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Facilitated spinal neuropeptide signaling and upregulated inflammatory mediator expression contribute to post-fracture nociceptive sensitization

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Abstract

Tibia fracture induces exaggerated substance P (SP) and CGRP signaling and neuropeptide-dependent nociceptive and inflammatory changes in the hindlimbs of rats similar to those seen in complex regional pain syndrome (CRPS). Inflammatory changes in the spinal cord contribute to nociceptive sensitization in a variety of animal pain models. This study tested the hypothesis that fracture induced exaggerated neuropeptide signaling up-regulates spinal inflammatory mediator expression, leading to post-fracture hindlimb nociceptive sensitization. At 4 weeks after performing tibia fracture and casting in rats, we measured hindlimb allodynia, unweighting, warmth, edema, and spinal cord neuropeptide and inflammatory mediator content. The antinociceptive effects of intrathecally injected neuropeptide and inflammatory mediator receptor antagonists were evaluated in fracture rats. Transgenic fracture mice lacking SP or the CGRP RAMP1 receptor were used to determine the effects of neuropeptide signaling on post-fracture pain behavior and spinal inflammatory mediator expression. Hindlimb allodynia, unweighting, warmth, edema, increased spinal SP and CGRP, and increased spinal inflammatory mediator expression (TNF, IL-1, IL-6, CCL2, NGF) were observed at 4 weeks after fracture in rats. Fracture induced increases in spinal inflammatory mediators were not observed in fracture mice lacking SP or the CGRP receptor and these mice had attenuated post-fracture nociceptive sensitization. Intrathecal injection of selective receptor antagonists for SP, CGRP, TNF, IL-1, IL-6, CCL2, or NGF each reduced pain behaviors in the fracture rats. Collectively, these data support the hypothesis that facilitated spinal neuropeptide signaling up-regulates the expression of spinal inflammatory mediators contributing to nociceptive sensitization in a rodent fracture model of CRPS.

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Keywords

fracture; complex regional pain syndrome; cytokines; nerve growth factor; substance p; calcitonin gene-related peptide; pain

1. Introduction

Population based studies indicate that distal limb fracture is the most common cause of CRPS [11,36] and we have developed a tibia fracture model in the rat and mouse that closely resembles CRPS. Distal tibia fractured rats treated with 4-weeks cast immobilization exhibited paw allodynia, unweighting, increased spinal Fosimmunoreactivity, increased hindpaw skin temperature, edema, increased spontaneous protein extravasation, facilitated neuropeptide signaling, regional periarticular bone loss, keratinocyte proliferation and epidermal thickening, and increased keratinocyte expression of TNF, IL-1, IL-6, and NGF inflammatory mediators in the affected skin [13,14,25,27,34,35,46,47].

Using the rat fracture model of CRPS we have shown that up-regulated expression of neuropeptides such as substance P (SP) and calcitonin gene-related peptide (CGRP) and their cognate receptors [46] leads to inflammation and pain sensitization by the activation of neuropeptide receptors on the surface of cutaneous keratinocytes, mast cells, and vascular endothelial cells, causing mast cell degranulation [26], vasodilatation and protein extravasation [46], keratinocyte proliferation and activation, and keratinocyte expression of high levels of inflammatory cytokines (TNF, interleukin 1 beta (IL-1), IL-6) and nerve growth factor (NGF) [15,25,27,34,35,47]. Experimental maneuvers blocking these inflammatory mediators can inhibit the nociceptive and vascular changes that develop over the first 4 weeks in the CRPS fracture model [26,27,34,35,47]. Similarly, clinical studies in CRPS patients have reported therapeutic responses with treatments that inhibit innate immune mechanisms, but these treatments are not effective for every patient [8,17,21,40]. Utilizing bilateral skin punch biopsies from 55 CRPS patients, we recently demonstrated unilateral mast cell and keratinocyte proliferation, epidermal thickening, and keratinocyte inflammatory cytokine expression (TNF, IL-6) in early (< 3 month duration) CRPS affected skin [4]. This study also found that increased keratinocyte and mast cell proliferation gradually resolved over time, despite persistent pain symptoms. These results are consistent with skin blister fluid studies indicating a gradual resolution of cutaneous inflammatory mediator (IL-6, TNF) expression over time in CRPS affected skin [49]. Collectively, these data support the hypothesis that cutaneous keratinocyte and mast cell mediated innate immune responses contribute to the development of post-traumatic CRPS, but cannot completely account for the inflammation and pain observed in early and particularly in chronic CRPS.

It has been hypothesized that neuropeptides released from the central terminals of spinal primary sensory afferent neurons can initiate glial activation and inflammatory mediator expression contributing to chronic pain [42]. Interestingly, several studies have reported elevated cerebrospinal fluid levels of IL-1 and IL-6 in chronic CRPS patients [2,3], but other investigators have been unable to confirm these findings [30]. Elevated cerebrospinal fluid

levels of cytokines and neurotrophic factors have also been observed in other chronic pain states [6]. We postulated that facilitated spinal neuropeptide signaling up-regulates the expression of spinal inflammatory mediators contributing to the development of CRPS and the current study tests this hypothesis in the rodent fracture model of CRPS.

2. Materials and methods

All experiments were approved by our Institutional Animal Care and Use Committee (IACUC) and followed the animal subjects guidelines of the International Association for the Study of Pain. Adult (9-month-old) male Sprague Dawley rats (Simonsen Laboratories) and adult (3-month-old) male C57BL/6 wild type mice, SP deficient mice ($TAC1^{-/-}$, Jackson Laboratories), and CGRP RAMP1 receptor deficient mice ($RAMP1^{-/-}$, generated by Dr. Kazutake Tsujikawa at Dept of Immunology, Osaka University) were used in these experiments. Rats were housed individually in isolator cages with solid floors covered with 3 cm of soft bedding, while mice were housed for 4 to a cage. Food and water were provided *ad libitum*.

2.1. Surgery

Tibia fracture was performed in the rat under isoflurane anesthesia as we have previously described [13]. The right distal tibia was fractured using pliers with an adjustable stop (Visegrip, Petersen Manufacturing) that had been modified with a 3-point jaw. The right hind limb was then wrapped in casting tape so the hip, knee and ankle were flexed. The cast extended from the metatarsals of the hind paw up to a spica formed around the abdomen. To prevent the rats from chewing at their casts, the cast material was wrapped in galvanized wire mesh. The animals were given subcutaneous saline and buprenorphine (0.03mg/kg) immediately after procedure and on the first day after fracture for post-operative hydration and analgesia. At 4 weeks post-fracture the rats were anesthetized with isoflurane and the cast removed with a vibrating cast saw. The following day the rats underwent behavioral testing for hind paw von Frey allodynia, unweighting, edema, and warmth. The tibia fracture protocol in mice was similar to the procedure describe in rats, but a 6 inch straight hemostat was used for making the fracture [15]. In mice the cast was removed at 3 weeks post-fracture and behavioral testing was performed the following day. All animals used in this study had union at the fracture site at the time of cast removal.

