### ➢ General data

1	Title	
2	First author	
3	Year of publication	
4	Reviewer	

### > Methods:

	Domain	Description	Reviewers'
			judgment
			High risk/ low risk/ unclear <sup>1</sup>
1	Fish strain		
2	Fish species		
3	Fish gender		
4	Number of fishes per group		
5	Genetic background		
6	Sample size		
7	Follow up period		
8	Age or weight (at the beginning and the end of the experiment)		
9	Method of allocation to treatment group: i.e. randomly assigning animals to a specific group		
10	Blindness of assessor		
11	Description of the method of injury		
12	Description of the severity of injury		

<sup>&</sup>lt;sup>1</sup>Low risk: no bias; Unclear risk: not mentioned; High risk: bias

13	Description of the level of injury	
14	Regulations and ethics	
15	Description of statistical analysis of data	
16	Using of appropriate tests to prove the research question and hypothesis	
17	Definition of the control group	
18	Method and drugs used for anesthesia/antibiotics	
19	Bladder expression	
20	Method of killing	
21	Experimental unit	
22	Housing conditions	
23	Exclusion of animals from the experiment	

## > Results

# Primary Outcome – Micro

	Outcome measures		Time	Events
Second Protein MRN Protein   A L1.1, L1.2 L1.1, L1.2   GAP-43 NCAM   NCAM ZFNLRR   HMGB1 SOX1   SOX1 Contactin-1   PtneaPtena CRP1	PO1: Expression of growth related molecules	SeriesmRN AProteinAL1.1, L1.2GAP-43uitouuoudSOX1VMOD SOX1SOX1Contactin-1PtneaPtenaCRP1		

Legumain SOX2		Legumain			
	MVP				
		PO (protein zero)			
		MiR-133b			
	ý	mRNA	Protein		
bitor		Nogo-A			
	f inhi ules	Notch-her 1			
	iion o molec	LPA (lipopolys	sacharid acid)		
	press	PtnebPtenb			
	Ex	Rho			
	Bobo1,2,Slit3,Sema3,b,Sema 3h,Plexina,mmp9,etc				
	I	Degeneration	Apoptosis Necrosis		
		Regeneration/neurogenesis			
		(type of neurons; motor,			
su		Synaptic formation			
leuro		Neurite sprouting/growth			
PO2:N		Axonal changes	Number(total, regenerated) Remyelination Repatterning Pathway		
		Escape behavior	r of growth cone		
		Proliferation			
13 : ortive 11s	Astrocyte	Morphology	Glial bridge formation		
Jdns			Typical morphology		

		Apoptosis/necrosis/ death
		Proliferation
		Migration (infiltration)
		Reactivity
		Differentiation to neurons
		(progenitor cell)
	roc	Apoptosis/necrosis/ death
	dendı e	Proliferation
	gogoo yt	Migration (infiltration)
	Olig	Reactivity/morphology
		Apoptosis/necrosis/ death
	Schwann	Proliferation
		Migration (infiltration)
		Morphology
	Microglia/macr ophage	Apoptosis/necrosis/ death
		Proliferation
		Migration (infiltration)
		Reactivity/ Morphology
	Ependyma	Apoptosis/necrosis/ death
		Proliferation
		Migration
		Reactivity/ Morphology
		Differentiation to neurons
		(progenitor cell)

PO4: Functional recovery (swimming)		

# Secondary Outcome – Macro

Outo	come measures	time	Events
Out		unite	
S01	Hemorrhage/ spinal cord blood flow		
S02	Edema (Gray matter, white matter)		
S03	Vasospasm / Blood- SpinalCord-Barrier rupture		
S04	Necrosis/ thrombosis/ infraction		
SO5	Regeneration		
S06	Gliosis/ scar formation		
SO7	Degeneration/tissue loss		
S08	Injury volume / atrophy		

> Note:

1	Molecular (name, place of expression,	
	its effectiveness etc)	
2	Cellular (type, position and	
	effectiveness)	
3	Position of recovery in spinal cord	
4	Direction of recovery (e.g. caudal,	
	rostral,)	
5	inhibitory or activation mechanism of	
	molecule or cell	
6	Model of recovery evaluation such as	
	cellular, immunohistochemistry,	
	molecular (PCR), physical activity	
	(swimming)	
	<b>.</b>	

\*Limitations of the study