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Investigating pathways associated with intervertebral disc degeneration and back pain.

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Supervisor: Seguin, Cheryle A., *The University of Western Ontario* A thesis submitted in partial fulfillment of the requirements for the Doctor of Philosophy degree in Physiology and Pharmacology © Geoffrey J. Kerr 2020

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Abstract

Lower back pain (LBP) is one of the most common conditions worldwide, yet, current therapeutics are limited to symptomatic relief and do not directly treat the underlying cause of pain. This is largely due to an incomplete understanding of the biological pathways and tissues involved in LBP. While many tissues appear to be involved, intervertebral disc (IVD) degeneration is believed to be a major contributor.

The main aim of this thesis was to characterize the role of two environmental risk factors, mechanical loading and obesity, in the initiation of IVD degeneration and associated pain using the mouse as a preclinical model. We first investigated the effects of mechanical loading, specifically whole-body vibration (WBV), on joint health. In contrast to previous findings in CD-1 mice, we show that WBV delivered using parameters that model those used clinically does not affect joint health in C57BL/6 mice . These findings suggest that the response of joint tissues to WBV may vary in a strain-specific manner.

We next evaluated the impact of diet-induced obesity on IVD degeneration and associated pain. Ten-week-old male C57BL/6N mice were fed either a control, high-fat, or western diet for 12, 24 or 40 weeks. At endpoint, mice were assessed for behavioral indicators of pain and histological and molecular analyses were carried out on joint and neural tissues. We demonstrated that diet-induced obesity accelerates IVD degeneration; however, painrelated behaviors precede histological joint damage.

Finally, we examined the role of peroxisome proliferator-activated receptor delta (PPAR δ) as a potential mediator of IVD degeneration. We confirmed that PPAR δ was expressed and functionally active in the IVD and used male and female, cartilage-specific *Ppard* knockout

(*Ppard* KO) mouse to assess age- and obesity-associated IVD degeneration. Loss of PPARδ appeared to be protected against age-associated IVD degeneration; however, this protective effect was not observed with obesity-induced IVD degeneration or knee osteoarthritis.

Overall, this data highlights the complex interactions between modifiable and nonmodifiable risk factors as drivers of IVD degeneration and associated back pain. Furthermore, these studies highlight the importance of including sex as a biological variable and assessing pain and IVD degeneration, as they do not always correlate.

Keywords: Intervertebral disc degeneration; Back pain; Whole-body vibration; Obesity; Peroxisome proliferator activated receptor delta (PPARδ); Mouse models.

Summary for Lay Audience

Lower back pain (LBP) is one of the most common causes of disability worldwide, yet current treatments are limited to pain relief and do not directly treat the underlying cause of pain. This is largely due to an incomplete understanding of what tissues contribute to LBP and why pain occurs. Large-scale association studies in humans have identified several risk factors for the development of LBP including age, genetics, mechanical loading and obesity; however, how and why they cause the pain is largely unknown.

While many tissues appear to be involved in LBP, intervertebral disc (IVD) degeneration is believed to be a major contributor in many cases. IVDs are specialized joints located between the vertebrae in the spine that act as shock absorbers, helping to dissipate mechanical loading. Degeneration is a progressive biological process that results in the functional failure of the tissue and can result in pain. This research was designed to investigate the impact of mechanical loading, specifically whole-body vibration (WBV), and obesity on IVD degeneration and associated pain using the mouse as a model system. Furthermore, we investigate a potential biological mediator to explain how obesity impacts IVD degeneration and associated pain.

In our first study we demonstrate that the response of joint tissues to WBV vary based on genetic background of the animals. In our second study we demonstrated that diet-induced obesity does accelerate IVD degeneration and pain-related behavior in a mouse model; however, pain related behaviors precede structural IVD damage. Lastly, we assessed a potential biological mediator of IVD degeneration - peroxisome proliferator activated receptor delta (PPAR δ) – in aging and obesity. Using mice that had the PPAR δ gene deleted in joint tissues, we showed that loss of PPAR δ protected against age-associated IVD

degeneration; however, this protective effect was not observed with obesity-induced IVD degeneration.

Taken together, this thesis highlights the complex interactions between mechanical loading and obesity as drivers of IVD degeneration and associated back pain. Furthermore, these studies highlight the importance of including sex as a biological variable and assessing pain and IVD degeneration, as they do not always correlate.

Co-Authorship Statement

Comprehensive co-authorship statements are included at the beginning of each chapter.

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List of Abbreviations

Abbreviation	Full Name
Abca1	ATP binding cassette subfamily A member 1
Acan	Aggrecan
ADAMTS	A disintegrin and metalloproteinase with thrombospondin motifs
AF	Annulus Fibrosus
AF-1	Activating function-1
AF-2	Activating function-2
AGE	Advanced glycation end product
Angptl4	Angiopoietin-like 4
ATF3	Activating transcription factor 3
ATP	Adenosine triphosphate
BDNF	Brain-derived neurotrophic factor
BMD	Bone mineral density
BMI	Body mass index
CCN2	Cellular communication network factor 2
CEP	Cartilage endplate
CGRP	Calcitonin gene-related peptide
Cpt1a	Carnitine palmitoyltransferase 1A
CRP	C-reactive protein
СТ	Computed tomography
DMMB	Dimethylmethylene blue
DNA	Deoxyribonucleic acid

DRG	Dorsal root ganglia
ECM	Extracellular matrix
FFA	Free fatty acid
FXR	Farnesoid X receptor
GAG	Glycosaminoglycan
Gdf5	Growth differentiation factor 5
GFAP	Glial Fibrillary acidic protein
Gsta4	Glutathione S-transferase A4
HU	Hounsfield unit
IBA-1	Ionized calcium binding adaptor molecule 1
IL	Interleukin
Insig1	Insulin induced gene 1
IP	Intraperitoneal
ISO	International organization of standardization
IVD	Intervertebral disc
КО	Knockout
LBP	Lower back pain
LFC	Lateral femoral condyle
LTP	Lateral tibial plateau
LXR	Liver X receptor
MCP-1	Monocyte chemoattractant protein-1
MFC	Medial femoral condyle
МНО	Metabolically healthy obese

MMP	Matrix metalloproteinase
MSK	Musculoskeletal
MTP	Medial tibial plateau
NGF	Nerve growth factor
Noto	Notochord homeobox
NP	Nucleus pulposus
OA	Osteoarthritis
OARSI	Osteoarthritis research society international
Pdk4	Pyruvate dehydrogenase kinase 4
PPAR	Peroxisome proliferator-activated receptor
Ptgs2	Prostaglandin-endoperoxide synthase 2
RAGE	Receptor for advanced glycation end products
ROS	Reactive oxygen species
Rps29	Ribosomal protein S29
RXR	Retinoid X receptor
Scx	Scleraxis BHLH transcription factor
SPARC	Secreted protein acidic and cysteine rich
TNF	Tumor necrosis factor
TrkA	Tropomyosin-related kinase A receptor
TRPV	Transient receptor potential vanilloid
Txnip	Thioredoxin interacting protein
VDR	Vitamin D receptor
VEGF	Vascular endothelial growth factor

WBV Whole-body vibration

WT Wild type

Chapter 1

1 Introduction

1.1 What is pain?

The International Association for the Study of Pain defines pain as "an unpleasant sensory and emotional experience associated with actual or potential tissue damage or described in terms of such damage"¹. The ability to detect noxious and potentially dangerous stimuli is essential to avoid tissue injury and facilitate healing. While pain is unpleasant, it is essential for overall survival and wellbeing². This is highlighted in individuals that have congenital insensitivity to pain. These individuals cannot feel broken bones, heat from a fire or being cut. As a result, they are not aware of potentially life-threatening injuries and do not know when to take appropriate protective behaviors³.

1.1.1 Biological mechanisms of pain

Pain is a multidimensional, dynamic interaction between biological, psychological, and social factors that influence one another and can contribute to how a person interprets noxious stimuli signaling tissue injury⁴. At the biological level, nociception is the process by which information about actual or potential tissue damage is relayed to the brain⁵. These signals are initiated at specialized receptors called nociceptors, which are attached to thin myelinated A δ fibers and unmyelinated C-fibers that transmit the signal to the brain⁶. At the site of injury these nociceptors become activated by chemical activation or mechanical and thermal stimuli², leading to the opening of voltage-gated sodium and potassium

channels². The opening of these channels forms an action potential, which is then propagated to the dorsal horn of the spinal cord and ultimately the brain².

1.1.2 Acute vs. Chronic pain

While acute pain plays a physiological role in response to tissue damage, chronic pain is defined as "pain that persists past the normal time of healing", which, in the context of back pain, is typically over 3 months in duration ⁷. Chronic pain conditions are one of the most common reasons people seek medical care, and have been linked with decreased quality of life, decreased mobility, and the development of psychiatric conditions such as depression and anxiety^{8–11}. Chronic pain conditions have a large socioeconomic effect, affecting up to 40% of Canadians¹². Chronic pain costs the Canadian taxpayer more than cancer and heart disease combined, with direct healthcare costs at \$6 billion a year^{13,14} and indirect costs estimated at \$37 billion a year¹³. Uncontrolled pain is the most common cause of disability amongst working-age adults in Canada, and up to 60% of people living with chronic pain lose their jobs^{14,15}.