2.2. Drug administration

All drugs were injected intrathecally in 10ul of saline under isoflurane analgesia. The following drugs were used; LY303870 (20ug), a SP neurokinin 1 (NK1) receptor antagonist (Eli Lilly Company); CGRP₈₋₃₇ (20ug), a CGRP receptor antagonist (Sigma); etanercept (100μg), a TNF inhibitor (Pfizer); anakinra (100ug), an IL-1 receptor antagonist (Amgen); TB-2081 (20ug), an IL-6 receptor antagonist (a generous gift from Dr. Kenner Rice, National Institute on Drug Abuse); RS504393 (3ug), a C-C chemokine receptor type 2 (CCR2) antagonist (Sigma); muMab 911 (12.4ug), an anti-NGF antibody (Pfizer); and PD98059 (20ug), a selective inhibitor of extracellular signal-regulated kinase 1/2 (ERK1/2) mitogen-activated protein kinase (MAPK) phosphorylation (Sigma).

2.3. Hind paw allodynia and unweighting measurements

To measure mechanical allodynia in the rats and mice an up-down von Frey testing paradigm was used as we previously described [15]. Test animals were placed in a clear plastic cylinder (20 cm in diameter for rats, 10 cm in diameter for mice) with a wire mesh bottom and allowed to acclimate for 15 minutes. The paw was tested with von Frey fibers of sequentially increasing stiffness applied against the hind paw plantar skin at the same site, and pressed upward to cause a slight bend in the fiber and left in place for 6 sec. Withdrawal of or licking the hind paw after fiber application was scored as a response. When no response was obtained the next stiffest fiber in the series was applied to the same paw; if a response was obtained a less stiff fiber was applied. Testing proceeded in this manner until 4 fibers had been applied after “negative + positive or positive + negative” response. Estimation of the mechanical withdrawal threshold by data fitting algorithm permitted the use of parametric statistics for analysis [32]. Baseline von Frey withdrawal thresholds were identical in WT, *Tac1^{-/-}*, and *RAMP1^{-/-}* mice. An incapitance device (IITC Inc. Life Science) was used to measure hind paw unweighting. The rats and mice were manually held in a vertical position over the apparatus with the hind paws resting on separate metal scale plates and the entire weight of the animal supported on the hind paws. The duration of each measurement was 6 seconds and 10 consecutive measurements for rats were taken at 60-second intervals. Mice were tested 6 consecutive times. Eight readings (excluding the highest and lowest ones) for rats and 6 for mice were averaged to calculate the bilateral hind paw weight bearing values.

2.4. Hind paw volume

A laser sensor technique was used to determine the dorsal-ventral thickness of the hind paw, as we have previously described [13]. For laser measurements each animal was briefly anesthetized with isoflurane and then held vertically so the hind paw rested on a table top below the laser. The paw was gently held flat on the table with a small metal rod applied to the top of the ankle joint. Using optical triangulation, a laser with a distance measuring sensor was used to determine the distance to the table top and to the top of the hind paw dorsal skin over the midpoint of the third metatarsal and the difference was used to calculate the dorsal-ventral paw thickness. The measurement sensor device used in these experiments (Limab) has a measurement range of 200 mm with a 0.01 mm resolution.

2.5. Hind paw temperature

The temperature of the hind paw was measured using a fine wire thermocouple (Omega) applied to the paw skin, as previously described [13]. The investigator held the wire using an insulating Styrofoam block. Three sites were tested over the dorsum of the hind paw: the space between the first and second metatarsals (medial), the second and third metatarsals (central), and the fourth and fifth metatarsals (lateral). After a site was tested in one hind paw the same site was immediately tested in the contralateral hind paw. The testing protocol was medial dorsum right then left, central dorsum right then left, lateral dorsum right then left, medial dorsum left then right, central dorsum left then right, and lateral dorsum left then right. The six measurements for each hind paw were averaged for the mean temperature.

2.6. Quantitative real-time PCR in spinal cord tissues

At 4 weeks after fracture the rats were euthanized by CO₂ inhalation and the ipsilateral spinal cord (L4,5 lumbar enlargement) was collected after behavioral testing and frozen immediately on dry ice. In mice the spinal cord was collected at 3 weeks post-fracture. Total RNA was extracted using the RNeasy Mini Kit (QIAGEN) and the purity and concentration were determined spectrophotometrically. Next, cDNA was synthesized from 1 µg RNA using an iScript cDNA Synthesis Kit (Bio-Rad Laboratories). Real-time polymerase chain reactions (PCRs) were conducted using the SYBR Green PCR master mix (Applied Biosystems) and performed on an ABI 7900HT sequencing detection system (Applied Biosystems). To validate the primer sets used, we performed dissociation curves to document single product formation, and agarose gel analysis was conducted to confirm the size (Table 1). The data from real-time PCR experiments were analyzed as described in the manual for the ABI prism 7000 real-time systems. All results were confirmed by repeating the experiment 3 times.

2.7. Enzyme immunoassay for SP, CGRP, TNF, IL-1, IL-6 and NGF levels in spinal cord

At 4 weeks after fracture the rats were euthanized by CO₂ inhalation and the ipsilateral spinal cord (L4,5 lumbar enlargement) was collected after behavioral testing and frozen immediately on dry ice. The spinal cord tissue was cut into fine pieces in ice-cold phosphate buffered saline (PBS), pH 7.4, containing protease inhibitors (Sigma) followed by homogenization using a Dounce tissue grinder. Homogenates were centrifuged for 15 min at 12,000 g, 4°C. Supernatants were transferred to precooled Eppendorf tubes. Triton X-100 was added at a final concentration 0.01 %. The samples were centrifuged again for 5 min at 12,000 g at 4°C. The supernatants were aliquoted and stored at -80°C. TNF-α was detected by using a TNF-α EIA kit (BD Biosciences). IL-1β, IL-6 and CCL2 protein levels were measured by using EIA kits from R&D Systems. The NGF concentrations were determined by using the NGF Emax® ImmunoAssay System kit (Promega) according to the manufacturer's instructions. The OD of the reaction product was read on a microplate reader at 450 nm, and values were normalized per mg of protein assayed. The concentrations of TNF-α, IL-1β, IL-6, NGF and CCL2 proteins were calculated from the standard curve for each assay. Each protein concentration was expressed as pg/mg total protein. Total protein content was determined using a Coomassie Blue Protein Assay Kit (Pierce).

SP and CGRP content in the rat spinal cord was measured as we have previously described [46]. Briefly, samples were minced in 1 ml of 3:1 ethanol/0.7 M HCl and homogenized for 20 s. The homogenates were shaken for 2 h at 4°C and centrifuged at 3,000 g for 20 min at 4°C. The supernatants were lyophilized and stored at -80°C. The lyophilized products were reconstituted with enzyme immunoassay (EIA) buffer before assay. After rehydration, the extracts were assayed in duplicate using SP and CGRP ELISA (Cayman Chemical) according to the manufacturer's instructions.

2.8. Western blotting of spinal cord

At 4 weeks after fracture the rats were euthanized by CO₂ inhalation and the ipsilateral spinal cord (L4,5 lumbar enlargement) was collected after behavioral testing and frozen immediately on dry ice. Spinal cord samples were later homogenized in modified RIPA

buffer (50mM Tris-HCl, 150 mM NaCl, 1 mM EDTA, 1% Igepal CA-630, 0.1% SDS, 50 mM NaF, and 1 mM NaVO₃) containing protease inhibitors (Sigma). The homogenate was centrifuged at 10,000 g for 30 min at 4°C. Total protein concentration of the homogenate was measured using a Coomassie Blue Protein Assay (Bio-Rad) and BSA protein standard (Pierce). Equal amounts of protein (50 µg) were subjected to SDS-PAGE (Bio-Rad) and electrotransferred onto a polyvinylidene difluoride membrane (Millipore). The blots were blocked overnight with 5% non-fat dry milk or normal serum in Tris-buffered saline with 0.5% Tween-20 (TBST), then incubated with primary antibodies against β-actin, NK1, total ERK1/2 and phosphorylated ERK1/2 (Santa Cruz Biotechnology), and total p38 and phosphorylated p38 (Cell Signaling Technology) for 1 hr on a rocking platform at room temperature. After washing in TBST, the blots were incubated with HRP-conjugated secondary antibody (Santa Cruz Biotechnology) for 1 h at room temperature. The blots were then washed in TBST again, then the proteins were detected using ECL chemiluminescence reagent (GE Healthcare) and scanned by PhosphoImager (Typhoon, GE Healthcare). The band intensity was analyzed using ImageQuant 5.2 software (Molecular Dynamics) and normalized with the corresponding internal loading control band. The specific protein expression is expressed as protein/actin band intensity ratio to demonstrate the change of the specific protein after treatments.

