

Lipocalin 2 Plays an Immunomodulatory Role and Has Detrimental Effects after Spinal Cord Injury

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Lipocalin 2 (Lcn2) plays an important role in defense against bacterial infection by interfering with bacterial iron acquisition. Although Lcn2 is expressed in a number of aseptic inflammatory conditions, its role in these conditions remains unclear. We examined the expression and role of Lcn2 after spinal cord injury (SCI) in adult mice by using a contusion injury model. Lcn2 expression at the protein level is rapidly increased 12-fold at 1 d after SCI and decreases gradually thereafter, being three times as high as control levels at 21 d after injury. Lcn2 expression is strongly induced after contusion injury in astrocytes, neurons, and neutrophils. The Lcn2 receptor (Lcn2R), which has been shown to influence cell survival, is also expressed after SCI in the same cell types. Lcn2-deficient (*Lcn2*^{-/-}) mice showed significantly better locomotor recovery after spinal cord contusion injury than wild-type (*Lcn2*^{+/+}) mice. Histological assessments indicate improved neuronal and tissue survival and greater sparing of myelin in *Lcn2*^{-/-} mice after contusion injury. Flow cytometry showed a decrease in neutrophil influx and a small increase in the monocyte population in *Lcn2*^{-/-} injured spinal cords. This change was accompanied by a reduction in the expression of several pro-inflammatory chemokines and cytokines as well as inducible nitric oxide synthase early after SCI in *Lcn2*^{-/-} mice compared with wild-type animals. Our results, therefore, suggest a role for Lcn2 in regulating inflammation in the injured spinal cord and that lack of Lcn2 reduces secondary damage and improves locomotor recovery after spinal cord contusion injury.

Introduction

Spinal cord injury (SCI) is associated with an inflammatory response that contributes to delayed secondary damage. Activated astrocytes, microglia, and neurons contribute to the initial inflammatory response by producing a number of pro-inflammatory cytokines and mediators (Bartholdi and Schwab, 1997; Yang et al., 2004, 2005; Pineau and Lacroix, 2007; Pineau et al., 2010) that are cytotoxic for neurons and glial cells at the lesion site. In addition, some of these molecules are important for recruiting immune cells to the lesion site and play a role in modulating the wound-healing response after SCI (Popovich and Longbrake, 2008).

Lipocalin 2 (Lcn2), also known as neutrophil gelatinase-associated lipocalin (NGAL), is a siderophore-binding protein that is reported to play an important role in the innate immune

response to certain kinds of bacterial infection (Flo et al., 2004). Siderophores are bacterial molecules that bind iron and are internalized by a receptor-mediated mechanism. They allow bacteria to obtain iron, an element that is critical for their growth and survival, from within the host (Wandersman and Delepelaire, 2004). The biological functions of Lcn2 stem from its ability to bind iron-loaded siderophores (Goetz et al., 2002), thus depriving pathogenic bacteria of iron. Lcn2 has also been shown to exert its effects on mammalian cells through its receptor Lcn2R (24p3R). Cells that expressed Lcn2R were sensitized to Lcn2-triggered apoptosis through a mechanism dependent on the intracellular depletion of iron (Devireddy et al., 2005). Recent reports have suggested that Lcn2 may play a similar role in sensitizing microglia and astrocytes *in vitro* to cell death induced by cytotoxic agents such as nitric oxide (Lee et al., 2007, 2009). In the CNS, expression of Lcn2 has been reported in the choroid plexus in response to peripheral lipopolysaccharide (LPS) administration (Marques et al., 2008), as well as in the striatum after intrastriatal injections of LPS (Glezer et al., 2007). Its expression was also reported to be upregulated in a microarray analysis of SCI tissue (De Biase et al., 2005). However, what role Lcn2 plays in CNS inflammation and injury remains unknown.

Here we report that Lcn2 plays a detrimental role in a mouse model of spinal contusion injury. Using *Lcn2*^{-/-} mice, we show that Lcn2 contributes to inflammation and secondary damage in the injured spinal cord. *Lcn2*^{-/-} mice also show improved locomotor recovery as compared with wild-type controls. Our results

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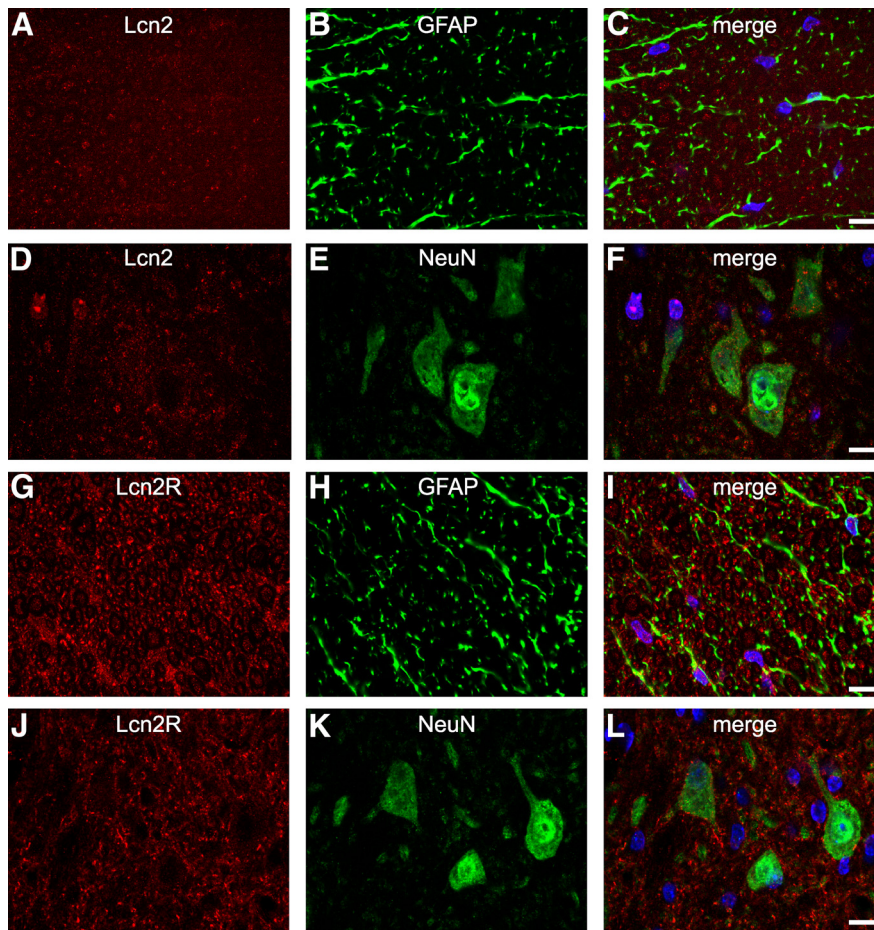


Figure 1. Lcn2 and Lcn2R expression in the uninjured spinal cord. Confocal images show double immunofluorescence labeling for Lcn2 (A–F) and Lcn2R (G–L) in the uninjured spinal cord. Representative micrographs show extremely low levels of Lcn2 and Lcn2R in astrocytes and neurons in the uninjured spinal cord. Scale bars, 10 μ m.

suggest that Lcn2 plays a detrimental role after SCI, likely through the modulation of diverse pathways.