While the precise mechanisms associated with the transition from acute to chronic pain are largely unknown, several potential mediators have been implicated including peripheral and central sensitization. Peripheral sensitization involves increased excitability of the primary afferent nociceptors, resulting in a decreased threshold for action potential generation or the generation of spontaneous action potentials without the presence of a noxious stimuli^{16,17}. While many biological mechanisms can contribute to peripheral sensitization, inflammatory mediators secreted at the site of injury can change the properties of nociceptors^{5,18}. Nerve growth factor (NGF) appears to be one key mediator

driving peripheral sensitization through both direct actions on the nociceptors and indirect actions on surrounding tissues. NGF is a neurotrophin released by many cell types, including inflammatory cells, at the site of tissue injury^{19,20}. NGF exerts its actions through binding with its receptor, Tropomyosin-related kinase A receptor (TrkA), activating a series of downstream signaling pathways. NGF-TrkA activation on nociceptors leads to rapid sensitization of the neuron through intracellular signaling cascades that results in increased expression and phosphorylation of channels and receptors involved in pain transduction^{21,22}. NGF-TrkA activation also leads to a prolonged increase in the expression of neuropeptides, receptors and ion channels involved in pain transmission ²³.

Central sensitization can also contribute to chronic pain. Central sensitization is caused by increased excitability of second order neurons within the spinal cord that transmit the nociceptive signals from the primary afferent neurons to the brain^{17,24}. This increased excitability within the central nervous system can contribute to the lack of a direct correlation between nociceptor activation and painful experience¹⁷. Several mechanisms contribute to central sensitization including increased neuronal membrane excitability, facilitated synaptic transmission, and disinhibition of second order neurons²⁵. While neuropeptides released from nociceptive neurons contribute to central sensitization, glial cells (astrocytes and microglia) within the spinal cord also play an important role in this process²⁶. Astrocytes are the most abundant cell type within the brain and perform a variety of different functions including, but not limited to, regulating synaptic transmission and neuroinflammation, these cells undergo a phenotypic switch to a so called "state of activation" marked by increased production of the astrocyte-specific marker glial fibrillary

acidic protein (GFAP), and an enlargement of the astrocyte processes²⁶. Activated astrocytes have decreased glutamate reuptake and release neuromodulators that can alter synaptic transmission within the spinal cord²⁶. Microglia, which are the resident macrophages in the spinal cord, are also activated in chronic pain conditions and contribute to central sensitization²⁶. Activated microglia release pro-inflammatory cytokines that further modulate synaptic activity by altering the activity and expression of ion channels and receptors²⁶. While both peripheral and central sensitization are potential mediators of chronic pain, it should be noted that pain is a biopsychosocial experience that has sensory, affective, and cognitive components that contribute to the interpretation of nociceptive signals, and whether the signals coming into the brain are perceived as pain⁵.

1.2 Back pain is a problem

The most recent *Global Burden of Disease* study reported back pain as the single most common cause of disability worldwide, with a lifetime prevalence of over 84% in developed countries²⁹. While 90% of people recover from acute episodes of lower back pain (LBP) within a few months, some will not recover and instead develop chronic LBP^{30,31}. Chronic LBP is debilitating physically as well as socially. People suffering from LBP report feeling hopelessness, loss of identity and social isolation, which can lead to depression and suicide ideation^{32,33}. Like other chronic pain conditions, chronic LBP has a large impact at the societal level as well. In the United States, LBP costs the economy an estimated \$100 billion a year in both direct (primary care, pharmaceuticals, etc.) and indirect costs (lost wages, reduced productivity)³⁴. Several tissues appear to be involved in chronic LBP, including muscles, ligaments, facet joints, nerve roots, vertebral bones, and

the intervertebral discs (IVDs)³⁵. Of these, pain arising from IVD degeneration – termed discogenic back pain - is believed to be the major contributor to pain in an estimated 40% of cases^{36,37}. Current therapeutic interventions focus on providing symptomatic relief, but do not address the underlying cause of the pain³⁸, or restore the mechanical function of the tissue³⁹. Altered mechanical function of the tissue can accelerate IVD degeneration in adjacent spinal levels, which can further contribute to pain and disability^{40,41}. Our ability to treat IVD degeneration is limited due to an incomplete understanding of molecular processes that regulate IVD homeostasis and disease.

1.3 The Spine

In humans, the spine has 3 main functions: protect the spinal cord and nerve roots, provide structural support to maintain balance and upright posture, and enable flexible motion⁴². Anatomically, the spine consists of an osseous vertebral column surrounded by musculature and connective tissues that aid in the overall function of the structure (**Figure 1.1**). The basic functional unit of the spine is called a motion segment, consisting of a single intervertebral disc, the inferior and superior vertebrae and associated connective tissue⁴³.

1.3.1 Vertebral bone

In humans, the vertebral column is composed of 24 presacral vertebrae separated by 23 intervertebral discs (IVDs), subdivided into three anatomical regions: cervical (7 vertebrae, 6 IVDs), thoracic (12 vertebrae, 12 IVDs) and the lumbar region (5 vertebrae, 5 IVDs)⁴⁴. In mice, the vertebral column is composed of 7 cervical, 13 thoracic, 6 lumbar,



Figure 1.1: Schematic depiction of the major components of the spine.

The spine is composed of several different tissues that are essential for its overall function. These include osseous vertebrae, fibrocartilaginous intervertebral discs, muscles, and ligaments. The spine also serves structurally to surround the spinal cord. The spinal cord is a component of the central nervous system that transmits motor and sensory information between the brain and the peripheral nervous systems.

4 sacral, and 28 caudal (tail) vertebrae⁴⁵. While the osseous features of each vertebrae are generally preserved in the different anatomical areas, their morphology differs slightly and contribute to the biomechanical stability and flexibility of the vertebral column⁴⁴.

1.3.2 The Intervertebral Disc

The intervertebral disc (IVD) is a fibrocartilaginous joint that links vertebral bodies along the spine and allows for spine flexibility and stability under axial compression⁴⁶. Anatomically, the IVD is a heterogeneous structure composed of three distinct, yet interdependent tissues. At the core of the IVD lies the gelatinous nucleus pulposus (NP), surrounded by the fibrocartilaginous annulus fibrosus (AF), and cartilage endplates that anchor the IVD to adjacent vertebrae (**Figure 1.1**). Each tissue is structurally and mechanically unique, but closely coupled with one another to contribute synergistically to the function of the IVD⁴⁷.

1.3.2.1 Nucleus Pulposus

At the center of the IVD lies the nucleus pulposus (NP), a gelatinous tissue composed of proteoglycans and water held together by an irregular network of type II collagen⁴⁸. Although the NP contains several different types of proteoglycans such as decorin, versican, and biglycan, the major proteoglycan in the NP is aggrecan^{48,49}. These proteoglycan core proteins are covalently bound by glycosaminoglycans (GAGs), such as chondroitin sulfate and keratin sulfate, that act to draw water into the NP thereby contributing to its osmotic properties and ability to dissipate compressive loads^{48,50,51}. Aggrecan also has the ability to form proteoglycan aggregates, in which many aggrecan molecules interact with hyaluronan via link protein, which limits their diffusion through

the tissue, allowing for an even distribution of load across the NP^{51,52}. Within the NP there are two different cell types that contribute to extracellular matrix (ECM) production: notochord cells and cartilage-like NP cells. During development, notochord cells are the major cell type in the NP and produce a proteoglycan rich ECM^{53,54}. In humans, the number of notochord cells within the NP steadily decrease with age, and the NP is populated instead by cartilage-like NP cells⁵⁵ that secrete an altered ECM ⁵³. Functionally, the role of the NP is to dissipate compressive loading between adjacent vertebral segments of the spine. This is facilitated through the osmotic pressure created by the anionic side chains attached to proteoglycan, which act to draw in water to this tissue^{46,47}. The NP's ability to dissipate compressive loading is dependent on the radial forces of the surrounding annulus fibrosus⁵⁶.

1.3.2.2 Annulus Fibrosus

The fibrocartilaginous annulus fibrosus (AF) encapsulates the gelatinous NP. The major component of the AF ECM is collagen, with lower levels of proteoglycans compared to the NP⁵⁷. In the AF, the collagen fibers run in parallel to one another within distinct lamella. In humans, the AF is arranged in 15-25 concentric lamellae, where collagen fibers of adjacent lamellae travel in opposite directions at oblique angles, to form an angle-ply structure providing tensile strength^{56,58,59}. The AF is typically divided into the outer AF and the inner AF, which have different ECM components. The outer AF is composed largely of type I collagen and elastic fibers, helping the tissue to dissipate tensile load^{56,60}. This is in contrast to the inner AF, which serves as a transition between the fibrocartilaginous outer AF and the NP, and has a higher type II collagen and proteoglycan

content within the interlamellar matrix than the outer AF; a feature that provides resistance to compressive loading^{56,61}. Similar to the ECM composition, the cellular morphology and biochemical profiles of the cells are different based on anatomical location within the AF⁶². Cells in the inner AF have more of a rounded, chondrocyte-like morphology compared to the spindle-shaped, fibroblast-like cells of the outer AF⁶². The difference in cellular profiles across this tissue are thought to arise from variations in mechanical loading seen across the tissue, where cells of the inner AF are exposed to compression and tension and the cells of the outer AF are exposed to predominantly tension^{63,64}. Functionally, the role of the AF is to support the radial forces exerted by the NP⁶³, which requires the tissue to be able to resist tensile loading and depends on the microarchitecture of the collagen network⁴⁷.

1.3.2.3 Cartilage endplates

The cartilage endplates (CEP) are thin layers of hyaline cartilage at the superior and inferior ends of the IVD⁵⁷. Similar to articular cartilage, the CEP is composed mostly of water, followed by type II collagen and proteoglycans^{57,65}. The collagen fibrils within the CEP are arranged differentially based on location within the CEP, but merge with collagen fibers of the AF at the margins of the IVD⁵⁶. The cells of the CEP are very similar in morphology and density to articular chondrocytes, yet, no zonal arrangement is seen within the CEP⁶⁶. The main role of the CEP is to act as a semi-permeable barrier. While the outer lamellae of the AF is vascularized, the NP is completely avascular and depends on diffusion through the CEP for nutrient supply⁶⁷. The vertebral bone underlying the CEP is highly vascularized, allowing for the diffusion of small molecules such as oxygen and glucose into the IVD⁶⁷. With age, the CEP can become calcified, thereby impairing the diffusion of solutes into the IVD; a process thought to drive IVD degeneration^{67,68}.