2.9. Tissue processing and immunofluorescence confocal microscopy

At 4 weeks post-fracture the rats were perfused with 4% paraformaldehyde (PFA) in phosphate buffered saline (PBS), pH 7.4, via the ascending aorta; the lumbar spinal cord was removed and post-fixed in 4% PFA overnight, then the tissues were treated with 30% sucrose in PBS at 4°C before embedding in OCT (Sakura Finetek). Following embedding, 20-µm thick slices were made using a cryostat, mounted onto Superfrost microscope slides (Fisher Scientific), and stored at -80°C.

Frozen sections were permeabilized and blocked with PBS containing 10% donkey serum and 0.3% Triton X-100, followed by exposure to the primary antibodies overnight at 4°C in PBS containing 2% serum. Upon detection of the first antigen, primary antibody from a different species against the second antigen was applied to the sections and visualized using an alternative fluorophore-conjugated secondary antibody. Sections were then rinsed in PBS and incubated with fluorophore-conjugated secondary antibodies against the immunoglobulin of the species from which the primary antibody was generated. After three washes, the sections were mounted with anti-fade mounting medium (Invitrogen). Images were obtained using confocal microscopy (Zeiss LSM/710, Carl Zeiss) and stored on digital media. The sources of primary antibodies were as follows: monoclonal mouse anti-phospho ERK, 1:100 (clone E4, Santa Cruz Biotechnology), monoclonal mouse anti-GFAP, 1:1000 (Clone GA5, EMD Millipore), polyclonal rabbit anti-Iba1, 1:1000 (WAKO). The sources of secondary antibodies were as follows: donkey anti-mouse IgG (1:400) conjugated with cyanine dye 3, or donkey anti-rabbit IgG (1:400) conjugated with fluorescein (FITC) secondary antibodies (Jackson ImmunoResearch Laboratories), incubated with respective primary antibodies. Control experiments included incubation of slices in primary and secondary antibody-free solutions both of which led to low intensity non-specific staining patterns in preliminary experiments (data not shown).

2.10. Statistical Analysis

Statistical analysis was done using Prism 4.02 (GraphPad Software). Sample sizes were based on a power analysis of preliminary and previously published data generated from using each of the proposed assays in fracture animals. Based on this analysis we calculated that the proposed experiments would require 8 animals per cohort to provide 80% power to detect 25% differences between groups. Animals were randomized to experimental groups using computer generated random numbers and all testing was performed in a blinded fashion when possible. No animals were excluded after enrollment into the experimental cohorts. Normal distribution of the data was confirmed using the D'Agostino-Pearson omnibus normality test. All data was evaluated using an analysis of variance (ANOVA) followed by Bonferroni *post hoc* multiple comparison testing. For simple comparisons of two means, two-tailed t-tests were performed. Data are presented as the mean \pm standard error of the mean (SEM) and a value of $P < 0.05$ was considered statistically significant.

The hindpaw von Frey mechanical nociceptive threshold, temperature, and thickness were analyzed as the difference between the fracture side (right, R) and the contralateral untreated side (left, L). Right hindpaw weight bearing data were analyzed as a ratio between twice the right hindpaw weight bearing and the sum of the right (R) and left (L) hindpaw weight bearing values ($(2R/(R + L)) \times 100\%$).

3. Results

3.1. Exaggerated neuropeptide signaling in the lumbar spinal cord after fracture

At 4 weeks after tibia fracture in rats there was increased expression of TAC1 (SP gene), TACR1 (SP neurokinin1 receptor gene), CALCA (alpha-CGRP gene) and CALCB (beta-CGRP gene) mRNA in the L4 and L5 spinal cord segments ipsilateral to the fracture (Fig. 1A). No changes were observed in mRNA expression for CALCRL (calcitonin receptor-like receptor, CGRP receptor gene) and RAMP1 (receptor activity modifying protein 1, CGRP receptor chaperone gene). SP and CGRP protein levels, as measured by enzyme immunoassay, also increased in the L4,5 spinal cord segments at 4 weeks post-fracture (Fig. 1B-C). Detected by western immunoblot assay, Fig. 1D illustrates that protein levels of the SP neurokinin 1 (NK1) receptor increased 2-fold in spinal cord after fracture. These data demonstrate the up-regulated expression of SP and CGRP signaling components in the lumbar spinal cord at 4 weeks post-fracture.

3.2. Inhibition of spinal SP and CGRP signaling reverses allodynia and unweighting

The anti-nociceptive effects of a single intrathecal injection of the SP NK1 receptor antagonist LY303870 and of the CGRP receptor antagonist CGRP₈₋₃₇ were evaluated in fracture rats. At 4 weeks after fracture hind paw von Frey nociceptive thresholds were reduced, but intrathecal LY303870 and CGRP₈₋₃₇ were both capable of reversing this mechanical allodynia (Fig. 2A). Similarly, intrathecal injection of LY303870 or CGRP₈₋₃₇ after fracture partially restored post-fracture hind paw weight bearing (Fig. 2B). These data indicate that spinal neuropeptide signaling contributes to nociceptive sensitization at 4 weeks post-fracture.

3.3. Cytokine induction in the lumbar spinal cord after fracture

The TNF, IL-1, IL-6, NGF and CCL2 mRNA and protein levels in spinal cord were measured by real-time PCR and EIA in fracture rats. Figure 3 demonstrates that the expression of these inflammatory mediators is up-regulated in the lumbar spinal cord at 4 weeks post-fracture.

3.4. Inhibition of inflammatory mediator signaling reverses allodynia and unweighting

To test the hypothesis that post-fracture up-regulated inflammatory mediator expression in lumbar spinal cord (Fig. 3) contributes to hind paw nociceptive sensitization, we intrathecally injected 4 weeks post-fracture rats with selective cytokine receptor antagonists (TNF- α receptor antagonist etanercept, IL-1 receptor inhibitor anakinra, IL-6 antagonist TB2081, or CCL2 receptor antagonist RS504393) or with the anti-NGF antibody muMab 911. As shown in Figure 4, all these inhibitors/antagonists partially reversed hind paw allodynia and unweighting in the fracture rats. These data indicate that up-regulated cytokine and nerve growth factor expression in the spinal cord can contribute to nociceptive sensitization at 4 weeks post-fracture.

