Cytokine mRNA Expression in Painful Radiculopathy

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Abstract: Inflammatory cytokines contribute to lumbar radiculopathy. Regulation of cytokines for transient cervical injuries, with or without longer-lasting inflammation, remains to be defined. The C7 root in the rat underwent compression (10gf), chromic gut suture exposure (chr), or their combination (10gf+chr). Ipsilateral C7 spinal cord and dorsal root ganglia (DRG) were harvested at 1 hour after injury for real-time PCR analysis of IL-1β, IL-6, and TNF-α. Cytokine mRNA increased after all 3 injuries. TNF-α mRNA in the DRG was significantly increased over sham after 10gf+chr (P < .026). Spinal IL-1β was significantly increased over sham after 10gf and 10gf+chr (P < .024); IL-6 was significantly increased after 10gf+chr (P < .024). In separate studies, the soluble TNF-α receptor was administered at injury and again at 6 hours in all injury paradigms. Allodynia was assessed and tissue samples were harvested for cytokine PCR. Allodynia significantly decreased with receptor administration for 10gf and 10gf+chr (P < .005). Treatment also significantly decreased IL-1β and TNF-α mRNA in the DRG for 10gf+chr (P < .028) at day 1. Results indicate an acute, robust cytokine response in cervical nerve root injury with varying patterns, dependent on injury type, and that early increases in TNF-α mRNA in the DRG may drive pain-related signaling for transient cervical injuries.

Perspective: Inflammatory cytokine mRNA in the DRG and spinal cord are defined after painful cervical nerve root injury. Studies describe a role for TNF-α in mediating behavioral sensitivity and inflammatory cytokines in transient painful radiculopathy. Results outline an early response of inflammatory cytokine upregulation in cervical pain.

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Key words: Cytokine, pain, radiculopathy, mRNA, PCR.

Nearly one-third of Americans suffer from pain, with as many as 13% of them experiencing neck pain.13,23,56 Painful cervical radiculopathy can result from nerve root compression by disc herniation or transient loading by the vertebrae.3,17,21,41,64 Inflammation and proinflammatory mediators contribute to painful lumbar radiculopathy.8,22,28,36,37 Early cytokine production and release have been studied extensively for sustained nerve root compression, cytokine profiles in the dorsal root ganglion (DRG) and spinal cord for transient painful cervical nerve root compression have not been defined.

An inflammatory response is produced in the nervous system after radiculopathy, including infiltration and activation of immune cells, cytokine upregulation, and growth factor release.12,36,49 Spinal inflammatory cytokines also contribute to neuropathic pain.11,14,15,22,30,48,49,58 Cytokines, including TNF-α, IL-1β, IL-6, and IL-10, are synthesized and released by astrocytes, microglia and macrophages and act throughout the nervous system in both paracrine and autocrine signaling. Cytokine production can indirectly induce the expression of a variety of pain mediators and directly activate glial cells in the central nervous system (CNS), leading to spinal sensitization.2,12,31,45,52 Spinal glial activation is produced in lumbar radiculopathy and exhibits a relationship with behavioral hypersensitivity.5,6,10,22,36,39,60,62 Recent work from our lab demonstrates that mechanical allodynia and spinal glial activation are similarly produced after a transient nerve root compression in the cervical spine and both responses are sensitive to the severity of compression and the presence of chemical irritation.26,28,46,47 Although relationships among injury parameters, mechanical allodynia, and spinal glial activation have been described for cervical radiculopathy, it is not known whether painful cervical in-
juries also modulate inflammatory cytokine expression in the DRG and/or spinal cord, and if these responses are specific to the type of injury.

Cytokines are differentially regulated in the DRG and spinal cord in radiculopathy. Inflammatory cytokine genes including TNF-\(\alpha\), IL-1\(\beta\), and IL-6 increase in the DRG after root injury, whereas mRNA for these cytokines increases in both the DRG and spinal cord, it peaks earlier (day 1) in the DRG. Spinal TNF-\(\alpha\) mRNA and protein also increase immediately after painful trauma to the nerve roots. TFN-\(\alpha\) provides positive-feedback for its own production, and initiates the production and release of other cytokines. Despite the known early upregulation of TNF-\(\alpha\) in pain and its direct involvement in the cytokine cascade, it is not known if, and how, TNF-\(\alpha\) mRNA is modulated immediately after insult to the cervical nerve root and whether inflammatory cytokine upregulation in the DRG and spinal cord are mediated by TNF-\(\alpha\).