1.3.3 Muscle and connective tissue

Muscle and connective tissues (i.e. ligaments and tendons) connect to the vertebral bodies at a number of anatomical landmarks and play a major role in movement as well as supporting the axial skeleton in an upright posture^{69,70}. Muscle specifically plays a very important role in supporting the spine against forces seen in everyday life⁷¹. Muscular dysfunction, whether due to muscle weakness or impaired neurological coordination, contributes to decreased stability of spinal structures, leading to gradual degeneration of joints and soft tissue over time, potentially contributing to LBP^{71,72}. Clinically, atrophy of paraspinal muscles (e.g. multifidus muscle) has been associated with chronic LBP⁷³, however, it is difficult to determine whether this is a cause or consequence of pain.

1.3.4 Innervation & Vascularization

In the lumbar spine, the vertebral bodies, muscles and ligaments are innervated by several different nerves including the sinuvertebral nerves, ventral rami, dorsal rami, and gray rami^{74,75}. While lumbar IVDs receive innervation from some of the same neural sources (sinuvertebral nerve, dorsal rami)⁷⁶, the presence of nerves is restricted to the outermost layers of the AF in healthy tissues^{77,78}. The lumbar spine receives its blood supply from several sources, including the lumbar, middle sacral, and branches of the iliolumbar arteries⁷⁹. While the tissues surrounding the IVD are well vascularized, only the outmost layers of the AF receive direct vascular supply⁸⁰.

1.4 Intervertebral Disc Degeneration

IVD degeneration has been defined as an aberrant, cell-mediated response leading to progressive structural failure⁸¹. It involves a progressive cascade of events, beginning with changes to the IVD microenvironment and eventually resulting in structural breakdown and functional impairment of the tissue ³⁹. Disc degeneration is characterized by increased ECM breakdown and abnormal (fibrotic) matrix synthesis leading to reduced hydration, loss of disc height, and decreased ability to absorb load^{50,82}. As mentioned previously, IVD degeneration is believed to be the primary contributor to chronic LBP in an estimated 40% of cases³⁶. IVD degeneration is distinguished from disc herniation, which is associated with a tear of the AF (often associated with acute injury), leading to extrusion of the NP into the space surrounding the IVD also resulting in pain⁸³.

1.4.1 Risk factors for IVD degeneration

Risk factors for IVD degeneration include non-modifiable risk factors such as genetics, age and sex, as well as modifiable/environmental risk factors such as obesity, mechanical loading, and smoking. It should be noted that these risk factors are not mutually exclusive, and individuals can have multiple risk factors contributing to disease progression. Moreover, IVD degeneration is likely a group of diseases with distinct etiologies yet a common radiographic presentation.

1.4.1.1 Non-modifiable risk factors

Genetic predisposition is the largest risk factor for IVD degeneration, followed by age^{50,84,85}. With the advance in genetic technologies, genome-wide association studies

have highlighted several gene variants as risk factors for the development of IVD degeneration. Many of these genes encode for proteins involved in ECM homeostasis including collagen I, collagen IX, aggrecan, matrix metalloproteinase-3 (MMP-3) amongst many others^{86,87}. Polymorphisms in the coding region of these matrix components may lead to altered protein products, functional impairment, and ultimately accelerated IVD degeneration. For example, polymorphisms within aggrecans coding region are believed to decrease the number of GAG binding sites, impairing the hydrophilic nature of the molecule. This is thought to contribute to IVD degeneration through decreased hydration of the IVD, reducing its ability to dissipate compressive load⁸⁸. While polymorphisms within intronic and exonic regions of single genes have been associated with IVD degeneration, it is believed that IVD degeneration is likely polygenic, where minor variations within several genes work synergistically to contribute to disease progression^{89–} ⁹². In addition to genetic predisposition, age and sex are also established risk factors for IVD degeneration^{93,94}. In humans, IVD degeneration accelerates linearly during the fifth decade of life⁹⁴, with potential causes being loss of nutrition caused by CEP calcification, cellular senescence, and accumulation of degraded ECM^{50,67,93}. In addition, females show an increased risk of developing IVD degeneration (Odds Ratio [OR], 1.69 when adjusted for age) compared to males, although biological mechanisms underlying this association are currently unknown⁹⁵.

1.4.1.2 Modifiable risk factors

Life-style related risk factors have been shown to have a large impact on IVD health. These risk factors include mechanical loading, obesity, shift work and smoking^{81,96,97}. Mechanical

loading plays a major role in IVD cell function⁴⁷, and mechanical overloading can promote processes associated with IVD degeneration including induction of catabolic gene expression and cell death⁹⁸. In humans, increased loading is associated with IVD degeneration, through occupational hazards⁹⁹ or recreational sports such as weightlifting¹⁰⁰. Increased mechanical loading on the IVD is also seen in obesity, which is a risk factor for both structural IVD degeneration and LBP^{101,102}. However, this link between obesity, IVD degeneration, and LBP is likely multifactorial, involving both mechanical and metabolic components¹⁰³. The exact mechanisms by which these modifiable risk factors contribute to IVD degeneration are still being explore, but their impact can be reduced through life-style modifications and education.

1.4.2 Pathogenesis of IVD degeneration

Cells of the IVD are necessary to produce, maintain and repair the extensive ECM needed for the IVD to be functional. With IVD degeneration, cells undergo substantial biological changes including altered cellular phenotype, senescence, and apoptosis, that may contribute to disease progression⁵³. Cellular senescence – the irreversible arrest of cell proliferation¹⁰⁴ – is increased in IVD degeneration¹⁰⁵. Alongside reduced proliferative capacity of these cells, there are also alterations in cellular function and phenotype. In human IVD cells, markers of senescence positively correlate with increased expression of matrix degrading enzymes such as MMP13 and ADAMTS5, which mediate matrix degradation in IVD degeneration¹⁰⁶. While senescence naturally occurs during aging through telomere shortening, it can be accelerated by stress signals such as the proinflammatory cytokine interleukin 1 (IL-1), which is also associated with IVD
degeneration^{107,108}. However, it is unlikely that all phenotypic differences associated with IVD degeneration are due to cellular senescence, as reversible changes can be caused by environmental signals such as mechanical loading⁵³. During IVD degeneration, cells produce increased levels of inflammatory and catabolic factors, decrease synthesis of normal ECM components (type II collagen, aggrecan), increase synthesis of fibrotic matrix components (type I collagen), and alter secretion of growth factors that can potentiate the degenerative process⁵³. In addition to alterations in the phenotype of remaining IVD cells, IVD degeneration is also associated with increased levels of cell apoptosis¹⁰⁹. Together these factors decrease the number of properly functioning cells needed to maintain and repair the ECM, potentially accelerating IVD degeneration.

Alterations in cellular function associated with IVD degeneration is marked by increased production of factors that accelerate ECM degradation, including matrix metalloproteinases (MMPs) and 'a disintegrin and metalloproteinase with thrombospondin modifs' (ADAMTS)¹¹⁰. MMP-1,-2,-3,-7,-12,-13 are upregulated in human IVD degeneration^{110–112}, and degrade a number of ECM components, including collagens type I and II, as well as non-collagenous proteins such as aggrecan¹¹³. ADAMTS-4, and -5 are also upregulated in IVD degeneration^{113,114}. Unlike the MMPs, ADAMTS enzymes demonstrate a narrow substrate specificity; ADAMTS-4, and -5 primarily cleave aggrecan, and their increased activity leads to the accumulation of fragmented proteoglycans within the IVD¹¹⁴. While not directly investigated in the IVD, these proteoglycan fragments potentiate inflammatory processes in articular cartilage through activation of toll-like receptors, and may therefore likewise accelerate IVD degeneration¹¹⁵. In addition to matrix degrading enzymes, IVD cells intrinsically produce and secrete inflammatory cytokines

and neurotrophins such as IL-1 β , IL-6, Tumor Necrosis Factor alpha (TNF α), NGF, which are all elevated with IVD degeneration^{78,108,116}. Aside from their pro-nociceptive roles on nerve roots^{93,117}, these inflammatory cytokines can potentiate pathways leading to increased expression of many of the catabolic enzymes mentioned above including MMP-1,-3,-13 and ADAMTS-4,-5¹⁰⁸. As a result of ECM breakdown and an altered secretion of matrix components, the biomechanical properties of the IVD are impaired^{50,118}. As the NP becomes less hydrated and more fibrous, it is no longer able to dissipate compressive loads effectively, resulting in increased transfer of load to the AF¹¹⁸. Increased AF mechanical loading in turn leads to structural degeneration of the AF¹¹⁹.

Lastly, IVD degeneration is associated with vascular and neural ingrowth into the normally avascular and aneural tissue^{120,121}. It has been suggested that this neural ingrowth is an important contributor to discogenic pain, which is pain originating from the IVD^{120,122}. While the mechanisms responsible for nerve ingrowth into the IVD are largely unknown, ECM breakdown and neurotrophins are believed to play an important role⁷⁸. In healthy IVDs, aggrecan has been shown to inhibit nerve ingrowth, however, when IVDs degenerate, loss of intact aggrecan may result in nerve ingrowth^{78,123}. This appears to be coupled with increased expression of proinflammatory cytokines such as IL-1 β , TNF α and IL-4 inducing the expression of the NGF, which can directly stimulate nociceptive afferents and cause further ingrowth of nociceptive nerve fibers into the IVD⁷⁸.

