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Robust Axonal Regeneration Occurs in the Injured CAST/Ei Mouse CNS

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### Authors

Omura, T  
Omura, K  
Tedeschi, A  
et al.

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## Robust Axonal Regeneration Occurs in the Injured CAST/Ei Mouse CNS

### Highlights

- Long-distance CNS axon regeneration in mammals is possible without genetic manipulation
- Pre-injured CAST/Ei sensory neurons grow on diverse growth-inhibitory substrates
- CNS regeneration occurs in CAST/Ei mice through Activin signaling
- Enhancing Activin signaling confers CNS regenerative capacity in C57/B6 mice

### Authors

Takao Omura, Kumiko Omura, ...,  
Michael Costigan, Clifford J. Woolf

### Correspondence

michael.costigan@childrens.harvard.edu  
(M.C.),  
clifford.woolf@childrens.harvard.edu  
(C.J.W.)

### In Brief

Omura et al. screened for neuronal growth on myelin and find CAST/Ei inbred mice are uniquely able to regenerate axons in the injured CNS through Activin signaling. Enhancing Activin signaling confers a CNS-regenerative capacity in normally non-regenerative mouse strains.

### Accession Numbers

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# Robust Axonal Regeneration Occurs in the Injured CAST/Ei Mouse CNS

Takao Omura,<sup>1</sup> Kumiko Omura,<sup>1</sup> Andrea Tedeschi,<sup>1</sup> Priscilla Riva,<sup>1</sup> Michio W. Painter,<sup>1</sup> Leticia Rojas,<sup>1</sup> Joshua Martin,<sup>1</sup> Véronique Lisi,<sup>2</sup> Eric A. Huebner,<sup>1</sup> Alban Latremoliere,<sup>1</sup> Yuqin Yin,<sup>1</sup> Lee B. Barrett,<sup>1</sup> Bhagat Singh,<sup>1</sup> Stella Lee,<sup>1</sup> Tom Crisman,<sup>3</sup> Fuying Gao,<sup>3</sup> Songlin Li,<sup>4</sup> Kush Kapur,<sup>1</sup> Daniel H. Geschwind,<sup>3</sup> Kenneth S. Kosik,<sup>2</sup> Giovanni Coppola,<sup>3</sup> Zhigang He,<sup>1</sup> S. Thomas Carmichael,<sup>4</sup> Larry I. Benowitz,<sup>1</sup> Michael Costigan,<sup>1,5,6,\*</sup> and Clifford J. Woolf<sup>1,6,\*</sup>

<sup>1</sup>F.M. Kirby Neurobiology Center, Boston Children's Hospital and Harvard Medical School, Boston, MA 02115, USA

<sup>2</sup>Neuroscience Research Institute, Department of Molecular, Cellular, and Developmental Biology, University of California, Santa Barbara, Santa Barbara, CA 93106, USA

<sup>3</sup>Departments of Psychiatry and Neurology, Semel Institute for Neuroscience and Human Behavior, David Geffen School of Medicine, University of California, Los Angeles, Los Angeles, CA 90095, USA

<sup>4</sup>Department of Neurology, David Geffen School of Medicine, and Multiple Myeloma Research Consortium, Semel Institute for Neuroscience and Human Behavior, University of California, Los Angeles, Los Angeles, CA 90095, USA

<sup>5</sup>Anaesthesia Department, Boston Children's Hospital and Harvard Medical School, Boston, MA 02115, USA

<sup>6</sup>Co-senior author

\*Correspondence: [michael.costigan@childrens.harvard.edu](mailto:michael.costigan@childrens.harvard.edu) (M.C.), [clifford.woolf@childrens.harvard.edu](mailto:clifford.woolf@childrens.harvard.edu) (C.J.W.)

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## SUMMARY

Axon regeneration in the CNS requires reactivating injured neurons' intrinsic growth state and enabling growth in an inhibitory environment. Using an inbred mouse neuronal phenotypic screen, we find that CAST/Ei mouse adult dorsal root ganglion neurons extend axons more on CNS myelin than the other eight strains tested, especially when pre-injured. Injury-primed CAST/Ei neurons also regenerate markedly in the spinal cord and optic nerve more than those from C57BL/6 mice and show greater sprouting following ischemic stroke. Heritability estimates indicate that extended growth in CAST/Ei neurons on myelin is genetically determined, and two whole-genome expression screens yield the Activin transcript *Inhba* as most correlated with this ability. Inhibition of Activin signaling in CAST/Ei mice diminishes their CNS regenerative capacity, whereas its activation in C57BL/6 animals boosts regeneration. This screen demonstrates that mammalian CNS regeneration can occur and reveals a molecular pathway that contributes to this ability.

## INTRODUCTION

Axons do not regenerate after an injury to the adult mammalian CNS. Two conditions appear to be required for successful re-growth of injured CNS axons: an intrinsic capacity of the neurons to grow and an ability to overcome growth-inhibitory environments. Many factors inhibit growth in the CNS (Filbin, 2003; Fitch and Silver, 2008; Fournier et al., 2001; Silver et al., 2014), and eliminating some of these can promote regeneration (Allilain et al., 2011). In the mature CNS the intrinsic growth capacity of

intact neurons is repressed to stabilize synaptic circuitry (Sun and He, 2010) and is not generally re-engaged after axonal injury (Cho et al., 2013). To experimentally increase intrinsic neuronal growth in the CNS, studies have genetically inhibited growth-suppressing genes (Sun et al., 2011), exposed CNS neurons to growth-promoting factors (Yin et al., 2006), and altered the neuronal growth state by a pre-conditioning axonal lesion (Kadoya et al., 2009; Mills et al., 2007; Neumann and Woolf, 1999). The data from many studies suggest that both the presence of extrinsic inhibition and the lack of an intrinsic growth capacity after injury are important in preventing axonal re-growth in adult CNS (Benowitz and Yin, 2007).

Neurons with axons in the peripheral nervous system retain an ability to regenerate following injury by inducing a robust intrinsic growth response; in addition, the PNS environment is growth permissive (Hoffman, 2010). Pre-conditioning PNS neurons by a peripheral axonal injury primes them through massive transcriptional changes to regenerate more vigorously and faster in the face of a second injury because of an induction of networks of regeneration-associated genes (Tedeschi, 2012). Such pre-conditioned growth can be detected in vitro both on permissive substrates (Smith and Skene, 1997; Richardson and Issa, 1984) and to a more limited degree on myelin, the latter enabling the central axons of pre-conditioned neurons to begin to grow in the CNS (Kadoya et al., 2009; Mills et al., 2007; Neumann and Woolf, 1999).

Studies of the transcriptional response in pre-conditioned dorsal root ganglia (DRG) neurons have implicated regeneration-promoting roles for many transcription factors (*c-Jun*, *Atf3*, *Klf* family, *Stat3*, *Sox11*, and *Smad1*) and regeneration-associated genes (*Gap43*, *Cap23*, *Arg1*, *Sprr1a*, *Hspb1*, and tubulins; reviewed in Sun and He, 2010; Tedeschi, 2012). In retinal ganglion cells (RGCs) a growth-priming phenomenon analogous to pre-conditioning can be initiated by intraocular inflammation using agents such as zymosan, which activates expression of the neutrophil- and macrophage-derived growth factor oncomodulin to increase neuronal expression of regeneration-associated genes (Benowitz and Popovich, 2011; Kurimoto et al., 2013).

Using an adult primary neuron cell-based assay in genetically diverse inbred mouse strains, we have investigated whether the capacity for adult sensory neurons to regenerate on an inhibitory substrate after a pre-conditioning lesion has a heritable component, with the intention of then using this for identifying signaling pathways that may enable growth of injured CNS neurons. Our results identify a particular mouse strain with the capacity to regenerate injured axons in the CNS to a far greater extent than other inbred strains and also identify Activin signaling as a major component of this ability.

## RESULTS

### In Vitro Screening for Heritability of DRG Neuron Axonal Growth on Central Myelin

We examined nine diverse inbred mouse strains for a potential differential capacity to initiate adult axonal growth in a CNS-like environment. To accomplish this we screened adult primary sensory neurons obtained from the lumbar L4 and L5 DRGs of genetically distinct mouse strains (A/J, C3H/HeJ, C57BL/6J, DBA/2J, 129S1/SvImJ, NOD/LtJ, NZO/HILtJ, CAST/EiJ, and WSB/EiJ) for their ability to extend neurites when grown in culture on CNS myelin over 24 hr, as a model for axonal growth in the face of the inhibitory CNS environment. The genetic diversity of the nine strains, which are founder members of the Collaborative Cross, has been well documented (Chesler et al., 2008). The uninjured central branch of DRG neurons can regenerate to some extent in the spinal cord in spite of the inhibitory growth environment when the peripheral axons of these cells are pre-injured (Kadoya et al., 2009; Mills et al., 2007; Neumann and Woolf, 1999). Consequently, the axonal outgrowth screens on myelin were performed using both naive (DRG neurons from uninjured mice) and pre-conditioned neurons (DRG neurons pre-injured by a sciatic nerve crush 5 days before culture).

