

IN VITRO INVESTIGATION OF CYTOKINE-MEDIATED NUCLEUS PULPOSUS DEGENERATION

NSF Summer Undergraduate Fellowship in Sensor Technologies

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ABSTRACT

Degeneration of the lumbar intervertebral discs is strongly implicated as a cause of low back pain, and may also lead to impaired mobility. A lack of understanding of the pathomechanisms that underlie degeneration limits our ability to develop biological treatments that both alleviate painful symptoms and restore function. The process of degeneration is characterized by up-regulation of pro-inflammatory cytokines—particularly interleukin 1 beta (IL1 β) and tumor necrosis factor alpha (TNF α)—within the central nucleus pulposus. Furthermore, the increased production of these cytokines is not matched by increasing amounts of their inhibitory regulators, resulting in an imbalance of catabolic and anabolic activity. In this study, we developed an in-vitro model of the nucleus pulposus that was used to investigate the effects of IL1 β and TNF α on composition and mechanical function. In addition, we examined the capacity of IL1 receptor antagonist (IL1ra) and soluble TNF receptor 1 (sTNFR1), inhibitors of IL1 β and TNF α respectively, to mitigate cytokine-mediated functional and compositional changes. Our results demonstrated that short-term exposure to IL1 β , but not TNF α , causes loss of matrix components that significantly compromises mechanical function, suggesting that IL1 β plays a more direct role than TNF α in driving matrix degradation in the nucleus pulposus. Our results also demonstrated that IL1ra can effectively prevent compositional and functional changes induced by IL1 β , highlighting its therapeutic potential.

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1. INTRODUCTION

Disorders of the lower back impact our society, both physically through individual impairment and financially through medical expenses. The most common cause of musculoskeletal impairment is low back pain. The prevalence of low back pain among the general U.S. population is very high, reaching 25 percent [1]. From a financial viewpoint, back disorders result in lost productivity and increased health care costs. Since low back pain is a common disorder negatively affecting society, much medical research has focused on the causes and treatment of low back pain.

Intervertebral discs (IVDs) are a critical component of the spine. These discs are pads of fibrocartilage that both transfer and distribute compressive loads between vertebral bodies, and permit spinal movement. Intervertebral disc degeneration is strongly implicated as a cause of low back pain [4-7]. Current treatments for painful disc degeneration, such as spinal fusion, are aimed at allaying painful symptoms, without restoring function. The limit of these treatment options is in part due to an incomplete understanding of the biological mechanisms involved. The central nucleus pulposus (NP) is crucial to disc function and health, and is the focus of this research [8,9]. Our objective was to investigate the initiation and progression of IVD degeneration and potential therapies using a new in vitro NP model. A complete understanding of the fundamental mechanisms associated with disc degeneration is necessary for developing novel treatments of low back pain.

2. BACKGROUND

2.2 Intervertebral Discs

2.2.1 Disc Anatomy

The spine consists of vertebral bones interconnected with intervertebral discs. The discs are soft tissue structures that provide flexibility and integrity to the spine. IVDs both transfer and distribute compressive loads between vertebrae, and permit spinal movement. Proper function is based on IVD structure. Three regions comprise each disc: an outer annulus fibrosus (AF); superior and inferior cartilaginous end plates; and the central NP [8, 9] (Figure 1).