3.5. Neuropeptide signaling contributes to post-fracture vascular and nociceptive changes

Transgenic mice were used to test the hypothesis that neuropeptide signaling mediates the post-fracture development of hind paw nociceptive sensitization and vascular changes. Figure 5 demonstrates that at 3 weeks post-fracture wildtype (WT) mice developed hindpaw von Frey allodynia, unweighting, warmth and edema, but SP deficient (SP KO, TAC1^{-/-}) mice and CGRP RAMP1 receptor deficient (RAMP1 KO, RAMP1^{-/-}) mice had significantly less allodynia, unweighting and warmth than WT fracture mice. SP deficient mice also failed to develop hind paw edema post fracture, but RAMP1 deficient fracture mice had a similar degree of hind paw edema as WT fracture mice (Fig. 5D). These results confirm that neuropeptide signaling is required for the development of post-fracture hind paw nociceptive sensitization, warmth, and edema.

3.6. Neuropeptide signaling contributes to post-fracture inflammatory mediator expression in the lumbar spinal cord

Figure 6 illustrates the post-fracture lumbar spinal cord expression of TAC1, TACR1, CALCA, CALCB, CALCRL, RAMP1, TNF, IL-1, IL-6, CCL2, and NGF mRNA in wildtype, SP deficient, and the CGRP RAMP1 receptor deficient mice. Compared to no fracture wildtype controls, the 3 weeks post-fracture wildtype mice exhibited a 2 to 5-fold increase in the spinal cord expression for all these genes. The SP deficient fracture mice had up-regulated TACR1, CALCA, TNF, and NGF expression, compared to no fracture wildtype controls, but this increase was less than the increase observed in the fracture wildtype mice. The increase in IL-6 expression in fracture SP deficient mice was same as that observed in the fracture wildtype mice. As expected, there was no TAC1 mRNA in the SP deficient mice and there was no post-fracture increase in spinal CALCB, CALCRL, RAMP1, IL-1, or CCL2. At 3 weeks after fracture, there was increased expression of TAC1, CALCA, TNF and CCL2 in the RAMP1 deficient mice, compared to control wildtype mice, but this increase was less than the increase observed in the fracture wildtype mice. TACR1,

CALCB, CALCRL, IL-1, IL-6 and NGF expression was unchanged in the fracture RAMP1 deficient mice, compared with no fracture wildtype controls. As expected, there was no RAMP1 mRNA in the RAMP1 deficient mice. These data indicate that neuropeptide signaling contributes to the post-fracture amplification of neuropeptide signaling and up-regulated inflammatory mediator expression in the lumbar spinal cord.

3.7. Spinal ERK1/2 activation contributes to post-fracture nociceptive sensitization and up-regulated TNF expression in spinal cord

Western immunoblot was used to evaluate phosphorylation of the mitogen-activated protein kinases (MAPK) extracellular signal-regulated kinase1/2 (ERK) and p38 in the L4,5 spinal cord segments ipsilateral to fracture. At 4 weeks post-fracture in rats there was a 50% increase in phosphorylated ERK, compared to controls (Fig. 7G), but no change was observed in p38 phosphorylation (data not shown). To elucidate the involvement of ERK in post-fracture behavioral changes and spinal inflammatory mediator expression, we intrathecally injected 4 weeks post-fracture rats with PD98059, a selective ERK phosphorylation inhibitor. PD98059 treatment reduced fracture-induced hindpaw allodynia and unweighting (Fig. 7A-B), partially blocked the post-fracture increase in TNF mRNA expression, but had no effects on IL-1, IL-6 and NGF expression at 4 weeks after fracture (Fig. 7C-F). Intrathecal injection of the SP NK1 receptor antagonist LY303870 only partially inhibited fracture-induced phosphorylation of ERK, but the CGRP receptor antagonist CGRP₈₋₃₇ completely reversed ERK phosphorylation (Fig. 7G). To identify the cell types in which ERK phosphorylation occurred, sections of the lumbar cord were immunostained for glial cell markers and phosphorylated ERK. Representative confocal images in Figure 8 demonstrate ERK activation was frequently observed Iba1-positive microglia and occasionally in GFAP-positive astrocytes in rat spinal cord dorsal horn at 4 weeks after fracture. These results suggest that CGRP induced phosphorylation of ERK in spinal microglia and astrocytes can stimulate spinal TNF expression and induce nociceptive sensitization after fracture.

4. Discussion

Neurogenic inflammation is mediated by sensory afferent release of SP and CGRP, neurotransmitters that activate their receptors in the dermal vasculature to induce extravasation and vasodilatation. Electrically evoked extravasation and vasodilatation responses are enhanced in CRPS patients, and when SP is microdialyzed in the skin there is an exaggerated extravasation response [24,44]. Furthermore, serum levels of SP and CGRP are elevated in CRPS patients [5,7,37]. Similarly, after fracture in rats we have observed increased SP and CGRP expression in the sciatic nerve and serum, up-regulated SP NK1 receptor expression in the endothelial cells and keratinocytes of the paw skin, and enhanced SP evoked extravasation and edema responses in the fracture limb [46]. Treating fracture rats with an NK1 receptor antagonist reduced hindpaw allodynia, warmth, and edema, providing evidence that exaggerated SP signaling contributes to the development of CRPS-like changes [13]. We also observed that fracture mice developed hindpaw allodynia, unweighting, warmth, and edema at 3 weeks post-fracture, but that in SP deficient fracture mice allodynia and unweighting were attenuated and there was no warmth and edema [15].

Fracture mice lacking the CGRP RAMP1 receptor had a similar presentation. Collectively, these data support the hypothesis that neuropeptide signaling is amplified in the skin and vasculature of CRPS patients and in the fracture rats and mice, contributing to CRPS-like changes.

At 4 weeks post-fracture the expression of SP and its NK1 receptor were up-regulated in the ipsilateral lumbar spinal cord (Fig. 1). There was also a post-fracture increase in the spinal expression of CGRP, but no change was observed in the expression of the CGRP receptor dimers CALCRL and RAMP1 (Fig. 1). Although the primary sources of dorsal horn SP and CGRP are the small sensory fiber terminals, other potential sources for up-regulated intrinsic spinal neuropeptide mRNA expression are spinal interneurons expressing SP [9] and ventral motor neurons expressing CGRP [12]. Previously we observed a post-fracture increase in SP and CGRP mRNA and protein expression in the ipsilateral L4,5 dorsal root ganglia and sciatic nerve, respectively [46]. Collectively, these data suggest that primary afferent and intrinsic neuropeptide signaling in the spinal cord is enhanced at 4 weeks post-fracture. Similarly, several previous studies using inflammatory and neuropathic pain models have also observed exaggerated spinal neuropeptide signaling [1,22,28]. Intrathecal injection of either a SP or CGRP receptor antagonist partially reversed hindpaw von Frey allodynia and unweighting in the 4 week fracture rats (Fig. 2), but had no effect on the contralateral von Frey thresholds, providing evidence that spinal neuropeptide signaling contributes to post-fracture nociceptive sensitization but not to basal pain thresholds.

When SP or CGRP is dialyzed through normal [48] or CRPS affected skin [24] there is no immediate painful response, leading us to postulate that SP and CGRP act as intermediate mediators in the development of post-traumatic pain. Increased levels of TNF and IL-6 are observed in experimental blister fluid or skin biopsies from the affected limbs, but not the contralateral limbs of CRPS patients [16,18,23] and we recently demonstrated activated keratinocytes expressing TNF and IL-6 in CRPS affected skin [4]. TNF, IL-1, IL-6, and NGF levels are increased in the fracture hindpaw skin and fracture rats treated systemically with the TNF inhibitor etanercept, the IL-1 receptor antagonist anakinra, or the NGF antibody tanezumab had reduced allodynia and hindlimb unweighting, indicating the contribution of cytokine and growth factor signaling in the development of trauma induced chronic pain [27,34,35].