Once the process of IVD degeneration has started, it is difficult to stop. A recent review describes IVD degeneration as a vicious circle, where abnormal interactions between biomechanics, cellular processes, and ECM components result in a positive feedback loop resulting in accelerated IVD degeneration (**Figure 1.2**)¹¹⁹. An alteration in any of these



Figure 1.2: Proposed feed forward model of intervertebral disc degeneration with contributing risk factors.

Intervertebral disc homeostasis is dependent on interactions between cells, extracellular matrix, and biomechanics of the spine. Alterations in any of these factors alter the others and can initiate the degenerative cycle. For an example, upregulation of catabolic genes (i.e. *Mmp3*, *Mmp13*, *Adamts4*, *Adamts5*) from the cells of the IVD – perhaps caused by aberrant mechanical loading or obesity - would lead to the degradation of the extracellular matrix, thereby altering the biomechanical properties of the IVD. Alterations in tissue mechanics would then alter the mechanical loads experienced by cells, causing further alterations to cellular function. Figure modified from¹¹⁸; Asterix indicate risk factors assessed in the current thesis.

elements may initiate IVD degeneration, ultimately leading to structural and functional failure of the tissue¹¹⁸.

1.4.3 The link between IVD degeneration and pain

Despite the clinical association between radiographic IVD degeneration and LBP, this link remains poorly understood. In fact, patients presenting with back pain often lack radiographic signs of IVD degeneration, and many individuals with radiographic IVD degeneration are completely asymptomatic^{124,125}. While the difference between symptomatic and asymptomatic IVD degeneration is not well understood, it has been suggested that nerve in-growth may cause discogenic back pain^{126–128}. During IVD degeneration, both NP and AF cells release pro-nociceptive factors including IL-1, IL-6, TNF α , and NGF^{108,129,130}, which can act as nociceptive triggers. In addition to nerve ingrowth to the disc and direct activation of nociceptors, it is believed that IVD degeneration contributes to LBP in indirect mechanisms as well¹²⁸. With IVD degeneration there are structural and function changes within the IVD that impact adjacent tissues and may provoke pain indirectly¹²⁸. For example, in advanced IVD degeneration the significant loss of IVD height can cause foraminal or central stenosis leading to compression of the nerve roots or spinal cord¹²⁸. Functional failure of the IVD may also lead to motion segment instability⁴³ leading to compensation from surrounding tissues¹²⁸. This functional compensation can lead to muscle strain/spasm, or degeneration of other tissues in the spine such as the facet joint, which may also contribute to pain directly^{128,131}.

1.4.4 Treatment for IVD degeneration

There are no treatments to modify or prevent the progression of IVD degeneration. Current therapeutic options address the primary clinical concern with IVD degeneration – back pain - using lifestyle, pharmacological, and surgical interventions¹³². Recent clinical guidelines suggest non-pharmacological therapies such as exercise and cognitive behavioral therapy as first-line treatments, with limited pharmacological and surgical interventions¹³². Despite these recommendations, surgical interventions (i.e. spinal fusions) for chronic LBP is on the rise despite a lack of evidence of effectiveness^{132,133}. The current lack of disease-modifying treatments for IVD degeneration and associated LBP is largely due to an incomplete understanding of the pathophysiology of the disease, and the interplay between the tissues involved.

1.5 Similarities between IVD degeneration and osteoarthritis

The IVD shares several similarities with articular cartilage, and many of the pathways associated with disc degeneration (detailed above) have also been implicated in the pathogenesis of osteoarthritis (OA)¹³⁴. OA is a heterogeneous disease that can affect multiple joints – most commonly the knee - resulting in pain and loss of function in patients ^{135,136}. It is characterized by a progressive loss of articular cartilage, subchondral bone sclerosis and synovial hyperplasia¹³⁷. Clinically, OA is typically diagnosed based on radiographic evidence of joint space narrowing, osteophyte formation, and sclerosis¹³⁸. Interestingly, these OA features are also detected in IVD degeneration¹³⁹, which has driven speculation about a potential link between the two pathologies. At the cellular level, NP cells resemble articular chondrocytes and produce a similar ECM rich in type II

collagen and aggrecan, while AF cells resemble those of the fibrous connective tissue comprising of the articular joint capsule¹³⁴. Similarities exist between the degenerative processes in IVD degeneration and OA including alterations to the ECM, biomechanics, genetic factors, inflammation, symptoms and risk factors, as recently reviewed¹³⁴. Despite these similarities, it should not be supposed that OA and IVD degeneration are identical. There are differences in both the developmental origins and biomechanical properties of each tissue. The IVD is formed by two different structures during development, the notochord and sclerotome^{140,141}. The notochord serves as the embryonic precursor to the NP¹⁴¹, while the sclerotome contributes to the AF and adjacent vertebral bones¹⁴⁰. In contrast, the articular joints of the appendicular skeleton are formed later in development, through the condensation of mesenchymal precursor cells¹⁴². While there are key similarities between these two disease states, each involves distinct tissue types and may differ in potential pathogenic pathways.

1.6 Mechanobiology of the IVD

In order for humans to remain upright, muscles of the posterior trunk contract and exert high levels of mechanical load, especially in the lumbar spine¹⁴³. As the "shock absorber" of the spine, cells within the IVD are exposed to a variety of different loading modalities such as compressive and tensile load¹⁴³. Mechanical loading is a key regulator of IVD homeostasis; multiple studies have highlighted the complexity of the cellular and molecular events that it triggers in IVD cells (as reviewed in¹⁴³). The effects of mechanical loading on IVD cell physiology are dependent on both cell type and the loading parameters applied (including load type, magnitude, frequency and duration). Several reviews have recently

described the biological response of NP and AF cells to various mechanical loads^{47,98,143,144}. This body of work describes the physiological zone in which loading promotes matrix synthesis and growth factor secretion to maintain disc health⁹⁸. Outside of this zone – either underloading or overloading - the cellular response shifts towards catabolism, marked by the induction of matrix-degrading enzymes, inflammatory cytokines, and cell apoptosis⁹⁸.

1.6.1 Mechanical loads seen by the IVD

Different compartments of the IVD experience different parameters of mechanical load and respond to these loads through different mechanisms. Due to overall anatomy of the spine and surrounding musculature and ligaments, IVDs are never completely unloaded. Instead, IVDs experience a constant level of mechanical stimulation known as preload⁵⁹. In addition to this preload, daily activity such as walking or lifting exerts additional mechanical forces on the spine that are dissipated by the IVD¹⁴³. In response to mechanical loading, the osmotic properties of the NP enable to absorb compressive loads on the spine. The consequent intradiscal pressure exerted by the NP is then dissipated by the tension that develops between the lamellae of the AF⁸¹.

1.6.2 Mechanotransduction

The process by which cells sense mechanical stress and convert it into downstream biological activity is known as mechanotransduction. Mechanotransduction occurs in three coupled, yet distinct, steps: first, the mechanical stimulus is sensed by mechanoreceptors; second, these receptors activate signal transduction pathways; and third, these pathways

induce a cellular response¹⁴⁵. While mechanotransduction pathways are not fully defined in the IVD, several targets including integrins, purinergic receptors, and transient receptor potential vanilloid 4 (TRPV4) ion channels have been implicated in the response of NP and AF cells to mechanical loading¹⁴⁶.

1.6.3 IVD tissue response to aberrant mechanical loads

While mechanical loading is essential for IVD health, overloading or underloading can lead to catabolic processes associated with IVD degeneration⁹⁸. Underloading of the spine is experienced during prolonged immobilization or in zero gravity and has been shown to accelerate IVD degeneration^{147,148}. On the opposite end of the spectrum is overloading, experienced for example during weightlifting or whole-body vibration, which likewise has been shown to induce IVD degeneration¹⁴⁷. *In vitro*, exposure to aberrant mechanical loading induces the expression of matrix-degrading enzymes, inflammatory cytokines, and cell apoptosis in both NP and AF cells⁹⁸.

1.6.3.1 Whole-body vibration

Occupational exposure to high-amplitude, low-frequency whole-body vibration - typically created by the operation of heavy equipment – has been shown to contribute to the development of IVD degeneration and back pain¹⁴⁹. Despite this, whole-body vibration (WBV) platforms designed to deliver low-amplitude, high frequency mechanical vibration are being used clinically based on their reported ability to increase bone density¹⁵⁰ and muscle strength¹⁵¹. Within the last decade, WBV has been integrated into physical therapy regimens for a variety of musculoskeletal (MSK) disorders, including osteoporosis¹⁵⁰, back

pain¹⁵², and OA¹⁵³. In addition to their clinical use, WBV platforms are marketed by the health and wellness industry as "no-workouts" to replace traditional resistance training. However, a dichotomy exists surrounding the safety of WBV: WBV is being promoted as a safe and effective treatment for MSK conditions, yet epidemiological studies have established a strong association between workplace exposure to WBV and the development of MSK conditions ¹⁴⁹. Recognizing the potential of WBV to cause tissue damage, the International Organization for Standardization (ISO) implemented guidelines indicating the maximum daily limit workers can be exposed to WBV without being at risk for injury¹⁵⁴. However, there are no regulation associated with use of commercially available WBV platforms, which are typically able to deliver vibration far greater than what ISO-2631 guidelines deem as acceptable¹⁵⁵.