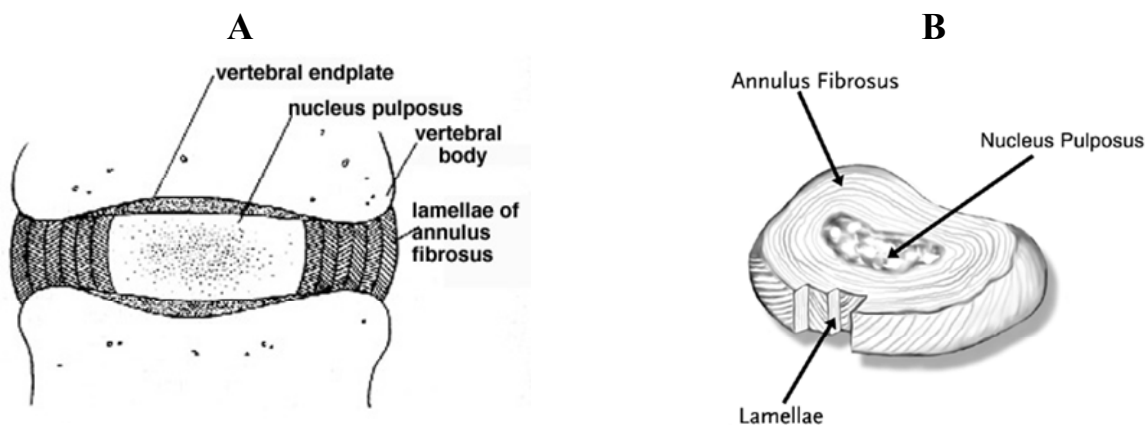


Figure 1. The structure of the intervertebral disc: (A) sagittal and (B) transverse sections. (Sources: http://www.medscape.com/viewarticle/405642_2 and <http://www.chiropractic-help.com/L4-Lumbar-Spine.html>)

2.2.2 Nucleus Pulposus

The NP is a pressurized gel of randomly distributed collagen II fibrils in a hydrated extrafibrillar matrix rich in proteoglycans [10, 11] (Figure 2). In compression under spinal axial loads, the NP is confined peripherally by the AF, which generates a region of uniform hydrostatic pressure in the disc. The NP pressure further enables the even distribution of the axial compressive loads between vertebral bodies [8, 9].

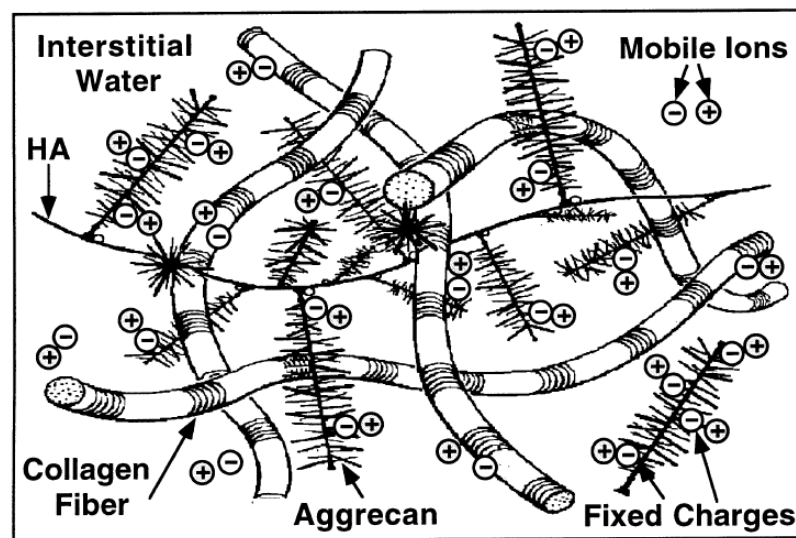


Figure 2. This NP matrix schematic shows collagen fibers interwoven with proteoglycans. GAG are the side chains found on proteoglycans. The negative charges are associated with GAG and the positive charges are the sodium ions in water. (Source: Mow et al, 1998, Int. J. Solids Structures)

2.3 Human Disc Degeneration

2.3.1 Alterations in NP Structure, Composition, and Mechanics

Human disc degeneration starts in the NP with associated alterations in structure, composition, and mechanics. One of the earliest indicators of disc degeneration is loss of glycosaminoglycan (GAG) content in the NP, followed by water loss [14-16]. Changes in matrix composition impair mechanical function. Such comprised functional changes result in reduction in NP pressure and alterations in motion segment stiffness, predisposing the tissue to injury and stress [12, 13]. These NP changes initiate a cascade expanding to other disc structures, particularly the AF. The reduced NP pressure causes an inward bulging of the AF, which is distinctly visible in the middle image of Figure 3 [12]. A loss of distinction between the nucleus pulposus and annulus fibrosus is also characteristic of disc degeneration, as seen in Figure 4. As

degeneration proceeds, progressive alterations in mechanics and composition occurs in the IVD structures, including decreased disc height decreases and the formation of tears [17] (Figure 3).