The TNF, IL-1, and NGF inhibitors we tested in the fracture rats are large molecules that don't cross the blood-brain barrier, suggesting that the pronociceptive effects of these inflammatory mediators occur at the nociceptor level in hindpaw skin. Pro-inflammatory cytokines and NGF can immediately evoke spontaneous firing and sensitization in sensory afferents [10,20,31]. Furthermore, we have observed that intraplantar injection of these mediators into normal hindpaw skin rapidly induces nociceptive sensitization and have identified proliferating keratinocytes as the primary cellular source of these mediators in the fracture hindpaw [25]. These convergent data support the premise that post-fracture keratinocyte proliferation and inflammatory mediator expression can induce nociceptive sensitization.

Previously we observed that SP and CGRP applied to keratinocytes in vitro stimulated keratinocyte expression of TNF, IL-1, IL-6, and NGF via cell surface receptor activation [38]. After plantar injection of SP in normal rats we observed a gradual increase in keratinocyte expression of TNF, IL-1, IL-6, and NGF that resolved after 48 hours, temporally correlating with the development and resolution of allodynia [45]. In another study we observed that wildtype fracture mice expressed elevated levels of TNF, IL-1, IL-6, and NGF in the hindpaw skin, but that in SP deficient fracture mice and CGRP RAMP1 receptor deficient fracture mice only IL-6 levels increase in the hindpaw skin [15]. These results demonstrate that neuropeptide signaling directly induces keratinocyte expression of inflammatory mediators in vitro and in vivo.

In the current study, at 4 weeks post-fracture TNF, IL-1, IL-6, CCL2, and NGF expression was elevated in the lumbar spinal cord (Fig.3). These inflammatory mediators are up-regulated in spinal cord tissue in a variety of neuropathic and inflammatory pain models [19,33,42]. Intrathecal injections of specific antagonists or inhibitors for each of these inflammatory mediators partially reduced allodynia and unweighting in the fracture hindpaw (Fig. 4), confirming their association with nociceptive sensitization in the fracture model.

Transgenic fracture mice deficient for SP or the RAMP1 receptor were utilized to examine the relationship between neuropeptide signaling, nociceptive sensitization, and expression of spinal inflammatory mediators. As previously noted, the SP and RAMP1 receptor deficient fracture mice had attenuated hindpaw allodynia and unweighting at 3 weeks post-fracture, compared with wildtype fracture mice (Fig. 5) [15]. Post-fracture increases in the spinal expression of SP and CGRP and of their cognate receptors were reduced in the SP and CGRP receptor deficient mice (Fig. 6A-F). These results suggest that exaggerated post-fracture neuropeptide signaling in the spinal cord can up-regulate intrinsic neuropeptide and neuropeptide receptor expression, further amplifying neuropeptide signaling in the cord. Furthermore, SP and RAMP1 deficient mice had reduced post-fracture increases in spinal TNF, IL-1, IL-6, CCL2, and NGF, indicating a crucial role for neuropeptide signaling in post-fracture spinal cord inflammatory mediator expression (Fig. 6G-K). Additional support for this hypothesis comes from the observation that exaggerated spinal SP signaling contributes to allodynia and elevated spinal TNF levels in rats exhibiting opiate withdrawal allodynia [41].

Spinal ERK phosphorylation was increased in fracture rats, and this increase was inhibited by intrathecal administration of the CGRP receptor antagonist CGRP₈₋₃₇ (Fig. 7G). Intrathecal injection of an ERK phosphorylation inhibitor (PD98058) partially reversed post-fracture pain behavior (Fig. 7A,B) and partially blocked the increase in TNF, but not IL-1, IL-6, or NGF expression (Fig. 7 C-F). Immunostaining for phosphorylated ERK and the microglial marker Iba-1 or the astrocyte marker GFAP in the dorsal horn of the lumbar cord demonstrated extensive ERK phosphorylation in microglia and occasionally in astrocytes (Fig 8). Macrophages and microglia are both immune cells derived from monocytes and it has been reported that the in vitro activation of macrophage NK1 receptors induces phosphorylation ERK and stimulates production of pro-inflammatory mediators such as CCL2 [39].

It has been proposed that intense nociceptive afferent activation triggers neurotransmitter release in the dorsal horn, thus inducing spinal glia to express and release inflammatory mediators [42,43]. Interestingly, the intensity of pain in the first week after wrist fracture is the best predictor of the development of CRPS in the ensuing 4 months [29]. The results of the current study support the hypothesis that intense post-fracture nociceptive signaling induces neuropeptide release in the spinal cord, with resultant amplification of spinal neuropeptide signaling and inflammatory mediator expression. We postulate that inflammatory mediator expression in the spinal cord, acting in concert with the inflammatory mediators released by the cutaneous keratinocytes and mast cells in the injured limb, can initiate the development of regional nociceptive sensitization in the fracture model of CRPS. The spinal cord inflammatory changes observed after fracture may also play a critical role in the maintenance of nociceptive sensitization in chronic CRPS. Over time keratinocyte and mast cell proliferation and inflammatory mediator expression return to normal levels in most CRPS patients [4,49], but in some patients the pain and nociceptive sensitization persist, perhaps due to the residual over-expression of spinal inflammatory mediators [2,3]. Future studies in the fracture model looking at changes over time in the expression of inflammatory mediators in skin and spinal cord may further our understanding of the post-fracture temporal evolution of neuro-keratinocyte and neuro-glial inflammatory signaling and its nociceptive sequelae.

To our knowledge this is the first study to examine spinal cord neuropeptide signaling and inflammatory mediator expression in a CRPS model. The current study has several limitations, including those inherent to using animal models of pain and the translational value of the data acquired. Clinical investigations are required to determine whether up-regulated spinal neuropeptide signaling and inflammatory mediator expression contribute to CRPS. The results of this study do provide preclinical support for CRPS treatments that inhibit spinal nociceptive processing, such as NMDA antagonists, intrathecal drugs, and spinal cord stimulation.

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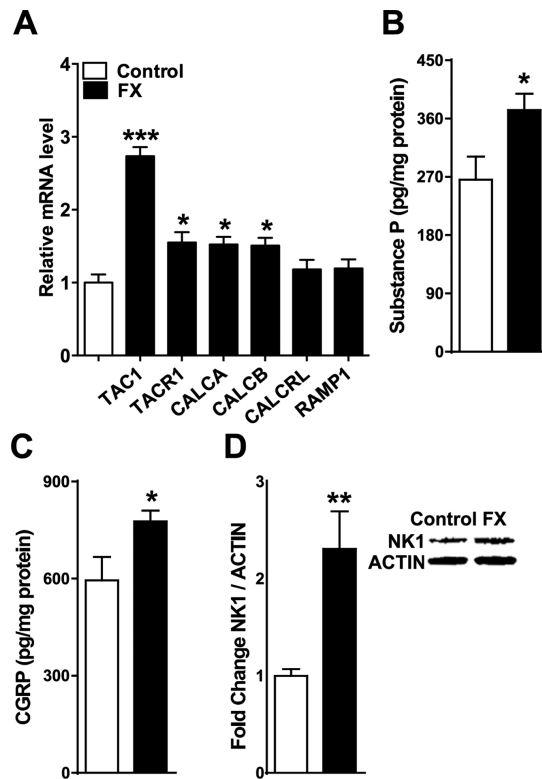


Figure 1. Distal tibia fracture facilitated spinal neuropeptide signaling in the ipsilateral L4,5 spinal cord segments at 4 weeks post-fracture (FX, n = 8), compared to control rats (Controls, n = 8)

(A) Tibia fracture caused increased spinal expression of the substance P (SP) gene (TAC1), the SP neurokinin1 (NK1) receptor gene (TACR1), the α -calcitonin gene-related peptide (α -CGRP) gene (CALCA), and the β -CGRP gene (CALCB). Fracture had no effect on the spinal expression of the CGRP calcitonin receptor-like receptor (CRLR) gene (CALCRL) or on the CGRP receptor activity-modifying protein (RAMP1) gene (RAMP1). Substance P (B) and CGRP (C) spinal protein levels, as measured by enzyme immunoassay, also increased at 4 weeks post-fracture. A western immunoblot assay was used to demonstrate a 2-fold increase in SP NK1 receptor protein levels after fracture (D). Values are means \pm SE. One-way ANOVA ($p < 0.001$) with Bonferroni *post hoc* testing (A) and t-tests (B-D) * $P < 0.05$, ** $P < 0.01$ and *** $P < 0.001$ for FX vs Control.

