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# Kallikrein Cascades in Traumatic Spinal Cord Injury: *In Vitro* Evidence for Roles in Axonopathy and Neuron Degeneration

Maja Radulovic, B.S.<sup>1</sup>, Hyesook Yoon, Ph.D.<sup>2</sup>, Nadya Larson, Ph.D.<sup>2</sup>, Jianmin Wu, Ph.D.<sup>2</sup>, Rachel Linbo, D.PT.<sup>2</sup>, Joshua E. Burda, Ph.D.<sup>1</sup>, Eleftherios P. Diamandis, M.D.<sup>3</sup>, Sachiko I. Blaber, B.S.<sup>4</sup>, Michael Blaber, Ph.D.<sup>4</sup>, Michael G. Fehlings, M.D., Ph.D.<sup>5</sup>, and Isobel A. Scarisbrick, Ph.D.<sup>1,2,6,\*</sup>

<sup>1</sup>Neurobiology of Disease Program, Mayo Medical and Graduate School, Rochester MN 55905

<sup>2</sup>Department of Physical Medicine and Rehabilitation, Mayo Medical and Graduate School, Rochester MN 55905

<sup>3</sup>Department of Pathology and Laboratory Medicine, Mount Sinai Hospital, Toronto, ON, Canada M5G 1X5

<sup>4</sup>Department of Biomedical Sciences, Florida State University College of Medicine, Tallahassee, FL 32306

<sup>5</sup>Department of Surgery, Toronto Western Research Institute, Toronto, ON, Canada M5T 2S8

<sup>6</sup>Department of Neurology, Mayo Medical and Graduate School, Rochester MN 55905

# **Abstract**

Kallikreins (KLKs) are a family of 15 secreted serine proteases with emerging roles in neurological disease. To illuminate their contributions to the pathophysiology of spinal cord injury (SCI) we evaluated acute through chronic changes in the immunohistochemical appearance of six kallikreins, KLK1, KLK5, KLK6, KLK7, KLK8 and KLK9 in post-mortem human traumatic SCI cases, quantified their RNA expression levels in experimental murine SCI, and assessed the impact of recombinant forms of each enzyme toward murine cortical neurons *in vitro*. Temporally and spatially distinct changes in kallikrein expression were observed with partially overlapping patterns between human and murine SCI, including peak elevations (or reductions) during the acute and subacute periods. KLK9 showed the most robust changes and remained elevated chronically. Importantly, a subset of kallikreins, KLK1, KLK5, KLK6, KLK7 and KLK9 were shown to be neurotoxic toward primary neurons *in vitro*. Kallikrein immunoreactivity was also observed in association with swollen axons and retraction bulbs in the human SCI materials examined. Together, these findings demonstrate that elevated levels of a significant subset of

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<sup>\*</sup>Correspondence to: Isobel A. Scarisbrick Ph.D., Neurobiology of Disease Program, 642B Guggenheim Building, Mayo Clinic Rochester, 200 First St., SW., Rochester, MN 55905, Tel: 507-284-0124, Fax: 507-266-4716, Scarisbrick.Isobel@mayo.edu.

kallikreins are positioned to contribute to neurodegenerative changes in cases of CNS trauma and disease and therefore represent new targets for the development of neuroprotective strategies.

#### Keywords

axonal injury; degeneration; inflammation; neural injury; spinal cord injury

# INTRODUCTION

Kallikreins are emerging as important mediators of disease, including neurological conditions such as Alzheimers (1–5), frontotemporal dementia (6), Parkinsons (3, 7), multiple sclerosis (MS) (8, 9) and post-polio syndrome (10). Alterations in proteolytic cascades are also an important component of the response of the CNS to trauma, including SCI in which severely damaging pathophysiological events occur secondary to the primary physical injury (11). These events include a complex cascade of vascular, cellular and chemical responses such as edema, ischemia, inflammation, ionic imbalance, free radical formation, astrogliosis, demyelination, excitotoxicity and apoptosis (12–15). We and others have become particularly interested in the role of extracellular proteolysis in secondary injury since secreted proteases offer great potential for therapeutic targeting.

The inventory of proteases with likely involvement in SCI is long and growing and includes multiple members of the matrix metalloprotease family (16), cysteine proteases such as calpain (17) and serine proteases including thrombolytic enzymes (18, 19), plasminogen activators (20, 21) and tissue kallikreins (22, 23). The current effort focuses on defining the involvement of tissue kallikreins in traumatic SCI, with an emphasis on those which are expressed at significant levels in the CNS (24–26) or which have been linked to pathology in other neurological conditions (6, 8, 27–30). The classical kallikreins have been recognized for some time and include true "tissue kallikrein", KLK1, which cleaves Lys-bradykinin to liberate kinins that have been implicated in inflammation and tumor cell invasion as well as KLK2 and KLK3 (prostate specific antigen) which are well recognized serum biomarkers of prostate cancer. Over the past decade, 12 additional kallikrein encoding genes were identified alongside these on human chromosome 19q13.4 and together these form the largest contiguous cluster of serine proteases in the human genome (31).

The urgent need to further understand the roles kallikreins play in nervous system injury and disease is highlighted by the growing list of neurological disorders in which kallikrein expression is dynamically regulated (32). For example, elevations in KLK1 occur in stroke (33) and hippocampal sclerosis (34). KLK8 is elevated in the hippocampus in Alzheimers disease (35), while KLK6 is reduced in Alzheimers plaques (1–4), CSF (36) and serum (5). KLK6 and KLK7 are also reduced in the CSF of patients with frontotemporal dementia, while KLK10 is elevated (6). KLK1 and KLK6 are each elevated in the serum of patients with progressive MS, a disease stage characterized by progressive neural injury (9). KLK6 is also localized to Lewy bodies in Parkinsons (7) and is elevated in the CSF of patients with post-polio syndrome (10). Less is known regarding the activity of kallikreins in CNS trauma, although elevations in KLK6 have been reported in human and rodent SCI (22–24)

and this protease promotes neurite retraction, neuron degeneration, oligotoxicity and astrogliosis *in vitro* (8, 9, 28, 30). KLK8 is elevated in a crush model of murine SCI and notably KLK8 deficient mice show reductions in SCI-related axon injury and motor deficits (37).

Despite a growing recognition of the potential involvement of kallikreins in neurological disease, little is known regarding their pathophysiological actions. To begin to address this gap we made a parallel examination of the appearance of six kallikreins in human traumatic SCI and in a murine SCI model and determined the impact of the recombinant forms of each enzyme on neurite stability and neuron survival *in vitro*. The findings presented point to the involvement of multiple kallikreins in the response of the human spinal cord to injury at acute and chronic time points, with a subset likely to participate directly in neuron injury, including neuron and neurite loss. The results of these studies therefore point to kallikreins as targets for the development of new neuroprotective strategies relevant to traumatic SCI and other neurological disorders.

#### MATERIALS AND METHODS

# **Human Spinal Cord Injury Material**

To determine the scope of action of kallikreins in human traumatic SCI we made a comprehensive analysis of the immunohistochemical appearance of six kallikreins in noninjured post-mortem human spinal cord material (n=3) and in cases of traumatic SCI ranging from 0 to 7704d post-injury (n=22). Specifically, the SCI cases included were classified as immediate acute (0 to 1d, n=3), acute (>1d to < 14d, median 4d (Range 2 to 8d), n=7), subacute (y 14d to 63d, median 30d (Range 14 to 63d), n=6) and chronic (>1 yr, median 2133d (Range 413 to 7704d), n=6) time points post-traumatic injury (Table 1). These injury intervals correspond to the immediate acute (1d), acute (3, 7), subacute (14, 21, 30d) and what could be considered more chronic time points (60d) examined in murine experimental SCI. A total of 22 post-mortem cases of traumatic SCI were included in these studies with specimens obtained from the Mayo Clinic archives. All cases were confirmed as traumatic involving contusion, compression or transection of the spinal cord by examination of patient records, including autopsy reports. Cases where injury was a result of vascular events or myelopathy of non-traumatic origin were excluded. In addition, the extent of injury in each case was evaluated in hematoxylin and eosin (H&E), in addition to Luxol fast blue/periodic acid Schiff's (LFB/PAS) stained sections, and only cases in which injury could be clearly ascertained, that is disruption of the spinal cord parenchyma, hemorrhage, inflammatory infiltrates, demyelination and/or cyst-like areas were included in subsequent studies to evaluate kallikrein expression levels. 77% of the SCI cases were male and the median age was 28.2 yr (Range 3.9 to 74 yr). 72.7% of cases were cervical, involving one or more segments C1 to C5 and the remaining cases were thoracic ranging from T1 to T7. 63% of cases were determined to be a result of motor vehicle accident, 14% falls and 14% sportsrelated injury (diving or go-cart), 4.5% gunshot wound and 4.5% due to unspecified trauma. Three cases with no history of CNS trauma were examined as uninjured-controls (median age 41, range 37-41; 67% male) and included death due to acute leukemia, liver disease or melanoma. The use of human materials in these studies was approved by the Mayo Clinic

Institutional Review Board and all clinical follow-up was obtained by review of medical records.