Materials and Methods

Animals. *Lcn2*^{+/+} and *Lcn2*^{-/-} mice on a C57BL/6 background (Flo et al., 2004; Huang et al., 2009) between 2 and 4 months of age were used for spinal cord contusion injury (60 kdyn force, 500–700 μ m tissue displacement) using the Infinite Horizons impactor (Precision Scientific Instrumentation). Mice were deeply anesthetized with ketamine, xylazine, and acepromazine (100, 50, and 3 mg/kg), a single laminectomy was done, and the contusion injury was performed at the T10 thoracic level (Ghasemlou et al., 2005). Motor control was evaluated using the nine-point Basso Mouse Scale (BMS) (Basso et al., 2006) by two observers trained in the Basso laboratory at The Ohio State University. Consensus scores were then recorded. These analyses were performed by observers blinded to experimental groups. All procedures were approved by the McGill University Animal Care Committee and followed the guidelines of the Canadian Council on Animal Care.

Western blotting. A 5-mm-long piece of spinal cord centered at the lesion site was removed at 1, 3, 7, 14, 21, and 28 d postinjury (dpi) ($n = 3$ mice per group) after intracardiac perfusion with 10 ml of 0.1 M phosphate buffer. Total protein was extracted using radioimmunoprecipitation assay buffer containing 1% Nonidet P-40 (Sigma), 1% sodium deoxycholate (BDH Chemicals), 2% SDS, 0.15 M sodium phosphate, pH 7.2, 2 mM EDTA, and a mixture of protease inhibitors (Roche Diagnostics). Twenty micrograms of total protein were loaded onto a 4–12% SDS-PAGE (Novex Bis-Tris gel; Invitrogen) and then transferred to polyvinylidene fluoride membrane (Millipore). Membranes were

blocked with 3% nonfat milk in PBS–Tween 20 (0.05%) and incubated at 4°C overnight with goat anti-mouse Lcn2 (1:400; R&D Systems). Blots were washed and incubated with peroxidase-conjugated IgG (1:200,000; Jackson ImmunoResearch) and detected with Western Lightning Chemiluminescence Reagent Plus (PerkinElmer Life and Analytical Sciences). Blots were subsequently reprobbed with rabbit anti- β -actin (1:400; Sigma-Aldrich) to confirm equal loading. Experiments were done in triplicate.

Quantitative real-time reverse transcription-PCR. Approximately 5 mm of spinal cord tissue containing the lesion site was removed at 1, 3, 7, and 21 dpi. Total RNA was isolated from SCI tissue as described previously (Rathore et al., 2008) using the RNeasy Lipid Tissue kit (Qiagen), and quantitative real-time reverse transcription (QRT)-PCR was performed using the MX4000 (Stratagene) as per the manufacturer's protocol for all mRNA expression analyses. The relative expression between the samples was calculated based on the threshold cycle (C_T) value, and was standardized to the amount of a housekeeping gene, *glyceraldehyde 3-phosphate dehydrogenase (GAPDH)*, using the $\Delta\Delta C_T$ method (Livak and Schmittgen, 2001): $\Delta\Delta C_T = (C_{T,Target} - C_{T,GAPDH}) - (C_{T,Target} - C_{T,GAPDH})_{(calibrator\ avg)}$. Results are shown as the mean (fold increase over uninjured controls) \pm SEM, $n = 4$.

Flow cytometry. Immune cells from the injured spinal cord were analyzed by flow cytometry at 2 dpi as described previously (Stirling and Yong, 2008). Briefly, the spinal cord samples were mechanically disrupted using a Dounce homogenizer and passed through an 80 μ m cell strainer (BD Falcon). Cells were then collected by mild centrifugation and resuspended in PBS containing 1% fetal bovine serum. Samples were divided,

and cells alone and isotype-matched control samples were generated to control for nonspecific binding of antibodies and for autofluorescence. Isotype control antibodies were purchased from BD Biosciences PharMingen and included phycoerythrin (PE)-labeled rat IgG2b, allophycocyanin (APC)-labeled rat IgG2b, and peridinin chlorophyll-a protein (PerCP)-conjugated rat IgG2b. After cell counts, the samples were Fc blocked (BD Biosciences) to minimize background staining. The following antibodies were purchased from BD Biosciences PharMingen and were optimized and used to determine different populations of resident microglia and blood-derived leukocytes: APC-conjugated rat anti-mouse CD45 (leukocyte common antigen, Ly-5), PE-conjugated rat anti-mouse Ly-6G and Ly-6C (Gr-1), and PerCP-conjugated rat anti-mouse CD11b. After 30 min of incubation with combinations of antibodies at 4°C, the samples were washed and fixed in 1% paraformaldehyde. The samples were then immediately analyzed with a FACSCalibur Flow Cytometer (BD Biosciences). Equal numbers of cells were analyzed for all conditions. Density plots were created using CellQuest software (BD Biosciences), and cells alone and isotype controls were used to set quadrants. CD45-positive gates ensured that only leukocytes and resident microglia were included in the analysis. The proportions of blood-derived leukocytes (neutrophils and monocytes) (CD45^{high}:CD11b), neutrophils (CD45^{high}:Gr-1^{high}), and resident microglia (CD45^{low}:CD11b) were determined from density plots (CD45, y -axis; CD11b or Gr-1, x -axis) using CellQuest software and compared between the different groups ($n = 4$).

