Supplemental Information

Combination of In Situ Lcn2 pRNA-RNAi

Nanotherapeutics and iNSC Transplantation

Ameliorates Experimental SCI in Mice

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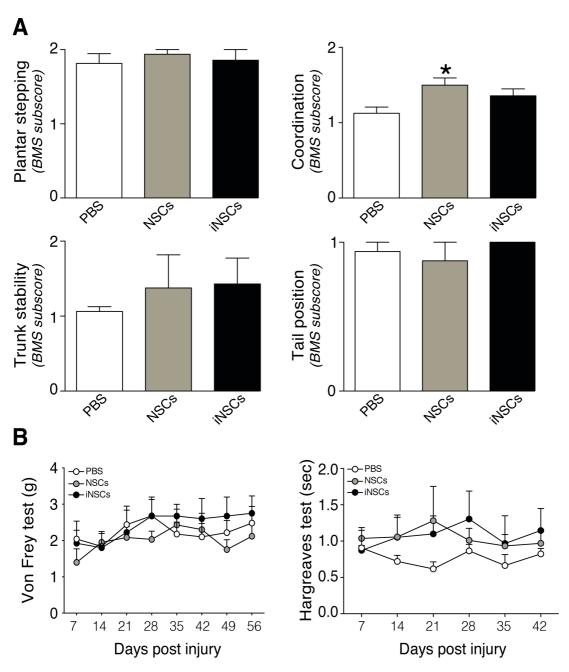


Figure S1. Evaluation of fine locomotion after NSC transplantation and assessment of allodynia.

(A) BMS sub-score parameters (plantar stepping, coordination, trunk stability, tail position) at 56 dpi. Data are mean values (\pm SEM) from n \geq 4 mice per group. *p< 0.05, vs. PBS group. (B) Mechanical allodynia (left) and thermal allodynia (right) by Von Frey test and Hargreaves test, respectively. Data are mean values (\pm SEM) from n \geq 4 mice per group. BMS = Basso Mouse Scale; dpi = days post injury.

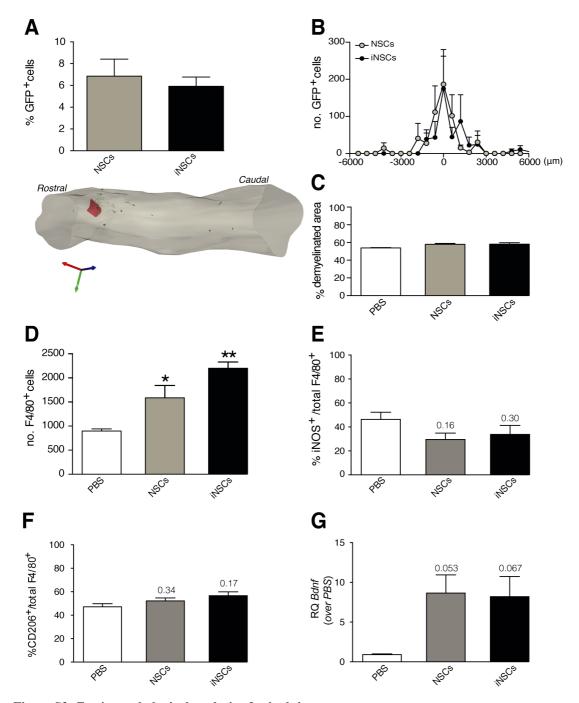


Figure S2. Ex vivo pathological analysis of spinal tissue.

(A) Proportion of surviving fGFP-NSCs or fGFP-iNSCs at 56 dpi over total transplanted cells. Data are mean values (± SEM) from n=3 mice per group. Representative 3D reconstruction depicting the lesion volume in red and fGFP-iNSCs as green dots. (B) Evaluation of migratory capabilities of transplanted cells. Distance from the lesion epicentre (0) is represented in μm. Data are mean values (± SEM) from n=3 mice per group. (C) Quantification of Luxol Fast Blue demyelinated area. Data are mean % percentage of demyelinated area over the total spinal cord section area (± SEM) from n=3 mice per group. (D) Quantification of F4/80-positive cells at 56 dpi. Data are mean numbers (± SEM) from n=3 mice per group. *p< 0.05 and **p< 0.01, vs. PBS. (E) iNOS-positive cells over the total number of F4/80-positive cells at 56 dpi. (F) CD206-positive cells over the total number of F4/80-positive cells at 56 dpi. Data in E and F are mean % of double positive cells over the total number of F4/80-positive cells (± SEM) from n≥5 mice per group. (G) qRT-PCR of Bdnf mRNA expression levels, expressed as a Relative Quantification (RQ) versus the Gapdh reference gene. Data are mean mRNA expression (vs. PBS) (± SEM) from n=3 mice per group. dpi = days post injury.

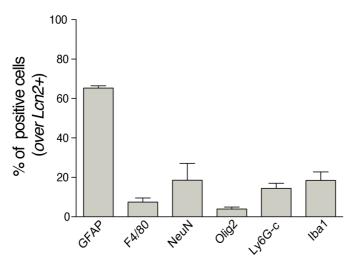


Figure S3. Lcn2 expression in spinal tissue.

Relative percentages of cell types expressing Lcn2 in the spinal cord at 21 dpi. Data are expressed as mean % of double positive cells over the total number of Lcn2-positive cells (± SEM) from n=3 mice. dpi = days post injury.

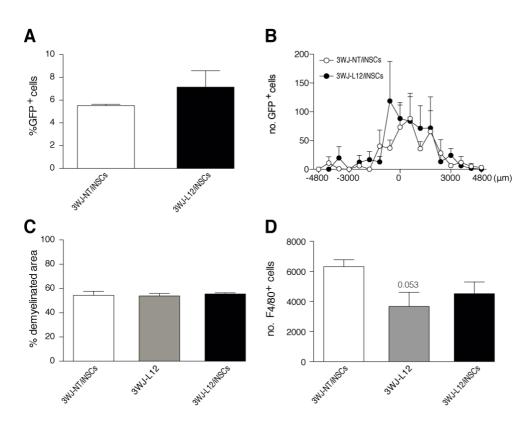


Figure S4. Ex vivo pathological analysis of spinal tissue from combinatorial treatment groups.

(A) Surviving fGFP-iNSCs in the spinal cord at 56 dpi over total transplanted cells. Data are mean % (\pm SEM) from n=3 mice per group. (B) Evaluation of migratory capabilities of transplanted cells. Distance from the lesion epicentre (0) is represented in μ m. Data are mean values (\pm SEM) from n=3 mice per group. (C) Quantification of Luxol Fast Blue demyelinated area. Data mean % of demyelinated area over the total area of the spinal cord section (\pm SEM) from n=3 mice per group. (D) Quantification of F4/80-positive cells at 56 dpi. Data are as mean values (\pm SEM) from n \geq 4 mice per group. Dpi = days post injury.