#### Kallikrein Immunohistochemistry in Human Spinal Cord Injury

Human kallikreins were examined in 5 µm sections obtained from paraffin embedded SCI autopsy material using immunohistochemical techniques as detailed in our prior studies (8, 22, 38, 39). Endogenous peroxidase activity was quenched in de-paraffinized sections by treatment with 0.3% H<sub>2</sub>O<sub>2</sub> in methanol and sections rehydrated through graded ethyl alcohols. Sections were then subject to steam heat-induced antigen retrieval using Antigen Retrieval Citra (BioGenex, San Ramon, CA, USA). Primary antibodies specific for each human kallikrein were applied to tissue sections overnight at 4 °C. KLK1 was detected using a mouse monoclonal antibody M01-H00003816 (Novus Biologicals, Inc. Littleton, CO, USA). KLK5 was detected with an affinity purified rabbit polyclonal antibody PA1-4238 (Thermo Fisher Scientific Inc., Rockford, IL). KLK6 was detected using either a monoclonal antibody (MSP-3-3) or a rabbit polyclonal antibody (KLK6-Rb008) each of which produced identical staining patterns as previously described (22, 28). KLK7 (6, 40) and KLK8 (41) were detected using rabbit polyclonal antibodies generated and validated as we have previously described (26). KLK9 was detected using a rabbit polyclonal antibody PAB-10236 (Orbigen, San Diego, CA, USA). Following incubation with primary antibodies, sections were washed in PBS and species appropriate AffiniPure F(ab')<sub>2</sub> fragment-specific biotin-conjugated secondary antibodies with minimal cross-reactivity to human serum proteins (Jackson ImmunoResearch, West Grove, PA) were applied for 1 hr at RT. Antibody binding was then visualized using HRP-conjugated streptavidin with 3',3'diaminobenzadine tetrahydrochloride as the substrate. Processing of tissue sections with primary antibody replaced with normal serum was used to control for non-specific binding of secondary antibodies (42). Parallel sections were stained with H&E to identify areas of tissue injury, LFB/PAS to determine areas of demyelination, Bielschowsky's silver stain to examine axon integrity and Masson's Trichrome to visualize collagen. The appearance of GFAP-immunoreactive (-IR) astrocytes (rabbit polyclonal, Z0334, Dako Carpenteria, CA) and CD68-IR macrophages of monocyte or microglial origin (43) (mouse monoclonal, PG-M1, Serotec, Raleigh, NC) were also evaluated in sections parallel to those used to examine kallikrein immunoreactivity.

#### Semi-Quantitative Assessment of Kallikrein Immunoreactivity in Human Spinal Cord Injury

The immunohistochemical appearance of kallikreins across the post-injury survival intervals examined was evaluated using a well validated semi-quantitative scale that considers both the intensity and relative abundance of immunoreactivity (44, 45). Kallikrein immunoreactivity (KLK-IR) was graded separately in the spinal cord white matter or in the gray matter without knowledge of the kallikrein being considered or the post-injury interval. In a subset of cases, only spinal cord white matter was available for assessment (see Table 1). Our analysis of KLK-IR was focused on the spinal level of injury indicated on the pathology report with tissue injury confirmed in parallel H&E stained sections. Regions of spinal cord pathology were further verified in parallel sections stained for LFB/PAS, Mason's Trichrome, GFAP or CD68. In some cases a second adjacent block with equivalent injury was also stained and scored as indicated in Table 1. In each case, the region of spinal

cord examined and KLK-IR scores were arrived at by consensus by two different observers (MR and IAS) using a multi-headed microscope. Intensity of immunoreactivity in each case was graded on a scale of 0 to 3 (no staining, weak, moderate staining, strong staining) and the percent area stained graded on a scale of 1 to 4, in 25% increments. A grade of 4 was only given if both cells and the extracellular matrix were immunoreactive. An overall immunoreactivity score was tabulated in each case as the product of intensity and percent area stained, with a maximal score of 12 as we and others have previously described (45–47). The mean and standard error of KLK-IR scores at the different post-injury intervals examined was then determined and compared to non-traumatic "uninjured" control cases using a Student's t-test or the Rank Sum test when data were not normally distributed (47). P < 0.05 was considered statistically significant.

# Murine Clip Compression Model of Traumatic Spinal Cord Injury

The potential significance of each kallikrein to traumatic SCI was further evaluated by quantitative real-time RT-PCR analysis of the expression of the murine ortholog in each case in a murine contusion-compression SCI model. Experimental SCI was generated in twelve-week old female (19-23 g) C57BL/6J mice (Jackson Laboratories) using a modified aneurysm clip (FEJOTA<sup>TM</sup> mouse clip, 8g closing force generating "moderate injury") (48). This model of SCI includes not only an initial contusion, but also a persistent compression phase that results in microcystic cavitation, degenerating axons, and robust astrogliosis, all hallmarks of human traumatic SCI (49, 50). Prior to compression, mice were deeply anesthetized with Rompun and Ketaset (10 mg/kg) and a laminectomy performed at T9 to T11. Injury was induced at the level of T10 by extradural application of the clip for a period of exactly 1 min, which produces mechanical contusion to both dorsal and ventral aspects of the spinal cord. Before and after surgery, saline was administered subcutaneously to replace lost blood volume. Pain was minimized by administration of Buprenorphine (0.05 mg/kg, Hospira, Lake Forest, IL) subcutaneously every 12 hr for 96 hr post-surgery. To avoid infection, Baytril (10mg/kg, Bayer Health Care, Shawnee Mission, KS) was administered intraperitoneally in a prophylactic fashion for the first 72 hr post-surgery. Bladders were manually voided twice daily even after spontaneous recovery of function until the endpoint of each experiment. Any mice with signs of infection or which were moribund were immediately euthanized and not included in the study. All animal experiments were carried out with careful attention to animal comfort and in strict adherence to NIH Guidelines for animal care and safety and were approved by the Mayo Clinic Institutional Animal Care and Use Committee.

To determine immediate acute (1d), acute (3, 7d), subacute (14, 21, 30d), and more chronic (60d) compression related changes in kallikrein RNA expression, mice were allowed to recover to each of experimental endpoint (n=3 per time point, except 14d n=4). Mice receiving surgery on a given day were randomized across two or more endpoints. In each case, the level of injury, as well as one segment above and below, was collected into a single tube and snap frozen prior to RNA isolation. Mice were evaluated across seven categories of locomotor recovery using the Basso Mouse Scale the day after surgery and weekly thereafter until experimental endpoints were reached (51). Age and gender matched uninjured mice served as controls. Due to the focus of these studies on a limited amount of spinal cord

tissue at the injury epicenter, samples from at least 3 mice at a given time point were pooled to facilitate RNA isolation (52). RNA was isolated using RNA STAT-60 (Tel-Test, Friendswood, TX) and stored at -80 °C until the time of analysis.

# Quantitative Assessment of Murine Kallikrein Expression in Response to Compression Injury

Expression levels of the murine orthologs of human kallikreins 1, 5, 6, 7, 8 and 9 were determined by real-time RT-PCR using primers specific to each kallikrein and designed to span at least two exons (Table 2). According to the new nomenclature, non-human forms of each kallikrein are designated in lower case (Klk) (53). Accuracy of amplification was verified by analysis of melting peaks. An estimate of the overall abundance of each kallikrein RNA was made based on a standard curve created by amplification of plasmids containing the gene of interest serially diluted to known copy number in the same PCR experiment. At least 3 individual SCI epicenters at each time point were pooled (52) and 0.25 µg examined in triplicate by real time RT-PCR, with control samples and all 7 time points post-injury examined in the same PCR run. A no RNA control was included in every PCR experiment to exclude reagent contamination. The copy number of each kallikrein determined in control and injured spinal cord was normalized to the constitutively expressed gene glyceraldehyde phosphate dehydrogenase (GAPDH) amplified in parallel and the data were expressed as a percent of uninjured control (23, 25, 30, 54, 55).

#### **Cortical Neuron Culture**

To determine the potential contribution of KLK1, KLK5, KLK6, KLK7, KLK8 and KLK9 to neural injury and their likely impact on the microenvironment available for neural repair we examined the effects of the recombinant forms of each enzyme on neurite stability and neuron survival *in vitro*. In addition to these kallikreins already linked to the CNS (25), we also examined 3 kallikreins (KLK2, KLK3 and KLK13) not densely expressed in the CNS, but which extravasate across a damaged blood brain barrier. Murine cortical neurons were isolated from embryonic day 16 C57BL6/J mice and plated at a density of  $2 \times 10^3$  cells/mm² on poly-L-lysine (10 µg/ml) coated 12 mm² glass cover slips. Cortical neurons were grown in defined media consisting of Neurobasal A media supplemented with 2% B27 supplement, 2mM Glutamax, 0.45% glucose and 50 U/ml penicillin-streptomycin (Invitrogen, Carlsbad, CA) (9). Neurons were allowed to mature for 96 hr *in vitro* and then treated with recombinant human kallikreins (10 µg/ml, 300 nM) for an additional 24 hr prior to fixation with 2% paraformalydehyde. Neurons were stained for neurofilament H protein (rabbit polyclonal, AHP245 Serotec) using standard avidin-biotin histochemistry as described above.