Raising further concern, clinical trials have shown conflicting reports regarding the ability of WBV to slow bone loss, and its effectiveness in reducing back and OA-associated pain. In young women (15-20 years old) with low bone mineral density (BMD) daily WBV therapy increased cortical BMD in the femoral mid-shaft¹⁵⁶; however, the same researchers subsequently reported that older males and females (mean age = 82) showed no significant differences in BMD following exposure to WBV¹⁵⁷. Clinical trials investigating the effectiveness of WBV to treat back pain reported improvements in patient self-reported pain^{158,159}; however, these studies did not directly assess joint health. Based on the lack of rigorous research-based evidence, a recent analysis cautioned against the use of WBV to treat back pain¹⁵². Clinical trials evaluating the effectiveness of WBV for the treatment of joint pain in knee OA have also shown conflicting results. A number of studies reported that WBV reduced pain intensity^{153,160}, while other studies report no significant

improvement in pain intensity in knee OA patients following WBV¹⁶¹. The lack of rigorous evidence-based research surrounding the efficacy and safety of WBV is concerning given their prevalent use in physiotherapy clinics and gyms.

Given this conflicting evidence, our group carried out several studies using the mouse as a preclinical model to directly assess the effects of WBV on skeletal health. These studies demonstrated that exposure of mice to WBV using vibrational parameters used clinically induces significant IVD degeneration and OA-like damage to the knee in as little as 4-weeks^{162–164}. In addition to gross histological damage, WBV induced inflammation within the IVD marked by upregulation of $II-1\beta$ followed by upregulation of matrix degrading enzymes associated with IVD degeneration including *Mmp-3,-13* and *Adamts5*¹⁶⁵. These findings highlight the need for more comprehensive studies investigating the effects of WBV on joint health.

1.7 Metabolism of the IVD

As the largest avascular tissue in the human body, the IVD has a unique metabolic profile compared to other tissues. Aside from the outermost lamellae of the AF, the majority of the nutrients are supplied to the IVD through diffusion from the CEPs⁶⁷. As such, several factors including blood supply to the adjacent tissues, loading and matrix composition affect nutrient supply to the IVD ⁶⁷. The IVD has low cellularity and an extensive ECM that impacts nutrient transport. The major components of this matrix are collagen fibers and negatively charged proteoglycans that act as semi-permeable barrier to molecules entering the disc. While small molecules such as oxygen and glucose are able to diffuse through, the ECM limits the movement of large molecules such as growth factors and

cytokines^{67,166}. As a result of cellular metabolism and limits of diffusion, large concentration gradients of oxygen, glucose and lactic acid are observed within the IVD⁶⁷. The center of the NP is an oxygen and glucose poor environment, with high levels of lactic acid^{67,167}. The main source of ATP for AF and NP cells is through anerobic glycolysis^{168,169}, allowing NP cells to survive up to 13 days in the absence of oxygen^{170,171}. While the role of glucose and glycolytic production of ATP has been well characterized in the IVD, the role of lipid metabolism in IVD homeostasis is unknown. Of note, mediators of lipid metabolism impact articular cartilage health both negatively and positively, and have been shown to alter the progression of OA^{172,173}.

1.8 Animal models of IVD degeneration and back pain

As with many other MSK diseases, animal models are often used to understand the pathophysiology of IVD degeneration and associated back pain¹⁷⁴. A range of animal models are used to study IVD degeneration, from large animals (e.g. sheep, goats) to small rodents, that each have unique strengths and weaknesses. A common criticism of basic spine research is the use of quadrupedal animals. It is often assumed that humans experience higher mechanical loading on their IVDs due to upright posture. However, muscular contraction is the major contributor to axial compression along the spine and quadrupedal animals in fact experience higher relative compressive loading on their spines compared to bipedal animals^{174–176}. Spinal tissues from large animal models such as sheep, and goats have a similar size, anatomy, mechanical loading and cellularity to those from humans¹⁷⁵, and have been used extensively for models of surgically-induced IVD degeneration¹⁷⁵. Despite these advantages, cost and associated limitations to sample size

in experiments is a major deterrent for research involving large animals¹⁷⁷. Most basic research investigating IVD degeneration use small animal models. Rodents are most frequently used based on their ease of use, low cost and the ability to manipulate their genome¹⁷⁴. Despite the size difference between human and mouse IVDs, both species have similar geometry, biomechanics and extracellular matrix composition^{178–180}. Aside from their low housing costs and short generational times, one of the largest advantages of mouse models is the relative ease of genetic manipulation. Use of the *Cre/LoxP* recombinase system allows for modification of the murine genome in either a global or tissue specific manner^{181–183}. A number of studies using genetically modified mice have highlighted the importance of matricellular and ECM components including CCN2, SPARC, and collagen type IX in IVD degeneration and back pain^{184–186}.

In addition to genetic manipulation, mouse models have been used extensively for pain research¹⁸⁷. Given the discordance between IVD degeneration and back pain, it is important to simultaneously assess both in preclinical models of disease¹⁸⁸. While pain is a complex phenomenon, and limitations of mouse models should be acknowledged, their use has provided a number of promising therapeutic targets for the treatment of chronic pain¹⁸⁹. To assess pain, several behavioral metrics have been developed for various modalities of pain^{187,188}. Generally, these tests assess for behavioral indicators of evoked or spontaneous pain and are used extensively in the pain field¹⁸⁷. In addition to behavioral metrics, mouse models also allow for the assessment of biological indicators associated with pain, such as neuroinflammation¹⁹⁰. The use of animal models has provided researchers with valuable knowledge about potential mechanisms contributing to IVD degeneration and LBP and may potentially yield future therapeutic targets.

Obesity – traditionally defined as a body mass index over 30 – is a worldwide epidemic. Obesity substantially increases the risk of developing metabolic, cardiovascular, neurological and MSK diseases¹⁹¹. With the prevalence of obesity nearly tripling over the last 30 years¹⁹¹, it poses a large public health concern. In the US, it is estimated that 21% of the total healthcare expenditure is used to directly treat obesity and obesity-related comorbidities¹⁹². At the individual level, obesity not only decreases life expectancy¹⁹³ but also decreases quality of life and leads to disability, mental illness and unemployment^{191,194}.

1.9.1 Metabolic syndrome

Although several factors contribute to obesity including hormonal, nutritional, psychological and metabolic factors, obesity is fundamentally caused by an imbalance in energy input and expenditure^{191,195} resulting in an accumulation of adipose tissue. Evolutionarily, the ability to store energy for long periods of time was essential for survival. However, in today's industrial world where food is abundant and lifestyles are sedentary, fat accumulation has negative health implications. As mentioned previously, obesity – or more specifically the accumulation of abdominal fat – is related to the development of several metabolic diseases, together termed metabolic syndrome¹⁹⁶. Components of metabolic syndrome include abdominal obesity/large waist circumference, type II diabetes mellitus, hypertension, and dyslipidemia^{191,196,197}. Together, these conditions have a tremendous impact on overall systemic health, and increase the risk of developing cardiovascular¹⁹⁸, renal¹⁹⁹ and MSK²⁰⁰ pathologies. Although the pathogenesis

of metabolic syndrome is complex, abdominal obesity appears to be a key causative factor²⁰¹. Adipose tissue not only stores and releases fatty acids, but is also is a major endocrine organ that synthesizes and secretes proteins and hormones termed adipokines (e.g. leptin, adiponectin, visfatin, resistin) and inflammatory cytokines (e.g. TNF- α , IL-6)^{202,203}. Increased release of these molecules is associated with the development of many aspects of metabolic syndrome including insulin sensitivity, hypertension and dyslipidemia independently^{204,205}. However, this link between obesity and metabolic syndrome appears to be multifactorial, as increased adiposity is also associated with increase plasma levels of free fatty acids (FFAs), which can independently contribute to insulin sensitivity, hypertension and dyslipidemia²⁰⁶.

1.9.2 Musculoskeletal health

Obesity impacts several MSK tissues, including muscle, bone, and cartilage. Clinically, obesity is associated with an increased risk of sarcopenia, osteoporosis, OA, LBP and IVD degeneration^{207,208}. While the mechanisms underlying these associations are not fully elucidated, obesity and obesity-associated diseases likely contribute through direct actions on the tissue of interest, as well as indirectly through effects on supporting tissues. A recent review by Zhuo *et al*²⁰⁰ highlighted the differential impact of each component of metabolic syndrome in the context of OA. A common indirect mechanism believed to impact joint health is cardiovascular complications – postulated to lead to subchondral bone ischemia and impaired nutrition of the joint. Hypertension¹⁶², dyslipidemia^{79,209} and type II diabetes^{210,211} can lead to microvascular narrowing, atherosclerosis and endothelial dysfunction^{79,209,210}, potentially resulting in subchondral bone ischemia. Lack of blood

supply could compromise nutrient exchange ultimately leading to cartilage degeneration^{200,212}. In humans, hypertension, dyslipidemia, and type II diabetes are also independently associated with increased risk of IVD degeneration^{213–215}. While the exact mechanisms have not been determined, impaired nutrient exchange has been implicated as a mediator of IVD degeneration⁶⁷, and may be caused by these conditions. Aside from these indirect effects, components of metabolic syndrome may impact joint health directly through abnormalities in lipid/glucose metabolism and release of systemic factors^{200,216}. Dyslipidemia may contribute to IVD degeneration through alterations in the lipid metabolism pathway. In chondrocytes, lipid metabolism regulates viability, matrix synthesis and inflammation²¹⁷. While the role of lipid metabolism in the IVD is not well characterized, dysregulation of metabolic pathways in hypertriglyceridemia is associated with IVD degeneration, caused by IVD cell apoptosis and matrix catabolism²¹⁴. Hyperglycemia resulting from type II diabetes is thought to contribute to IVD degeneration through increased production of reactive oxygen species (ROS) and advanced glycation end products (AGEs)²⁰⁰. *In vitro*, hyperglycemic conditions induce ROS production in both articular chondrocytes²¹⁸ and NP cells²¹⁹, leading to matrix degeneration and apoptosis in both tissues^{219,220}. AGEs are products of non-enzymatic glycation of proteins that accumulate with uncontrolled hyperglycemia²²¹. In humans, AGEs accumulate in IVDs of people with type II diabetes²²², which is thought to contribute to IVD degeneration through increased protein crosslinking, causing increased mechanical stiffness of the IVD²²³. Supporting this, dietary consumption of AGEs in mice was associated with increased IVD degeneration and altered biomechanics of spinal motion segments²²⁴. While biomechanical changes may play a role in AGE-induced IVD degeneration, AGEs may also activate the

receptor for advanced glycation end products (RAGE), which has been found to activate pro-inflammatory and catabolic processes in chondrocytic cells^{225,226}.