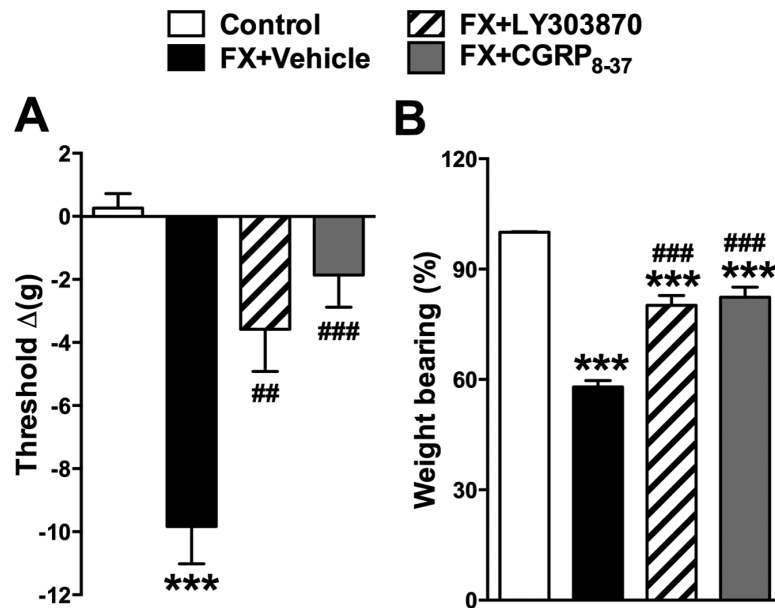


Figure 2. Spinally administered neuropeptide receptor antagonists reduced allodynia and unweighting at 4 weeks post-fracture

Fracture (FX) rats were intrathecally injected with 20 μ g of the SP NK1 receptor antagonist LY303870 or 20 μ g of the CGRP CRLR receptor antagonist CGRP₈₋₃₇. Both LY303870 and CGRP₈₋₃₇ reduced hind paw mechanical allodynia (A) and unweighting (B) at 30 min after injection, compared to FX + Vehicle. Values are means \pm SE. One-way ANOVA ($p < 0.001$) with Bonferroni *post hoc* testing ** $P < 0.01$ and *** $P < 0.001$ for FX + Vehicle (n=8), FX + LY303870 (n=8), or FX + CGRP₈₋₃₇(n=8) vs Control (n=8), ### $p < 0.001$ for FX + LY303870 (n=8), or FX + CGRP₈₋₃₇ (n=8) vs FX + Vehicle (n=8).

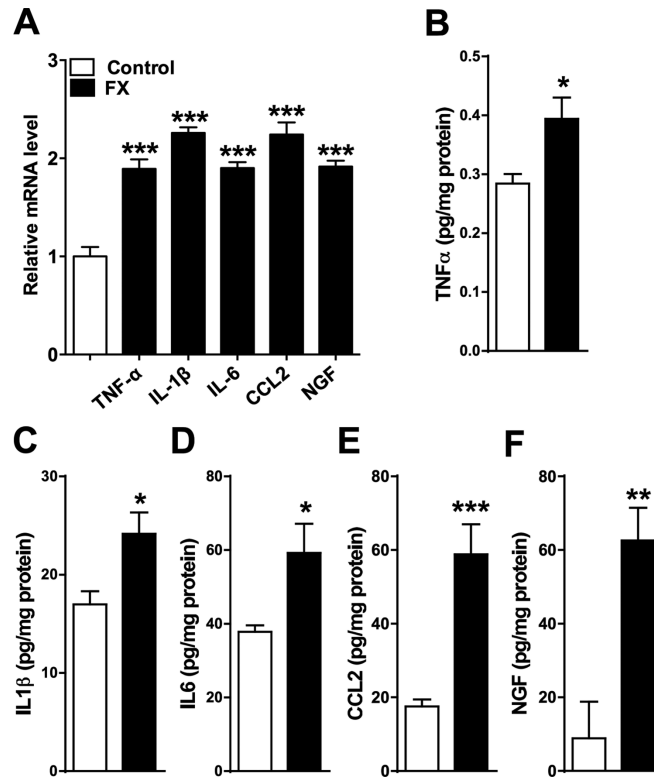


Figure 3. Fracture caused an increase in spinal cord inflammatory mediator expression

Tumor necrosis factor- α (TNF- α), interleukin-1 β (IL-1 β), interleukin-6 (IL-6), chemokine (C-C motif) ligand 2 (CCL2), and nerve growth factor (NGF) gene expression measured by real-time PCR (A) and protein levels determined by enzyme immunoassay (B-F) increased in rat spinal cord post-fracture. Values are means \pm SE. One-way ANOVA ($p < 0.001$) with Bonferroni *post hoc* testing (A) and t-tests (B-F) * $P < 0.05$, ** $P < 0.01$ and *** $P < 0.001$ for FX (n=8) vs Control (n=8).

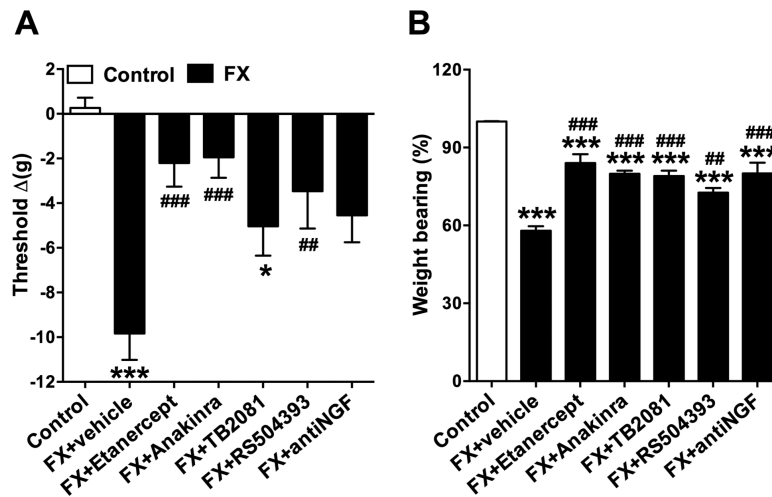


Figure 4. Spinally administered inflammatory mediator receptor antagonists/inhibitors reduced allodynia and unweighting at 4 weeks post-fracture