All human kallikreins were expressed using a baculovirus expression system and subsequently purified and activated as previously described in detail (30, 54, 56–60). 300 nM of KLK1 or KLK6 were previously shown to promote a dying back of cortical neurites in prior studies (9) and therefore, here we examined KLKs 2, 3, 5, 7, 8, 9 and 13 at parallel concentrations. This concentration is designed to be reflective of the elevated levels of kallikreins seen in SCI materials (25, 61) and is approximately 5-fold higher than the level of KLK6 detected in human CSF (26, 62).

Changes in the number of cortical neurons or neurite density in response to recombinant kallikrein were quantified from 5 digitally captured  $20\times$  microscopic fields encompassing the center and 4 poles of each cover slip, taken without knowledge of the treatment groups. Digital images were then used to make neuron counts. Neurite number was evaluated by superimposing a grid consisting of  $967.5~\mu\text{m}^2$  squares on to each digital image and counting processes which overlapped grid lines (8). Histograms shown represent the mean and standard error (s.e.) of counts across three independent cover slips treated in parallel within a given experiment and results were reproducible across at least 3 experiments utilizing separate murine cortical neuron preparations. Statistical differences between groups were compared using One Way ANOVA with the Student Newman Keul's (SNK) *post hoc* test, with P < 0.05 considered statistically significant.

#### **RESULTS**

### Regulated Expression of Kallikreins in Human Traumatic Spinal Cord Injury

KLK Expression in the Uninjured Human Spinal Cord—In the uninjured adult human spinal cord, overall levels of kallikrein-IR were highest in the case of KLK6, followed by KLK5, KLK7, KLK9, KLK1 and KLK8 (Figs. 1, 2). Consistent with previous reports, in white matter KLK6-IR was dense in glia with the cytological appearance of oligodendroglia that is round cell bodies, a paucity of cytoplasm and processes encircling axons (Fig. 2B) (38, 39). KLK1 could also be identified in glia with a round nucleus characteristic of oligodendroglia, but was more prominent in those with a triangular shape and tapering processes resembling astrocytes in parallel GFAP-immunostained sections. KLK5, KLK7 and KLK9 were also prominent in astroglial-like cells. KLK8-IR was low when observed in white matter glia of the uninjured spinal cord.

At least a subset of kallikreins is known to be expressed by neurons (24, 39, 63). Here we document expression of KLK6, KLK5, KLK7, KLK1, KLK9, and KLK8, listed in the relative order of abundance, in neurons across the dorsal and ventral horns of the uninjured adult spinal cord, with prominent expression in ventral horn motoneurons as seen in Fig. 2C. KLK5, KLK7, KLK9 and at least low levels of KLK8 were also readily observed in association with axons in uninjured spinal cord white matter (Fig. 2B).

## KLK Expression in Human Spinal Cord White Matter after Traumatic Injury—

Overall, in the human traumatic SCI cases examined there was a trend towards elevations in kallikrein expression in spinal cord white matter, although these elevations only reached the level of statistical significance in the case of KLK7 and KLK9 (Figs. 1, 3A, 4A, 5A, 6A, B, 7A). In the first 24 hr after injury (0–1d, immediate acute), KLK7- and KLK9-IR showed minor elevations in white matter, otherwise either no change, or minor decreases in kallikrein expression were observed (Fig. 1, Fig. 3). In the acutely (>1d <14d) injured spinal cord white matter, elevated levels of immunoreactivity were detected for each KLK, with elevations in KLK7-IR reaching the level of statistical significance (Student's t-test, P=0.04, Figs. 1, 4). In the subacute ( 14d to 63d) cases examined, overall levels of immunoreactivity for KLKs 1, 5, 7 and 9 remained elevated relative to the uninjured spinal cord, although the overall changes observed did not reach the level of statistical significance (Figs. 1, 5, 6).

Swollen axons and retraction bulbs (64, 65) were immunoreactive for each KLK in the later subacute cases examined, including KLK1 and KLK6 which were not apparent in association with axons of the intact spinal cord (Figs. 2B, 6B). The distribution of axons was verified in parallel sections stained by Bielschowsky's silver stain. At chronic post-injury intervals, only KLK9-IR remained significantly elevated in spinal cord white matter (Student's t-test, P=0.02) while the other kallikreins examined were unchanged or slightly lower than that observed in the uninjured spinal cord (Figs. 1, 7). When primary antibodies were replaced with normal serum, immunoreactivity scores for either uninjured or traumatically injured spinal cord specimens were 0, reflecting the absence of staining (Fig. 5C).

KLK Expression in Human Spinal Cord Gray Matter after Traumatic Injury—In the gray matter of the injured spinal cord, there was an overall decrease in immunoreactivity for KLK1 and KLK5 over the time course examined with levels remaining below baseline in the most chronic cases examined (Figs. 1, 3B, 4B, 5B, 6C, 7B). Decreases in KLK1-IR in injured spinal cord gray matter were statistically significant at subacute time points (Student's t-test, P=0.04), while decreases in KLK5-IR were statistically significant at both acute (Rank Sum test P=0.03) and subacute (Student's t-test, P=0.02) time points. KLK9-IR was significantly reduced in the injured spinal cord gray matter at acute time points (Rank Sum test, P=0.04), but elevated by more than two-fold at chronic time points post-injury (Student's t-test, P=0.01, Figs. 1, 7B).

KLK-Immunoreactivity in Astrocyte-Like Cells in Human SCI—Spinal cord trauma-related changes in KLK-IR observed from acute through chronic time points could be related to co-ordinate changes in the cellular hallmarks of neural injury, including the appearance of GFAP-IR astrocytes. In the uninjured cases examined, moderate levels of GFAP-IR were observed in white matter, while levels of GFAP were very low or absent in gray matter (Fig. 2). As expected (66), increases in GFAP-IR cells and processes were observed from the acute through the most chronic time points examined (Figs. 4 to 7). In the cases examined in this study, increases in GFAP-IR predominated in spinal cord white matter, although clear increases in gray matter were observed in some cases. Each KLK was elevated in astrocyte-like cells over that detected at baseline, with KLK5 being more dense than KLK7 followed by KLK9, KLK6, KLK1 and KLK8 (Figs. 2 to 7). KLK5-, KLK7- and KLK9-IR appeared in many cases to be co-extensive with that for GFAP-IR, particularly in white matter at chronic stages after injury (Figs. 6A, 7A). The most prominent staining for KLKs in astrocyte-like cells occurred in the later subacute and chronic SCI cases examined (> 63d to 7704d, Figs. 6A, 7A).

**KLK-Immunoreactivity in Macrophage-Like Cells in Human SCI**—Macrophages, derived from hematogenous monocytes and resident microglia are reactive for the CD68 antigen and play important roles in the response of the spinal cord to injury (43, 67–69), including involvement in cytotoxicity as well as the wound healing process. In the uninjured spinal cord and within the first 24 hr after injury (immediate acute), CD68-IR cells and processes were largely absent or extremely sparse (Figs. 2, 3) (69). Increases in CD68-IR associated with clearly defined cell bodies was seen in the acute cases examined (1d to 14d,

Fig. 4), but was most prominent at subacute ( 14d) time points when staining included larger rounded macrophages as well as smaller densely CD68-IR cells with short processes, likely ramified microglia (Figs. 5 and 6). Monocytes/microglia immunoreactive for CD68 persisted in many cases out to the late chronic time points examined (Fig. 7). At subacute time points (Fig. 5), KLKs 6, 5 and 7 were prominent in rounded-macrophage-like cells prominent in the injured white matter, while KLK1 and KLK9 were seen at lower levels (Figs. 5).

#### Regulated Expression of Kallikreins in Murine Experimental SCI

The murine orthologs of each human kallikrein were examined in an experimental murine SCI model using quantitative real time RT-PCR. We note that antibodies specific to each of the murine kallikreins studied herein are not currently available. As expected based on our prior studies regarding the expression of Klk6 in rodent brain and spinal cord (22, 24) and kallikrein RNA and protein expression patterns documented in the human spinal cord (Fig. 1) (25, 26), of all the kallikreins examined Klk6 was by far the most abundant (Fig. 8). In the uninjured murine spinal cord, Klk6 RNA was detected at  $4.2 \times 10^7 \pm 2.7 \times 10^6$  copies in 0.25 µg of RNA. Klk1 was the next most abundant of those examined, being detected at 5.2  $\times 10^3 \pm 1.6 \times 10^2$  copies in a parallel amount of RNA. Significantly lower levels of each of the other kallikreins examined were observed in the uninjured spinal cord, each at approximately  $10^2$  copies in 0.25 µg of RNA with Klk9 being more abundant than Klk5, followed by Klk7 and Klk8.