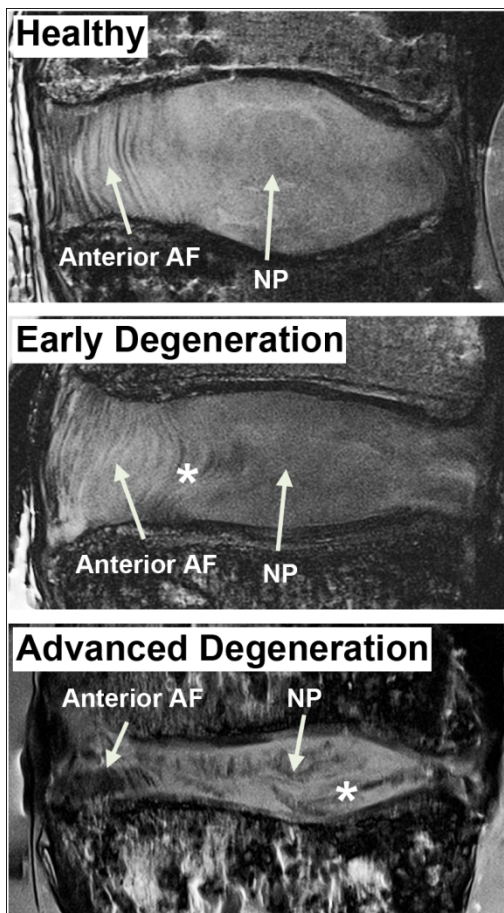


Figure 3. The MRI images of human IVDs shows the contrast between a healthy disc and the progressive effects of degeneration. (Source: Smith et al, 2010, Dis. Model. Mech.)



Figure 4. Intervertebral disc degeneration: healthy disc (left) and degenerate disc (right). (Source: <http://www.physiol.ox.ac.uk/EURODISC/>)

2.3.2 Cytokine-Mediated Matrix Degradation

Human disc degeneration is also associated with alterations in cellularity and biology. A lack sufficient nutrient supply to the NP affects the normal biological cellular pathways, beginning the degeneration process. As degeneration progresses, overall cell density decreases

primarily due to apoptotic response [18]. Consequently, the compositional and functional NP changes occur in an increasingly inflammatory microenvironment, characterized by the up-regulation of cytokines [19, 20]. The two key pro-inflammatory cytokines associated with disc degeneration are interleukin-1 β (IL1 β) and tumor necrosis factor α (TNF α). The expression of IL1 β is significantly greater in degenerate discs and TNF α is also increased but to a lesser extent [19]. In healthy IVDs, naturally occurring inhibitory proteins—interleukin-1 antagonist receptor (IL1ra) and soluble TNF receptor 1 (sTNFR1)—block the activity of catabolic cytokines—IL1 β and TNF α —respectively [19, 21, 22]. However, in disc degeneration, the increased cytokine levels occur without a concomitant increase in their associated inhibitors. The unmatched production of cytokines and inhibitors causes an imbalance in catabolic and anabolic events. Secondly, increased levels of catabolic enzymes, specifically MMP3, MMP13, and ADAMTS4, are characteristic of human disc degeneration [23]. The increases in both cytokines have been demonstrated to significantly increase MMP3, MMP13, and ADAMTS4 expression and decrease expression of NP matrix proteins including aggrecan and collagen II [24, 25]. Matrix degradation of the NP is a key event in IVD degeneration [24].

The up-regulated cytokines are targeted in order to prevent the cascading effects of NP matrix degradation. Therefore, inhibitors IL1ra and TNFR1 have promising therapeutic potential in preventing cytokine-mediated NP degradation and reducing further catabolic activity found in disc degeneration.

2.4 Pro-Inflammatory Cytokines

2.4.1 Association of IL1 with IVD

As a key event in IVD degeneration, matrix degradation in the nucleus pulposus has been the focus of many investigations. The first evidence of the possible involvement of IL1 in matrix degradation in degenerate discs was found in 1988 in in-vitro experiments using rabbit AF cells [26]. A 1997 study extended these observations of IL1 into human IVDs without fully localizing enzymes in the AF and NP [27]. According to more recent evidence, normal IVD cells show an expression of both isoforms—IL1 α and IL1 β —with a matched expression of the natural IL1 inhibitor, IL1ra [21]. In degeneration, IL1 β is up-regulated without increased IL1ra, leading to localized increased catabolic activity. A 2008 in-situ zymography investigation further showed that IL1 up-regulates enzyme activity and IL1ra inhibitor reduces induced catabolic enzyme activity [24].