Immunofluorescence and histological analyses. Animals were perfused with 0.1 M phosphate buffer, followed by 4% paraformaldehyde in 0.1 M phosphate buffer, pH 7.2. A 7 mm length of spinal cord tissue centered on the lesion epicenter was dissected out and cryoprotected, and 12- μ m-thick cryostat sections were obtained. Tissue sections were incubated with 0.1 M PBS

containing 2% normal goat serum and 1% ovalbumin to block nonspecific binding of antibodies. Tissue sections were incubated overnight at 4°C with primary antibodies at the following concentrations: goat anti-Lcn2 (1:200; R&D systems), rabbit anti-Lcn2R (1:100; Sigma/Atlas Antibodies), rat anti-gliofibrillary acidic protein (GFAP) (1:800; Invitrogen), rabbit anti-Iba1 (1:500; Wako Chemicals), and rat anti-7/4 (1:500; Serotec) at room temperature. After washing, sections were then incubated for 1 h at room temperature with Alexa-568-conjugated donkey anti-rabbit IgG (1:200; Invitrogen), Alexa-488-conjugated donkey anti-rat IgG (1:200; Invitrogen), or Alexa-568-conjugated donkey anti-goat (1:200; Invitrogen). Nuclear staining for confocal analysis was done by incubating sections with TOPRO-3 (1:5000, Invitrogen). Sections were mounted with Slowfade Gold antifade media (Invitrogen), and a coverslip was applied. For confocal microscopy, immunostained sections were viewed with a spinning disk confocal system (PerkinElmer) connected to an EclipseTE2000 (Nikon). Image capture and analysis were conducted with MetaMorph (Molecular Devices). For a negative control for Lcn2 antibody, we stained spinal cord section from Lcn2 knock-out mice. No staining was detected, as shown in Results. The Lcn2 receptor antibody was generated against a 60-mer recombinant protein fragment of Lcn2R. The antibody was validated using an antigen microarray against 384 different antigens as well as by immunostaining by the company that generated the antibody (Atlas Antibodies). We were, however, unable to obtain the recombinant protein for immunodepletion, and there are no knock-out mice available.

To detect ventral horn neurons, tissue sections were stained by incubating tissue in 0.1% cresyl violet (Sigma-Aldrich) for 10 min or in 0.1% Luxol fast blue overnight and cleared in 0.05% lithium carbonate to stain for myelin. This was followed by dehydration through ascending alcohols and Hemo-De (Thermo Fisher Scientific). All bright-field and immunofluorescent microscopy was conducted on the Axioskop 2 Plus (Zeiss), by using a QImaging Retiga 1300 camera and BIOQUANT image analysis system for image capture and analysis. For the analysis of the GFAP-positive area in the spinal cord cross sections, the pixel density function of the BIOQUANT image analysis system was used to analyze the total cross-sectional area in stained sections. For assessment of myelin sparing, the total extent of myelin staining in the tissue section was quantified using the pixel count function in BIOQUANT and normalized to the pixel count for the entire cross section. Sparing myelin was presented as a percentage of myelin at the lesion site compared with the percentage of myelin in intact spinal cord.