Fracture (FX) rats were intrathecally injected with either; 1) vehicle, 2) the TNF inhibitor etanercept (100 μ g), 3) the IL-1 receptor antagonist anakinra (100 μ g), 4) the IL-6 receptor antagonist TB-2081 (20 μ g), 5) the chemokine (C-C motif) receptor type 2 (CCR2) antagonist RS504393 (3 μ g), and 6) the anti-NGF antibody muMab 911 (12.4 μ g). Behavioral testing was performed at various time intervals after intrathecal injection based on preliminary pilot time course studies identifying the peak analgesic effects for each drug (etanercept 3 h, anakinra 30 min, TB2081 15 min, RS504393 and muMab 911 1 h). All receptor antagonists/inhibitors reduced post-fracture hind paw allodynia (A) and unweighting (B). There were no effects on hindpaw temperature or thickness. Values are means \pm SE, n=8 per cohort. One-way ANOVA ($p < 0.001$) with Bonferroni *post hoc* testing * $P < 0.05$. ** $P < 0.01$ and *** $P < 0.001$ for FX + vehicle, FX + etanercept, FX + anakinra, FX + TB2081, FX + anti-NGF antibody or FX + RS504394 vs Control, ## $p < 0.01$ and ### $p < 0.001$ for FX + etanercept, FX + anakinra, FX + TB2081, FX + anti-NGF antibody, or FX + RS504394 vs FX + vehicle.

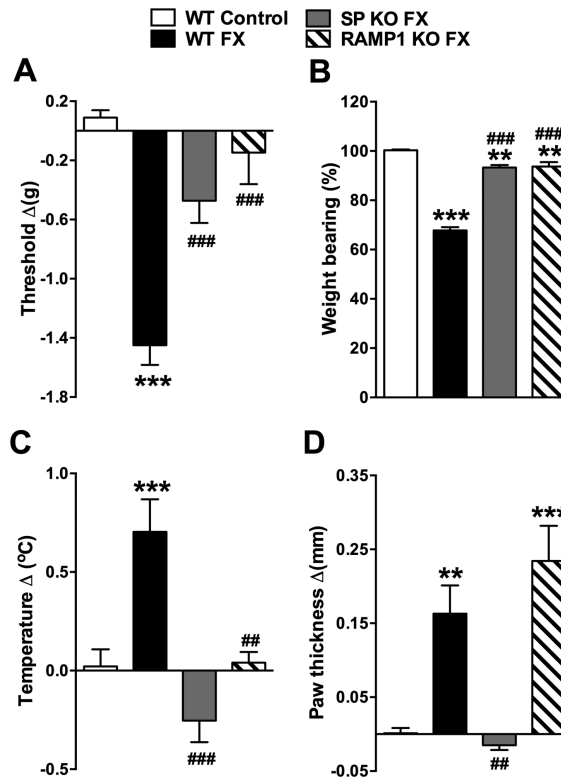


Figure 5. Substance P deficient (SP KO) and CGRP RAMP1 receptor deficient (RAMP1 KO) fracture (FX) mice had reduced hindpaw allodynia, unweighting, warmth, and edema, compared to wildtype (WT) fracture mice

WT fracture mice exhibited hind paw von Frey allodynia (A), unweighting (B), warmth (C) and edema (D) at 3 weeks post-fracture. The SP deficient fracture mice had attenuated hind paw allodynia (A) and unweighting (B), and failed to develop hind paw warmth (C) or edema (D). Similarly, the RAMP1 receptor deficient fracture mice had no von Frey allodynia (A), attenuated unweighting (B), and no warmth (C), but the development of post-fracture hindpaw edema was unaffected by the loss of CGRP signaling (D), compared to WT FX mice. Values are means \pm SE, n=8 per cohort. One-way ANOVA ($p < 0.001$) with Bonferroni *post hoc* testing ** $P < 0.01$ and *** $P < 0.001$ for WT FX, SP KO FX, or RAMP1 KO FX vs WT Control, ## $p < 0.01$ and ### $p < 0.001$ for SP KO FX, or RAMP1 KO FX vs WT FX.

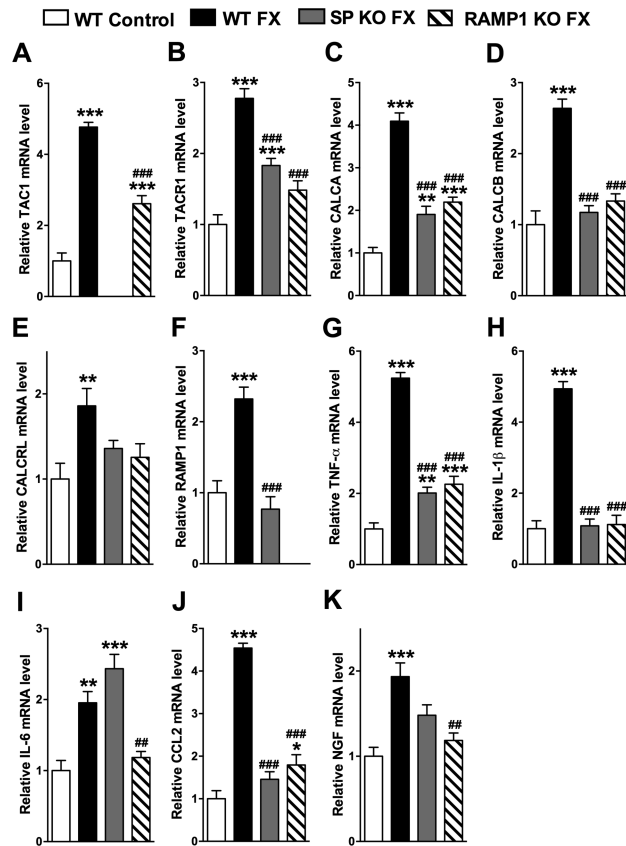


Figure 6. Substance P deficient (SP KO) and CGRP RAMP1 receptor deficient (RAMP1 KO) fracture (FX) mice failed to develop exaggerated spinal neuropeptide signaling and inflammatory mediator expression at 3 weeks post-fracture, measured by real-time PCR

Similar to the changes observed in 4 weeks post-fracture rats (Figs. 1,3), in wild type (WT) fracture (FX) mice there was increased gene expression of (A) SP (TAC1), (B) SP NK1 receptor (TACR1), (C, D) CGRP (CALCA and CALCB), (E) CGRP CRLR receptor (CALCRL), (F) CGRP RAMP1 receptor (RAMP1), (G) TNF- α , (H) IL-1 β , (I) IL-6, (J) CCL2, and (K) NGF, compared to WT mice with no fracture (WT Control). Substance P deficient fracture (SP KO FX) mice had attenuated post-fracture spinal cord changes compared to WT FX mice, and there was no increase in the expression of (D) CALCB, (E) CALCRL, (F) RAMP1, (H) IL1 β , and (K) CCL2 in the SP KO FX mice, compared to WT control mice. CGRP RAMP1 receptor deficient fracture (RAMP1 KO FX) mice also exhibited attenuated post-fracture spinal cord changes, compared to WT FX mice, and there was no increase in (B) TACR1, (D) CALCB, (E) CALCRL, (H) IL-1 β , (I) IL-6, and (J) NGF mRNA levels, compared to WT controls. (A) TAC1 mRNA was not detected in the TAC1 deficient mice and (F) RAMP1 mRNA was not measurable in the RAMP1 receptor deficient mice. Values are means \pm SE, n=8 per cohort. One-way ANOVA ($p < 0.01$ for graph E, $p < 0.001$ for all other graphs), with Bonferroni *post hoc* testing * $P < 0.05$, ** $P < 0.01$ and *** $P < 0.001$ for WT FX, SP KO FX, or RAMP1 KO FX vs WT Control, # $p < 0.05$, ## $p < 0.01$ and ### $p < 0.001$ for SP KO FX, or RAMP1 KO FX vs WT FX.