Studies investigating the role of adipokines have also highlighted the importance of secreted factors in obesity-associated pathologies. For example, leptin-deficient mice become obese, yet they do not develop knee OA, suggesting leptin may play a key role in obesity-induced OA²²⁷. In the IVD, exposure of NP cells to adipokines, such as leptin and resistin, promotes catabolic metabolism associated with increased expression of matrix remodeling enzymes such as MMP and ADAMTS genes^{228,229}.

1.9.3 Pain

In addition to the induction of degenerative joint diseases, obesity is also associated with chronic pain conditions, such as fibromyalgia, headaches and abdominal pain²³⁰. While the underlying mechanisms linking obesity and chronic pain remains unknown, it has been suggested that systemic immune and endocrine alterations play a role in the altered pain response²³¹. Obesity is considered a low-grade chronic systemic inflammatory state with elevated levels of IL-6 and C-reactive protein (CRP) that may contribute to pain^{231,232}. Pain is a biopsychosocial phenomenon⁴, so, in addition to these biological factors, obesity is also associated with depression²³³ and poor sleep quality²³⁴. These conditions are both known to contribute to pain through various mechanisms^{235,236}. A summary of the proposed mechanisms linking obesity with IVD degeneration and associated pain can be seen in **Figure 1.3**.



Figure 1.3: Proposed mechanism underlying the association between obesity/ metabolic syndrome, IVD degeneration, and pain.

The link between obesity, IVD degeneration and back pain is multifactorial, involving both direct effects on the IVD and indirect effects on the nervous system and other tissues. Obesity may directly contribute to structural IVD degeneration through increased mechanical loading, systemic inflammation and adipokine signaling. Metabolic dysregulation associated with obesity leading to diabetes, hypertension and dyslipidemia may compound these direct effects and potentiate further degeneration. Indirectly, obesity and obesity-related pathologies may also alter pain-processing pathways contributing to back pain. Furthermore, damage/dysfunction of adjacent tissues (i.e. vasculature, muscle, vertebral bone, facet joints) caused by obesity may lead to both structural IVD degeneration and back pain. Figure based on content provided in¹⁹⁹.

1.10 Nuclear Receptors

1.10.1 Structure

Nuclear receptors are a large family of ligand-regulated transcription factors activated by binding of steroid hormones, such as estrogen and progesterone, and lipid-soluble signals, including retinoic acid, oxysterols, and thyroid hormone, to alter transcription of downstream target genes²³⁷. Although nuclear receptors have different, and often opposing actions, they share a common structure. All nuclear receptors contain 4 domains, including the N-terminal domain that is of variable size and sequence, a conserved DNA binding domain, a variable hinge domain, and a ligand binding domain ²³⁷. Some nuclear receptors contain an activation function-1 (AF-1) domain at the N-terminal domain, which is a ligand-independent domain that assists in the recruitment of coactivators²³⁸. The centrally located DNA binding domain is the most conserved segment of nuclear receptors; it contains two zinc fingers that binds to hormone response elements on the promoter of downstream target genes²³⁹. The hinge domain connects the DNA binding domain to the ligand-binding domain, which is a 12 helixed structure containing a ligand-dependent activation-function 2 (AF-2) domain²³⁷. A major characteristic of the ligand binding domain is a hydrophobic cavity that allows lipophilic ligands to be shielded from the hydrophilic environment²³⁷. These pockets allow selectivity for specific ligands through variations in size and amino-acid composition²⁴⁰. Binding of the ligand stabilizes the surface elements of the ligand binding domain and allows for coactivator/corepressor binding, altering downstream gene transcription^{237,241,242}.

Nuclear receptors are divided into 4 main classes. Class I nuclear receptors are normally located in the cytoplasm bound to chaperone proteins, but through ligand activation are translocated to the nucleus where they can bind DNA and affect gene transcription²⁴³. Class I nuclear receptor ligands are steroid hormones such as estrogen, testosterone and corticoids²⁴⁴. Class II nuclear receptors on the other hand, remain in the nucleus regardless of ligand-activation²⁴³. Class II nuclear receptors are commonly found as heterodimers with retinoid X receptor (RXR), and include Peroxisome Proliferator Activated Receptors (PPARs), Liver X receptor (LXR), Farnesoid X receptor (FXR) and Vitamin D receptor (VDR)²⁴⁵. Class III nuclear receptors are similar to class II, however they typically exist as homodimers on their response elements²⁴³. Class IV nuclear receptors also have a similar mechanism of action to class II nuclear receptors, but typically are found as monomers on DNA²⁴³.

Many class II receptors respond to dietary lipids such as fatty acids (PPARs), sterols (LXR), bile acid (FXR), serving as "lipid sensors"²⁴⁶. Activation of these receptors occurs at physiological lipid concentrations influenced by dietary intake and leads to changes in transcription of genes involved in lipid metabolism, storage, transport and elimination²⁴⁶. While ligand binding is the major activator of nuclear receptors, their activity can also be modulated by posttranslational modifications, including phosphorylation, ubiquitination, and SUMOylation²³⁹.

1.10.2 Peroxisome Proliferator Activated Receptors (PPARs)

The PPARs are a subfamily of nuclear receptors whose ligands include fatty acids and their metabolites, which are of interest as they are dysregulated in obesity^{200,247}. As a type II

nuclear receptor, PPARs form heterodimers with RXR, and when bound to their ligands they recruit co-activators or co-repressors to either activate or repress downstream gene expression depending on specific gene and cellular context^{243,248}. There are three PPAR isoforms (alpha, delta, gamma) that play distinct roles in energy balance and metabolism²⁴⁹. The two most studied isoforms, PPAR α and PPAR γ , are highly expressed in the liver and adipose, respectively. Agonists of these receptors are currently being used clinically for the treatment of a variety of obesity-associated conditions including dyslipidemia and type II diabetes mellitus^{250,251}. Studies examining the role of PPARy in murine cartilaginous tissues have shown that a cartilage-specific knockout of PPARy leads to spontaneous OA development²⁵². In contrast to the tissue-specific expression of other PPARs, PPAR δ has broad expression patterns and affects glucose and lipid metabolism, cell differentiation, proliferation, apoptosis and immune regulation²⁵³. Recent studies in mice have shown PPARδ agonism improves insulin resistance, fat burning, and muscle endurance^{247,254,255}; however, PPARS agonism has also been shown to induce catabolic processes within articular cartilage^{172,256}. In a murine organ culture model, activation of PPAR δ lead to proteoglycan degradation in articular cartilage through increased expression of proteases including *Mmp-3* and *Adamts-5*^{172,256}. Moreover, mice with cartilage-specific deletion of PPAR δ are protected from cartilage damage in a post-traumatic model of OA¹⁷². While the biological mechanisms behind the role of PPAR δ in articular cartilage are not fully understood, it appears to contribute to the regulation of lipid metabolism and the cellular response to oxidative stress²⁵⁶. Although the role of PPARs in the IVD have not been directly investigated, PPAR expression is detected in both the NP and AF²⁵⁷. The similarities between articular cartilage and IVD biology and pathophysiology suggest that PPARs may be intriguing candidates as potential regulators of IVD degeneration.

1.11 Overall objectives and hypothesis

Using the mouse as a preclinical model, this thesis focuses predominantly on examining the role of two important environmental risk factors in the initiation and progression of IVD degeneration and associated pain: mechanical loading and obesity.

1.11.1 Objective #1

Investigate whether the detrimental effects of WBV on joint health are altered by genetic background in mouse models.

1.11.1.1 Rationale #1

Previous research by our group demonstrated that exposure of mice to protocols of WBV that model those used clinically (30 minutes/day, 5 days/week; 45 Hz, 0.3 g) induced damage to the IVD and knee joint within two weeks, that progressed in severity with continued daily exposure to WBV (i.e. 4, 8 weeks)^{162,164,165}. These studies were however limited to skeletally mature (10-week old) male mice of a single outbred stock (CD1). Due to the diverse human population using WBV platforms we aimed to determine whether the response of joint tissues to WBV was consistent between mice of different genetic backgrounds.

1.11.1.2 Hypothesis # 1

The effects of whole-body vibration on joint health will differ based on genetic background.

1.11.2 Objective #2

Determine whether chronic exposure of mice to a high-fat or high-fat/high-sugar western diet will accelerate age-related IVD degeneration and/or back pain.

1.11.2.1 Rationale #2

Throughout the world, the prevalence of obesity is on the rise. In 2014 over 25% of Canadians were obese, resulting in an enormous toll on our health care system of over \$4 billion spent annually on direct costs alone, let alone costs associated with obesity-associated co-morbidities²⁵⁸. In addition to being a major risk factor for the development of type II diabetes, cardiovascular disease and metabolic syndrome²⁵⁹, obesity also increases the risk of IVD degeneration and lower back pain^{260,261}. Despite this association there is very little known about the pathogenesis of obesity induced IVD degeneration. Previous research demonstrated that obesity induced by a high-fat diet accelerates OA in mouse and rat models^{262–266}. In addition to structural changes in the joint, the high-fat diet also induces symptomatic characteristic of OA, such as hyperalgesia²⁶³. To date, the effect of diet-induced obesity on IVD degeneration and back pain has not been directly studied.