These assays revealed that DRG neurons from CAST/Ei mice were capable of producing substantially higher levels of growth on a myelin substrate than all the other strains assayed, measured both as numbers of neurons with neurites and longest axonal process per neuron. The axonal growth in this strain surpassed the average growth of the other strains under naive conditions (2-fold increase,  $p < 0.01$ , one-way ANOVA; post hoc Sidak's, for naive CAST/Ei against all other naive strains) and was markedly greater than the average growth of the other strains after pre-conditioning (3.5-fold increase,  $p < 0.0001$ , one-way ANOVA; post hoc Sidak's, for conditioned CAST/Ei against all other pre-injured strains; Figures 1A–1C). Pre-conditioning increased growth on myelin in all the strains, but the relative increase in effect size was much greater for neurons from the CAST/Ei strain (Table S1). The low-power micrographs presented in Figure 1 show the extent of this growth in pre-injured CAST/Ei neurons, a finding that was replicated multiple times (see Experimental Procedures).

The sensory neurons from the CAST/Ei mice also exhibited elevated levels of growth compared to those from C57BL/6 mice when grown on chondroitin sulfate proteoglycans (CSPGs), an inhibitory substrate found in glial scars (Dickendesher et al., 2012; Fisher et al., 2011; Shen et al., 2009). Pre-conditioned CAST/Ei neurons grew far more robustly than pre-conditioned

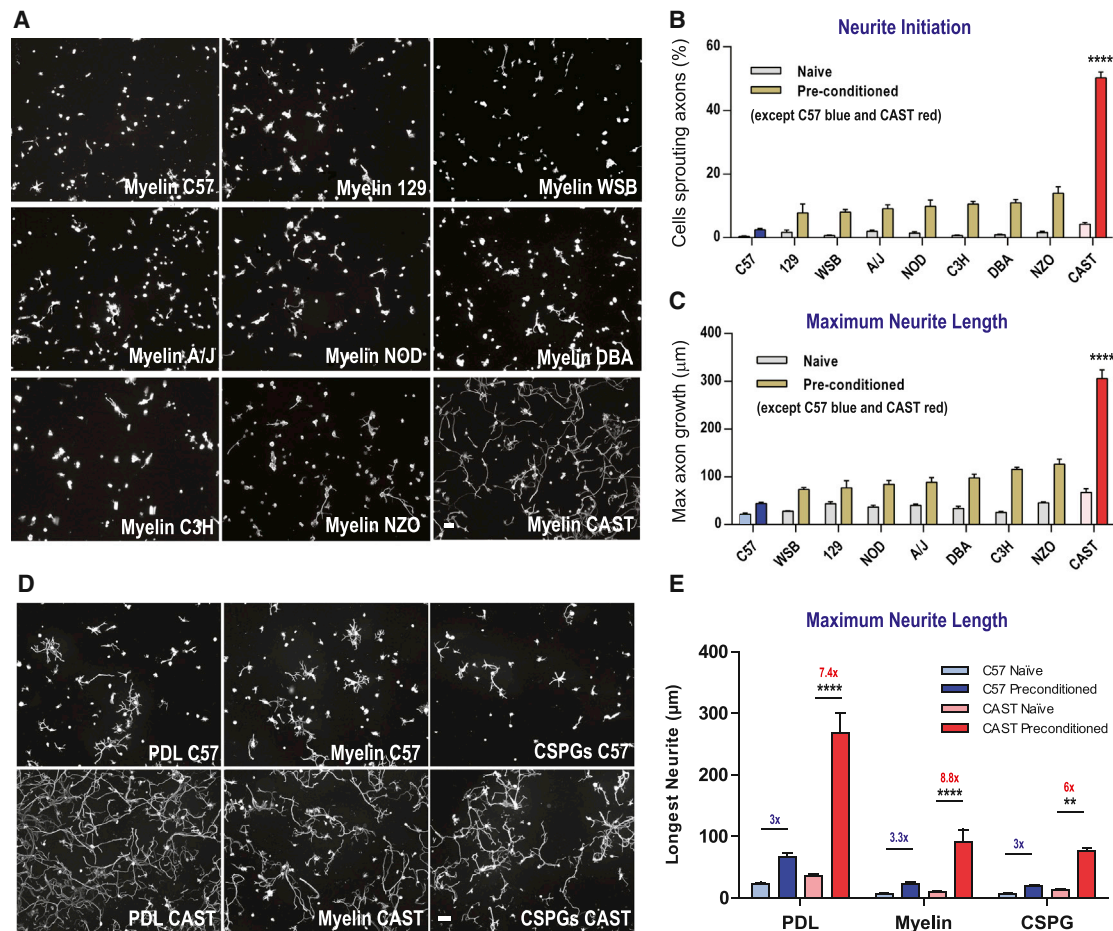
C57BL/6 neurons on both CSPGs (4-fold increase,  $p < 0.05$ , 56.7 [17.0]; CI, 8.3–105.1; three-way ANOVA; post hoc Sidak's) and on myelin (as an independent replicate of the earlier findings; 4.1-fold increase,  $p < 0.001$ , 68.7 [17.0]; CI, 20.3–117.1; three-way ANOVA; post hoc Sidak's; Figure 1E). These results show that injury-conditioned CAST/Ei DRG neurite growth is strikingly greater than that of conditioned C57BL/6 on two major CNS-inhibitory substrates, myelin and CSPGs, indicating a broad reduction in responsiveness of sensory neurons from this strain to potent CNS inhibitory cues.

In contrast, pre-conditioned growth on permissive substrates (dissociated DRG neurons on laminin, whole adult DRG explants on matrigel, and also within the PNS in vivo after a sciatic nerve crush) was broadly similar between CAST/Ei and C57BL/6 (1.3-fold CAST/Ei pre-conditioned versus C57BL/6 pre-conditioned at 12 hr on laminin;  $p < 0.001$ ; one-way ANOVA; post hoc Sidak's; Figure S1B). These results imply that DRG neurons from CAST/Ei mice activate a growth program when injured that specifically enables extensive axonal regeneration on CNS inhibitory substrates, and that this differs from the generalized “pre-conditioning effect” present across all the lines, which promotes regeneration strongly on permissive substrates or in the PNS, but only to a much more limited extent on myelin or in the CNS.

### Growth Phenotype of Injured CAST Neurons in the CNS

To determine whether the growth capacity of injured CAST/Ei DRG sensory neurons on myelin and CSPGs translates into enhanced growth in the CNS, we assayed three different animal models of CNS regeneration: dorsal column injury, optic nerve (ON) injury, and ischemic stroke. We first assayed regeneration of the centrally projecting DRG sensory axons in CAST/Ei or C57BL/6 mice after a dorsal column injury at T10, with and without a pre-conditioning lesion to the sciatic nerve. This is an established model of DRG axonal growth in the CNS, in which the pre-conditioning enables some regeneration of central axons of DRG neurons in the injured spinal cord to occur (Kadoya et al., 2009; Mills et al., 2007; Neumann and Woolf, 1999). Both strains received either a sham or sciatic nerve pre-conditioning injury a week prior to the dorsal column injury at T10. At 6 weeks after the dorsal column injury the sciatic nerve was re-exposed and injected with dextran 3000 (micro-Ruby) to retrogradely label ascending dorsal column fibers, and these were measured 2 weeks later.

Axonal growth in the dorsal column after a lesion was very limited in both C57BL/6 and CAST/Ei mice that had not undergone a pre-conditioning lesion; however, the pre-conditioning of sensory neurons prior to the spinal cord injury resulted in growth into and beyond the lesion site in both strains (Figures 2A, 2B, S2, and S3). In these mice the dorsal columns from the conditioned CAST/Ei mice contain significantly more regenerating axonal trajectories quantified up to 3 mm rostral of the injury site than pre-injured C57BL/6 mice ( $p = 0.004$ ; differences determined by permutation tests for CAST/Ei and C57BL/6 axonal growth profiles; Figure 3E). Overall pre-conditioned CAST/Ei sciatic dorsal column fibers regenerated  $2,974 \pm 89 \mu\text{m}$  rostral from the lesion site compared to  $789 \pm 271 \mu\text{m}$  in C57BL/6 (a 3.5-fold difference,  $p < 0.01$ , 2,184.9 [557.1] CI 826.3–3,543.6; two-way ANOVA; post hoc Sidak's; Figure S4). In two