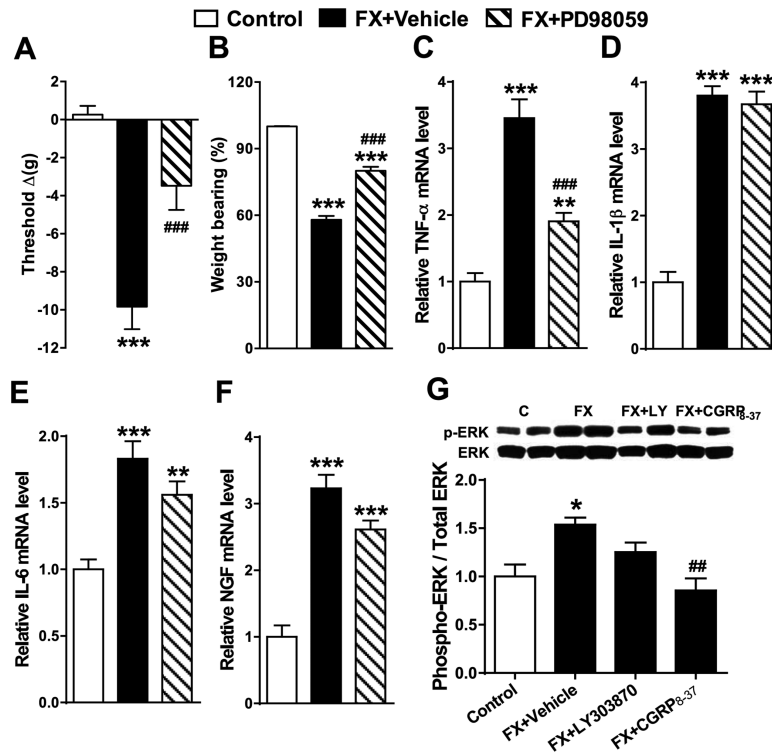


Figure 7. Spinal extracellular signal-regulated kinase 1/2 (ERK1/2) mitogen-activated protein kinase (MAPK) phosphorylation increased after fracture and contributed to upregulated spinal TNF- α expression and hindpaw nociceptive sensitization

At 4 weeks post-fracture rats underwent intrathecal injection with vehicle (FX + Vehicle) or the ERK1/2 MAPK phosphorylation inhibitor PD98059 (20 μ g). Peak analgesic effects for hindpaw von Frey allodynia (A) and unweighting (B) were observed at 1 hour post-injection. PD98059 intrathecal injection also partially inhibited post-fracture increases in spinal TNF- α mRNA levels (C), but had no effect on post-fracture increases in IL-1 β (D), IL-6 (E) and NGF (F), compared to FX + vehicle group. At 4 weeks post-fracture there was an increase ratio of phosphorylated (pERK) to total ERK1/2 (ERK) in the lumbar spinal cord, indicating ERK1/2 activation (G). Intrathecal injection of the SP NK1 antagonist LY303870 (20 μ g) in fracture rats (FX + LY) did not significantly inhibit ERK1/2 phosphorylation, but injection of the CGRP antagonist CGRP₈₋₃₇ (20 μ g) (FX + CGRP₈₋₃₇) completely blocked post-fracture induced phosphorylation of ERK1/2. Values are means \pm SE, n=8 per cohort. One-way ANOVA ($p < 0.001$) with Bonferroni *post hoc* testing * $P < 0.05$, ** $P < 0.01$ and *** $P < 0.001$ for FX + Vehicle, or FX + PD98059 vs Control, ### $p < 0.001$ for FX + PD98059, or FX + CGRP₈₋₃₇ vs WT FX.

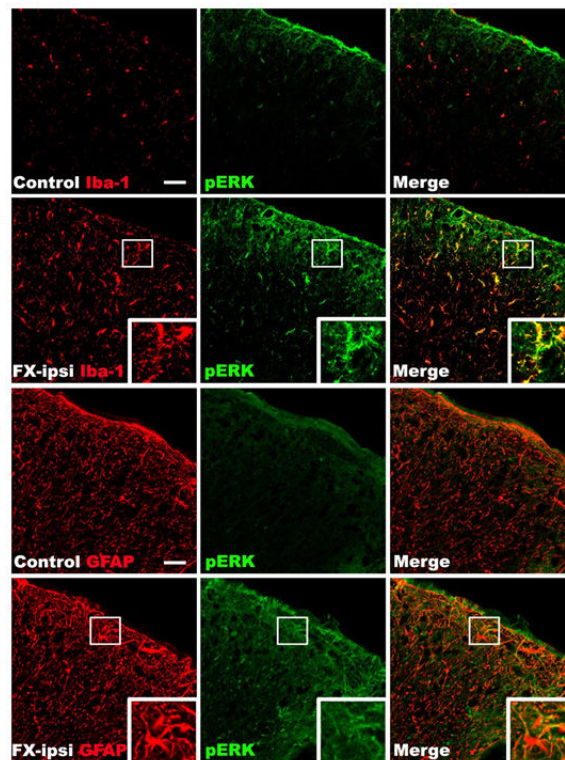


Figure 8. Representative confocal images of co-immunostaining for glial cell markers and phosphorylated ERK1/2 (pERK) in the lumbar spinal cord at 4 weeks after fracture
 Top and third row panels are from a normal control rat, second and fourth row panels are from a fracture rat. Minimal pERK (green) immunostaining was seen in the spinal cord from normal control. In contrast, a substantial increase in pERK immunostaining was observed in the cord at 4 weeks after fracture. Double labeling demonstrates that ERK is activated in the Iba1-positive microglia (red) and GFAP-positive astrocytes (red) in the spinal cord dorsal horn after fracture (n = 4 per cohort). FX: fracture, ipsi: ipsilateral. Scale bar = 50 μ m.

Table 1

Primers used for real-time PCR

Gene	GenBank Accession #	Forward primer	Reverse primer	Product Size(bp)
TAC1	NM_012666	tttcagaggaaatcgggtccaac	ggcattgcctccttgatttggtca	83
TACR1	NM_012667	ctggaagaggagccttgtg	ctgagacggaaaggaacagc	205
CALCA	NM_017338	agaagagatcctgcaactgcca	ggcacaagttgtccttcaccaca	94
CALCB	NM_138513	cccagaagatcctgcaac	agttcctcagaccgaaggt	158
CALCRL	NM_012717	tcattgtggtgctgtgtt	aatggaccatggatgatgt	176
RAMP1	NM_031645	ggcaacaagattggctgtt	aatggggagcacaatgaaag	154
TNF- α	NM_012675	ctcccagaaaagcaagcaac	cgagcaggaatgagaagagg	210
IL-1 β	NM_031512	agtctgcacagttcccac	agacctgactggcagagga	230
IL-6	NM_012589	cacaagtccggagaggagac	acagtcatcatcgctgttc	168
NGF	XM_227525	acctctcggacctctgga	gtcctggctgtgtcttat	168
CCL2	NM_031530	tcccactctgctgcttctta	agcaaaggctgctgctcatagt	86
18S	NR_046237	tcaacttcgatggtagtcgccgt	tccttggatgtgtagccgttct	108