Results

Expression of Lcn2 is markedly upregulated after SCI

We assessed the expression of Lcn2 in the uninjured spinal cord by double immu-

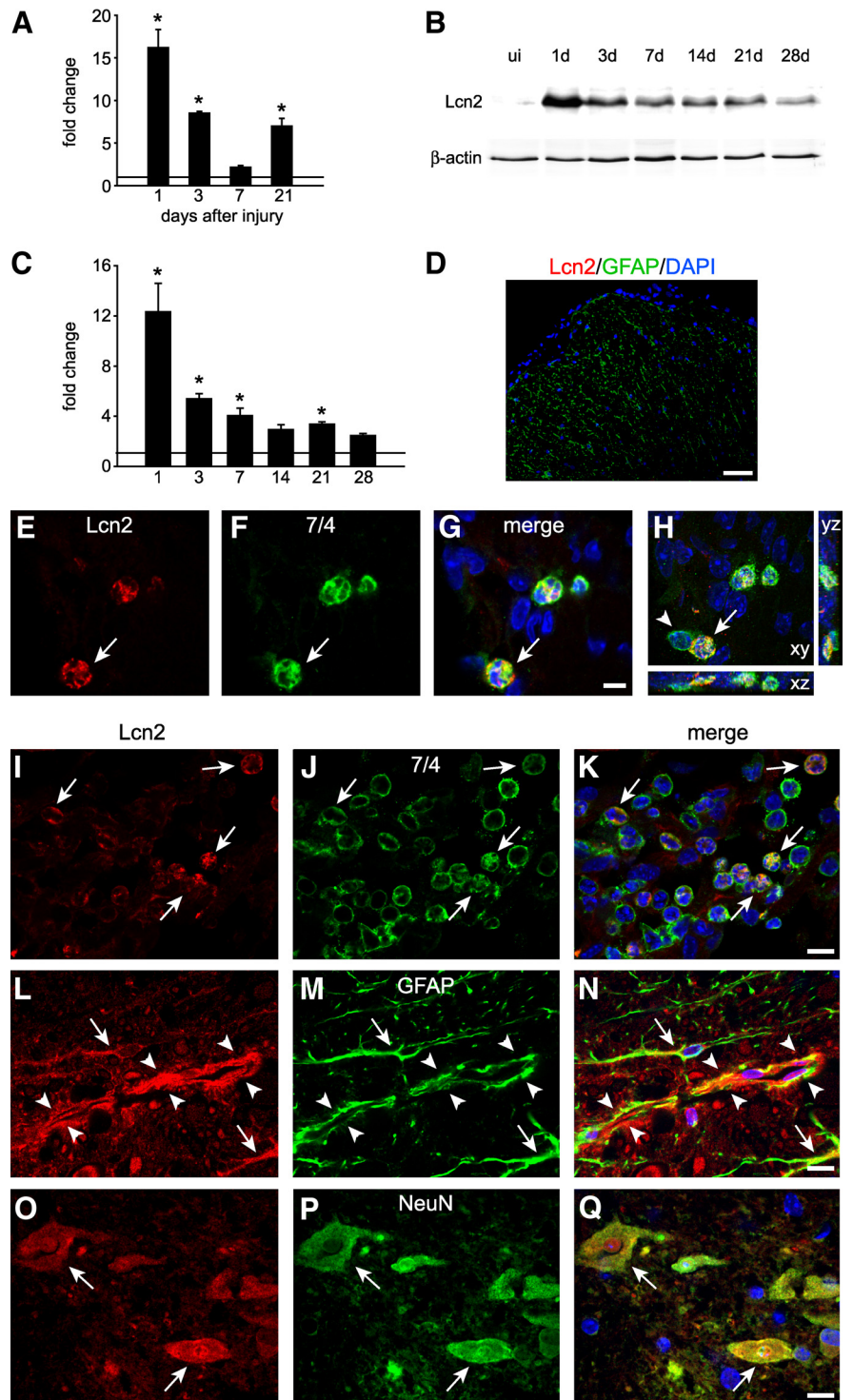


Figure 2. Lcn2 expression after SCI. **A**, Changes in Lcn2 mRNA expression as determined by QRT-PCR in the injured spinal cord at various times after injury. Lcn2 mRNA levels are significantly upregulated above controls at 1, 3, and 21 dpi ($*p < 0.05$, 2-sample unequal variance *t* test, $n = 3$ for each time point). **B**, Representative Western blot showing Lcn2 protein in spinal cord homogenates from uninjured naive (ui) mice and at several time points after SCI. β -Actin was used as a loading control. Note the marked increase immediately after SCI and its gradual decrease thereafter. **C**, Densitometric quantification of Western blot data ($*p < 0.05$, 2-way ANOVA, *post hoc* Tukey; $n = 3$ per group). **D**, Negative staining control shows no anti-Lcn2 labeling (red) in Lcn2^{-/-} spinal cord tissue, double labeled with anti-GFAP (green); nuclei are stained with DAPI. **E–G**, Single-plane confocal images (0.6 μ m thickness) of double immunofluorescence labeling of spinal cord tissue sections 1 d after SCI stained for Lcn2 (red) and 7/4 for neutrophils (green); arrows indicate double-labeled cells. **H**, A full z-stack image (14 μ m) showing double labeling for Lcn2 and 7/4 in the *x–y*, *x–z*, and *y–z* planes. **I–K**, Single-plane confocal images showing double immunofluorescence labeling of infiltrating 7/4-positive neutrophils expressing Lcn2 (arrows) in the injured spinal cord at 3 dpi. Similar confocal images showing that GFAP-positive astrocytes (**L–N**) as well as NeuN-positive neurons (**O–Q**) also express Lcn2 at 1 dpi (arrows indicate double-labeled cells; arrowheads point to a longitudinal profile of a blood vessel that shows Lcn2 labeling of the endothelial cells lining the inner aspect of the GFAP staining). Scale bars: **D**, 50 μ m; **E–G**, 5 μ m; **I–Q**, 10 μ m.

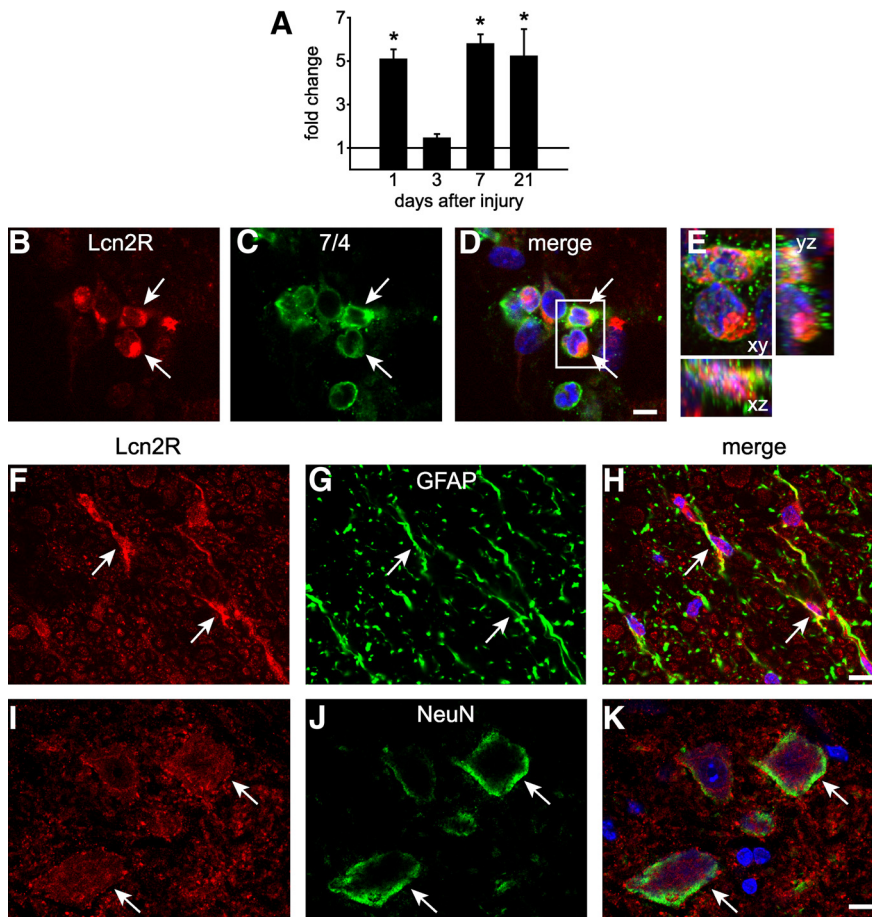


Figure 3. Lcn2 receptor expression after SCI. **A**, Graph showing changes in Lcn2R mRNA expression as detected by QRT-PCR. Expression levels are increased at 1, 7, and 21 dpi ($*p \leq 0.05$; 2-way ANOVA, *post hoc* Tukey). **B–D**, Single-plane confocal images (0.6 μm thickness) showing double labeling for Lcn2R and 7/4 (arrows indicate double-labeled cells) at 1 dpi. **E**, A full z-stack image (14 μm) of the area in the box in **D** showing double labeling for Lcn2R and 7/4 in the *x–y*, *x–z*, and *y–z* planes. **F–K**, Single-plane confocal images showing GFAP-positive astrocytes (arrows) and NeuN-positive neurons at 1 dpi. Scale bars: **B–D**, 5 μm ; **E–J**, 10 μm .

