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**F**. BrdU<sup>+</sup> cell population breakdown. EGFP<sup>+</sup>CC1<sup>+</sup> oligodendrocytes, EGFP<sup>neg</sup>GFAP<sup>+</sup> astrocytes and NG2<sup>+</sup> cells comprise about half of all BrdU<sup>+</sup> cells in the chronic animal.

Figure <u>6</u>. *In vivo* treatment with glial growth factor 2 and basic fibroblast growth factor spinal cord injury results in increased numbers of EGFP<sup>+</sup>CC1<sup>+</sup> oligodendrocytes.

SCI Mice were injected subcutaneously with 0.02mg/kg FGF2 and 0.8mg/kg GGF2 for 8 days, beginning at 1 DPI, and sacrificed on day 9. NG2 immunoreactivity was increased in GGF2 + FGF2 mice (**A**), compared to saline-injected controls (**B**). EGFP<sup>+</sup>CC1<sup>+</sup> oligodendrocytes were also increased in GGF2 + FGF2 treated mice (**C**). **D**. Quantification of NG2<sup>+</sup> cells indicating a significant increase in EGFP<sup>neg</sup>NG2<sup>+</sup> cells. **E**. Quantification of CC1<sup>+</sup> cells indicating a significant increase in EGFP<sup>+</sup>CC1<sup>+</sup> oligodendrocytes; bars represent mean  $\pm$  SEM. Two-way ANOVA was performed with Bonferrroni's post-hoc analysis.

## SUPPLEMENTAL MATERIAL

## **<u>Supplemental Figure</u>** 1. Histopathology and functional deficits after spinal cord injury in the CNP-EGFP mouse

**A**. Representative sections depicting eriochrome (myelin, blue), hemotoxylin (nuclei, purple) and eosin/phyloxine (cytoplasm, pink) stain showing an uninjured control spinal cord section (top) and sections from 1, 3, and 7 days post-injury (top to bottom, respectively), every 1mm from 2mm rostral to 2mm caudal. Scale bar =  $500 \mu m$ .

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**B**. Total Spared White Matter, measured as a percentage of the total cross-sectional area of the spinal cord at the epicenter, decreases with time after injury, and is significantly decreased at the injury epicenter as early as 1 day. Results represented as mean  $\pm$  SEM. One-way ANOVA with Tukey's post-hoc analysis, \* p<0.05; \*\* p<0.01 vs. Control. p<0.05; t p<0.01 vs. 1 day, # p<0.05 vs. 3 days. N = 8 animals for all timepoints.

C. Hindlimb functional recovery after spinal cord injury. Animals exhibit significant hindlimb paralysis, slight movement at the ankle joint, no plantar placement (BMS score 1) at 1 day post-injury, and recover to weight supported plantar stepping with little coordination (BMS score 5-6) by 7 days. Continued recovery occurs with further significant increase at 35-42 days. Symbols represent mean  $\pm$  SEM. N = 8 animals for acute timepoints, N = 6 for chronic timepoints. Two-way ANOVA with Bonferroni's post-hoc analysis was performed. \* = p<0.01 vs. 1 day, # = p<0.05vs. 7 days.

**D**. CNP-EGFP cells (green) are distributed in both grey and white matter in uninjured control tissue. The white dotted line delineates grey matter. By 1 day at the injury epicenter, EGFP cells are visibly lost dorso-centrally. This loss continues at 7 days and is maintained at 42 days post-injury. GM = gray matter, WM = white matter. Sections are 10 $\mu$ m coronal sections from thoracic level 8 of CNP-EGFP transgenic mice. Scale bar = 250 $\mu$ m.

E. Upper panel:  $CC1^+$  oligodendrocytes and  $GFAP^+$  astrocytes (asterisks) in the uninjured ventral-lateral white matter. Inset shows a  $GFAP^+$  cell neighboring a  $CC1^+$  cell.  $CC1^+$  cells are always EGFP^+.  $GFAP^+$  cells are always EGFP^-. Scale bars = 10  $\mu$ m. Lower panel: EGFP<sup>+</sup>CC1<sup>+</sup> oligodendrocytes and EGFP<sup>-</sup>GFAP<sup>+</sup> astrocytes (asterisks) at 1

day post-injury. After injury,  $CC1^+$  cells are always EGFP<sup>+</sup> and GFAP<sup>+</sup> cells are always EGFP-. Scale bars = 10  $\mu$ m.

**F-G.** Quantification of DAPI<sup>+</sup> nuclei, and of EGFP<sup>+</sup>CC1<sup>+</sup> oligodendrocytes and EGFP<sup>neg</sup>GFAP<sup>+</sup> astrocytes after SCI. Most oligodendrocyte and astrocyte loss occurs at the epicenter by 1 day after injury; loss at the epicenter is significant at all time points. Represented as Mean  $\pm$  SEM. One-Way ANOVA. \* = p<0.05, \*\* = p<0.01, \*\*\* = p<0.001 vs. Control. However, this loss is not reflected in the density of DAPI<sup>+</sup> nuclei. Black boxes in the ventral-lateral white matter indicate where cell counts were performed.