1.11.2.2 Hypothesis # 2

Chronic consumption of a high-fat or high-fat/high-sugar western diet will accelerate IVD degeneration and/or behavioral indicators of back pain in the mouse model.

1.11.3 Objective # 3

Determine the role of PPAR δ in age- and obesity-associated IVD degeneration and investigate sex as a variable in the mouse model system.

1.11.3.1 Rationale # 3

Based on previous research demonstrating that loss of PPAR δ in articular cartilage protects from surgically-induced and age-associated OA^{172,267}, we sought to determine the role of PPAR δ in the IVD and is contribution to both age- and obesity-associated IVD degeneration and pain. Previous studies by our lab using genome-wide transcriptome analysis demonstrated that many nuclear receptor pathways are activated in ageassociated IVD degeneration, including the PPAR signaling pathway²⁵⁷, highlighting the potential importance of PPAR in age-associated IVD degeneration. Given that its endogenous ligands are dysregulated in obesity²⁴⁷, and the detrimental role of PPAR δ activation in articular cartilage¹⁷², we aimed to determine whether similar findings would be seen between the different models of IVD degeneration.

This study also sought to address potential sex-related differences in IVD degeneration and back pain in our model of diet-induced disc degeneration. Numerous pre-clinical and clinical studies have shown that IVD degeneration and back pain are sexually dimorphic, where females are disproportionately more affected by spinal pathologies and back pain than males^{95,268}.

1.11.3.2 Hypothesis # 3

Loss of PPAR δ in IVD cells will delay the progression of age- and obesity-related IVD degeneration and back pain.

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Chapter 2

C57BL/6 mice are resistant to joint degeneration induced by whole-body vibration.

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2 C57BL/6 mice are resistant to joint degeneration induced by whole-body vibration.

2.1 Co-authorship Statement

G.J. Kerr performed most of the experiments, contributed to study design and wrote the manuscript. J.K. Branch assisted with the *in vivo* vibration and sectioning of the lumbar spines (**Figure 2.1**). M.R. McCann assisted with histopathological scoring for IVD degeneration (**Figure 2.1**). A. Ratneswaran, and M.A. Pest conducted histopathological scoring on the knee joints (**Figure 2.2**). Drs. D.W. Holdsworth, F. Beier, S.J. Dixon and C.A. Séguin contributed to the study design and editing of the manuscript. All authors read and approved the submitted version of the manuscript.

2.2 Chapter Summary

Objective: Whole-body vibration (WBV) platforms are commercially available devices that are used clinically to treat numerous musculoskeletal conditions based on their reported ability to increase bone mineral density and muscle strength. Despite widespread use, there is an alarming lack of understanding of the direct effects of WBV on joint health. Previous work by our lab demonstrated that repeated exposure to WBV using protocols that model those used clinically, induces intervertebral disc (IVD) degeneration and osteoarthritis-like damage in the knee of skeletally mature, male mice of a single outbred strain (CD-1). The present study examined whether exposure to WBV induces similar deleterious effects in a genetically different strain of mouse (C57BL/6).

Design: Male 10-week-old C57BL/6 mice were exposed to vertical sinusoidal WBV for 30 min/day, 5 days/ week, for 4 or 8 weeks using previously reported protocols (45 Hz, 0.3 g peak acceleration). Following WBV, joint tissues were examined using histological analysis and gene expression was quantified using real-time PCR (qPCR).

Results: Our analyses show a lack of WBV-induced degeneration in either the knee or IVDs of C57BL/6 mice exposed to WBV for 4 or 8 weeks, in direct contrast to the WBV-induced damage previously reported by our lab in CD-1 mice.

Conclusions: Together with previous studies from our group, the present study demonstrates that the effects of WBV on joint tissues vary in a strain-specific manner. These findings highlight the need to examine genetic or physiological differences that may underlie susceptibility to the deleterious effects of WBV on joint tissues.

2.3 Introduction

Platforms that deliver whole-body vibration (WBV) are currently used in the clinical setting based on their reported ability to increase bone density¹ and muscle strength². Within the last decade, WBV has been integrated into physical therapy regimens for a variety of musculoskeletal disorders, including osteoporosis¹, back pain³, and osteoarthritis⁴. In addition to their clinical use, WBV platforms are promoted in the health and wellness industries as "no-work workouts" equivalent to traditional resistance training. The proposed use of WBV as a treatment for musculoskeletal conditions appears contradictory to epidemiological studies establishing an association between workplace exposure to WBV and the development of several conditions, including back pain⁵. Recognizing the potential of WBV to induce tissue damage, the International Organization for Standardization (ISO) implemented guidelines defining the maximum daily limit of WBV that workers can be exposed to without risk for injury⁶. Conversely, commercially available WBV platforms are currently unregulated, with some able to deliver accelerations that exceed current ISO-2631 recommended guidelines⁷.

Raising further concern regarding the use of WBV, clinical trials show conflicting reports regarding the ability of WBV to increase bone mineral density^{1,8}. Similarly, clinical trials investigating the effectiveness of WBV for knee or back pain have failed to show conclusive evidence, with only a small proportion of trials showing benefits based on assessments of patient self-reported pain^{3,9}. A cause for concern is that the widespread use of WBV has not been validated with rigorous research-based evidence addressing the broad range of parameters used and potential harmful effects.

Investigating the effects of WBV on joint tissues, our research has demonstrated that exposure of mice to protocols of low-amplitude, high-frequency WBV that model those used clinically induced intervertebral disc (IVD) degeneration and osteoarthritis-like damage in the knee of skeletally mature, male mice, of a single outbred strain $(CD-1)^{10-12}$. Given these strikingly deleterious effects of WBV, the present study aimed to determine if the response of joint tissues to WBV was consistent in age- and sex-matched C57BL/6 mice representing a distinct genetic background.

2.4 Methods

2.4.1 Whole body vibration

Based on WBV parameters used in clinical protocols and our previous studies^{10–12}, 10week-old male C57BL/6 mice (Charles River) were subjected to vertical sinusoidal vibration (45Hz, peak-to-peak amplitude 74 µm, peak acceleration 0.3 g) 30 minutes/day, 5 days/week for 4 or 8 weeks. Age- and sex-matched controls were housed in identical chambers on a non-vibrating sham platform to replicate handling and environmental conditions. Following WBV, mice were returned to conventional housing and monitored daily. No differences in total body mass were detected between groups. Mice were euthanized with sodium pentobarbital 24 hours after final exposure to WBV. The Council on Animal Care at Western University approved all procedures, in accordance with the Canadian Council on Animal Care and the ARRIVE guidelines.

2.4.2 Gene expression analysis

Thoracic IVDs (T10-T15) were isolated from mice exposed to 8 weeks WBV and sham controls, placed in TRIzol (Life Technologies) and homogenized using a PRO250 tissue homogenizer (PRO Scientific). RNA was extracted according to manufacturer's instructions, quantified using a NanoDrop 2000 spectrophotometer (Thermo Scientific), and 0.5 µg was reverse transcribed into complementary DNA (cDNA) (iScript; Bio-Rad). Gene expression was assessed by real-time PCR (qPCR) using the Bio-Rad CFX384. PCR analyses were run in triplicate using 120 ng of cDNA per reaction and 310 nM forward and reverse primers with 2x SsoFast EvaGreen Supermix (Bio-Rad) using previously optimized PCR parameters and primers¹¹. Transcript levels were calculated using $\Delta\Delta$ Ct, with data normalized for input based on Ribosomal protein S29 (*Rps29*) and expressed relative to non-vibrated sham controls.

2.4.3 Histological analysis

Intact lumbar spine segments (L1-L5) and knees were isolated, fixed and paraffin embedded as previously described¹¹. Spines were sectioned sagittally and knees were sectioned coronally, at a thickness of 5 μ m using a microtome (Leica Microsystems). Spines sections were stained using 0.1% Safranin-O/0.05% fast green, and knee sections were stained with 0.04% Toluidine Blue. Sections were imaged on a Leica DM1000 microscope, with Leica Application Suite (Leica Microsystems). To evaluate IVD degeneration, mid-sagittal sections were scored using the modified Thompson grading scheme to assess IVD health based on criteria specific to each tissue compartment (scores 1-4, where a lower score corresponds to a more healthy tissue), as previously reported^{10,11}. Knee joint health was assessed using the murine Osteoarthritis Research Society International (OARSI) histopathological scale¹³. For each mouse, the four quadrants of the knee joint were evaluated in 10 serial sections taken 75 μ m apart to include most of the weight-bearing area of the femorotibial joint. For each quadrant of each section individual scores were averaged between three independent blinded observers and summed across the 10 sections for each mouse. The whole joint score was calculated as the sum of the quadrant scores for each mouse.

2.4.4 Statistical analysis

Data are from experiments conducted with n = 5-6 mice per group at each time point. For qPCR analyses, data from mice exposed to WBV were compared to non-vibrated sham controls using a parametric Welch's t-test. At each time point, histopathological scores from the IVD or knee joint were compared using a non-parametric Mann-Whitney U-test. Differences were accepted as statistically significant at P < 0.05.