no fluorescence labeling and confocal microscopy. There was minimal Lcn2 staining in the uninjured spinal cord (Fig. 1A–F). After SCI, Lcn2 mRNA was significantly induced ~ 16 -fold higher than control levels at 1 d after contusion injury as determined by QRT-PCR (Fig. 2A). The mRNA expression levels progressively decreased from 3 through 7 dpi, reaching levels approximately twofold higher than control levels at day 7, and increasing to levels approximately sevenfold higher than control levels at 21 dpi. Densitometric quantification of Western blot analysis showed a 12-fold increase in Lcn2 protein at 1 dpi and a gradual decrease thereafter, but the levels still remained at approximately threefold higher than control levels at day 21 (Fig. 2B, C). We next assessed the cellular localization of Lcn2 after SCI using double immunofluorescence labeling and confocal microscopy. Double immunofluorescence staining at 1 dpi showed that 7/4-positive neutrophils close to the lesion site expressed Lcn2 (Fig. 2E–H). Confocal single-plane images showed Lcn2 expression in 7/4-positive neutrophils at 1 dpi (Fig. 2E–G). Colocalization is seen further in the complete z-stack images with *x–y*, *x–z*, and *y–z* planes (Fig. 2H). Single-plane images from 3 dpi tissue also showed expression of Lcn2 in neutrophils at this time point (Fig. 2I–K). Single-plane images from 1 dpi also showed Lcn2 expression in GFAP-positive astrocytes (Fig. 2L–N). Neurons in the ventral gray matter of the injured spinal cord were also positive for Lcn2

(Fig. 2O–Q). At 1 dpi, elongated profiles that are likely to be endothelial cells were also positive for Lcn2 (Fig. 2L–N, arrowheads). Lcn2 was not detected in oligodendrocytes or Iba1-positive microglia (data not shown).

We next assessed the expression of Lcn2R, also known as BOCT (brain-type organic cation transporter), after contusion injury. Lcn2R levels are low in the uninjured spinal cord as determined by double immunofluorescence labeling and confocal microscopy (Fig. 1G–L). The mRNA expression of the Lcn2 receptor as determined by QRT-PCR is significantly increased in spinal cord tissue after contusion (Fig. 3A). Immunofluorescence labeling showed that after SCI, Lcn2R was expressed in neutrophils (Fig. 3B–D), astrocytes (Fig. 3E–G), and weakly in neurons (Fig. 3H–J).

Lcn2 deletion improves locomotor recovery after SCI

The effects of Lcn2 deficiency on locomotor recovery after SCI were assessed using the BMS (Basso et al., 2006). The BMS analysis is specifically designed to assess open-field locomotor recovery in mice after contusion injury. This analysis shows that *Lcn2*^{−/−} mice had significantly better recovery of locomotor function than did wild-type mice starting at day 5 after injury (Fig. 4A). By 10 dpi, *Lcn2*^{−/−} mice had an average score of 4.05 ± 0.43 , which corresponds to occasional stepping, whereas *Lcn2*^{+/+} mice had an average score of 3.0 ± 0.61 , which corresponds to paw placement, but no stepping. By 28 dpi, *Lcn2*^{−/−} had an average score of 4.9 ± 0.39 , whereas *Lcn2*^{+/+} animals had an average score of 3.7 ± 0.36 . *Lcn2*^{−/−} mice also had significantly better improvement of finer aspects of locomotor control as reflected by higher BMS subscores than *Lcn2*^{+/+} mice. BMS subscores can only be scored in mice that are frequently stepping and attain a score of 5. Accordingly, wild-type mice had a subscore of zero, while the Lcn2-null mice had an average subscore of 1.2 ± 0.29 (SEM) at 28 dpi.

Lcn2^{−/−} mice have increased neuron and tissue sparing after SCI

Given the proposed role for Lcn2 in promoting cell death, we assessed neuronal survival in the ventral horn of *Lcn2*^{−/−} and *Lcn2*^{+/+} mice at 28 dpi. Quantification of neurons in cresyl violet-stained tissue sections showed that *Lcn2*^{−/−} mice have significantly more neurons at 500 and 1000 μm rostral and 1000 μm caudal to the injury site (Fig. 4B–D). Using GFAP staining as a marker to label astrocytes, we examined cross sections of the spinal cord at the epicenter and 200 μm rostral and caudal to the lesion site at 3 and 7 dpi. Our results show that there is significantly increased tissue sparing in *Lcn2*^{−/−} mice as compared with wild-type injured controls at 3 and 7 dpi. The data for the 7 dpi are shown in Figure 5, A and B. Similar results were seen at 3 dpi (data not shown). We also assessed myelin sparing at the lesion