The expression of Klk1, Klk5, Klk6, Klk7, Klk8 and Klk9 RNA were each dynamically regulated in the murine spinal cord in response to experimental compression injury over the 60d post-injury interval examined (Fig. 8). Transcriptional alterations in each kallikrein showed clear differences in direction, timing and magnitude, although examination of RNA from whole spinal cord preparations does not permit us to comment on any changes specific to gray or white mater. Overall, increased levels of expression were the predominant pattern observed. Increases in RNA expression were observed in the case of Klk1, KLlk6, Klk7, Klk8 and Klk9 (P < 0.05 SNK) and in each case this reached the highest level by 7d after injury. The magnitude of the change in RNA expression from baseline was most robust in the case of Klk8 and Klk9, each of which reached from 10- to more than  $1 \times 10^3$ -fold base line levels during the first 7d after injury (P 0.001, SNK). Klk8 expression remained significantly elevated through day 21 after injury, but only Klk9 continued to be elevated at the 30 and 60d endpoints examined (P 0.001, SNK). By contrast, Klk5 RNA expression showed significant decreases at 1 and 3d after injury (P < 0.002 SNK). After transient increases at 1 and 7d after injury, Klk7 also showed significant decreases in RNA expression at 21 through the 60d endpoint examined (P < 0.05 SNK).

To facilitate interpretation of the changes in kallikrein expression in terms of other experimental SCI models, we evaluated Basso Mouse Scale scores at 1d after injury and weekly thereafter. Functional deficits reflected in BMS scores induced by the 8 g Fejota clip contusion compression model were largely parallel to those seen in BL6 mice in prior studies in which a moderate injury was induced by the Infinite Horizon or Ohio State

University impactor model (51). BMS scores (mean  $\pm$  s.e.) at each endpoint examined were: 1d 1  $\pm$  0.5; 3d 3  $\pm$  0.3; 7d 2  $\pm$  0.7; 14d 2.9  $\pm$  1; 21d 3.7  $\pm$  0.4; 30d 4  $\pm$  1.1; and 60d 5  $\pm$  1.

To put the changes in kallikrein RNA expression into the context of well characterized pathogenic events associated with SCI, levels of GFAP, MBP, PLP and the proinflammatory cytokines (IL-6, TNFα and IL-1β) were examined in the same RNA preparations (Fig. 8). Alterations in kallikrein expression were paralleled by significant elevations in IL-6, which showed a  $12 \times 10^3$ -fold increase at 1d and remained  $4 \times 10^3$ -fold elevated through 14d post injury (P < 0.001, SNK, Fig. 7). Another pro-inflammatory cytokine, TNF $\alpha$ , was also significantly elevated by 1d (4.5 × 10<sup>3</sup>-fold elevated), peaked at day 14 (12.8  $\times$  10<sup>3</sup>-fold elevated) and remained elevated out to the most chronic 60d endpoint examined (5.6  $\times$  10<sup>3</sup>-fold elevated, P 0.001, SNK). Peak elevations in IL-1 $\beta$  were also observed at 1d 49-fold elevated P < 0.001), showed a second peak in expression at 14d 20-fold elevated, P < 0.001) and remained significantly elevated at the 60d time point 10fold, P = 0.01). As expected, the murine contusion-compression model examined also triggered robust and sustained elevations in GFAP RNA expression, which were observed as early as 1d (2-fold) and reached a peak of 4-fold baseline levels at 21d (P 0.013, SNK). Significant elevations in GFAP RNA expression persisted through 60d (P 0.001, SNK). Across the same time points (1-60d) there was a significant decrease in PLP RNA expression, which remained at approximately 50% below baseline levels from the acute through chronic time points examined (P 0.001, SNK). In parallel, MBP RNA levels were also reduced at 3 and 7d post-injury (P 0.04, SNK), but returned to uninjured levels on days 14 through 30. By 60d however, MBP RNA levels were again significantly reduced reaching approximately 50% of that observed at baseline (P 0.001, SNK).

#### Kallikrein-Elicited Neural Injury In Vitro

Given the abundance of several kallikreins across the spinal cord white and gray matter, differential regulation in cases of traumatic SCI (Figs. 1–8), and potential to be involved in kallikrein-specific activation cascades (57, 59) we examined the effects of elevated levels of the recombinant forms of each enzyme, in addition to KLKs 2, 3 and 13 on neuron number and neurite integrity of mature murine cortical neurons in vitro (Fig. 9). KLKs 1, 5, 6, 7 and 9 each promoted significant loss of neurons after 24 hr treatment in vitro (P 0.03, SNK) and these kallikreins, in addition to KLK2, significantly reduced neurite numbers (P 0.001, SNK, Fig. 8). By contrast, KLK3, KLK8 and KLK13 did not promote significant neural injury-related changes in vitro. Indeed, there was significant increase in neuron process density in KLK3 treated cultures (P = 0.04, SNK). KLK-induced neural injury was most dramatic in the case of KLK7 and KLK9, each of which reduced neuron survival by 60% or more (P < 0.001, SNK). KLK1, KLK5 and KLK6 reduced neuron survival by approximately 40% relative to control cultures (P 0.03, SNK). Each of these kallikreins, in addition to KLK2, also significantly reduced neurite density by 40% or more (P < 0.001, SNK), with KLK7 and KLK9 exerting the most vigorous effects. KLK8 and KLK13 did not alter neuron survival or neurite density at the concentrations and time points examined.

# **DISCUSSION**

These studies document regulated patterns of expression of multiple members of the kallikrein family of secreted serine proteases (KLKs 1, 5, 6, 7, 8 and 9) in human spinal cord trauma. The most widespread changes in kallikrein expression occurred during the acute and subacute periods, with KLK9 remaining highly elevated even at late chronic post-mortem intervals. Importantly, despite differences in the evolution of the kallikrein locus across species (70), the same general patterns of kallikrein expression were observed in a complementary clinically relevant murine SCI model, pointing to conserved pathophysiological mechanisms. In addition, five of the six kallikreins studied in detail promoted injury to cortical neurons and their processes *in vitro*. Taken together, these studies define kallikreins as important regulators of the SCI microenvironment and as potential new targets for the design of therapies to reduce axon injury and create an environment favorable to repair across the acute through chronic SCI continuum.

Given the dynamic alterations documented in the expression of KLKs 1, 5, 6, 7, 8 and 9 in both human and rodent traumatic SCI, including the association of each with injured axons, we made a comprehensive assessment of the effects of the recombinant forms of these proteases, in addition to KLKs 2, 3 and 13, toward primary murine cortical neurons *in vitro*. Results demonstrate that elevated levels of KLKs 1, 2, 5, 6, 7 and 9 are capable of promoting a significant injury to neurons and/or their processes *in vitro*. These findings are consistent with prior studies pointing to the neurotoxic effects of KLK1 and KLK6 (9) and reveal the potential for KLKs 2, 5, 7 and 9 to promote neural injury at parallel concentrations for the first time. Taken with previously published reports implicating deregulated expression of multiple kallikreins in Alzheimers (1–5), fronto-temporal dementia (6), Parkinsons (3, 7), post-polio-syndrome (10), glioblastoma multiforme (47, 71) and multiple sclerosis (8, 9), it will be important in future studies to examine the consequences of a wider range of kallikrein concentrations and to define their parallel or differential effects towards multiple neural cell types.

Taken with the substantial elevations in kallikrein expression at one or more of the post-SCI intervals examined, additional efforts to fully resolve the mechanism(s) by which they alter neuron and neurite numbers and their potential utility as targets for neuroprotection and neural repair will be needed. The possibility that a subset of kallikreins exert their neurotoxic effects at least in part by activation of G-protein linked protease activated receptors (PARs) is supported by *in vitro* studies demonstrating that KLKs 5 and 6 activate one or more PARs *in vitro* (23, 28, 30, 61, 72–74). Importantly, the ability of KLK6 to elicit mitogen activated protein kinase signaling and to promote injury of cultured cerebellar granule neurons was recently shown to be dependent on both PAR1 and PAR2 (23). Whether other kallikreins activate neuronal PARs and the roles played in neural injury remains an important avenue for future investigation.

