Supplementary Materials: Cellular Changes in Injured Rat Spinal Cord Following Electrical Brainstem Stimulation

Antibody	Manufacturer	Host	Dilution			
Primary						
BrdU	Novus	Rat	1:100			
BrdU	Roche	Mouse	1:100			
NeuN	Millipore	Mouse	1:1000			
Blbp	Abcam	Rabbit	1:100			
Sox2	Santa Cruz	Rabbit	1:500			
GFAP	Invitrogen	Rat	1:500			
CD68	Millipore	Mouse	1:500			
Double cortin	Millipore	Guinea Pig	1:100			
APC	Abcam	Mouse	1:500			
Arginase-1	Invitrogen	Rat	1:200			
iNOS	Novus	Rabbit	1:500			
NG2	Millipore	Mouse	1:500			
Secondary (488, 594, 647 nm)						
Mouse	Alexa Fluor	Goat	1:10,000			
Rabbit	Alexa Fluor	Goat	1:10,000			
Rat	Alexa Fluor	Goat	1:10,000			

Table S1. List of antibodies and concentrations used.

Table S2. Univariate and multivariate regression analyses with all three spinal cord subregions included in analyses. All cell counts are measured at the level of the lesion, without pooling of cell counts from either of the spinal cord subregions. On the left are shown mean (\pm SE) counts for the immune and neuroglial progenitor markers. P values are estimated for individual comparisons using bootstrap analysis. Bolded are regressors with *p* < 0.05. On the right are shown counts of markers co-stained with BrdU. At the bottom of the table are shown the results of multivariate stepwise regression analysis, which yielded a significant model associated with stimulation for the regressors on the left side of the table, but not for those on the right side of the table. Abbreviations: Stim., stimulation; NS, not significant.

Immune/Precursor Cells			Proliferation				
Marker	Control	Stim.	Р	Markers	Control	Stim.	Р
				BrdU	587 ± 50	519 ± 49	NS
CD68	300 ± 39	224 ± 28	0.043	CD68/BrdU	88 ± 20	71 ± 20	NS
CD68/iNOS	16 ± 7	30 ± 10	NS				
CD68/Arg-1	21 ± 6	36 ± 11	NS				
GFAP	187 ± 29	185 ± 29	NS	GFAP/BrdU	55 ± 6	67 ± 10	NS
NG2	181 ± 41	252 ± 64	NS	NG2/BrdU	49 ± 13	59 ± 17	NS
Sox2	384 ± 61	250 ± 42	0.016	Sox2/BrdU	74 ± 11	64 ± 4	NS
Blbp	181 ± 38	349 ± 67	0.000	Blbp/BrdU	14 ± 2	22 ± 6	0.047
Stepwise regression model:			Stepwise regression model:				
NRM LFS ≈ Blbp + CD68			N/A				
$(F_2 = 51.3, p < 0.001)$							