Although many studies have quantified inflammation in the DRG and spinal cord in neuropathic pain, there is no characterization for cervical radiculopathy. Similarly, it remains to be seen if proinflammatory cytokines are increased after transient compression of the nerve root and whether these cytokines are influenced by a chemical insult. A goal of this study is to quantify changes in inflammatory cytokine mRNA in the DRG and spinal cord after 2 cervical nerve root compression injuries. A separate investigation tests whether allodynia produced by these injuries can be ameliorated by immediate antagonism of TNF-\(\alpha\) and if it also modulates cytokine mRNA.

Methods

Two studies were performed to define cytokine mRNA expression immediately after painful insults to the cervical nerve root and to utilize a cytokine antagonist to evaluate if TNF-\(\alpha\) modulates behavioral hypersensitivity. In the first study, cytokine (IL-1\(\beta\), IL-6, and TNF-\(\alpha\)) mRNA in the DRG and spinal cord was characterized at an early time point (1 hour) after cervical nerve root injury; either applied compression, chronic gut suture exposure, or their combination, in separate investigations. Based on those findings, a second study was performed using the TNF receptor-1 to neutralize TNF-\(\alpha\) and evaluate its role in modulating allodynia in those 3 models. Experiments were performed using male Holtzman rats (Harlan Sprague-Dawley, Indianapolis, IN), weighing 250 to 350 grams at the start of the study. All experimental procedures were approved by the University of Pennsylvania Institutional Animal Care and Use Committee and carried out according to the guidelines of the Committee for Research and Ethical Issues of the IASP. Animals were housed with a 12-12 hour light-dark cycle and free access to food and water.

Cytokine Characterization Study

Rats underwent 1 of 4 previously reported surgical procedures: (1) 10gf compression of the right C7 dorsal nerve root proximal to the DRG for 15 minutes (10gf), (2) exposure of the nerve root to 4 pieces of 3-0 chromic gut suture material (chr), (3) 10gf compression of the dorsal nerve root in addition to placement of 4 pieces of 3-0 chromic gut suture (10gf+chr), or (4) exposure of the nerve root with no additional insult (sham). At 1 hour after the surgical procedure, the C7 spinal cord on the side ipsilateral to the compressed nerve root was harvested (n = 6, 10gf; n = 6, chr; n = 7, 10gf+chr) for analysis of cytokine mRNA (IL-1\(\beta\), IL-6, and TNF-\(\alpha\)) using real-time reverse transcriptase polymerase chain reaction (RT-PCR). For some rats, the C7 ipsilateral and contralateral DRGs (n = 6 each group) were also harvested. Tissue from normal, unoperated rats (n = 5), and from rats receiving sham surgery (n = 5 spinal cord samples; n = 6 DRG samples) were also used for comparison for both DRG and spinal cord assays.