**Figure 1. Screening for Axonal Growth in an Inhibitory Central Growth Environment**

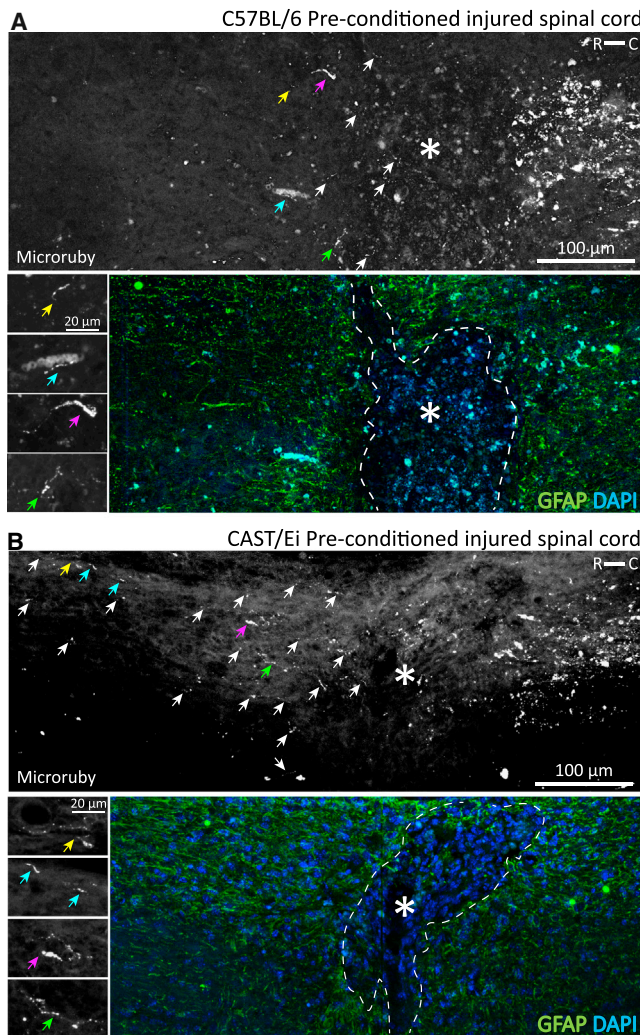
(A) Representative photomicrographs of pre-conditioned axonal growth for nine screened mouse strains (A/J, C3H/HeJ, C57BL/6J, DBA/2J, 129S1/SvImJ, NOD/LtJ, NZO/HILtJ, CAST/EiJ, and WSB/EiJ) on myelin. (B) Maximal level of axonal sprouting achieved by naive and pre-conditioned DRG neurons grown on myelin for 24 hr in these mouse strains. (C) Maximal level of axonal outgrowth achieved in naive and pre-conditioned DRG neurons by a sciatic nerve crush injury 5 days prior to growth on myelin for 24 hr. (D) Representative photomicrographs of pre-conditioned DRG axonal growth for C57BL/6 and CAST/Ei DRG neurons on PDL, myelin, and CSPGs, respectively. (E) Quantification of growth of naive and pre-conditioned C57BL/6 and CAST/Ei DRG neurons on a neutral substrate, poly-d-lysine, and on inhibitory substrates, myelin and CSPGs. Alternate image capture and counting schemes were used between data given in (A)–(C), (D), and (E) (see [Experimental Procedures](#)). All error bars represent SEM. Statistical analysis of (B) and (C): one-way ANOVA; post hoc Sidak's corrected,  $p < 0.0001$  for pre-conditioned CAST/Ei against all other conditions. For (E): three-way ANOVA; post hoc Sidak's. \*\* $p < 0.01$ ; \*\*\*\* $p < 0.0001$ . Scale, 100  $\mu\text{m}$ . For growth of C57BL/6 and CAST/Ei DRG neurons in the PNS see [Figure S1](#).

out of seven CAST/Ei animals, axonal growth reached 5 mm beyond the crush site. The limited extent of growth rostral to the injury site in C57BL/6 mice was consistent with previous results ([Seiffers et al., 2007](#)).

We next investigated whether CAST/Ei neurons that lie entirely within the CNS would also show enhanced regenerative growth. To do this, we assayed RGC axon regeneration after ON crush using cholera toxin B fragment (CTB) as an axonal tracer. Axon regeneration under baseline conditions was minimal in both strains 2 weeks after ON crush. Zymosan is commonly used to condition these cells in a manner analogous to a priming nerve injury for DRG neurons ([Yin et al., 2003](#)), with a dose of 36  $\mu\text{g}$  zymosan optimal for promoting RGC regeneration in C57BL/6

mice ([Kurimoto et al., 2010](#); [Yin et al., 2006](#)). To compare relative regeneration between the two strains we tested the effects of two reduced doses of zymosan applied 5 days before and immediately after the ON crush, in order to induce a lower-than-maximal growth in C57BL/6 mice.

In C57BL/6 animals 6.4  $\mu\text{g}$  and 25  $\mu\text{g}$  of zymosan produced modest and dose-dependent levels of axonal regeneration measured 14 days after the ON injury ([Figures 3A and 3B](#)) ([Kurimoto et al., 2010](#); [Yin et al., 2006](#)). The same doses in CAST/Ei mice, however, produced substantially more extensive axonal regeneration, with axons observed as far as 4,000  $\mu\text{m}$  from the crush site reaching close to the optic chiasm. Differences were determined by permutation tests for contrast based on average



**Figure 2. CAST versus C57 Axonal Regeneration in the Adult Spinal Cord**

(A and B) Regeneration of central axons of pre-conditioned DRG neurons rostral to a dorsal column injury. A small level of growth is present in C57BL/6 mice (A), but it is significantly greater in CAST/Ei mice (B). Growth quantified in Figure 3. Arrows identify retrogradely labeled axons from the PNS. The white dashed line indicates the border of the lesion. Asterisk indicates the lesion epicenter (R, rostral; C, caudal). The images are automated tile-scanned confocal images using a Zeiss LSM 700 confocal microscope, and two consecutive sections (20 μm each) were overlaid and aligned. Scale, 100 μm. See also Figures S2 and S3.

difference of regenerating axonal trajectories; at 6.4 μg zymosan CAST versus C57, mean difference, 118.02,  $p = 0.012$ ; at 25 μg zymosan CAST versus C57, mean difference, 93.28,  $p = 0.031$  (Figures 3C and 3D, quantified in Figure 3F). As robust growth occurred in CAST/Ei RGCs at a low dose of zymosan (6.4 μg), one at which C57BL/6 RGCs respond minimally, these results suggest that CAST/Ei CNS neurons are either primed more easily for re-growth than those from C57BL/6 mice or overcome inhibitory substrates better.

Next, we assayed whether CAST/Ei mice exhibit a greater axonal sprouting ability than C57BL/6 mice following an

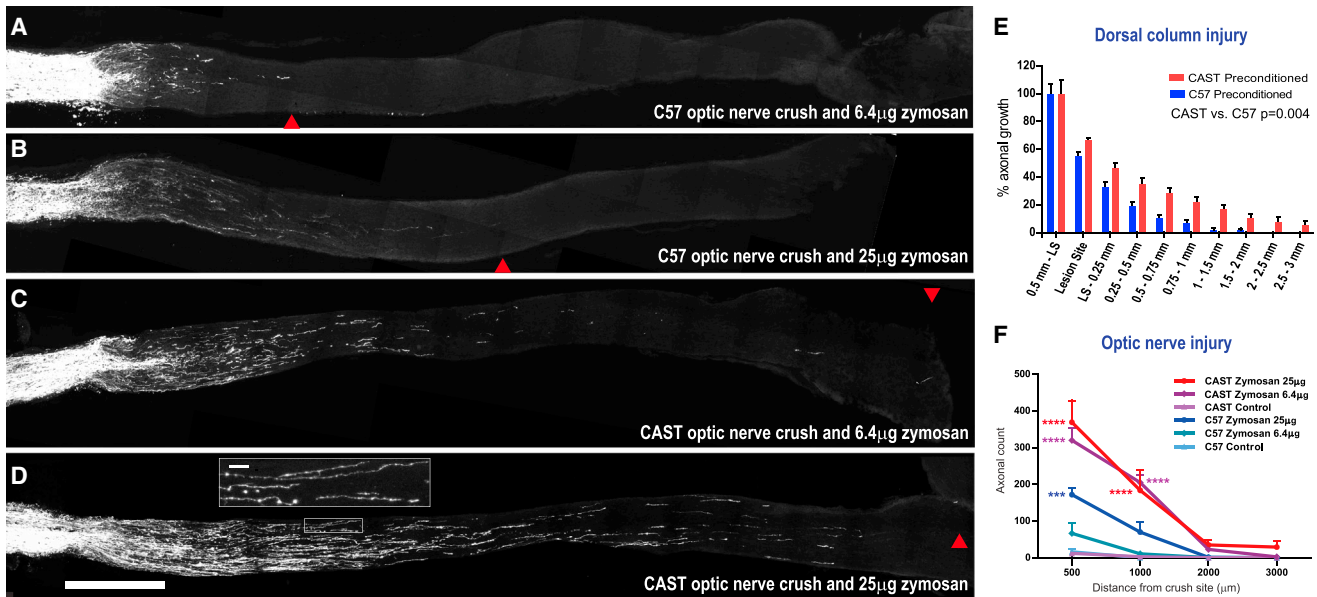
ischemic stroke produced by middle cerebral artery occlusion, where neuronal growth is impeded by myelin and CSPGs. To determine the extent of axonal sprouting we mapped cortical motor system connections in naive CAST/Ei and C57BL/6 mice 1 month after stroke, a time point that coincides with the formation of new connections (Clarkson et al., 2013; Li et al., 2010; Overman et al., 2012). In C57BL/6 animals, motor cortex projections showed only a limited degree of axonal sprouting (Figures 4A and 4B). However, sprouting in CAST/Ei mice extended from the motor cortex into the premotor cortex, as well as to posterior and lateral areas corresponding to the first and second somatosensory cortices (Figures 4C and 4D), with axonal projections > 5 mm from cell bodies (43% area increase in CAST/Ei stroke  $16.55 \pm 1.9 \text{ mm}^2$  versus C57BL/6 stroke  $11.58 \pm 1.0 \text{ mm}^2$ ,  $p < 0.05$ , Student's *t* test, two-tailed, Figure 4E) (Clarkson et al., 2013; Li et al., 2010; Overman et al., 2012). Pre-conditioning by axonal injury or zymosan administration was not necessary in this model to achieve enhanced axonal regeneration in the CAST/Ei mice, perhaps because the inflammation intrinsic to this model is sufficient to prime the cortical neurons for growth, and possibly, in part, because the axons do not have to traverse a physical lesion, as in the spinal cord and ON CNS injury models.