While increases in KLK8 were also observed in human and rodent SCI, recombinant KLK8 did not elicit direct loss of cortical neurons or neurites *in vitro*, nor did KLK3 or KLK13, at the concentrations and time points examined. Of interest, KLK3, better known as prostate specific antigen, acted in a nerve growth factor-like fashion promoting an increase in

cortical neuron processes. While the increases in neurite density elicited by KLK8 in the present study did not reach statistical significance, a prior study showed KLK8 enhances neurite density in cultured hippocampal neurons (75). In this regard, it is important to highlight the substantial prior work indicating that KLK8 plays a key role in neuroplasticity. For example, KLK8 deficient mice show reduced early-LTP and have abnormal CA1 synapses (76, 77). Importantly, we show that KLKs 1, 5, 6, 7 and 9 are expressed across both neurons and glia of the intact human spinal cord and therefore future efforts to determine whether their regulated proteolytic actions may likewise be involved in neural plasticity, or other key neurophysiologic activities, is needed. While KLK2 and KLK3 are not densely expressed in the intact CNS (25), and their expression with injury has not been studied, each could enter the damaged brain or spinal cord in areas of increased vascular permeability. In this case, further study regarding the potential neurite outgrowth promoting effects of KLK3, and the ability of KLK2 to promote neural injury, will also be important future directions.

We note that only the direct effects of kallikreins toward neurons were examined and therefore these studies do not exclude a wider range of action in vivo, including modification of extracellular matrices or other CNS proteins and cell types that may indirectly influence neurite integrity and neuron viability. Several kallikreins are reported or predicted to degrade extracellular matrix (ECM) proteins, including basement membrane components, such as fibronectin, laminin and heat denatured collagen (8, 56, 78–80). In fact, the role of KLK8 in promoting neural plasticity occurs in part by its activities at the neuronextracellular matrix interface, including cleavage of synaptic L1 (81), EphB2 (82), and Neuregulin 1 (83). Our prior studies indicate that KLK6 hydrolyzes laminin to decrease neurite outgrowth and aggrecan to promote neurite extension in vitro (22). Deregulated expression of KLK6 with CNS injury is therefore positioned to modulate the balance of permissive and inhibitory cues that regulate axon outgrowth and the potential for nerve regeneration. Also, KLK6 readily degrades myelin proteins, including MBP and myelin oligodendrocyte glycoprotein (8, 72) and therefore may contribute to axon injury and conduction block in SCI by promoting demyelination. Adding to the complexity of the potential consequences of elevated (or reduced) kallikrein expression at sites of CNS pathology, KLKs may participate in activation of metalloproteases (23, 84) or fibrinolytic enzymes (57-59, 85), which themselves may promote neurotoxic effects.

The kallikrein expression patterns described in the intact and injured spinal cord do not distinguish the relative amounts of inactive pro-kallikrein compared to that which N-terminal cleavage of a 3 to 37 amino acid pro-peptide results in the release of enzymatically active protein (57). While tools to distinguish between pro- and active kallikreins are yet to be developed, we have previously made a comprehensive analysis of activation cascades among the 15 kallikreins (57, 59). Since KLK9 not only showed the most robust and long-lasting elevations in expression across the cases of human and murine SCI examined, but was also among the most neurotoxic, we summarize the potential for the other kallikreins examined to be involved in KLK9-activation cascades (Fig. 10). For example, KLKs 2, 3, 5, 6 and 13 may directly activate pro-KLK9. In addition, KLKs 5, 6, 7 and 8 may indirectly activate pro-KLK9. While not examined here, KLKs 4, 12 and 14 also activate KLK9.

Interestingly, KLK9, as well as KLK1 do not activate other kallikreins (59). Taken together these data reveal for the first time multiple avenues for kallikrein enzymatic activation in the intact and injured CNS.

Dense kallikrein staining associated with astrocyte-like cells at sites of SCI points to novel potential roles for these unique enzymes in the process of injury-induced astrogliosis. For example, KLKs 1, 5, 7 and 9 were localized to astrocyte-like cells in the intact human spinal cord and each of these, in addition to KLK6, were elevated in swollen/hypertrophic astrocytes that were prominent in the injured spinal cord from acute through chronic time points, particularly in white matter. Supporting the likely role of at least a subset of KLKs in the process of astrogliosis, KLK6 activates astrocyte PAR1 and PAR2 to promote Ca<sup>2+</sup> flux and mitogen activated protein kinase signaling (61). In addition, KLK6 promotes robust astrocyte stellation and IL-6 secretion in vitro, in part by activation of PAR1 (28). Given the prominent association of several kallikreins with astrocyte-like cells in human SCI, a model is suggested whereby elevations in astrocyte kallikrein act in an autocrine or paracrine fashion to promote key aspects of astrogliosis, including stellation and IL-6 secretion. As this line of research proceeds, it will be necessary to determine the respective roles of individual or combinations of kallikreins in astrogliosis in order to delineate whether their effects are part of the beneficial wound healing response and/or may contribute to the formation of an astroglial nerve regeneration barrier (66, 86).

The inflammatory response in spinal cord trauma is complex and includes both cellular and chemical mediators (67, 69, 87–90). Corresponding to what has been reported, monocytes and microglia reactive for the lysosomal marker CD68 were observed from acute through the most chronic time points of human SCI examined and were most abundant from the subacute period onward (43, 69). In parallel sections, KLKs 1, 5, 6, 7 and 9 were identified in association with enlarged rounded macrophages. Notably, KLKs 1, 5, 6 and 7 have been previously implicated in inflammation (25, 91, 92) and results here support such roles in context of traumatic CNS injury. Kallikrein 6 has been previously localized to monocytes and microglia in human and rodent SCI (22) as well as in MS lesions and in viral and autoimmune MS models (8, 29, 54, 55). The association of at least low levels of KLK9 with rounded macrophages in SCI, points to potential roles for this protease in inflammatory events as well. Further supporting the involvement of a subset of kallikreins in inflammatory cascades secondary to SCI, peak elevations in KLKs 1, 6, 7, 8 and 9 seen across the acute and subacute periods in experimental murine SCI corresponded to parallel increases in the pro-inflammatory cytokines IL-6, IL-1β and TNFα, which are known to be produced by monocytes, microglia and other reactive cells, including T cells and astrocytes at sites of SCI (93, 94). The meaning of co-ordinate expression of kallikreins and cytokines is not yet known, but regulatory roles are possible based on prior studies demonstrating KLK6 promotes secretion of IL-6 from astrocytes (28) and KLK5 stimulates production of TNFa and IL-8 from keratinocytes (91). Future studies will be needed to determine how kallikreins affect the balance of cellular and chemical mediators of CNS inflammation (95-97) and whether, like the much better studied protease matrix metalloprotease 9 (50, 98), select kallikreins may serve as targets for therapeutic intervention.

Taken together, results spanning human SCI and a complementary murine experimental SCI model indicate that multiple kallikreins participate in the response of the spinal cord to trauma with possible roles in axon injury, astrogliosis and inflammatory events occurring from acute through more chronic stages. For the subset of kallikreins shown to be elevated post-injury and capable of promoting neural injury *in vitro*, that is KLKs 1, 5, 6, 7 and 9, additional efforts to evaluate whether they can be targeted to prevent secondary injury cascades and promote spinal cord repair and plasticity will be of particular interest.

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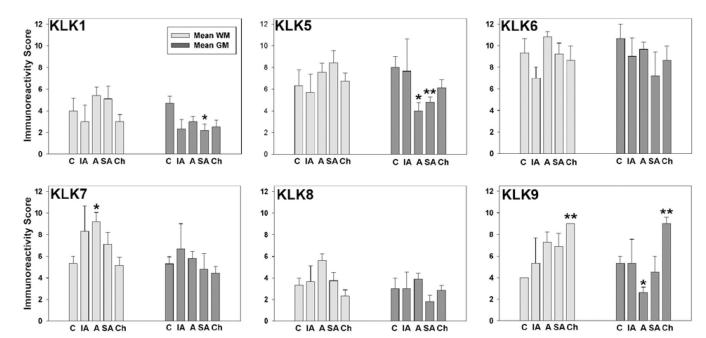
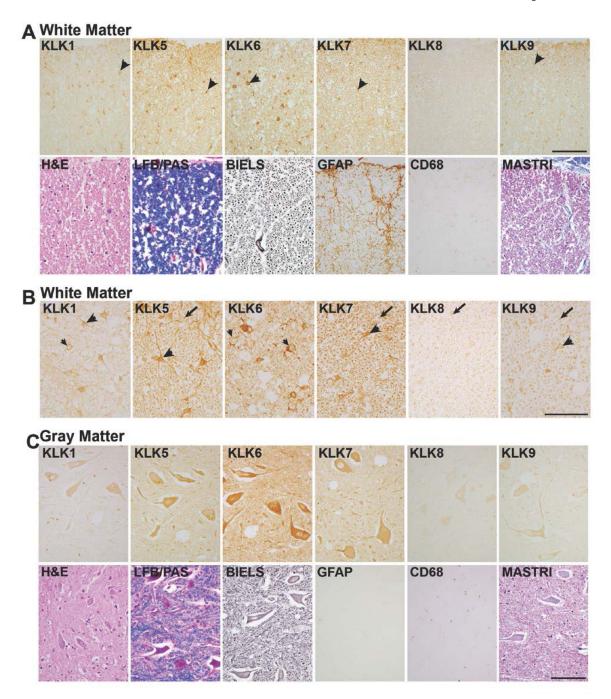