TNF-\(\alpha\) Receptor Treatment Study

In a separate study, soluble TNF- receptor 1 (sTNF-R1-PEG; gift from Amgen, Inc., Thousand Oaks, CA) was administered at the site of the compressed nerve root in the 10gf, chr, and 10gf+chr injury models. For this study, 200 \(\mu\)g sTNF-R1 in 40 \(\mu\)L was administered directly to the injured nerve root at the time of injury, according to other neuropathic pain studies. In addition, at 6 hours after the initial surgery, the C7 nerve root was again exposed and a second, identical dose of sTNF-R1 was administered directly to the injured nerve root. Spinal cord and DRG samples were harvested at day 1 after either surgery alone (10gf, chr, 10gf+chr, sham) or surgery with sTNF-R1 treatment (10gf+sTNF-R1, chr+sTNF-R1, 10gf+chr+sTNF-R1). Each rat was tested for bilateral mechanical allodynia at day 1 after injury to assess behavioral hypersensitivity. In this study, C7 spinal cord on the side ipsilateral to the nerve root injury was harvested after behavior testing was performed (n = 4, sham; n = 4, 10gf; n = 5, 10gf+sTNF-R1; n = 6, chr; n = 6, chr+sTNF-R1; n = 4, 10gf+chr; n = 4, 10gf+chr+sTNF-R1). In addition, for a subset of rats, the ipsilateral and contralateral C7 DRGs were harvested at day 1 from each group (n = 4, sham; n = 3, 10gf; n = 5, 10gf+sTNF-R1; n = 6, chr; n = 6, chr+sTNF-R1; n = 3, 10gf+chr; n = 4, 10gf+chr+sTNF-R1). DRG and spinal cord tissue from normal rats (n = 5) were used as controls for the cytokines probed (IL-1\(\beta\), IL-6, and TNF-\(\alpha\)).

Specific Methods

Surgical Procedures

All procedures were performed under inhalation anesthesia (4% halothane for induction, 2% for maintenance). Surgical procedures have been detailed previously. Briefly, rats were placed in a prone position, and an incision was made in the skin from the base of the skull to the second thoracic vertebra. A C6/C7 hemilaminectomy and partial facetectomy were performed on the right side to expose the C7 dorsal root. Procedures for nerve root compression (10gf) involved transient compression of the C7 dorsal nerve root for 15 minutes using a 10gf microvascular clip (World Precision Instruments,
Pellet was centrifuged at 12,000 rpm for 10 minutes at room temperature. The pellets were washed with 75% ethanol, and then dissolved in RNAse free water and stored at 80°C until use.

For reverse transcription, 1 to 2 μg of total RNA was treated with 1 unit of amplification grade DNase I (Invitrogen, Carlsbad, CA). The DNase I treated RNA was subject to reverse transcription according to the manufacturer’s instructions using SuperScript III reverse transcriptase (Invitrogen), random hexamer primer (Invitrogen) and RNase inhibitor RNaseOUT (Invitrogen) at 50°C for 1 hour. Single-strand synthesized cDNA was used for real-time PCR. Specific primer sequences are listed in Table 1. Sybr-green real-time PCR was performed in an ABI-7300 system (Applied Biosystems, Foster City, CA). Each reaction contained a total of 25 μL of Sybr-green master reagent (Applied Biosystems), cDNA, and 900nM of each primer. The PCR conditions were as follows: 50°C for 2 minutes, 95°C for 10 minutes, followed by 40 cycles of 95°C for 15 seconds and 60°C for 60 seconds. A no cDNA template control was included in each run. All samples and no-template controls were assayed in duplicate. The target gene expression was normalized against the cyclophilin-A gene. Samples were normalized to results from normal tissue. The relative target gene expression was analyzed using the ΔΔCt method.

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### Cytokine Characterization Study

For the cytokine characterization study, a one-way ANOVA was performed to compare differences in mRNA between groups (10gf, chr, 10gf+chr, sham, normal) for each cytokine (IL-1β, IL-6, and TNF-α) for the spinal cord and DRG samples, separately. For the sTNF-R1 treatment study, a one-way ANOVA tested for differences in mechanical allodynia between sham, treated and untreated groups for each injury paradigm. t tests were used to detect differences in cytokine mRNA between injury and treatment (10gf, 10gf+sTNF-R1), (chr, chr+sTNF-R1) and (10gf+chr, 10gf+chr+sTNF-R1), with separate tests for the spinal cord and DRG. For all tests, significance was defined at P < .05 (SYSTAT, Richmond, CA).

## Results

### Cytokine Characterization Study

All proinflammatory cytokines probed in this study were detected in the ipsilateral DRG of all groups at 1 hour. Sham procedures increased all 3 cytokines in the ipsilateral DRG over normal levels, but none of these changes were significant. IL-1β mRNA was significantly increased over normal levels by the 10gf and 10gf+chr treatments and was highest in the 10gf+chr+sTNF-R1 group.