#### Heritability Estimates of Axonal Growth Phenotype

We next wished to understand what molecular signaling pathways might operate in CAST/Ei mouse neurons to enable regeneration in a CNS environment and if these are genetically determined. To estimate the genetic contribution to the observed regenerative variance across strains we estimated narrow-sense heritability, defined as the additive contribution of genetic factors to the observed phenotype from normalized phenotypic data across all nine strains (Table S2). Pre-conditioned growth on myelin showed a high degree of heritability, 72% ( $p = 2.2 \times 10^{-14}$ ) for axonal length, suggesting that most of the inter-strain axonal growth variation can be attributed to genetic factors.

#### Genome-wide Expression Profiling of Naive and Pre-conditioned DRGs across All Nine Strains

Given the high heritability, we undertook two bioinformatic strategies to identify candidate genes and pathways that may contribute specifically to the CAST/Ei pre-conditioned growth on myelin phenotype. In the first, we performed a genome-wide microarray expression profiling screen of naive and pre-conditioned DRG tissue across all nine mouse strains (Table S3). We then correlated this with the degree of DRG axonal growth on myelin, and ranked the resulting genes by Pearson correlation coefficient (Table S4). In the second strategy, we assayed global transcript expression in naive and pre-conditioned DRGs by RNA-seq from three strains chosen for their differing regenerative capacity (C57BL/6, low; DBA/2, mid-range; and CAST/Ei, high). This global expression dataset was then compared to naive and pre-conditioned axonal growth on myelin using an independent screening assay (see Experimental Procedures) to determine the genes that most closely correlated to growth. Remarkably, in both bioinformatic analyses, *Inhba* was revealed to be the transcript most strongly correlated with ability of the pre-injured DRG neurons to grow on myelin (Figures 5A,



**Figure 3. CAST versus C57 Axonal Regeneration in the Adult CNS**

(A–D) Optic nerve injury treated with low-dose (6.4 μg) zymosan in C57BL/6 results in minimal RGC re-growth (A), but the same dose initiates robust growth in CAST/Ei (C). Medium-dose zymosan (25 μg) stimulates growth in C57BL/6 optic nerves (B); however, CAST/Ei RGC growth is much greater (D). Scale, 500 μm. Red arrowhead signifies maximal axonal growth in that section. Inset, magnified image of the regenerating fibers. Scale, 25 μm. These images were obtained using a Zeiss Spinning Disk Confocal at a magnification of 20×, then automatically tiled using Velocity Image Analysis Software (PerkinElmer).

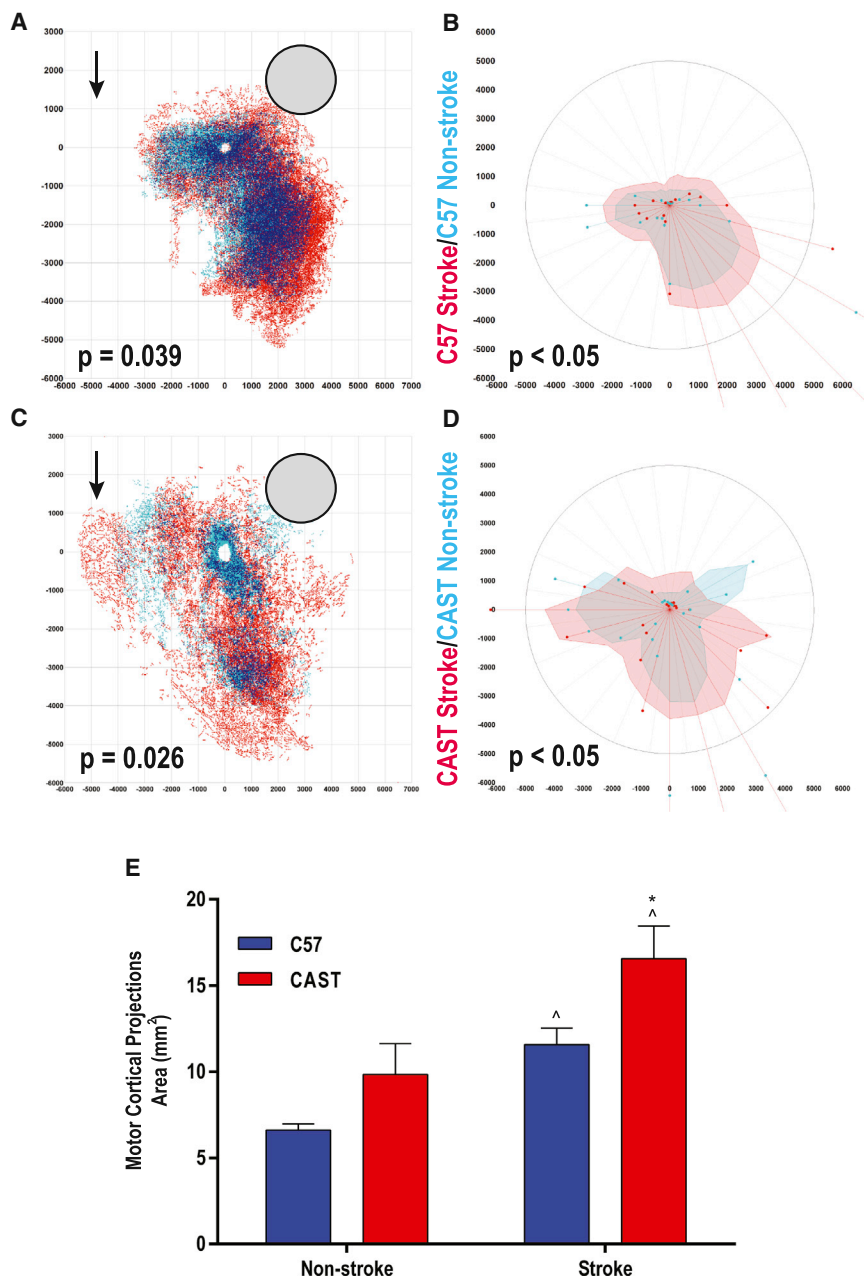
(E) Quantification of percentage of labeled axons crossing the injury site rostral to the lesion site in pre-conditioned adult C57BL/6 and CAST/Ei spinal cords, respectively.

(F) Quantification of numbers of axons reaching defined distances from the crush site in the optic nerve injury model of central regeneration. C57BL/6 and CAST/Ei 25 μg or 6 μg zymosan versus control comparisons; \*\*\*p < 0.001, \*\*\*\*p < 0.0001 (two-way ANOVA, post hoc Sidak's). All error bars represent SEM. For quantification of longest axonal growth in the injured spinal cord see [Figure S4](#).

5C, and 5D). The strain-dependent regulation of *Inhba* indicated by the profiling studies ([Figures 5A](#) and [5E](#)) was confirmed in independent samples across time by qRT-PCR ([Figure 5F](#)). As two independent genome-wide screens identified Activin signaling as being most highly correlated to CAST/Ei active growth on myelin, we chose to investigate this cascade further as detailed below. However, this is unlikely to be the only signaling system that contributes to the CAST/Ei CNS growth phenotype, since we identified two sets of other highly correlated transcripts ([Figures 5A](#) and [5D](#)) from the full screen data.

Overall, the DRG gene expression levels in naive mice and the transcriptional changes induced by sciatic nerve crush were very similar across the nine strains, with relatively few transcripts showing large strain-dependent expression differences across the dataset. To illustrate this, we performed a weighted gene co-expression network analysis (WGCNA) ([Oldham et al., 2008](#)) on the full mouse array dataset. Briefly, absolute Pearson correlation coefficients between each transcript and every other gene in the expression analysis were computed, weighted, and used to determine the topological overlap, a measure of connection strength, or “neighborhood sharing,” in the network. A set of “expression networks” were then constructed where the members of each network have high topological overlap. Groups of genes with very similar expression patterns across large datasets are often functionally connected being part of the same tissue, cell type, or biological pathway ([Oldham et al., 2006, 2008](#)).

Across the mouse array dataset the largest of these expression networks (turquoise, n = 1,824 unique transcripts) represented genes commonly expressed and regulated after nerve injury between the strains ([Table S5](#)). This gene set also overlapped substantially with that group of transcripts that is regulated in the rat DRG following a sciatic nerve injury, and previously implicated as contributing to the conditioning response (p value =  $7.1 \times 10^{-52}$  at the 7-day time point) ([Chang and Jovanis, 1990](#); [Kadoya et al., 2009](#)). The turquoise “mouse pre-conditioning response” network did not include *Inhba*, which is uniquely induced in CAST/Ei relative to marginal changes in expression in the other strains. In addition, *Inhba* is not upregulated in the rat DRG following sciatic nerve injury ([Blesch et al., 2012](#); [Costigan et al., 2010](#)). Moreover, when the degree of regulation of annotated regeneration-associated genes ([Sun and He, 2010](#); [Tedeschi, 2012](#)) between C57BL/6 and CAST/Ei was analyzed, we found virtually no difference ([Figure S5](#)). We interpret these data as indicating that those elements of the pre-conditioning transcriptional response that facilitate accelerated regeneration in the PNS are largely uniform across all nine mouse strains, in keeping with the minimal difference we found in growth rates between C57BL/6 and CAST/Ei on permissive substrates or in the PNS ([Figure S1](#)). Our screen of pre-injured DRG neuronal growth on central myelin has assayed a completely different genomic program, one which enables primed CAST/Ei injured neurons to grow robustly on CNS growth-inhibitory substrates.