2.4.2 Association of TNF α with IVD

TNF α was first described in IVDs in relation to sciatic pain [24]. Other studies have found that TNF α is present in both normal and degenerate IVDs [28]. However, the role of TNF α in disc degeneration is under debate with conflicting evidence. According to an in-vitro NP model study, using normal bovine NP cells on a porous calcium-based substrate, TNF α induced a loss of matrix molecules and decreased matrix synthesis [29]. However, a recent in-situ study of human IVDs reported no effect of TNF α on matrix-degrading activity. In this study, gene and protein expression data indicated TNF α up-regulation, and further evidence showing an absence of TNF receptor expression [24].

2.4.3 Roles of Cytokines

Previous studies have established an association of both cytokines IL1 β and TNF α in normal and degenerate IVDs. However, the roles of these cytokines in initiating NP matrix changes is under debate. Previous studies have also shown matrix changes due to increases in these cytokines, but not the functional significance of NP compositional changes.

3. METHODS

3.1 Cell Isolation and Preculture

Mature NP cells were isolated from bovine caudal discs. Isolated cells were expanded in monolayer, in Dulbeccos Modified Eagle Medium (DMEM) containing 10% fetal bovine serum and 1% penicillin/streptomycin/Fungizone (PSF). Cells were expanded through two passages for this study.

3.2 NP Constructs and Treatment

NP cells were suspended in chemically defined media (CM) and combined 1:1 with sterile type VII agarose in phosphate buffered saline (PBS) at room temperature. Chemically defined media consisted of DMEM supplemented with 1x PSF; 0.1 μ M dexamethasone; 50mg/mL ascorbate 2-phosphate; 40mg/mL L-proline; 100mg/mL sodium pyruvate; and 1X6.25 μ g/mL insulin, 6.25 μ g/mL transferrin, 6.25ng/mL selenious acid, 1.25mg/mL bovine serum albumin (BSA), and 5.35 μ g/mL linoleic acid. NP Cells were seeded in agarose gels (4mm diameter X 2.254mm thick) at a density of 20X10⁶/ml and allowed to gel for 20 minutes. NP constructs were precultured for 6 weeks in CM containing 10ng/ml transforming growth factor beta 3 (TGF β 3). The preculture period allowed for the development of a functional matrix. Samples were then removed and cultured for an additional two weeks in media without TGF β 3. Samples were divided into treatment groups (n=7) designated for IL1 treatment or TNF treatment (Table 1). After 3 days of treatment, samples were evaluated for histology, mechanics, and biochemistry.

IL1 Treatment Groups (n=7)	TNF Treatment Groups (n=7)
rh IL1 β , 10ng/ml	rh TNF α , 10ng/ml
rh IL1 β , 10ng/ml + rh IL1ra, 100ng/ml	rh TNF α , 10ng/ml + rh TNFR1, 100ng/ml
rh IL1ra, 100ng/ml	rh TNFR1, 100ng/ml
Control (no treatment)	Control (no treatment)

Table 1. Treatment groups for both IL1 and TNF with dosages of cytokine and/or inhibitor.

3.3 Histology

For histology, samples were fixed in 4% paraformaldehyde, dehydrated in a graded series of ethanol, and embedded in paraffin. Samples were sectioned at 7 μ m-thickness from the

middle of the construct. Sections were stained with Alcian Blue (pH 1.0) for sulfated proteoglycans and Picrosirius Red (0.1% w/v in saturated picric acid) for collagens.

3.4 Mechanics

Samples from each treatment group were mechanically tested in confined compression ($n=5$). The confined compression test was used to replicate the physiological conditions of the NP. The mechanical testing device consisted of an acrylic chamber fixed above a porous platen, in a bath filled with culture medium. The in-vitro NP construct was peripherally confined by the chamber in order to model the in-vivo NP confined by the AF. An impermeable ceramic indenter attached to a mechanical testing device applied compression. First, samples were equilibrated under a static preload of 0.02 N held for 500 seconds. Samples were subjected to a stress relaxation test, consisting of 10% strain (calculated from post-creep thickness values) applied at 0.05%/s, followed by relaxation to equilibrium for 10 minutes. Aggregate modulus, H_A , was calculated as the final equilibrium stress divided by the applied strain. The aggregate modulus was used as a measure of stiffness of the tissue at equilibrium. Hydraulic permeability, k_0 , was calculated using a linear biphasic theory, assuming material isotropy [30]. The hydraulic permeability models the rate at which fluid flows out of the construct.