<table>
<thead>
<tr>
<th>Cytokine</th>
<th>Gene Symbol</th>
<th>Primer Sequence</th>
<th>GENE PRIMER SEQUENCE</th>
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<tr>
<td>IL-1β</td>
<td>forward 5’-CACCTCTCAAGCAAGCAGC-3’</td>
<td>reverse 5’-GGTTCCATGGTAGTCAAC-3’</td>
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</tr>
<tr>
<td>IL-6</td>
<td>forward 5’-AGTCGGAGGCTTAATTACATATGGTC-3’</td>
<td>reverse 5’-TGCATGCAACACTCTTCTTCT-3’</td>
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<tr>
<td>TNFα</td>
<td>forward 5’-ATCATCTTCTCAAAAACTCGAGTACAA-3’</td>
<td>reverse 5’-CTGCTCTTGTCTGTGTT-3’</td>
<td></td>
</tr>
<tr>
<td>Cyclophilin A</td>
<td>forward 5’-TATCGCAGCTCAGAGCTGAGTG-3’</td>
<td>reverse 5’-CTTCTCTGGTTGCTTCCATTCC-3’</td>
<td></td>
</tr>
</tbody>
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For tissue harvest, rats were deeply anesthetized with sodium pentobarbital (65 mg/kg) followed by transcardiac perfusion with 200 mL of phosphate buffered saline (PBS). After perfusion, the ipsilateral and contralateral C7 DRGs were identified, harvested and immediately frozen. The cervical spinal cord was exposed by laminectomy, the C7 segment of the cervical spinal cord was harvested, separated into ipsilateral and contralateral sections, and frozen immediately on dry ice. Total RNA was then isolated using Trizol reagent (Life Technologies). For reverse transcription, 1 to 2 μg of total RNA was treated with 1 unit of amplification grade DNase I (Invitrogen, Carlsbad, CA). The DNase I treated total RNA was subject to reverse transcription according to the manufacturer’s instructions using SuperScript III reverse transcriptase (Invitrogen), random hexamer primer (Invitrogen) and RNase inhibitor RNaseOUT (Invitrogen) at 50°C for 1 hour. Single-strand synthesized cDNA was used for real-time PCR. Specific primer sequences are listed in Table 1. Sybr-green real-time PCR was performed in an ABI-7300 system (Applied Biosystems, Foster City, CA). Each reaction contained a total of 25 μL of Sybr-green master reagent (Applied Biosystems), cDNA, and 900nM of each primer. The PCR conditions were as follows: 50°C for 2 minutes, 95°C for 10 minutes, followed by 40 cycles of 95°C for 15 seconds and 60°C for 60 seconds. A no cDNA template control was included in each run. All samples and no-template controls were assayed in duplicate. The target gene expression was normalized against the cyclophilin-A gene. Samples were normalized to results from normal tissue. The relative target gene expression was analyzed using the ΔΔCt method.

**Table 1. Specific Primer Sequences**

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<tr>
<td>TNFα</td>
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<td>reverse 5’-CTGCTCTTGTCTGTGTT-3’</td>
</tr>
<tr>
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<td>reverse 5’-CTTCTCTGGTTGCTTCCATTCC-3’</td>
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### Behavioral Assessment

Each rat was evaluated for bilateral forepaw mechanical allodynia postoperatively until the time of tissue harvest, as previously described. Before surgery, rats were acclimated to the tester and environment. Allodynia was measured for each unoperated rat before the surgical procedure to provide an assessment of baseline and unoperated control values. A single tester performed all testing and was blinded to the surgical procedures. For each testing session, after 20 minutes of acclimation, rats were stimulated on the plantar surface of each forepaw using 3 von Frey filaments (1.4, 2, and 4g; Stoelting, Wood Dale, IL). Each testing session had 3 rounds of 10 stimulations, separated by 10 minutes. Total withdrawals were recorded for each forepaw of each rat and averaged for each group.