<u>Supplemental Figure 2. Nestin and S100beta are expressed in subsets of</u> EGFP<sup>+</sup>NG2<sup>+</sup> cells in the spinal cord and after spinal cord injury.

**A.** Example of a nestin (blue)<sup>+</sup> EGFP<sup>+</sup> (green) NG2<sup>+</sup> (red) cell in the ventrolateral white matter in injured tissue at 3 DPI. Scale bar =  $10\mu m$ .

**B.** Quantitative analysis for EGFP<sup>+</sup>NG2<sup>+</sup>nestin<sup>+</sup> cells in the control animal and in the injured animal at 1, 3 and 7 DPI, at 1.5mm rostral, epicenter, and 1.5mm caudal. Cell counts were performed in the ventral-lateral white matter. A side-by-side comparison for EGFP<sup>+</sup>NG2<sup>+</sup> and EGFP<sup>+</sup>NG2<sup>+</sup>nestin<sup>+</sup> cells is shown for each timepoint. A subset of EGFP<sup>+</sup>NG2<sup>+</sup> cells express nestin. Immunoreactive cells were quantified for 5 animals every 0.5 mm from the injury epicenter, and at the injury epicenter in the spared ventral-lateral white matter. For each distance at each timepoint, three 10 µm-thick sections were counted, left and right hemispheres, for a total of six fields per animal. Therefore, thirty

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200 µm2 fields of view were quantified for each. One-Way ANOVA with Tukey's post-
hoc analysis for variance. * p<0.05, ** p<0.01, *** p<0.001 vs. control.
<u><b>C.</b></u> Example of S100 $\beta^+$ (blue) EGFP <sup>+</sup> (green) NG2 <sup>+</sup> (red) cells in the ventrolateral white
<u>matter in injured tissue at 3 DPI. Scale bar = <math>10\mu m</math>.</u>
<b>D.</b> Quantitative analysis for EGFP <sup>+</sup> NG2 <sup>+</sup> S100 $\beta$ <sup>+</sup> cells in the control animal and in the
injured animal at 1, 3 and 7 DPI, at 1.5mm rostral, epicenter, and 1.5mm caudal. Cell
counts were performed in the ventral-lateral white matter. A side-by-side comparison for
<u>EGFP<sup>+</sup>NG2<sup>+</sup></u> and EGFP <sup>+</sup> NG2 <sup>+</sup> S100 $\beta$ <sup>+</sup> cells is shown for each timepoint. A subset of
EGFP <sup>+</sup> NG2 <sup>+</sup> cells express S100β. Immunoreactive cells were quantified for 5 animals
every 0.5 mm from the injury epicenter, and at the injury epicenter in the spared ventral-
lateral white matter. For each distance at each timepoint, three 10 µm-thick sections were
counted, left and right hemispheres, for a total of six fields per animal. Therefore, thirty
200 µm2 fields of view were quantified for each. One-Way ANOVA with Tukey's post-
hoc analysis for variance. * p<0.05, ** p<0.01, *** p<0.001 vs. control.

## <u>Supplemental Figure 3. Olig2, Sox10 and Sox17 expression in EGFP<sup>+</sup>NG2<sup>+</sup> cells is</u> <u>altered after spinal cord injury.</u>

**A.** Examples of EGFP<sup>+</sup>NG2<sup>+</sup> cells (blue) expressing Olig2 (red) in acutely injured (top panels) and chronically injured (bottom panels) white matter. DAPI (white) labels nuclei. Scale bar =  $10\mu m$ .

**B-C.** Cell counts of EGFP<sup>+</sup>NG2<sup>+</sup>Olig2<sup>+</sup> cells in acutely injured (**B**; 1, 3 and 7 DPI) and chronically injured (**C**; 6 weeks DPI) at 1.5mm rostral, at the injury epicenter, and at

<u>1.5mm caudal. Cell counts were performed as described above. One-Way ANOVA with</u> Tukey's post-hoc test for variance. \* p < 0.05; \*\* p < 0.01 vs. control.

**D.** Examples of EGFP<sup>+</sup>NG2<sup>+</sup> cells (blue) expressing Sox10 (red, top panels) and Sox17

(red, bottom panels) in acutely injured animals. DAPI (white) labels nuclei. Scale bar = 10µm.

**E-F.** Cell counts of Sox10<sup>+</sup>EGFP<sup>+</sup>NG2<sup>+</sup> (**E**) and Sox17<sup>+</sup>EGFP<sup>+</sup>NG2<sup>+</sup> (**F**) cells in acutely injured (1, 3 and 7 DPI) at 1.5mm rostral, at the injury epicenter, and at 1.5mm caudal. Cell counts were performed as described above. One-Way ANOVA with Tukey's posthoc test for variance. \* p < 0.05; \*\* p < 0.01 vs. control; \*\*\* p < 0.001.