3.5 Biochemical Analysis

For the biochemical analysis, the glycosaminoglycans, collagen, and DNA content were determined using three assays. First, the samples were weighed for wet weight, dried overnight, and weighed again for dry weight. Dried samples were digested for 16 hours in papain (0.56U/mL in 0.1M sodium acetate, 10M cysteine hydrochloric acid, 0.05M ethylenediaminetetraacetic acid, pH 6.0) at 60°C. After digestion, the PicoGreen dsDNA assay was used to determine the DNA content. Digested samples were evaluated for GAG content using the 1,9-dimethylmethylene blue dye-binding (DMMB) assay against a standard curve of chondroitin-6-sulfate. After acid hydrolysis of the sample digests, the collagen content was determined using the orthohydroxyproline (OHP) assay. DNA content was reported as amount per construct, and GAG and collagen values are reported as percentages of wet weight.

3.6 Statistical Analysis

Differences in mechanics and composition between groups were assessed using ANOVAs and Student Newman-Keuls pair-wise post-hoc tests, with significance considered for $p < 0.05$.

4. RESULTS

4.1 Histology

Histological staining results were used to confirm the uniform distribution, particularly of GAG, in the extracellular matrix. Functionally mature constructs are shown in Figure 3. Alcian Blue staining for GAG was uniformly distributed and intense. Picrosirius Red staining for collagen was a more diffuse distribution, showing both pericellular and intercellular staining.

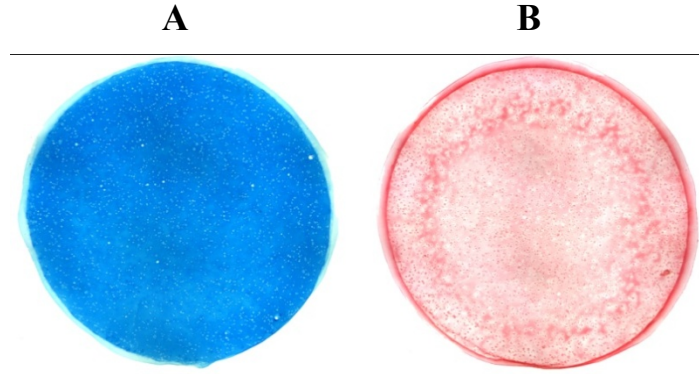


Figure 3. Histology staining of functionally mature constructs for (A) GAG using Alcian Blue and (B) collagen using Picrosirius Red. (Images taken by author)

4.2 Mechanics

Treatment with IL1 β resulted in a significant decrease of 33% in aggregate modulus, H_A , and a significant increase of 41% in hydraulic permeability, k_0 , compared with untreated controls (Figure 4, * $p < 0.05$). For groups treated with IL1 β + IL1ra or IL1ra only, aggregate modulus, H_A , and hydraulic permeability, k_0 , did not differ significantly from controls. For all samples in the TNF treatment groups, no changes in H_A and k_0 were found.