### Tissue Processing of Cytokine mRNA

For tissue processing, rats were deeply anesthetized with sodium pentobarbital (65 mg/kg) followed by transcardiac perfusion with 200 mL of phosphate buffered saline (PBS). After perfusion, the ipsilateral and contralateral C7 DRGs were identified, harvested and immediately frozen. The cervical spinal cord was exposed by laminectomy, the C7 segment of the cervical spinal cord was harvested, separated into ipsilateral and contralateral sections, and frozen immediately on dry ice. Total RNA was then isolated using Trizol reagent (Life Technologies, Rockville, MD) according to the manufacturer’s instructions. Bilateral DRGs were assayed as well as the ipsilateral spinal cord sample. Tissue was homogenized in 1 mL Trizol reagent (Life Technologies) using a pellet pestle motor. After phase separation with 1-bromo-3-chloropropane, the lysate was centrifuged at 12,000g for 15 minutes at 4°C. RNA was precipitated with isopropanol for 10 minutes at room temperature. The pellet was centrifuged at 12,000g for 10 minutes at 4°C and washed with 75% ethanol, and then dissolved in RNAse free water and stored at -80°C until use.

RNA concentration and quality were measured by a Nanodrop spectrometer (Nanodrop Technologies, Wilmington, DE). For reverse transcription, 1 to 2 μg of total RNA was treated with 1 unit of amplification grade DNase I (Invitrogen, Carlsbad, CA). The DNase I treated total RNA was subject to reverse transcription according to the manufacturer’s instructions using SuperScript III reverse transcriptase (Invitrogen), random hexamer primer (Invitrogen) and RNase inhibitor RNaseOUT (Invitrogen) at 50°C for 1 hour. Single-strand synthesized cDNA was used for real-time PCR. Specific primer sequences are listed in Table 1. Sybr-green real-time PCR was performed in an ABI-7300 system (Applied Biosystems, Foster City, CA). Each reaction contained a total of 25 μL of Sybr-green master reagent (Applied Biosystems), cDNA, and 900nM of each primer. The PCR conditions were as follows: 50°C for 2 minutes, 95°C for 10 minutes, followed by 40 cycles of 95°C for 15 seconds and 60°C for 60 seconds. A no cDNA template control was included in each run. All samples and no-template controls were assayed in duplicate. The target gene expression was normalized against the cyclophilin-A gene. Samples were normalized to results from normal tissue. The relative target gene expression was analyzed using the ΔΔCt method.
increased compared with normal for all injury groups (P < .003) (Fig 1A). All 3 types of injuries nearly doubled IL-6 mRNA in the DRG relative to sham, but only 10gf displayed significant increases over normal (P = .005) (Fig 1B). Last, TNF-α mRNA expression in the ipsilateral DRG was only increased after injury (Fig 1C). This increase was significant compared with normal for all types of nerve root insult (P < .042). IL-1β mRNA were both significantly increased compared with normal (P < .024). This increase was also significant compared with sham for TNF-α after 10gf+chr (***P < .026). Error bars indicate ±SD.

As with the DRG response, sham increased spinal cytokine mRNA at 1 hour for all cytokines probed, but not significantly over normal (Fig 2). IL-1β was significantly elevated over sham for the 10gf and 10gf+chr groups (P < .024) and over normal for all 3 injuries (P < .005) (Fig 2A). The trend was similar for IL-6 with all 3 injuries producing elevated mRNA over sham and normal; yet increases over sham and normal were only significant for 10gf (P < .024) (Fig 2B). Spinal TNF-α mRNA levels were significantly greater than normal after both 10gf and 10gf+chr (P < .019), but the increase was not significant for chr (Fig 2C).