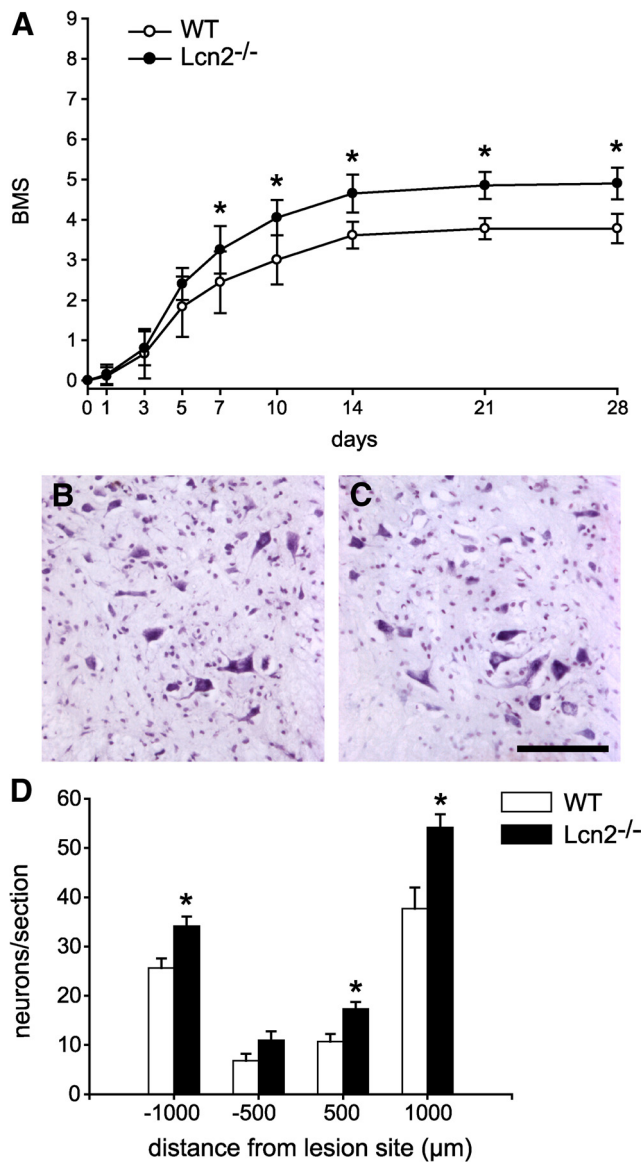


Figure 4. Improved locomotor recovery and neuronal sparing in *Lcn2*^{-/-} mice. **A**, Graph showing the BMS score of injured wild-type (WT) and *Lcn2*^{-/-} after SCI. *Lcn2*^{-/-} mice have significantly improved locomotor recovery from 5 dpi onwards, mean ± SD. *n* = 9 for WT, *n* = 10 for *Lcn2*^{-/-}. **p* < 0.05, two-way repeated-measures (RM)-ANOVA, *post hoc* Tukey. **B**, **C**, Micrographs of cresyl violet–stained sections of the spinal cord 1000 μm caudal to the lesion showing the ventral horn region of wild-type (**B**) and *Lcn2*^{-/-} (**C**) mice. Note the greater number of surviving large neurons in the *Lcn2*^{-/-} mouse. **D**, Quantification of neuronal survival at 500 and 1000 μm distances rostral and caudal to the lesion site, mean ± SEM. **p* < 0.05, two-tailed *t* test, *n* = 6 for each genotype. Scale bar, 100 μm.

site using Luxol fast blue staining of tissue sections. *Lcn2*^{-/-} mice (*n* = 6) showed a significantly higher percentage of myelin sparing (52.9 ± 4%) at the lesion epicenter than *Lcn2*^{+/+} animals (28.4 ± 4%; *n* = 4) (*p* ≤ 0.05).

Lcn2^{-/-} mice have altered cytokine expression and leukocyte infiltration

We next assessed the mRNA expression of MIP1-α, MCP-1, TNF-α, IL-1α, IL-1β, IL-6, and iNOS in the contused spinal cords of *Lcn2*^{+/+} and *Lcn2*^{-/-} mice at 1 dpi. The above chemokines and cytokines are reported to be expressed after SCI and contribute to the pro-inflammatory milieu that is present after SCI (Pineau and Lacroix, 2007). The relative mRNA expression

of all these pro-inflammatory mediators was significantly reduced in *Lcn2*^{-/-} mice (Fig. 6). Flow cytometry was used to quantify the infiltration of immune cells at the lesion site. Our expression data mentioned above indicate that *Lcn2* and its receptor are expressed in cells infiltrating into the spinal cord at 1 and 3 dpi, a time when the majority of immune cells are neutrophils and monocytes (Stirling and Yong, 2008). We therefore assessed whether neutrophil/monocyte infiltration was altered in *Lcn2*^{-/-} mice at 2 dpi. CD45 was used as a marker for peripheral immune cells and microglia to sort cells obtained from injured spinal cords from *Lcn2*^{-/-} and *Lcn2*^{+/+}. Further gating with CD11b made it possible to distinguish between leukocytes (CD45^{high}CD11b⁺) and microglia (CD45^{low}CD11b⁺). The leukocyte population was selected for further analysis and gated on Gr-1, a marker for both monocytes and neutrophils (Stirling et al., 2009). We detected two distinct populations of (CD45^{high}Gr-1⁺) cells, a Gr-1^{high} population that corresponds to neutrophils (Fig. 7C,F, R8), and a Gr-1^{low} population that corresponds to monocytes (Fig. 7C,F, R9). Our results show that, at 2 dpi, there are significantly fewer neutrophils in the injured spinal cord of *Lcn2*^{-/-} mice (49.6 ± 0.97% of the leukocyte population) than in *Lcn2*^{+/+} mice (59.5 ± 0.86% of the leukocyte population). In addition, there is a small but statistically significant increase in the monocyte population in *Lcn2*^{-/-} mice to 30.3 ± 0.82% compared with 25.1 ± 0.51% in *Lcn2*^{+/+} mice. These results suggest that *Lcn2* may play a role in leukocyte infiltration into the injured spinal cord.

Discussion

Lcn2 is a siderophore-binding protein that is thought to be involved in several inflammatory pathologies in non-CNS tissues (Bolignano et al., 2010). We showed that *Lcn2* plays a detrimental role in SCI in mice. *Lcn2* and its receptor are expressed in astrocytes, neurons, and infiltrating neutrophils after SCI. *Lcn2*^{-/-} mice have improved locomotor recovery after SCI and increased neuron and tissue sparing. In addition, our results show that *Lcn2*^{-/-} mice have decreased expression of certain pro-inflammatory mediators and altered neutrophil/monocyte infiltration into the injured spinal cord.

The role of *Lcn2* as part of the innate immune system defense against bacteria and pathogens is becoming better defined (Flo et al., 2004; Saiga et al., 2008; Chan et al., 2009; Nairz et al., 2009). However, its role in other inflammatory diseases and traumatic injuries has not been thoroughly examined. Our results suggest that, in the context of SCI, the observed increased expression of *Lcn2* is likely to be detrimental. The *Lcn2* receptor is expressed on neurons in the injured spinal cord, suggesting that they may be sensitive to *Lcn2*-mediated cell death, as has been shown for other cells (Devireddy et al., 2005). Such a role for *Lcn2* in neuronal loss would be consistent with our observation of increased neuronal sparing in *Lcn2*^{-/-} mice after SCI. Our results also show that *Lcn2* and its receptor are expressed in infiltrating leukocytes. The human form of *Lcn2* was first identified as neutrophil gelatinase-associated lipocalin (NGAL), so called because it covalently binds human gelatinase B (MMP9), stabilizes it, and promotes its proteolytic activity (Triebel et al., 1992). Whether murine *Lcn2* plays a similar role is disputed (Kjeldsen et al., 2000; Tveita et al., 2008; Berger et al., 2010). However, our own zymogram analysis of MMP9 activity after SCI showed no difference between *Lcn2*^{-/-} and *Lcn2*^{+/+} mice (data not shown). Our results indicate that there is a significant reduction in neutrophils in the spinal cord after SCI in *Lcn2*^{-/-} mice, which may contribute to the improved outcome. The role of neutrophils in SCI is debated; several studies