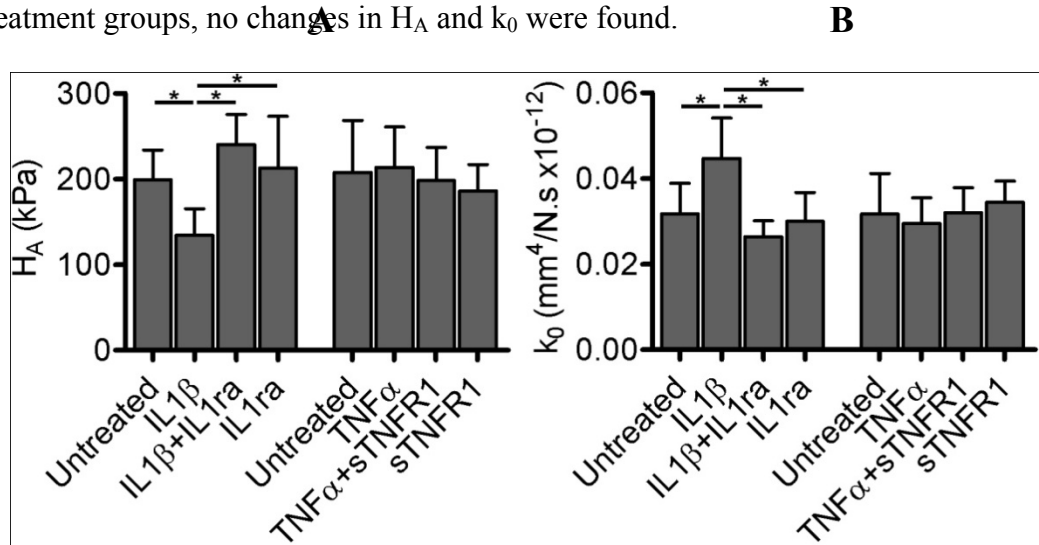


Figure 4. Mechanical testing results: (A) aggregate modulus and (B) hydraulic permeability.

4.3 Biochemical Analysis

For all treatment groups, no significant changes in DNA content were found. Samples treated with IL1 β showed a significant decrease in GAG (27%) relative to untreated controls. For samples treated with IL1 β + IL1ra or IL1ra alone, GAG was not significantly different from controls, but was significantly greater than IL1 β only group. In comparison to controls, there were no differences in GAG for all TNF treatment groups. Collagen followed a similar trend as for GAG, but no significant differences between treatment groups were found.

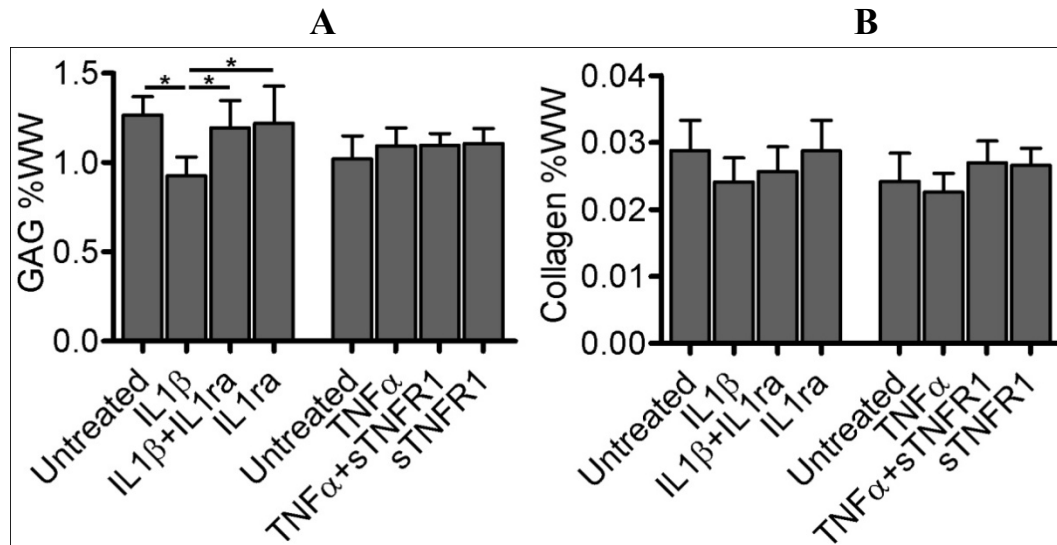


Figure 5. Biochemical analyses results: (A) GAG content as %/water weight and (B) collagen amount as %/water weight.

5. DISCUSSION

Intervertebral disc degeneration is the major cause of chronic low back pain for which current treatment methods have limited success. In intervertebral disc degeneration, changes in the extracellular matrix of the nucleus pulposus lead to reduction in function. Up-regulation of catabolic cytokines—IL1 β and TNF α —in the NP of degenerate discs are associated with alterations in composition [20]. However, there has been some debate as to which of these cytokines has a more direct role in initiating NP extracellular matrix changes [22, 29]. In the literature, the effects of TNF α in disc degeneration are under debate with some evidence showing minimal effects and others showing no effects. The results of this study support the idea that IL1 β plays a more direct role than TNF α [22].

Both mechanical and biochemical analyses demonstrate statistically significant effects of IL1 β treatment. Mechanical testing results for IL1 β groups showed decreased aggregate modulus and increased hydraulic permeability relative to untreated controls. Alterations in mechanical properties for IL1 β treatment followed similar trends as for physiological conditions of degenerate discs, while treatment with TNF α showed no difference. Further, biochemical results showed changes in composition corresponding to changes observed in mechanics. Treatment with IL1 β , but not TNF α , showed extracellular matrix changes, specially decreased GAG. This study demonstrated short-term exposure to IL1 β induced matrix changes that are functionally significant.