**TNF-α Receptor Treatment Study**

Mechanical allodynia was produced in the forepaws after all injuries (Fig 3). However, although ipsilateral allodynia was increased over sham after chronic exposure, this increase was only significant in the ipsilateral forepaw for testing with the 4g von Frey filament (P = .046) (Fig 3). In addition, no significant decreases were measured for treatment with sTNF-R1-PEG in the chr injury paradigm (Fig 3). Ipsilateral allodynia at day 1 after 10gf was significantly greater than sham for stimulation using all von Frey filaments (P < .007) (Fig 3). In general, treatment with the sTNF-R1 attenuated allodynia for the compression alone insult (10gf), but this was only significant for testing using the 1.4g von Frey filament (P < .005) (Fig 3). Administration of the sTNF-R1-PEG signifi-
Cytokine mRNA Expression in Painful Radiculopathy

Figure 3. Average mechanical allodynia in the ipsilateral and contralateral forepaws at day 1 for sham, 10gf, 10gf+sTNF-R1, chr, chr+sTNF-R1, 10gf+chr, or 10gf+chr+sTNF-R1. Allodynia is measured by the number of paw withdrawals for stimulation with each von Frey filament: (A) 1.4g, (B) 2g, and (C) 4g. *Significant difference from sham values. Treatment with sTNF-R1 significantly decreased allodynia compared with nontreatment responses for 10gf and 10gf+chr for testing with the 1.4g filament. This trend was conserved across filaments for 10gf+chr injury (A, B, C). Error bars indicate ±SD.

Figure 4. Quantification of cytokine gene expression in the ipsilateral DRG at 1 day after injury either with or without sTNF-R1 treatment: IL-1β (A), IL-6 (B), and TNF-α (C). Treatment with sTNF-R1 significantly decreased IL-1β and TNF-α mRNA compared with their matched untreated groups (P < .028). Error bars indicate ±SD.

Discussion

A transient nerve root compression (10gf) alone is sufficient to produce immediate (1 hour) significant eleva-
In the current study, only a slight increase in cytokine mRNA was noted after chr injury alone. This is in contrast to previous studies that show cytokine protein and mRNA increases in models of inflammatory injury to the DRG or nerve root. However, those models of inflammatory pain utilize compounds such as Complete Freund’s adjuvant or zymosan which exhibit immediate effects, whereas chronic gut suture resorbs slower and over a longer timeframe. It is likely that the 1 hour and 1 day time points studied here may be too early to detect the inflammatory effects produced from the chronic gut suture alone. Indeed, previously published work from our lab indicates that astrocytic activation is present by day 7 but not day 1 after chronic exposure, further implying that inflammation from chronic gut sutures increases over time.

Although allodynia can persist for 14 days after 10gf or 10gf+chr, it is necessary to probe early cytokine responses (both mRNA and protein) to fully define the temporal and spatial relationship of these cytokines, as they relate to pain initiation and maintenance in cervical radiculopathy. For example, mRNA levels in the spinal cord for all of these cytokines exhibit positive, significant correlations with imposed injury strain for ligation with silk suture, indicating their spinal responses are sensitive to previous studies that show cytokine protein and mRNA increases in models of inflammatory injury to the DRG or nerve root. However, those models of inflammatory pain utilize compounds such as Complete Freund’s adjuvant or zymosan which exhibit immediate effects, whereas chronic gut suture resorbs slower and over a longer timeframe. It is likely that the 1 hour and 1 day time points studied here may be too early to detect the inflammatory effects produced from the chronic gut suture alone. Indeed, previously published work from our lab indicates that astrocytic activation is present by day 7 but not day 1 after chronic exposure, further implying that inflammation from chronic gut sutures increases over time.

Although many cytokines have been investigated in several rat models of neuropathy, their interactions and signaling in pain are still poorly defined, especially for transient injuries. The present findings suggest that early upregulation of TNF-α transcription in the DRG, together with upregulation of genes for TNF-α, IL-1β, and IL-6 in the spinal cord, may trigger downstream pain-related signaling cascades. Indeed, although no significant differences were noted between mRNA levels of different injury groups, the proinflammatory cytokine responses were most robust after the combination of both compression and application of inflammatory material (10gf+chr) (Figs 4 and 5). This is consistent with other reports that this combined insult model has both greater allodynia and more robust spinal glial activation than that produced for compression alone.