FIGURE 1. KLK immunoreactivity scores in the intact and traumatically injured human spinal cord

Histograms show changes in kallikrein immunoreactivity (IR) scores seen in the uninjured spinal cord (C) and at immediate acute (IA, 0 to 1d), acute (A, >1d to < 14d), subacute (SA, > 14d to 63d), and chronic (Ch 413d to 2133d) time points post-injury. The most widespread changes in KLK-IR occurred during the acute and subacute periods, including significant elevations in KLK7 in white matter (\*P = 0.04, Student's t-test) and significant reductions in KLK1 (\*P = 0.04, Student's t-test), KLK5 (\*P = 0.03, Rank Sum test (Acute); \*\*P = 0.02, Student's t-test (Subacute), and KLK9 (\*P = 0.04, Rank Sum test (Acute) in gray matter. KLK9 was significantly elevated in both spinal cord white matter and gray matter at the chronic time points examined (\*\*P = 0.01, Student's t-test). Photomicrographs showing representative KLK-IR staining patterns in the uninjured spinal cord (Fig. 2) and at immediate acute (Fig. 3), acute (Fig. 4), subacute (Fig. 5, 6); and chronic (Fig. 7) time points are provided. Also, see Fig. 8 for comparison of changes in KLK expression in human SCI with those occurring in a murine experimental contusion-compression SCI model.



 $FIGURE \ 2. \ Differential \ expression \ of \ kallikreins \ in \ the \ white \ and \ gray \ matter \ of \ the \ uninjured \ human \ spinal \ cord$ 

A, In spinal cord white matter, KLKs 1, 5, 6, 7 and 9 were localized to glia (arrow heads, A). B, Higher power photomicrographs show localization of KLKs 1, 5, 7 and 9 in glia with a triangular shape resembling astrocytes (arrow heads). KLK6 predominated in glia with a round nucleus and thin rim of cytoplasm resembling oligodendroglia (small arrow heads). KLK1 was also seen a few oligodendrocyte-like cells (small arrow head). KLKs 5, 7, 8 and 9 were also localized to white matter axons (arrows). C, In spinal cord gray matter (ventral horn shown), each kallikrein was detected in neurons with highest levels of KLK6, 5 and 7

followed by KLK1 and KLK9 and finally, KLK8. Comparison of KLK-IR scores in the uninjured and injured spinal cord is provided in Fig. 1. Parallel sections were stained with H&E, LFB/PAS, Bielschowsky' silver stain (BIELS) and Masson's Trichrome (MASTRI), in addition to antibodies recognizing GFAP and CD68. GFAP-IR cells and processes were identified in uninjured spinal cord white matter, but were very low in uninjured gray matter. CD68-IR monocytes/microglia were very sparse or absent in both the white and gray matter of the uninjured spinal cord. The cervical spinal cord from a case in which death was due to acute leukemia is shown (Scale bar =  $100 \, \mu m$  A and C;  $50 \, \mu m$  B).

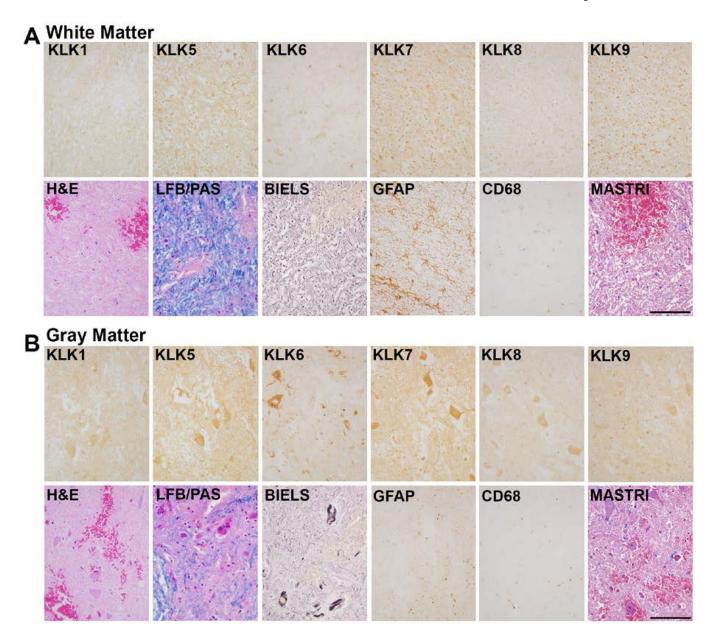


FIGURE 3. Expression of kallikreins in the immediate acute period after traumatic SCI A, In the immediate acute period after SCI (0–24 hr), minor elevations in KLK7 and KLK9 were seen in spinal cord white matter while levels of the other kallikreins examined were unchanged or decreased relative to those observed in the uninjured spinal cord (Fig. 1, 2). Similarly, very minor changes in kallikrein-IR were observed in spinal cord gray matter (B) in the immediate acute period (Fig. 1). Parallel sections were stained with H&E, LFB/PAS, Bielschowsky' silver stain (BIELS) and Masson's Trichrome (MASTRI), in addition to antibodies recognizing GFAP and CD68. Levels of GFAP-IR and CD68 in spinal cord gray and white matter in parallel sections were similar to those observed in the uninjured cases examined and patches of petechial hemorrhage were obvious in both H&E and Masson's Trichrome (MASTRI) stained sections. LFB/PAS and Bielschowsky's silver stain (BIELS)

are also shown. Case shown is C1 fracture dislocation as a result of a motor vehicle accident (Scale bar =  $100 \ \mu m$ ).

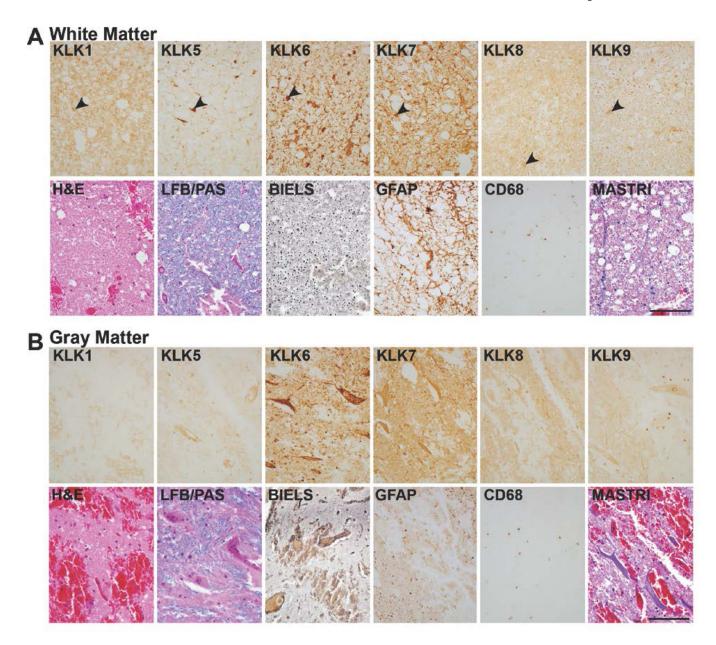


FIGURE 4. Expression of kallikreins in the acutely injured human spinal cord white and gray matter

A, At acute time points after traumatic injury (4d shown), elevations were observed in each of the kallikreins examined in white matter and there each was associated with glial-like cells (arrow heads, A). Elevations in white matter KLK7 were statistically significant (P = 0.04, Student's t-test, see Fig. 1). B, Overall, KLK-IR in the gray matter at acute time points showed either no change (KLK6 and KLK7) or decreases in immunoreactivity relative to uninjured spinal cord (Fig. 2). Reductions in gray matter KLK5 and KLK9 were statistically significant (P 0.04, Rank Sum test, see Fig. 1). Cytological and histopathological signs of injury evaluated in parallel sections showed patches of petechial hemorrhage in H&E and Masson's Trichrome (MASTRI) stained sections. Overall levels of GFAP and CD68-IR in the case shown were similar to controls although elevations were evident in other acute

cases examined, defined as >1d to <14d after injury. LFB/PAS and Bielschowsky's silver stain (BIELS) are also shown. Case shown is C4 fracture dislocation as a result of a motor vehicle accident (Scale bar =  $100~\mu m$ ).