In IVD degeneration, IL1 β is upregulated without increased levels of its natural inhibitor, IL1ra. Several recent studies have investigated the therapeutic potential of IL1ra in reducing the catabolic IL1 β cytokine activity for disc degeneration. The results of this study demonstrate that

IL1ra can effectively prevent NP matrix changes, particularly GAG loss, and the associated functional changes induced by IL1 β .

In investigating the initiation and progression of IVD degeneration, specifically factors affecting the NP, our study found IL1 β as a key cytokine regulating matrix degradation. As a key event in IVD degeneration, prevention of catabolic cytokine has practical therapeutic implications. IL1ra can effectively inhibit composition changes of the NP matrix, particularly GAG loss, and the associated functional effects induced by IL1 β . Significant inhibitory effects place IL1ra as a key therapeutic agent for restoring function in the impaired disc tissue. Results are also clinically significant for developing novel treatments for IVD degeneration.

6. RECOMMENDATIONS

The in vitro investigation of disc degeneration and potential therapies is necessary for designing future applications of cytokine antagonists. Future work will focus on gene expressions specific to the down-regulation of anabolic genes and up-regulation of catabolic genes associated with degeneration. Investigating gene expression will further quantify changes in catabolic enzyme activity and matrix synthesis that follow cytokine exposure. Changes in catabolic and anabolic activity will be contributed to alternations in composition and mechanics. Other future work includes using biodegradable polymeric microspheres to deliver therapeutic agents and evaluating therapeutic agents in an in vivo model of disc degeneration. Results of the current study demonstrated the direct role of IL1 β in disc degeneration and the inhibitory effects of IL1ra, which is important for novel treatment approaches.

7. ACKNOWLEDGMENTS

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9. APPENDIX

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Inhibition of Functional Matrix Degradation in a Cytokine-Mediated In-Vitro Model of Nucleus Pulposus Degeneration

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INTRODUCTION

Altered nucleus pulposus (NP) extracellular matrix composition and mechanical function are hallmarks of disc degeneration [1,2]. Early stage degeneration is characterized by decreasing aggrecan content, with the associated decrease in hydrostatic pressure impairing the ability of the NP to perform its most critical function: the even distribution and transfer of compressive loads between the vertebral bodies [3].

Changes in matrix composition occur within an increasingly inflammatory local microenvironment [1]. There is up-regulation of pro-inflammatory cytokines, particularly interleukin 1 beta (IL1 β) and tumor necrosis factor alpha (TNF α), without a concomitant increase in associated inhibitory regulators of these cytokines [1].

The direct functional consequences of cytokine-mediated extracellular matrix changes in the NP have not been described, in part due to the lack of an appropriate mimetic in-vitro culture model. In this study we developed an in-vitro model that was used to investigate the effects of IL1 β and TNF α on both NP extracellular matrix composition and mechanical function. In addition, we examined the capacity of IL1 receptor antagonist (IL1ra) and soluble TNF receptor 1 (sTNFR1), inhibitors of IL1 β and TNF α respectively, to mitigate cytokine-mediated functional and compositional changes.

METHODS

Cell Isolation, Preculture and Treatments: NP cells were isolated from mature bovine caudal discs and expanded through 2 passages. Cells were seeded in agarose gels (4mm diam. x 2.4mm thick) at a density of 20×10^6 /ml, and precultured for 6 weeks in chemically defined media containing 10ng/ml TGF β 3 [4]. Samples were then removed and cultured without TGF β 3 for an additional 2 weeks. Samples were then divided into cytokine and inhibitor treatment groups (Table 1). After 3 days of treatment, mechanical properties and biochemical composition were evaluated.

Treatment Group	IL1	TNF
Cytokine Only	rhIL1 β , 10ng/ml	rhTNF α , 10ng/ml
Cytokine + Inhibitor	rhIL1 β , 10ng/ml + rhIL1ra, 100ng/ml	rhTNF α , 10ng/ml + rhTNFR1, 100ng/ml
Inhibitor Only	rhIL1ra, 100ng/ml	rhTNFR1, 100ng/ml
Untreated Control	Control (no treatment)	Control (no treatment)

Histology: To confirm uniform matrix distribution in mature constructs, samples were processed into paraffin. Sections were stained with either picosirius red or alcian blue to demonstrate collagen and glycosaminoglycan (GAG) accumulation respectively.