For example, after either 10gf or 10gf+chr, TNF-α is increased in the DRG at 1 day; however for IL-1β this increase is present after 10gf+chr only (Figs 1 and 4). Although it appears that the increase in IL-1β in the DRG after 10gf+chr is due to TNF-α signaling (Fig 4A), this corresponding decrease is not noted for 10gf alone, implying varying responses to pharmacologic treatment for different injury paradigms.

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Tnf-α in neuropathic pain, based on a transgenic mouse model of conditional expression of Tnf-α in astrocytes under the GFAP promoter. Results from our lab also show that spinal glial activation and mechanical allodynia are increased when mechanical and chemical factors are combined at injury. Taken with results presented here, this could indicate a specific role for glial cells in upregulating cytokine mRNA after mechanical and chemical nerve root injury.

Only Tnf-α mRNA was significantly increased in the ipsilateral DRG at 1 hour after any of the injuries studied. This may suggest that the insults of mechanical injury and chromic gut suture elicit an early response of Tnf-α in the DRG as compared with IL-1β and IL-6. On the other hand, mRNA results in the ipsilateral spinal cord demonstrate different cytokines were increased significantly at this same time point in either 10gf or 10gf + chr group, suggesting that the type of injury may encode a specific profile for gene expression of proinflammatory cytokines in the spinal cord. Other aspects of the inflammatory response, such as glial activation and macrophage infiltration, are also variably activated for different types of injuries. After spinal nerve injury, macrophages infiltrate the endoneurial blood vessels adjacent to the DRG, and can produce and release Tnf-α and other proinflammatory cytokines. However, data from our lab demonstrate macrophage infiltration in the dorsal root and dorsal root ganglia at day 7 but not day 1 after injury. Further, macrophage infiltration is common in other models which demonstrate axonal degeneration and release of inflammatory cytokines. Tnf-α and IL-1β mRNA induction have been reported to peak earlier (at day 3) than IL-6 mRNA (at day 7) in the DRG and spinal cord after a painful chronic nerve constriction model, implying that Tnf-α and IL-1β have roles in initiating pain, while IL-6 may be responsible for its maintenance.

Studies quantifying cytokine protein and release after injury are important for defining a role for inflammatory cytokines in mediating nerve root pain. Assays at later time points would provide further insight into mechanisms of persistent pain in this transient injury. In addition, assays probing dorsal and ventral horn inflammation separately would offer information on regional patterns of cytokine changes. Further, defining the mechanical allodynia responses at the 1 hour time point would be valuable to place the cytokine mRNA changes in the context of behavioral hypersensitivity. Yet, in the current study such testing was not possible owing to the need to preserve the time period of 1 hour for tissue harvest after injury. Although mechanical allodynia was the only behavioral outcome probed for this injury model it provides a clinically relevant measure of behavioral hypersensitivity. Sustained sensitivity in the forepaw has been previously reported for this nerve root compression model; patients with cervical nerve injury can present with increased sensitivity to touch and heat stimuli extending from the spine into the forearms and fingers in some cases.

The current study indicates that mechanical compression of the cervical nerve root, both with and without chemical irritation, initiates very early increases in local
and central cytokine mRNA. Administration of sTNF-R1 at the site and time of injury decreased cytokine mRNA and attenuated mechanical allodynia. These findings suggest that TNF-α signaling mediates both downstream cytokine increases and behavioral hypersensitivity in cervical radiculopathy. Although inflammatory cytokine mRNA at day 1 was not decreased after 10gf+sTNF-R1, this could be due to the fact that these cytokines demonstrate only a transient increase in that model. Conversely, after 10gf+chr, inflammatory cytokine mRNA remains elevated at day 1. It may be possible that the inflammatory component of the chromic gut material introduced in the 10gf+chr injury dominates over the mechanical one present in either injury and that the sTNF-R1 was more active against the elements of the nociceptive cascade which are inflammatory-based than those from the mechanical component. Further studies outlining changes in cytokine protein and their specific cellular sources are required to further elucidate inflammatory signaling in cervical radiculopathy. Nonetheless, results presented here demonstrate both local and central inflammation after nerve root injury and provide evidence for a role for TNF-α in cervical radiculopathy.

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References

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