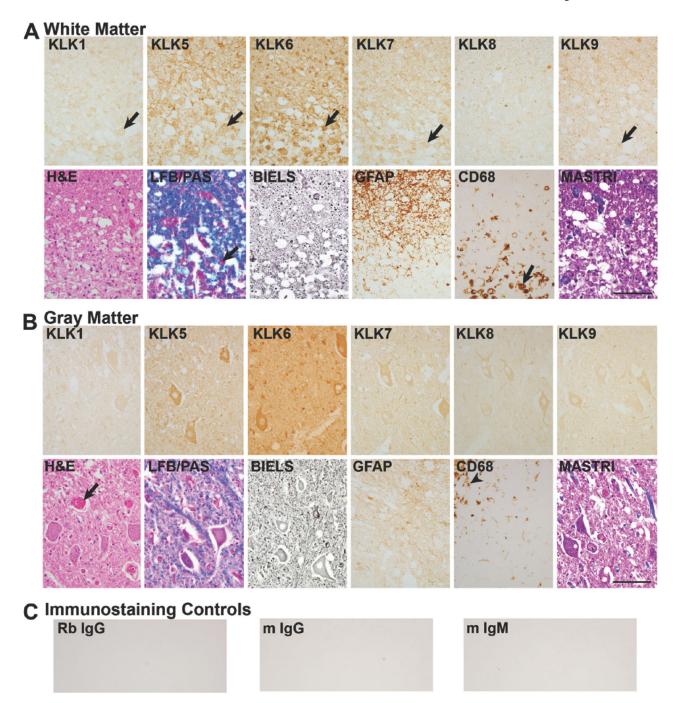


FIGURE 5. Expression of kallikreins in a case of subacute traumatic SCI

A, At subacute time points after traumatic injury (14d shown), elevations in KLK1, KLK5, KLK7 and KLK9 were observed in white matter, although these did not reach the level of statistical significance across the cases examined. Increases in CD68-IR monocytes/microglia were prevalent in white matter at subacute time points, which were either densely (KLK6), moderately (KLKs 5 and 7) or lightly (KLKs 1 and 9) KLK-IR in near adjacent sections (arrows, A). Inflammatory cells in white matter were also distinguished in LFB/PAS stained sections (arrows). Very clear increases in GFAP-IR were also observed in

white matter at subacute time points. B, Overall levels of KLK1 and KLK5 were significantly reduced in spinal cord gray matter across the subacute cases examined (P 0.04, Student's t-test, see Fig. 1). The cytoplasm of some anterior horn cells was associated with hypereosinophilia in parallel sections (arrow, B) and CD68 positive monocytes/microglia (arrow heads, B) were also observed in gray matter at subacute time points. C, Panels show the level of immunoreactivity achieved in the injured spinal cord when primary antibodies were replaced with normal serum and tissue processed with either rabbit IgG, or with mouse IgG or IgM secondary antibodies alone. Case shown is a T5 fracture dislocation, paraplegia. (Scale bar =  $100 \ \mu m$ ).

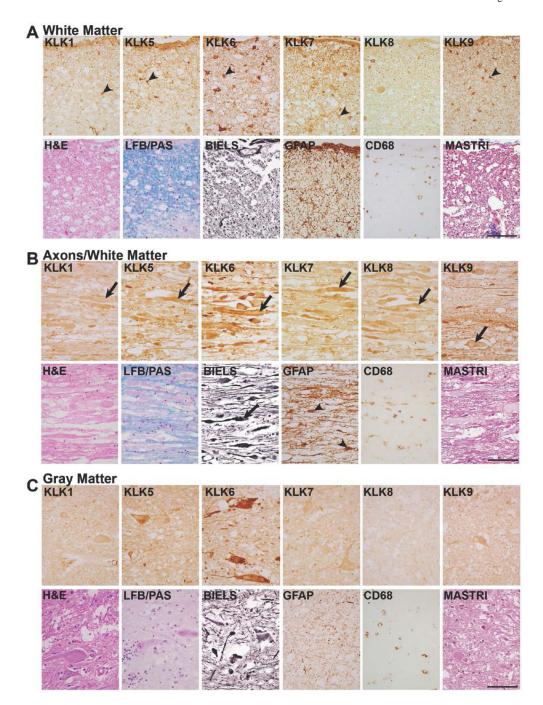


FIGURE 6. Elevated levels of kallikrein expression by axons in traumatic SCI

A, In subacute cases (63d shown), KLKs 1, 5, 6, 7 and 9 were associated not only with white matter glia (arrow heads, A), but along with KLK8 each was also clearly identified in swollen axons and axon ovoids seen in longitudinal sections taken from the same case (arrows, B). These injured/degenerating and swollen axons were also apparent in the near adjacent Bielschowsky's silver stained (BIELS) sections (arrows). Prominent GFAP-IR, including hypertrophic astrocytes (arrow heads, B) and CD68-IR monocytes were also seen in association with injured axons in white matter. C, Shows the overall reductions in KLK1

and KLK5 seen in gray matter during the subacute injury interval. GFAP-IR in gray matter was elevated relative to controls and CD68-IR monocytes/microglia were also evident. At this time point, numerous examples of explicit loss of LFB staining were seen both in the white and gray matter (see Fig. 1). Case shown is a C4 fracture dislocation as a result of a motor vehicle accident resulting in quadriplegia. (Scale bar =  $100 \mu m$ ).

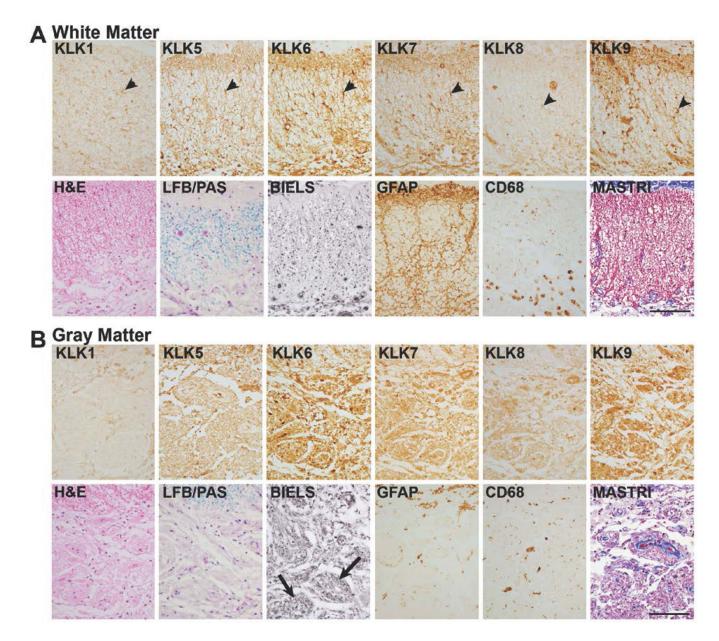


FIGURE 7. Expression of kallikreins in chronic SCI

A, At chronic time points after injury (1247d shown), overall levels of KLK9 were significantly elevated in spinal cord white matter (P = 0.02, Rank Sum test) and in gray matter (P = 0.01, Student's t-test), see Fig. 1. Changes in expression of the other kallikreins were also observed particularly in the cellular distribution of immunoreactivity in each case, although overall changes IR levels were not statistically significant. At this level of injury, the central core of gray matter was reorganized with some nerve fibers arranged in a fascicular fashion (arrows, B, BIELS). The density of LFB staining was reduced in both white and gray matter compared to uninjured controls (see Fig. 1). GFAP-IR and CD68-IR continued to be elevated in white matter even at very chronic stages. KLK1, 5, 6, 7 and 9 continued to be clearly associated with white matter glia, and KLK8-IR glia were also occasionally seen (arrow heads, A). Masson's trichrome (MASTRI) was used to examine

collagen. In both white and gray matter, blue stained collagen fibers were largely confined to blood vessels. Case shown is a C5 injury due to fall. (Scale bar =  $100 \ \mu m$ ).

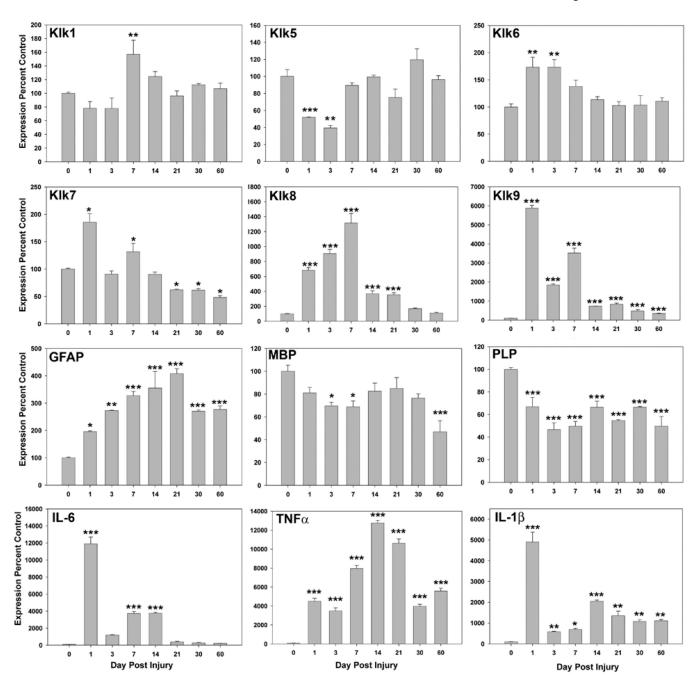


FIGURE 8. Experimental murine traumatic SCI promotes changes in kallikrein gene expression that parallel those seen at a protein level in the case of human SCI

Histograms show the change in kallikrein RNA expression seen in response to clip-compression SCI in mice at immediate acute (1d), acute (3, 7d), subacute (14, 21, 30d) and more chronic (60d) time points after compression injury, relative to that seen in the uninjured spinal cord (n=3 at each time point, except 14d n=4). Samples at each time point were combined and examined by PCR in triplicate. Co-ordinate changes in the expression of GFAP, myelin genes (MBP and PLP) and the pro-inflammatory cytokines IL-6, TNF $\alpha$  and IL-1 $\beta$  are also provided. RNA expression levels in each case were normalized to GAPDH to

control for loading and are expressed as a percent control (mean and s.e. shown, \* P < 0.05, \*\* P = 0.01, \*\*\* P = 0.001, SNK).