**Figure 4. CAST versus C57 Axonal Regeneration Post Cortical Stroke**

(A) Connections of forelimb motor cortex in C57BL/6 with and without stroke. Light blue dots indicate the location of projections in non-stroke, red in stroke, and dark blue dots are areas of dense overlap between the two conditions. Gray circle is site of tracer injection. The x and y axes represent micron distance from the center of the tracer injection. p value, Hotelling's inverse t test. (B) Polar distribution map in register with connective plot in (A) showing localization of sprouting in stroke versus control (Watson's U2 test;  $p < 0.05$ ). Shaded polygons represent the 70th percentile of the distances of labeled projections from the injection site in each segment of the graph; weighted polar vectors represent normalized distribution of the quantity of points in a given segment of control (blue) or stroke (red) graphs.

(C) Same conventions as in (A) for CAST/Ei mice ( $n = 5$  per group). Note the connections in the far premotor cortex, 4 to 5.5 mm anterior to the tracer injection site. These are not present in C57BL/6 after stroke (arrows).

(D) Same conventions as in (B). Overlay of control and post-stroke forelimb motor cortex projections.

(E) Area of motor cortex projections in C57BL/6 and CAST/Ei mice. Areas taken from 70th percentile of all distances in each radial segment from the injection site (B and D). Student's t test, two-tailed. ^  $p < 0.05$ , non-stroke versus stroke C57BL/6 ( $n = 5$  for both stroke and non-stroke) and CAST/Ei ( $n = 5$  for both stroke and non-stroke), respectively; \*  $p < 0.05$ , CAST/Ei stroke versus C57BL/6 stroke. All error bars represent SEM. See [Supplemental Experimental Procedures](#) for further details.

CAST/Ei DRGs than in C57BL/6 DRGs (13.2-fold), but is not regulated by nerve injury. In C57BL/6 DRGs, levels of this transcript are low in the naive state, but it slowly upregulates following injury. Both type II receptors are expressed at higher levels in naive CAST/Ei than C57BL/6 DRGs and were minimally regulated by injury in either strain (Figure 5F).

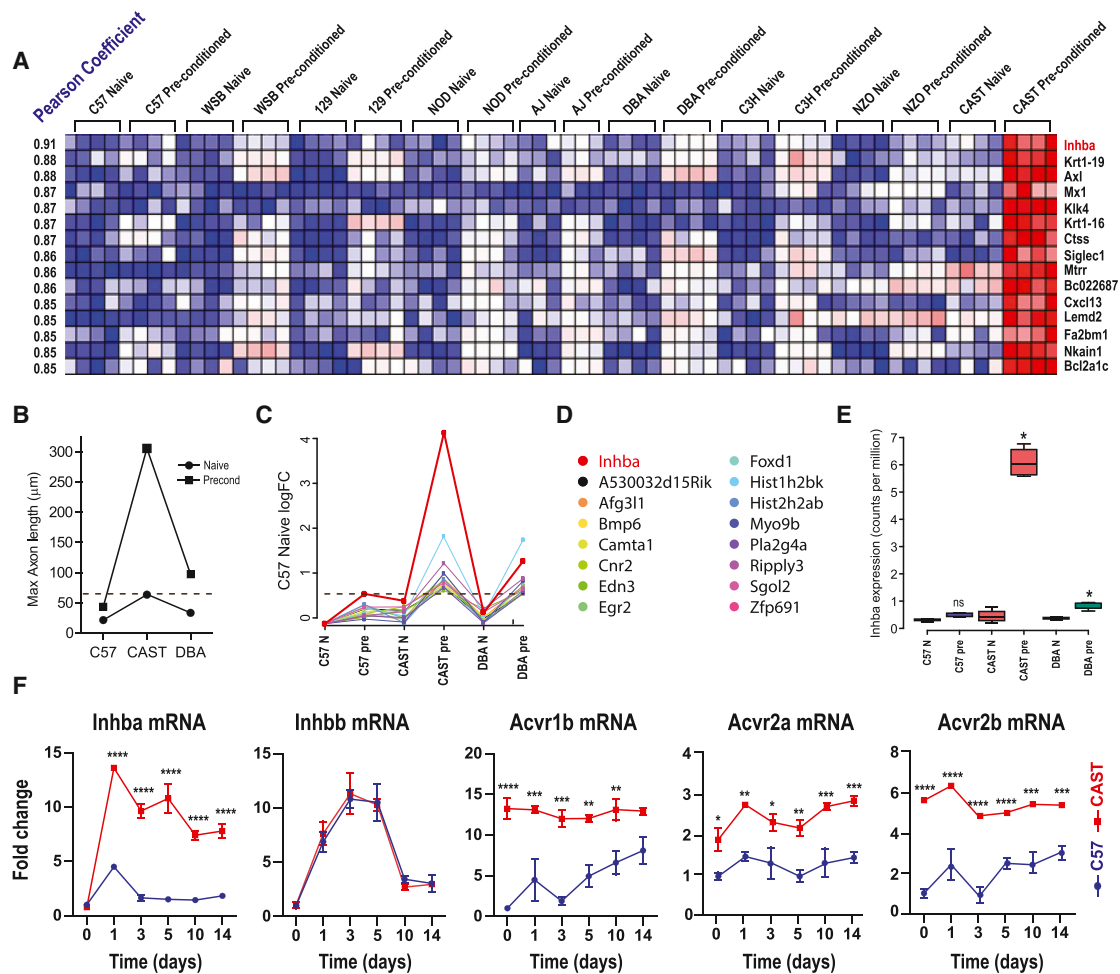
### Activin Signaling Components in C57 and CAST DRGs

*Inhba* mRNA encodes Activin- $\beta_A$ , and the Activin signaling cascade is a component of the TGF- $\beta$  superfamily. Activin comprises three functional protein forms, Activin-A ( $\beta_A\beta_A$ ), Activin-B ( $\beta_B\beta_B$ ), and Activin-AB ( $\beta_A\beta_B$ ). The Activin- $\beta_B$  transcript *Inhbb*, unlike *Inhba*, was upregulated to a similar extent in CAST/Ei and C57BL/6 DRGs following sciatic nerve crush injury (Figure 5F). Activin is signaled by one type I serine/threonine kinase receptor (*Acvr1b*, *Alk4*) and two type II co-receptors (*Acvr2a* and *Acvr2b*) (Harrison et al., 2003). These receptors form an active tetrameric complex composed of two of each receptor subtype. *Acvr1b* is expressed at higher levels in naive

Using in situ hybridization and immunohistochemistry we demonstrate that *Inhba* mRNA, the Activin- $\beta_A$  peptide, and all receptor subtypes are expressed in injured CAST/Ei DRG neurons. Furthermore, we confirm lack of *Inhba* mRNA induction and Activin- $\beta_A$  peptide expression in injured C57BL/6 DRG neurons (Figure S6).

Activin signaling promotes translocation of phosphorylated Smad2/3 from the cytoplasm to the nucleus to initiate downstream transcriptional signaling, and we found a large increase in nuclear phospho-Smad2/3 in pre-conditioned DRG neurons from CAST/Ei, but not C57BL/6, mice (Figure S5). Thus, pre-conditioned CAST/Ei, but not C57BL/6, neurons manifest





**Figure 5. Genome-wide Expression Analysis of CAST CNS Growth**

(A) Naive and pre-conditioned array data for the 15 genes whose DRG expression behavior most correlated with axonal growth on myelin. Each box represents data from a single array, each column a single DRG sample, each row a single gene, blue is less expressed, and red is most expressed. The transcripts are listed in order of Pearson correlation coefficient between expression and axonal growth, and the strains are ordered relative to pre-conditioned growth on myelin.

(B) The growth phenotype of each strain measured in a follow-up RNA-seq screen determined by maximal axonal length. Horizontal red checked line represents the minimal growth level required in the screen for strong growth in the CNS.

(C) Expression data of the 16 genes identified by RNA-seq screen. These data highlight the *Inhba* transcript expression (red line) as the most related to long-distance regeneration in the mammalian CNS.

(D) Each of the transcripts identified by the screen with the color indicator for the expression profiles shown in (C).

(E) Actual expression of the *Inhba* transcript measured by RNA-seq (units are counts per million reads); *Inhba* expression is significantly upregulated after pre-conditioning only in the DBA/2 and CAST/Ei strains (\* $p < 0.01$ , false-discovery rate  $< 0.01$ , edgeR). Data from two independent screens then place Activin signaling at the apex of the strong central axonal regeneration phenotype, most clearly seen in the CAST/Ei strain.