Mechanics: Constructs from each treatment group were tested in confined compression (n=5). The testing system consisted of an acrylic chamber fixed above a porous platen in a bath filled with culture medium. Compression was applied with an impermeable ceramic indenter attached to a mechanical testing system. Samples were first subjected to a 0.02N preload for 500s, followed by a stress relaxation test, consisting of 10% strain applied at 0.05%/s, followed by relaxation to equilibrium for 10mins. Aggregate modulus, H_A , was calculated as equilibrium stress/applied strain. Hydraulic permeability, k_0 , was determined using linear biphasic theory assuming material isotropy [5].

Biochemistry: Following mechanics, the same samples were weighed and digested in papain. Digests were assayed for DNA content (per construct), and sulfated GAG and hydroxyproline (collagen) normalized to wet weight.

Statistics: Differences in mechanics and composition between groups were assessed using ANOVAs and Student Newman-Keuls pair-wise post-hoc tests, with significance considered for $p < 0.05$.

RESULTS

Histology: Functionally mature constructs are shown in Fig 1. GAG staining was intense and uniform, while collagen staining was more diffuse.

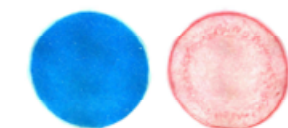


Figure 1. Functionally mature constructs: GAG (left) and collagen (right).

Mechanics: H_A increased from 2.5 ± 0.4 kPa (mean \pm SD) at day zero, to 199.23 ± 4.7 kPa following preculture. Treatment with IL1 β resulted in a significant decrease in H_A (33%), and an increase in k_0 (41%) compared with untreated controls (Fig 2, * $p < 0.05$). For samples treated with IL1 β +IL1ra, or IL1ra alone, H_A and k_0 were not significantly different from controls. No changes in H_A or k_0 were found for any of the TNF treatment groups.

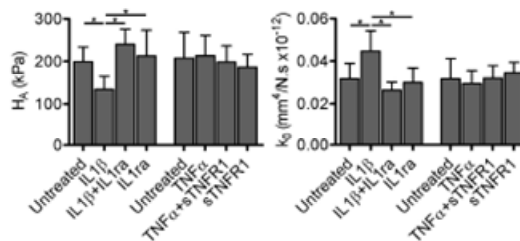
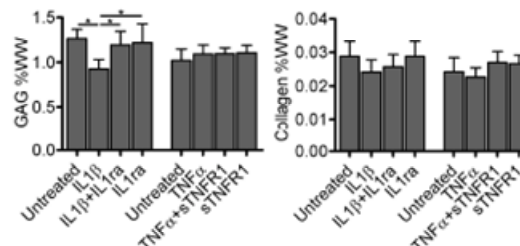


Figure 2. Mechanical properties

Biochemistry: Relative to untreated controls, GAG decreased significantly for samples treated with IL1 β (27%), but was not different from controls for all other treatments (Fig 3, left, * $p < 0.05$). Collagen showed a similar trend as for GAG, but there were no significant differences between groups (Fig 3, right). DNA did not change significantly for any of the treatments (not shown).



DISCUSSION

While both IL1 β and TNF α are upregulated in the NPs of degenerate discs [1], there has been some debate as to which of these cytokines plays a more direct role in mediating extracellular matrix changes [6,7]. The results of this study support the hypothesis that IL1 β plays a more direct role than TNF α [6]. Further, we have demonstrated that matrix changes, particularly decreased GAG, following short term exposure to IL1 β are functionally significant. Our results support the therapeutic potential of IL1ra for disc degeneration [8,9], by demonstrating that IL1ra can effectively prevent NP GAG loss and the associated functional changes induced by IL1 β .

The in-vitro culture model developed here presents a new platform for investigating biological mediators of disc degeneration and the associated compositional and functional changes. Future work will seek to quantify the changes in catabolic enzyme activity and matrix synthesis that follow cytokine exposure, and contribute to alterations in composition and mechanical properties.

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