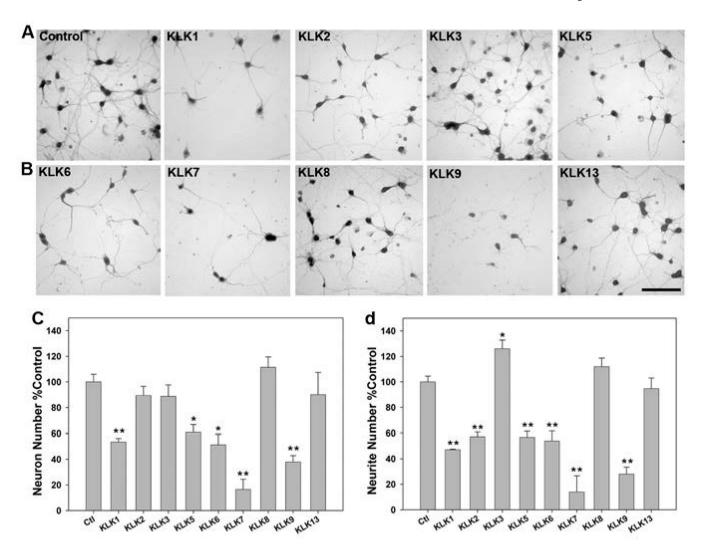
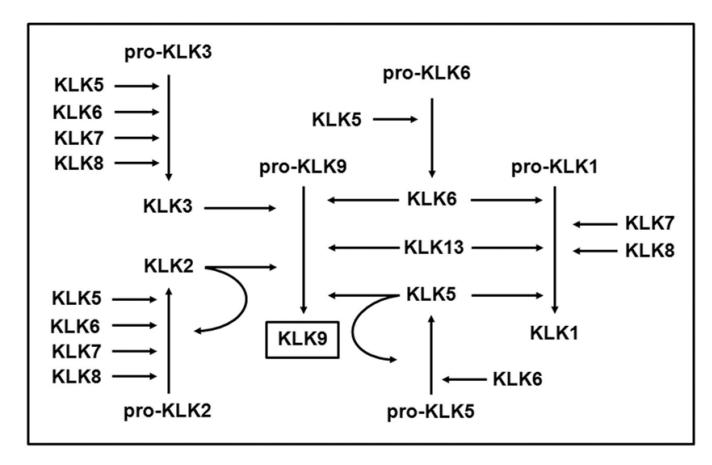


FIGURE 9. Differential activity of kallikreins in promoting neural injury in vitro

Murine embryonic day 15 (E15) cortical neurons were cultured on cover glass coated with poly-L-lysine for 96 hr prior to treatment with recombinant kallikreins (10 µg/ml (300 nM)) for 24 hr (A, B). Neurons were immunostained for Neurofilament H and neuron number (C) and neurite density (D) quantified. KLKs 1, 5, 6, 7 and 9 promoted significant loss of neurons and their processes. KLK2 also promoted neurite loss, while KLK3 increased the number of neurites counted. (\* P < 0.03, \*\* P 0.001, SNK). Histograms show the mean and s.e. of neuron and process counts across three independent cover slips treated in parallel and findings were reproducible across 3 experiments utilizing independent murine cortical neuron preparations. (Scale bar = 100 µm)



 ${\bf FIGURE~10.~Schematic~depicts~potential~kallikrein~activation~cascades~among~the~proteases~examined~in~the~current~study}$ 

The ability of each kallikrein to activate other members of the kallikrein family of proteases was established in our prior studies (57, 59). Here we present a theoretical scheme leading to the production of enzymatically active KLK9, which of six kallikreins examined was the most highly upregulated in the injured human and rodent spinal cord (Figs. 1 and 8) and exhibited robust neurotoxicity (Fig. 9).

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**TABLE 1** 

Summary of the human SCI and control cases examined

Injury Cause of Level Injury	C1 MVA	C3 MVA	C5 MVA	C1 MVA	C4 MVA	C MCA	C2 Go-cart	C1 MVA	C1-C4 MVA	T3 MVA	T5 UD	T5 MVA	T2 GSW	T7 Fall	C3 MVA	C4 MVA	C1 MVA	C5 Fall	C5 MVA	C5 Dive	T4 Fall	C4 Dive	N/A N/A	
Injury Inj Interval (d) Le	0	0	0	2	Α (	4	4 O	9	7 CI-	R 8	T T	Z2 T	30 T	30 T	37 C	63 C	413 C	1247 C	1264 C	3001 C	5315 T	7704 C	N/A	
Group	Imm. Acute	Imm. Acute	Imm. Acute	Acute	Acute‡	Acute	Acute#	Acute	Acute#	$Acute^{\dagger}$	Subacute#	Subacute <sup>†‡</sup>	$Subacute^{\dagger}$	$Subacute^{\dagger}$	Subacute#	Subacute	Chronic#	Chronic#	Chronic	Chronic	Chronic	Chronic	Control	
Sex	Щ	Σ	Σ	Σ	щ	Σ	$\mathbf{Z}$	Ц	щ	Σ	Σ	Σ	Σ	Σ	Σ	Σ	$\boxtimes$	щ	M	Σ	M	Σ	H	
Age (Yr)	29	19.6	35.8	4.4	43.8	19.8	11.7	3.9	18.4	36.9	54.5	T.TT	55.4	78.4	18.7	39.3	6	65.4	27.5	22.3	46.4	18.2	41	
Case	-	2	3	4	2	9	7	∞	6	10	11	12	13	41	15	16	17	18	19	20	21	22	23	

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Immediate (Imm.) Acute (0 to 1d); Acute (>1d to <14d); Subacute (>14d to 63d); Chronic (>413d to 2133d); MVA (motor vehicle accident); MCA (motorcycle accident); UD (cause of trauma undetermined); N/A (not applicable);

 $^{\dagger}\mathrm{IHC}$  available for white matter only;

 $^{\sharp}$ IHC examined across two adjacent blocks.

# TABLE 2

Summary of primers used for quantitative real time PCR analysis of murine kallikreins (Klk), GFAP, myelin genes (MBP and PLP) and pro-inflammatory cytokines (IL-6, TNF $\alpha$ , IL1 $\beta$ ).

Accession Number	Base Pairs	Forward Primer	Reverse Primer					
mKlk1 NM_010639	363–471	CAGGAGATATGGATGGAGGCAAAGA	CCCGGCACATTGGGTTTAC					
mKlk5 NM_026806.2	854–1007	GCGGAACAGACCAGGTGTCTA	CTTCAGCATTCTCAGAGAGGG					
mKlk6 NM_011177.1	749–949	CCTACCCTGGCAAGATCA	GGATCCATCTGATATGAGTGC					
mKlk7 NM_011872.2	143–245	ACCGCTGGACAAGGAGAAA	CTCCACAGTGAAGCTGATTGC					
mKlk8 NM_008940.2	1004–1236	GTGCGGAAGTGAAAATCTAT	GGTAGTGTAGCGGCAG					
mKlk9 NM_028660.3	766–884	AGGATCTGAGCCTTGT	TCATGCTAAGATGGACAGAC					
<b>GFAP</b> NM_010277.2	600–767	GCAGATGAAGCCACCCTGG	GAGGTCTGGCTTGGCCAC					
PLP NM_011123.2	377–537	TCTTTGGCGACTACAAGACCAC	CACAAACTTGTCGGGATGTCCTA					
IL-6 NM_031168.1	Amplicon Length: 78	The assay probe spans Exon Boundary: 2–3 (Applied Biosytems)	Assay ID: Mm00446190_m1					
<b>TNFa</b> NM_013693.2	Amplicon Length: 81	The assay probe spans Exon Boundary: 1–2 (Applied Biosytems)	Assay ID: Mm00443258_m1					
IL-1β NM_008361.3	110–224	TGACAGTGATGAGAATGACCTG Probe: /56-FAM/AGT TGA CGG /ZEN/ACC CCA AAA GAT GAA GG/3IABkFQ/(IDT)	CGAGATTTGAAGCTGGATGC Assay ID: Mm.PT.51.17212823					
GAPDH NM_008084.2	351–584	ACCACCATGGAGAAGGC	GGCATGGACTGTGGTCATGA					