(F) Time course of expression of *Inhba*, *Inhbb*, and Activin receptor subtype transcripts in ipsilateral L4/L5 DRG from naive mice and 1, 3, 5, 10, and 14 days post-sciatic nerve crush for both C57BL/6 and CAST/Ei. qRT-PCR; mean fold change  $\pm$  SEM normalized to naive C57BL/6 values. For statistical analysis, two-way ANOVA was performed, post hoc Sidak's (\* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$ , \*\*\*\* $p < 0.0001$ ). All error bars represent SEM. For Activin cascade member sites of expression in the DRG see Figure S5.

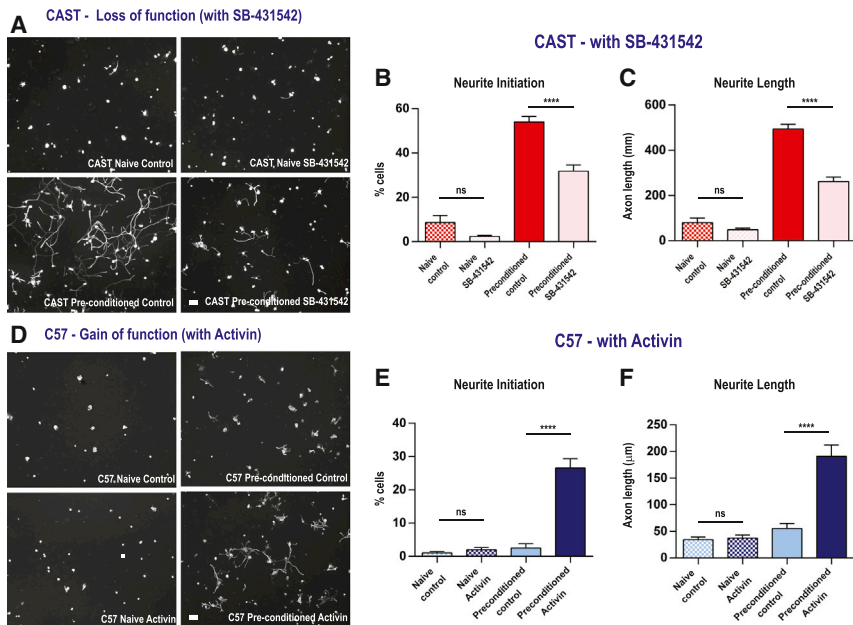
activation of downstream signaling components of the Activin pathway.

### Activin Gain and Loss of Function

To evaluate whether Activin signaling is necessary for the elevated axonal growth of CAST/Ei DRG neurons on myelin, we cultured pre-conditioned CAST/Ei DRG neurons for 24 hr (the same protocol used for screening inbred strains) in the presence or absence

of SB-431542, a potent Activin receptor antagonist (Laping et al., 2002). SB-431542 (100 nM) significantly reduced neurite initiation (42% reduction,  $p < 0.0001$ , 22.3 [2.6] CI, 15.6–28.9; two-way ANOVA, post hoc Sidak's) and maximal axon length (48% reduction,  $p < 0.0001$ , 232.7 [21.9] CI, 176.6–288.9; two-way ANOVA, post hoc Sidak's) relative to vehicle controls (Figures 6A–6C).

We also determined whether exogenous introduction of Activin ligands could promote axonal growth on myelin in C57BL/6



**Figure 6. Gain- and Loss-of-Function Assays of Activin Action in Promoting Central Axonal Growth**

(A–C) The Activin receptor, type 1B inhibitor SB-431542 (100 nM), significantly decreases neuronal growth of pre-conditioned CAST/Ei DRG neurons on myelin, quantified in (B) and (C). (D–F) A mixture of Activin-A, Activin-B, and Activin-AB peptides (10 ng/ml) significantly increased axonal growth in pre-conditioned C57BL/6 DRG neurons, quantified in (E) and (F). Neuronal counts average of four wells per group, run three independent times as counted in Figures 1A–1C. For (B), (C), (E), and (F): mean ± SEM. \*\*\*\* $p < 0.0001$  (two-way ANOVA, post hoc Sidak's). Scale, 100 µm. All error bars represent SEM.

fold increase in the number of axons 500 µm distal to the crush site relative to zymosan alone ( $p < 0.05$ , 112.4 [39.5] CI, 14.9–209.9; two-way ANOVA, post hoc Sidak's; Figure 7D), showing that Activin enhanced CNS axonal regeneration in

neurons. Dissociated pre-conditioned DRG neurons from C57BL/6 mice showed a 10.6-fold increase in sprouting ( $p < 0.0001$ , 24.1 [2.2] CI, 18.8–29.4; two-way ANOVA, post hoc Sidak's) and a 3.6-fold increase in neurite elongation ( $p < 0.0001$ , 135.6 [19.8] CI, 88.1–183.1; two-way ANOVA, post hoc Sidak's) on myelin when cultured in the presence of Activin A, B, and AB peptides (10 ng/ml; Figures 6D–6F). To monitor Activin downstream signaling in these cells, we assayed nuclear pSmad2/3 localization, finding low overall levels in C57BL/6 naive neurons (4.4% positive nuclei) with significantly more, but still relatively low, levels in pre-conditioned C57BL/6 neurons (21% positive nuclei,  $p < 0.05$ , 0.19 [0.05] CI, 0.03–0.35; Poisson regression, rate estimate on a log scale, post hoc Sidak's) that increased robustly when treated with Activin to 56% nuclear staining ( $p < 0.0001$ , 0.59 [0.06] CI, 0.39–0.79; Poisson regression, rate estimate on a log scale, post hoc Sidak's). Nuclear pSmad2/3 levels were relatively high in naive dissociated CAST/Ei DRG neurons at 46%, but increased further after pre-conditioning to 69% ( $p < 0.0001$ , 0.53 [0.08] CI, 0.27–0.79; Poisson regression, rate estimate on a log scale, post hoc Sidak's). In addition, the number of pSmad2/3-stained nuclei decreased from 69% to 45% when pre-conditioned CAST/Ei neurons were treated with SB-431542 (100 nM) ( $p < 0.0001$ , 0.56 [0.08] CI, 0.31–0.81; Poisson regression, rate estimate on a log scale, post hoc Sidak's; Figures 7A and 7B).

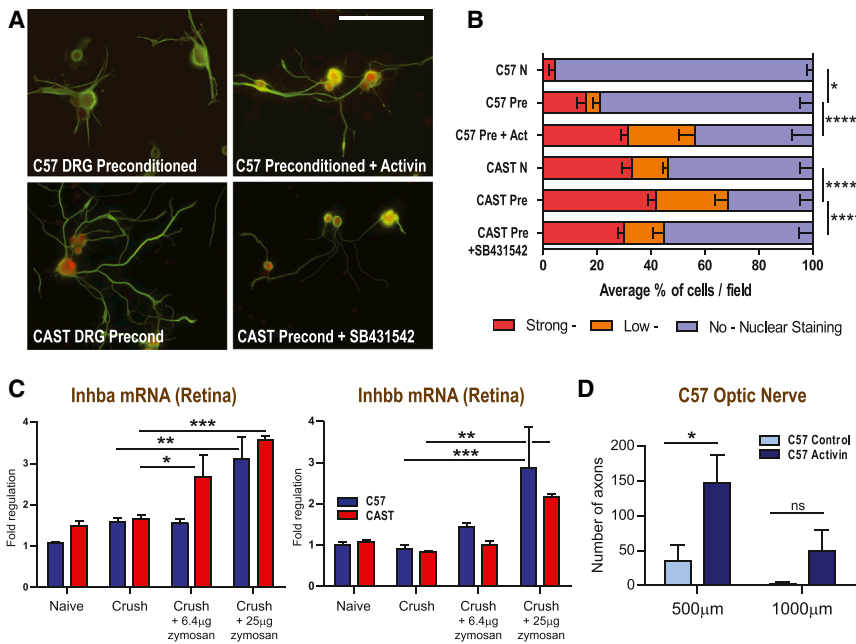
Zymosan at a dose (6.4 µg) that produces extensive growth only in CAST/Ei mice induced *Inhba* mRNA expression in the retina of this strain, but not in retina from C57BL/6 mice, whereas a higher zymosan dose (25 µg), which promotes regeneration in both strains, induced *Inhba* mRNA expression in both mouse lines (Figure 7C). To test if Activin signaling can boost RGC axonal regeneration in vivo we measured axonal growth in C57BL/6 ONs after crush where the retina had been treated with low-dose (6.4 µg) zymosan and then with 10 ng/ml of Activin immediately after ON crush. This combination resulted in a 4.2-

C57BL/6 mice after a low-dose zymosan priming. Finally, we assayed if Activin promoted CNS neuron growth in another species using rat RGCs in culture. Rat RGCs that had been injured and primed in vivo with low-dose zymosan (6.4 µg) showed a 2.8-fold increase in outgrowth when subsequently grown on PDL in vitro in response to the addition of Activin peptides (relative to control RGCs,  $p < 0.01$ , Student's t test, two-tailed; Figure S7).

