expression patterns to Noggin. To the right are gene names highlighted for function, orange indicates transcripts with a published role in axonal regeneration which have been linked to Hspb1 within (Ma et al., 2011).