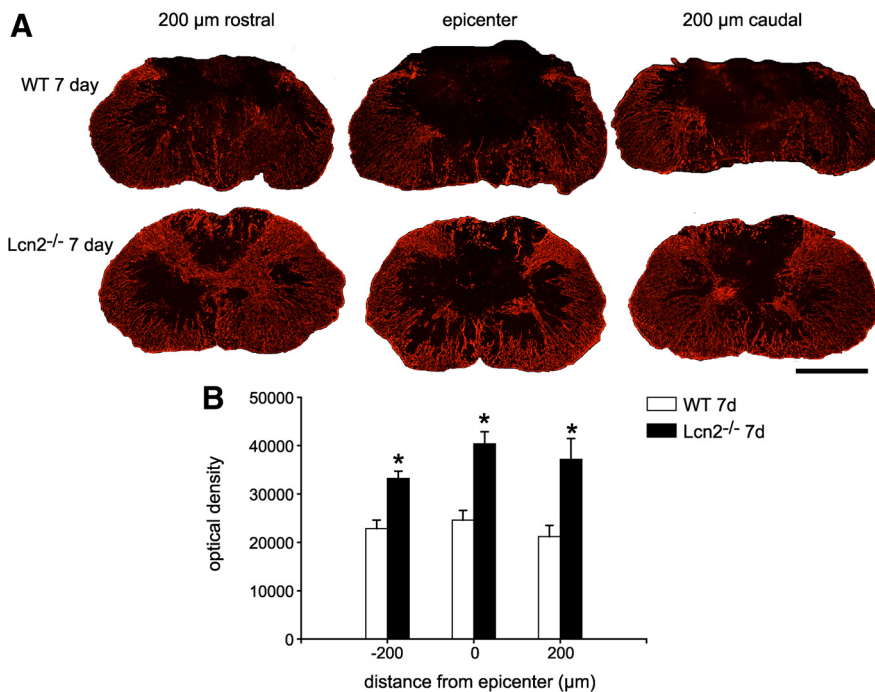


Figure 5. Increased tissue sparing in *Lcn2*^{-/-} mice after SCI. **A**, Immunofluorescence labeling for GFAP in cross sections of the spinal cord of wild-type (WT) and *Lcn2*^{-/-} mice at 7 dpi. Note the greater degree of tissue sparing in the *Lcn2*^{-/-} spinal cords than in the wild-type mice. Representative micrographs from the lesion epicenter and at 200 μ m rostral and caudal are shown. **B**, Quantification of GFAP immunofluorescence staining in entire cross section shows significantly greater tissue sparing in *Lcn2*^{-/-} mice than in wild-type control mice. Data are shown as mean \pm SEM. * p < 0.05, two-tailed t test, n = 3 each genotype. Scale bar, 500 μ m.

have suggested that neutrophils play a detrimental role after SCI (Taoka et al., 1997; Gris et al., 2004), while a recent report suggests that they may be beneficial after SCI (Stirling et al., 2009) through effects on wound healing and repair. Neutrophils have been shown to be largely insensitive to Lcn2-mediated death *in vitro* (Miharada et al., 2005), so the decrease in neutrophil numbers that we observed in the injured spinal cord in *Lcn2*^{-/-} mice may be due to either direct or indirect effects of Lcn2 on neutrophil recruitment. We also detected a small but significant increase in monocyte infiltration into the injured spinal cord in *Lcn2*^{-/-} mice. Whether this reflects an influx of a nonphlogistic (noninflammatory) type of monocytes, as has been reported under conditions in which there is resolution of inflammation (Serhan et al., 2008), is not yet known. A recent report has also suggested that certain monocyte populations can have beneficial effects on the injured spinal cord (Shechter et al., 2009). Consistent with these findings, we also observed a significant reduction in the expression of various pro-inflammatory mediators (MIP-1 α , MCP-1, IL-1 α , IL-1 β , TNF- α , and iNOS) in the injured spinal cord of *Lcn2*^{-/-} mice. Together, these data suggest that Lcn2 may play a role in the initiation of inflammation and leukocyte migration into the injured spinal cord. Whether this is a direct effect of Lcn2 on leukocyte migration or a secondary effect of decreased cytokine synthesis will require further study.

As yet, the mechanism of Lcn2 function is thought to stem from its ability to bind iron. Lcn2 cannot bind iron directly, but binds to siderophores that in turn bind iron. There are several bacterial siderophores, and, more recently, mammalian siderophores have been identified (Bao et al., 2010; Devireddy et al., 2010). The biological effects of Lcn2 on mammalian cells depend on its iron status. Lcn2 loaded with a siderophore/iron complex can deliver iron to cells (Yang et al., 2002; Devireddy et al., 2005), whereas Lcn2 without bound siderophore or iron has been

shown to deplete intracellular iron levels and cause apoptosis (Devireddy et al., 2005). The iron status of Lcn2 produced as a result of inflammation has not been directly assessed. Increased Lcn2 expression is an innate immune defense mechanism aimed at preventing bacteria from acquiring iron, and, in the injured spinal cord, it is likely an innate immune mechanism to prevent infections after injury. In the absence of any bacterial siderophore, however, it is conceivable that Lcn2 would bind its mammalian receptors and exert its iron-binding effects within host cells. One effect of intracellular iron depletion that has already been mentioned is apoptosis; however, intracellular iron levels have also been shown to regulate a number of other processes, including inflammation (Cheepsunthorn et al., 2001; Wang et al., 2008), cell motility (Zhang et al., 2009), and cell division (Yu et al., 2007).

Alternatively, Lcn2 bound to an unknown endogenous iron-binding protein may serve to transport iron to other cells. Whether this occurs *in vivo* and in the CNS is still unknown. Recent reports suggest that Lcn2 has bifunctional effects on microglia and astrocytes *in vitro* (Lee et al., 2007, 2009). The addition of Lcn2, as well as the forced expression of Lcn2 in

astrocyte cultures, caused an increase in GFAP expression and morphological changes consistent with reactive astrogliosis (Lee et al., 2009). On the other hand, Lcn2 sensitized astrocytes *in vitro* to various cytotoxic stimuli, including NO-mediated apoptosis, and promoted cell death. The suppressed expression of Lcn2 by antisense knockdown in astrocytes promoted cell survival (Lee et al., 2009). In the injured spinal cord, we did not detect Lcn2 and Lcn2R by immunofluorescence labeling in microglia, but high levels were seen in astrocytes. We also observed that the cross-sectional area of the spinal cord occupied by GFAP astrocytes was greater in *Lcn2*^{-/-} mice at 3 and 7 dpi than in injured wild-type controls, which may contribute to the increased tissue sparing seen in the knock-out mice. An important protective role for astrocytes in controlling or limiting the inflammatory response at early time points after spinal cord injury has been proposed previously (Okada et al., 2006; Sofroniew, 2009). The contusion injury model that we have used produces significant white and gray matter damage. Our results suggest that the absence of Lcn2 increases tissue sparing as assessed by GFAP staining and neuronal counts. Recombinant Lcn2 did not induce any loss of viability of primary cerebellar neurons *in vitro* assayed using the lactate dehydrogenase assay (data not shown), suggesting that the loss of neurons after SCI may be secondary to the inflammatory response. White matter sparing, which is associated with improved recovery from this kind of injury (Magnuson et al., 2005), is also increased in *Lcn2*^{-/-} mice, and these mice show greater sparing of myelin after SCI. Our immunostaining results, however, showed little or no expression of Lcn2 and its receptor in oligodendrocytes, suggesting that the increased myelin sparing we observed is likely an indirect effect of Lcn2 deficiency.