## DISCUSSION

Neurons in the adult mammalian PNS have the capacity, also found in the CNS of some lower organisms, to mount a regenerative response following axonal damage. In stark contrast, this regenerative ability has been considered to be essentially lost in the mammalian adult CNS. Two factors have been considered cardinal to achieving long-distance axonal growth in the adult mammalian CNS: removal of the growth-inhibitory environment and initiation of intrinsic molecular growth programs that enable active axonal growth. Our data show that CAST/Ei mouse neurons, following a pre-conditioning injury, are not inhibited by extrinsic inhibitory cues from the CNS extracellular environment in the way that other mouse strains are. The extent of the CNS regeneration in primed RGCs from CAST/Ei mice (as determined by maximal axonal growth) is similar to that found when tonic growth suppressors are eliminated (Gaub et al., 2010; Sun et al., 2011). The discovery of this unexpected growth capacity of injured neurons in the CNS in this strain allows molecular mechanisms required to initiate and sustain regeneration in the adult mammalian CNS to be teased out, and therapeutic implications of growth studies in these mice will aid drug discovery without removal of tumor suppressor mechanisms, which although effective, carry an oncogenic risk (Sun et al., 2011).

We designed a cell-based phenotypic screen to look for heritable differences in neuronal growth capacity, using pre-conditioned DRG neurons on myelin to model one of the very few



**Figure 7. Molecular Mechanisms of Activin Function**

(A) Representative cultured pre-conditioned C57BL/6 and CAST/Ei DRG neurons immunostained for pSmad2/3 and Tuj1 before and after Activin or SB-431542 treatment. Scale, 100 µm. (B) Quantification of nuclear pSmad2/3 in naive and pre-conditioned C57BL/6 and CAST/Ei DRG neurons treated with Activin or SB-431542.

(C) qPCR data showing that treating the mouse retina with 6.4 µg intraocular zymosan increases *Inhbba* mRNA expression only in CAST/Ei, while 25 µg induces regulation in both C57BL/6 and CAST/Ei animals. Intraocular injection of 6.4 µg and 25 µg zymosan showing similar *Inhbhb* regulation in C57BL/6 and CAST/Ei animals.

(D) A mixture of Activin-A, Activin-B, and Activin-AB increases ON regeneration in C57BL/6 mice treated with 6.4 µg of zymosan. Data quantified as numbers of optic nerve axons growing 500 µm and 1,000 µm from crush site in presence of PBS or Activin peptide mix, respectively. For (B): mean ± SEM. Statistical analysis Poisson regression, post hoc Sidak's. \* $p < 0.05$ , \*\*\*\* $p < 0.0001$ . For (C): mean fold change ± SEM normalized to naive C57BL/6 values, for comparisons shown \* $p < 0.05$ , \*\* $p < 0.01$ , \*\*\* $p < 0.001$  (two-way ANOVA, post hoc

Sidak's). For (D): counts are average of four sections per animal, five animals. Average numbers of axons reaching defined distances from the crush site in each animal ± SEM; \* $p < 0.05$  (two-way ANOVA, post hoc Sidak's). All error bars represent SEM. For in vitro gain-of-function assay using Activin peptides on rat retinal ganglion cells see Figure S7.

situations where axonal growth occurs in the CNS without genetic manipulation: namely when a prior peripheral nerve injury enables dorsal column fibers to regenerate into and beyond a spinal cord injury (Lu et al., 2004; Neumann et al., 2002; Neumann and Woolf, 1999). Our hypothesis was that differences in the growth of isolated DRG neurons across different mouse strains on a CNS inhibitory substrate might predict a differential capacity to grow within a CNS environment. Our screen revealed both that there is a large heritable component to the regeneration on myelin phenotype after pre-conditioning, and that CAST/Ei display a much higher level of growth in this circumstance than all the other strains. Furthermore, the growth of DRG neurons from this strain in vitro in this model accurately predicted increased growth in vivo of pre-conditioned dorsal column fibers after a spinal cord injury, RGC axons after ON crush when primed by zymosan, and enhanced spatial reorganization of projections arising from cortical neurons after a stroke.

To successfully regrow after axonal injury, PNS neurons need to realign their genomic structure through epigenetic mechanisms (Cho et al., 2013; Finelli et al., 2013; Kiryu-Seo and Kiyama, 2011; Puttagunta et al., 2014) to engage transcriptional programs with miRNAs (Li et al., 2012) and signaling pathways (Perry et al., 2012) to activate complex networks of transcription factors and repressors (Costigan et al., 2002, 2010; Tedeschi, 2012). This injury response leads to the upregulation of the many genes required for initiating, guiding, and sustaining axonal growth. Injuring the central axons of these DRG neurons, however, has no such effect (Ylera et al., 2009). Essentially, peripheral nerve injury reprograms adult sensory and motor neurons from a quiescent to an active intrinsic axonal growth mode, but these changes are not a dedifferentiation to an immature state,

since most of the genes induced are not developmentally regulated. The increased intrinsic growth in DRG neurons after a peripheral axonal injury is accompanied by somewhat reduced responses to inhibitory cues, such as central myelin (Qiu et al., 2002). As such large-scale transcriptional reprogramming does not occur after axonal injury in CNS neurons (Cho et al., 2013), this has been thought to be an important contributor to the failure of CNS regeneration. We found that both C57BL/6 and CAST/Ei neurons display increased growth in a permissive environment after pre-conditioning, in agreement with multiple previous studies in rats and mice (Richardson and Issa, 1984; Smith and Skene, 1997), and that the extent of CAST/Ei pre-conditioned growth in the PNS was not markedly different from that in C57BL/6 mice. In keeping with this, the major component of the transcriptional response to peripheral axonal injury (constituting pre-conditioning) was essentially identical across strains. Therefore, an augmented or different transcriptional growth response after conditioning could not account for the superior regenerative capacity found for DRG neurons from CAST/Ei mice on myelin, opening up the possibility that the pre-conditioned growth on myelin phenotype results from a more discrete process, specifically activated in this strain by injury, which may add to the more generic pre-conditioned response that is uniform across strains. CAST/Ei mice seem to engage, after pre-conditioning in the case of DRG neurons or zymosan priming for RGCs, a specific transcriptional mechanism distinct from the standard induction of RAGs, which enables them to grow well on myelin and CSPGs, and within the CNS in vivo.

Our bioinformatic analysis for differential transcript expression correlates of the growth on myelin phenotype showed that induction of Activin-βA was a prime candidate. In naive CAST

DRG neurons, expression of transcripts that encode Activin- $\beta$ A and Activin- $\beta$ B peptides (*Inhba* and *Inhbb*) is low, but the expression of the Activin receptors is high. Following axonal injury, expression of *Inhba* and *Inhbb* mRNAs are upregulated, with the increase in *Inhba* being specific to CAST/Ei mice and leading to a markedly greater neuronal pSmad2/3 activation. How Activin signaling is selectively enhanced after injury in the CAST/Ei strain relative to the others, and the transcriptional cascades downstream of pSmad2/3 that increase axon growth and/or reduce responses to CNS inhibitory cues, will now need to be elucidated. Future work, for example, by assaying growth variation on myelin in collaborative cross-recombinant backcross strains (Threadgill et al., 2011), could help define the complete genomic pathway required for this growth phenotype.

To test if Activin- $\beta$ A mechanistically contributed to the CNS regeneration phenotype, we used gain- and loss-of-function manipulations *in vitro* in DRG neurons and *in vivo* for RGCs. Antagonizing Activin/TGF- $\beta$  signaling diminished CAST/Ei axonal growth after injury toward the level found in C57BL/6 mice, whereas, conversely, Activin administered to pre-conditioned C57BL/6 neurons shifted regeneration capacity toward that present in CAST/Ei mice. The fact that Activin only induced active growth on myelin in pre-injured C57BL/6 DRG neurons means that other components induced by the conditioning injury need to be present for successful growth, components also provided by exposure to inflammation for RGCs, and we aim in the future to tease out what these are and how they interact with Activin.

A primordial function of Activin in the CNS is revealed by fly brain metamorphosis, a process that involves pruning of larval-specific dendritic and axonal processes and outgrowth of adult axonal structures. A genome-wide mutant screen for blockade of this growth plasticity resulted in only two genes, the *Drosophila* Activin serine kinase receptor (baboon) and dSmad2 (Zheng et al., 2003). In addition, the functional involvement of the two *Drosophila* Activin co-receptors and dActivin in neuronal remodeling has since also been shown (Zheng et al., 2003, 2006). Genome-wide studies have also found that Activin-A mediates tail regeneration in the leopard gecko (Gilbert et al., 2013) and fin re-growth in fish (Jazwinska et al., 2007). Activin can also promote embryonic chick DRG cell outgrowth (Fang et al., 2012), and its expression in the adult CNS can be strongly regulated after injury (Tretter et al., 2000). In the injured CNS Activin promotes dendritic complexity (Ishikawa et al., 2010), modifying spine morphology and increasing the number of synaptic contacts on dendritic spines (Shoji-Kasai et al., 2007). Activin signaling through Smad2/3 is therefore a conserved neural/tissue-regenerative pathway in flies, fish, and birds, and as we now show, the pathway is also activated in the injured adult mammalian nervous system in a mouse strain in which, uniquely, substantial central axon regeneration can occur.

In conclusion, we have found that injured neurons in the mammalian CNS can unexpectedly mount a large regenerative response and show a central role for Activin signaling in mediating this, providing potential therapeutic opportunity for stroke and other forms of traumatic brain injury where injury and a priming inflammation co-exist. Identification of the regenerative potential of injured CNS neurons in the CAST/Ei strain provides

an opportunity to begin to exhaustively interrogate molecular mechanisms that may enable CNS regeneration in mammals.

## EXPERIMENTAL PROCEDURES

Additional information can be found in [Supplemental Information](#).

### Mice

All strains of mice were purchased from Jackson Laboratory (Bar Harbor). All animal experiments except those detailed in [Figure 4](#) (stroke model) were conducted according to institutional animal care and safety guidelines at Boston Children's Hospital and Harvard Medical School. Stroke model animals were used in accordance with the University of California, Los Angeles Chancellor's Animal Research Committee.

### Sciatic Nerve Crush

Sciatic nerve crush was performed by exposing the left sciatic nerve at the level of sciatic notch and crushing with smooth forceps for 30 s.

### Primary Dissociated DRG Cultures

For pre-conditioned DRG cultures, ipsilateral L4/L5 DRGs were collected 5 days after a sciatic nerve crush. Appropriate L4/L5 DRGs from five mice (naive or pre-conditioned) were pooled and processed in one tube per experiment as in [Ma et al. \(2011\)](#) and [Seiffers et al. \(2007\)](#). Once dissociated, 3,000 cells were grown per chamber in an eight-well slide, coated with myelin (100 ng/well) or CSPGs (20 ng/well, Millipore), and 1,500 cells were grown on laminin (5  $\mu$ g/well, Invitrogen).

### Neurite Growth Assays

For quantification of sensory neuron culture experiments, neurons were plated in eight-well chamber slides. For data given in [Figures 1A–1C](#), nine non-overlapping low-power (4 $\times$ ) microscope images were taken per well, and at least four wells were assayed per condition. All neurons in these images were analyzed by NeuroMath software. Neurons with neurites exceeding 50  $\mu$ m were defined as sprouting cells, and the top 10% of the longest neurite lengths were quantified to obtain an average score; three independent experiments were performed. The values shown are the averages of these experiments and SEM ( $n = 3$ ). For data given in [Figures 1D](#) and [1E](#), all neurons in at least four chambers of an eight-well slide were counted. Images were collected with ImageXpress Micro High-Content Systems, and neurite outgrowth was automatically quantified using the MetaXpress software.

### Activin In Vitro Gain- and Loss-of-Function Experiments

For gain-of-function experiments, either 10 ng/ml of recombinant Activin A, B, and AB (R&D Systems) or PBS containing 0.1% BSA was added into myelin-coated eight-chamber slides with 3,000 naive or pre-conditioned C57BL/6 sensory neurons. For loss-of-function experiments, 3,000 naive and pre-conditioned CAST/Ei sensory neurons were cultured on myelin-coated wells for 24 hr in 100 nM SB-431542 (Tocris) in 1% final concentration ethanol. Vehicle control, 1% final volume ethanol.

For nuclear pSmad2/3 counts, C57BL/6 and CAST/Ei DRG naive and pre-conditioned DRG neurons cultured with or without the presence of either Activin or SB-431542 were assayed for nuclear pSmad2/3 expression by immunostaining (Cell Signaling, #8828). Cells were defined as either containing no, low, or strong pSmad2/3 signal within the nucleus. The percentages of the cells lacking nuclear pSmad2/3 were compared. The values given are the averages of at least three independent experiments counted blind to treatment.

### Spinal Cord Injury

Mice 6 weeks old were anesthetized with a mixture of ketamine/xylazine and received either sciatic nerve transection or sham operation. At 1 week later, a T10 laminectomy was performed, and the dorsal half of the spinal cord was crushed with #5 modified forceps (Dumont, Fine Science Tools). The forceps were deliberately positioned to cut dorsal column axons completely. At 6 weeks after spinal cord injury, axons were traced by injecting 2  $\mu$ l of 10% micro-Ruby tracer (3,000 MW, Invitrogen) into the left sciatic nerve. Mice were

kept for an additional 2 weeks before termination. For the quantification of dorsal column injury model, the retrogradely labeled axons in a series of regularly spaced sagittal sections (C57BL/6  $n = 5$ , CAST/Ei  $n = 7$ ) were imaged, and the number of regenerating axons at different distances from the lesion epicenter was normalized to the number of labeled axons caudal ( $-0.5$  mm/LS) to the lesion. The lesion site was defined according to GFAP immunostaining. Transverse sections of the spinal cord (3 mm caudal and 7–8 mm rostral to the lesion) were taken to confirm the completeness of the lesion and tracing efficiency among experimental groups. Mice with incomplete lesions were excluded from the analysis. Quantification was performed blinded to the treatment and the mouse strain.

### ON Injury

ON crush surgery and intraocular injections were performed under general anesthesia. The mice received either 6.4  $\mu\text{g}$  or 25  $\mu\text{g}$  of zymosan 5 days prior and immediately after ON crush into the vitreous. At 5 days prior to sacrifice, CTB (3  $\mu\text{L}$ , 1%; 103B, List Biological) was injected as an anterograde tracer. For C57BL/6 control  $n = 5$ ; 6.4  $\mu\text{g}$   $n = 4$ ; 25  $\mu\text{g}$   $n = 5$ . For CAST/Ei control  $n = 4$ ; 6.4  $\mu\text{g}$   $n = 4$ ; 25  $\mu\text{g}$   $n = 6$ . Quantification was performed blinded to the treatment and mouse strain.

### Quantification of ON Axon Regeneration

At 14 days after ON injury, mice were perfused with saline and 4% PFA. Regenerating axons were visualized by staining with primary antibodies to CTB (1:500, GWB-7B96E4, GenWay). Axons were counted manually in at least eight longitudinal sections per case at pre-specified distances from the injury site, and these numbers were converted into the number of regenerating axons at various distances (Kurimoto et al., 2010).

### ON Gain-of-Function Experiment

C57BL/6 mice were injected with 6.4  $\mu\text{g}$  of zymosan 5 days prior to and immediately after ON crush to prime RGCs. The animals were then injected with either 0.1% BSA or 10 ng/ml of Activin A, B, and AB immediately after ON crush. A total of five mice were assayed for both control and Activin-treated groups. Animals were sacrificed 14 days after ON injury, and axonal regeneration quantified. Quantification as above was performed blinded to the treatment and mouse strain.

### Stroke Model

Stroke was produced in the forelimb motor cortex as described (Clarkson et al., 2013; Li et al., 2010; Overman et al., 2012). The neuroanatomical tracer biotinylated dextran amine (BDA) was injected into the motor cortex anterior to the stroke. The connections labeled by BDA in the ipsilateral hemisphere were quantified in tangential cortical sections and statistically tested between stroke and control in CAST/Ei and C57BL/6 as described (Clarkson et al., 2013; Li et al., 2010; Overman et al., 2012). For further details see Supplemental Information.

### Microarray Analysis

Total RNA was extracted from naive and injured DRGs (six DRGs per sample, three mice) across nine strains and assessed with the Agilent Bioanalyzer (Agilent Technologies). A total of four biologically distinct replicates were run per sample for a total of 72 arrays. Total RNA (200 ng) was amplified, biotinylated, and hybridized to Illumina MouseRef-8 v2.0 Expression BeadChips. Raw data were analyzed using Bioconductor packages. Normalized expression values were used to compute Pearson correlations with phenotypic data.

### RNA-Seq Analysis

Reads were quality controlled using FastQC, mapped to the reference genome using Bowtie2 (Langmead et al., 2009), and counted using summarizeOverlaps from the GenomicRanges package, and genes were called differentially expressed using edgeR (Robinson et al., 2010). We searched for genes whose expression is significantly increased after pre-conditioning injury in the DBA/2 and CAST/Ei strain to specifically select for genes responsible for the medium-low to strong growth phenotype using methods described in Supplemental Information. This resulted in a list of 16 genes for which we plotted the expression fold change (compared to the C57BL/6 naive expression level) across all six conditions.

### Time Course of Expression of *Inhba*, *Inhbb*, and Activin Receptor Subtype Transcripts

Ipsilateral L4/L5 DRGs were dissected from naive mice and 1, 3, 5, 10, and 14 days post-sciatic nerve crush for both C57BL/6 and CAST/Ei. Each RNA sample was three animals pooled or six DRGs. A total of four distinct samples that were collected for naive ( $n = 4$ ) and three distinct samples for injured DRGs ( $n = 3$ ) for each time point were analyzed by QRT-PCR; mean fold change  $\pm$  SEM normalized to naive C57BL/6 values. For the quantification of mRNA expression of the retina in naive, post ON crush, and post ON crush with 6.4  $\mu\text{g}$  zymosan- and 25  $\mu\text{g}$  zymosan-treated C57BL/6 and CAST/Ei, three retinas were pooled per RNA sample. Three independent RNA samples were analyzed by qRT-PCR ( $n = 3$ ); mean fold change  $\pm$  SEM normalized to naive C57BL/6 values.

### ACCESSION NUMBERS

The accession number for the array and RNA-seq data reported in this paper is GEO: GSE55424.

### SUPPLEMENTAL INFORMATION

Supplemental Information includes Supplemental Experimental Procedures, seven figures, and five tables and can be found with this article online at <http://dx.doi.org/10.1016/j.neuron.2015.05.005>.

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