Increasing evidence suggests that Lcn2 plays an important role in the innate immune response, primarily as a defense against

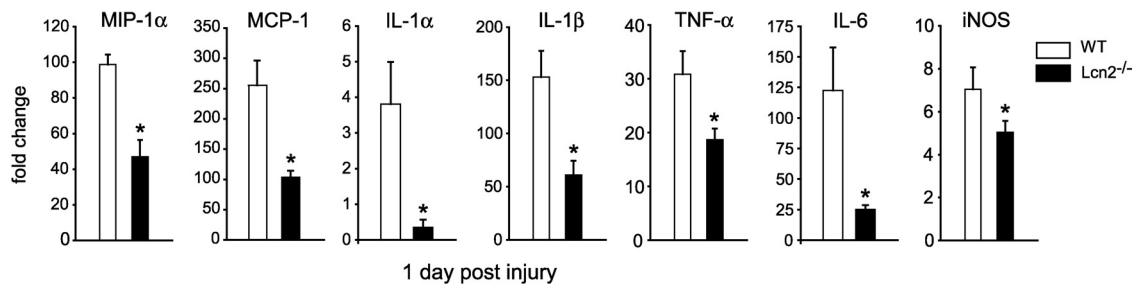


Figure 6. Reduced expression of pro-inflammatory mediators in *Lcn2*^{-/-} mice. The bar graphs show the relative mRNA expression of several pro-inflammatory mediators (MCP-1α, MCP-1, IL-1α, IL-1β, TNF-α, IL-6, and iNOS) in the spinal cord at 1 dpi as determined by QRT-PCR. The expression of all of these mediators is significantly reduced in *Lcn2*^{-/-} mice. Data are shown as mean ± SEM. **p* < 0.05, two-tailed *t* test, *n* = 4 each genotype.

bacterial infection. However, our results suggest that, in the setting of aseptic SCI, *Lcn2* may also have significant detrimental effects. We provide evidence that *Lcn2* and its receptor are expressed after SCI, and that *Lcn2* plays a detrimental role in SCI, contributing to neuronal and astrocyte loss, pro-inflammatory cytokine expression, and immune cell infiltration. Our results have implications for other inflammatory CNS conditions that are characterized by high levels of *Lcn2* expression.

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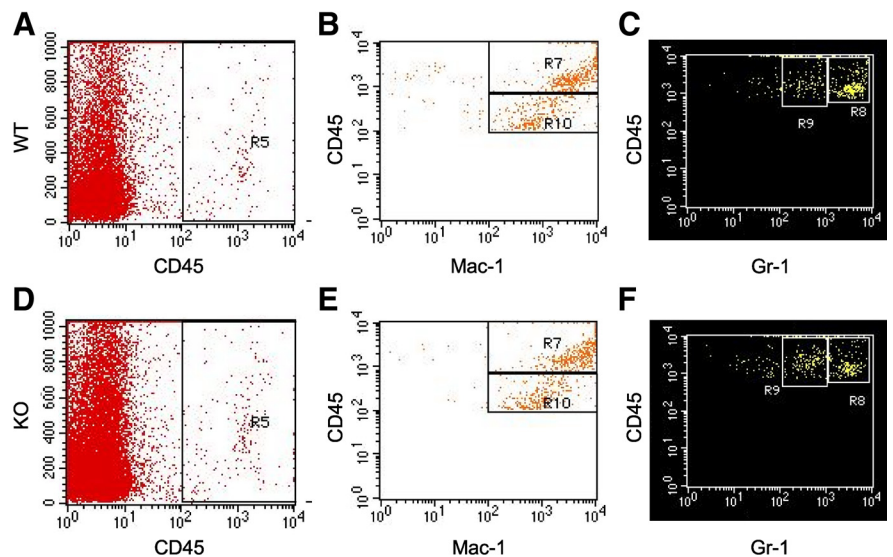


Figure 7. Altered numbers of neutrophils and monocytes in the injured spinal cord in *Lcn2*^{-/-} mice. Representative density plots of fluorescence-activated cell sorting analysis of cells isolated from injured spinal cord at 2 dpi. All CD45⁺ cells (**A**, **D**, **R5**) were further gated on Mac-1 labeling (**B**, **E**). In **B** and **E**, two distinct populations of Mac1⁺ cells can be observed, i.e., the CD45^{high} (box labeled **R7**) and the CD45^{low} (box labeled **R10**) populations. The CD45^{high} population is a mixed population of blood-derived immune cells, whereas the CD45^{low} population comprises resident microglia from the spinal cord. The blood-derived leukocyte population (**R7**) was further gated on Gr-1 labeling (**C**, **F**). Two Gr-1⁺ populations emerge, Gr-1^{high} (box labeled **R8**) and Gr-1^{low} (box labeled **R9**). The Gr-1^{high} population is composed of neutrophils, whereas the Gr-1^{low} population is made up of monocytes. There is a significantly smaller Gr-1^{high} population in *Lcn2*^{-/-} injured spinal cords and a significantly larger Gr-1^{low} population in *Lcn2*^{-/-} injured spinal cord than in wild-type controls [*p* < 0.05, 2-tailed *t* test, *n* = 6 for wild-type (WT), *n* = 5 for knock-out (KO)]. Gr-1^{high} in WT = 59.5 ± 0.86%, in KO = 49.6 ± 0.97%. Gr-1^{low} in WT = 25.1 ± 0.51%, in KO = 30.3 ± 0.82% (mean ± SEM).

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