2.5 Results

To examine the effects of repeated exposure to WBV on joint health, 10-week-old C57BL/6 mice were exposed to WBV for 30 min/day, 5 days/week for either 4 or 8 weeks, using parameters that model clinical use in humans (45 Hz, 0.3 g)¹¹. Importantly, these parameters were previously found to induce degeneration of both the IVD and knee joint in age- and sex-matched CD-1 mice, marked by histological hallmarks of tissue damage including annulus fibrosus degeneration in the IVD as well as meniscal damage and focal defects in the articular surface of the knee^{10–12}. In the C57BL/6 mice, we first assessed

IVD health. No discernable differences were detected in the histological appearance of lumbar IVDs in mice exposed to 4 or 8 weeks of WBV compared to their respective sham controls (**Figure 2.1A**). Accordingly, evaluation of IVD structure using the modified Thompson score showed no significant differences in degeneration grade between mice exposed to 4 or 8 weeks of WBV and their respective sham controls (**Figure 2.1B**). As a more sensitive measure of early degenerative changes in IVD tissues, we evaluated the cellular response to WBV using real-time PCR. Similar to histological evaluation, this analysis revealed no change in the expression of anabolic (*Acan, Col1a1, Col2a1, Sox9*) or catabolic (*Mmp3, Mmp13, Adamts4, Adamts5*) factors between mice exposed to 8 weeks of WBV and non-vibrated sham controls (**Figure 2-1C**).

We then assessed if exposure of C57BL/6 mice to WBV induced changes to knee joint tissues. Consistent with findings in the IVD, no discernable differences were detected in the knee joint of C57BL/6 mice exposed to 4 or 8 weeks of WBV compared to their respective sham controls (**Figure 2-2A**). In contrast to our previous studies in CD-1 mice¹⁰, exposure of C57BL/6 mice to WBV did not induce any observable meniscal damage, articular cartilage erosion or osteophyte formation. Evaluation of cartilage degeneration using the OARSI histopathological score supported these observations; no significant differences were found in the summed OARSI score between mice exposed to WBV and sham controls (**Figure 2-2B**).



Figure 2.1: Effect of repeated exposure to WBV on IVD health.

A. Representative sagittal sections of the lumbar spine stained with safranin-O/fast green from mice exposed to WBV (for 4 or 8 weeks, 30 minutes per day, 5 days per week, at 45 Hz and 0.3 g) and non-vibrated sham controls. Images are oriented with rostral portion of the lumbar spine on top and dorsal portion on the left. B. Histopathological scores generated using the modified Thompson grading scale demonstrated no significant differences in degeneration score corresponding to the annulus fibrosus (AF) or nucleus pulposus (NP) from mice exposed to either 4 or 8 weeks WBV compared to their respective non-vibrated sham controls. C. Real-time PCR analysis of IVD gene expression in mice exposed to 8 weeks WBV and sham controls. Expression of extracellular matrix genes (*Acan, Col1a1, Col2a1*), the transcription Sox9, and matrix remodeling enzymes (*Mmp3, Mmp13, Adamts4, Adamts5*) were determined relative to the expression of the housekeeping gene *Rps29* and normalized to that of non-vibrated sham controls. Data are presented as mean \pm 95% CI (n = 5-6 mice per group).





A. Representative coronal sections of the medial knee compartment stained with toluidine blue from mice exposed to WBV (30 minutes per day, 5 days per week, for 4 or 8 weeks at 45 Hz and 0.3 g) and non-vibrated sham control mice. Images are oriented with the medial femoral condyle located superiorly, and the medial tibial plateau inferiorly. B. Histopathological grading of the knee joints using the murine Osteoarthritis Research Society International (OARSI) scale. Joints from each mouse were evaluated using 10 serial sections (taken 75 μ m apart) and the summed OARSI scores are presented corresponding to the medial femoral condyle (MFC), medial tibial plateau (MTP), lateral femoral condyle (LFC) and lateral tibial plateau (LTP), which were then combined to generate the summed score for the whole joint (WJ). No significant differences in degeneration scores were detected between mice exposed to WBV and sham controls in any of the compartments investigated. Data are presented as mean ±95% CI (n = 5-6 mice per group).

Despite the widespread clinical and recreational use of WBV platforms, there is an alarming lack of understanding of the effects of WBV on overall joint health as well as the effects elicited in specific joint tissues. Previous studies from our group demonstrated the ability of WBV to induce IVD degeneration and osteoarthritis-like damage in the knee of skeletally mature male outbred CD-1 mice¹¹. After 4 weeks of repeated exposure to WBV, CD-1 mice showed degeneration in the annulus fibrosus of the IVD, marked by disruption of collagen organization, enhanced MMP-mediated collagen and aggrecan degradation, and increased cell death. Similarly, 4 of 5 CD-1 mice exposed to WBV for 4 weeks showed meniscal damage, and 2 of 5 showed focal damage to the articular cartilage associated with increased MMP-mediated matrix degradation¹¹. Subsequent studies from our group showed that WBV-induced damage to both the IVD and knee joint increased in severity when CD-1 mice were exposed to 8 weeks WBV^{10,12}. In direct contrast, the current study demonstrates that exposure of C57BL/6 mice to those same parameters of WBV did not induce degeneration-associated changes in either the knee or IVD after 4 or 8 weeks of exposure. Thus, the effects of WBV on joint tissues differ between different strains of mice, highlighting the need to examine factors that may determine susceptibility to the deleterious effects of mechanical loading on joint health.

Previous work has reported strain-specific differences in the attenuation of vibration loading in rodent models, as well as its downstream consequences on bone parameters^{14,15}. Studies investigating vibration frequency thresholds in rodents reported that the resonant frequency range differs between strains of mice¹⁴. At the resonant frequency, it is assumed

that there is maximal displacement and consequently strain experienced by skeletal structures, whereas frequencies outside of this range of vibration are attenuated by the body¹⁴. When anesthetized mice were exposed to vibration frequencies ranging from 0-600 Hz (at a constant magnitude of 0.3 m/s²), C57BL/6 mice showed the least amount of attenuation between 41-50 Hz, whereas CD-1 mice showed the least amount of attenuation between 51-60 Hz, suggesting difference in their resonant frequency ranges¹⁴. While the biological factors underlying these differences were not explored, it has been suggested that differences in baseline bone parameters between mouse strains may alter the mechanical forces experienced at the tissue level¹⁵. Studies investigating the response of trabecular bone to whole-body vibration (45 Hz, 0.25 g; 10 min/day, 3 weeks) demonstrated that low bone mineral density C57BL/6 mice showed significantly higher bone formation rates induced by WBV compared to mid-density BALB/cByJ mice¹⁵. Moreover, C3H/HeJ mice which showed the highest bone mineral density, showed no changes in bone formation rates following exposure to the same parameters of WBV¹⁵.

Clinical trials in humans report that women with lower body mass showed a greater inhibition of bone loss when exposed to WBV compared to women with higher body mass; since bone mineral density is positively correlated with body mass the authors suggested that women with low bone mineral density may be most responsive to mechanical loading to prevent bone loss¹. Attenuation of vibration may depend on bone density and other physiological factors including activity levels, body mass or joint structure, may alter the mechanical loading experienced at the tissue or cellular level, and potentially contribute to the differential response of joint tissues to WBV that we have shown between the CD-1 and C57BL/6 strains. Of note, the C57BL/6 mice used in the current study had an average

mass of 26 g at 18 weeks of age, compared to the CD-1 mice used in the previous study which had an average mass of 41 g at 18 weeks of age. Further studies are required to assess the relative contribution of biological mediators that might underlie the differences in the effects of WBV on joint tissues in different strains of mice, allowing for potential extrapolation to the clinical setting.

In summary, this study showed that repeated exposure of skeletally mature male C57BL/6 mice to WBV for 4 or 8 weeks did not alter overall health of the IVD and knee joints; in direct contrast to the WBV-induced joint degeneration we previously reported in CD-1 mice of the same age and sex. Although the results of the current study suggest that C57BL/6 mice are resistant to WBV-induced damage, it should be noted that different frequencies and accelerations may lead to differential responses. Previous studies in musculoskeletal tissues established that mechanical loading can induce both catabolic and anabolic responses, dependent on the load type, frequency, amplitude and exposure time¹⁶. Previous research by our group has shown the potential of WBV to induce anabolic effects in the IVD after acute exposure¹⁷. Together, these findings highlight the need for more rigorous investigations to examine the efficacy and safety of WBV in the diverse human populations currently using them, with emphasis on quantifying potential beneficial or detrimental effects on musculoskeletal tissues.

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Chapter 3

Diet-induced obesity leads to behavioral indicators of pain preceding structural joint degeneration in wild-type mice.

Chapter 3 is authored by Kerr, G.J, To, B., White, I., Millecamps, M., Beier, F., Stone, L.S., and Séguin, C.A. and is entitled "Diet-induced obesity leads to behavioral indicators of pain preceding structural joint damage in wild-type mice". This study is in preparation for submission.

3 Diet-induced obesity leads to behavioral indicators of pain preceding structural joint degeneration in wild-type mice.

3.1 Co-authorship Statement

G.J. Kerr performed most of the experiments, contributed to study design and wrote the manuscript. I. White assisted with scoring the tail suspension (Figure 3.2) and cold sensitivity assay (Figure 3.3). B. To conducted all histological and histomorphometry analysis on the knee joints and G. J. Kerr assisted with histopathological scoring. (Figure 3.5). I. White assisted with sectioning, staining and analysis of spinal cord sections (Figure 3.6). Manuscript was written by G.J. Kerr with suggestions from Dr. C.A. Séguin.

3.2 Chapter Summary

Introduction: Obesity is one of the largest modifiable risk factors for the development of musculoskeletal diseases, including intervertebral disc (IVD) degeneration and back pain. Despite the clinical association, no studies have directly assessed whether diet-induced obesity accelerates IVD degeneration, back pain, or investigated the biological mediators underlying this association. In this study we examine the effects of chronic consumption of a high-fat or high-fat/high-sugar (western) diet on the IVD and pain-associated outcomes.

Methods: Male C57BL/6N mice were randomized into one of three diet groups (chow control; high-fat; high-fat, high-sugar western diet) at 10-weeks of age and remained on the diet for 12, 24 or 40 weeks. At endpoint, animals were assessed for behavioral indicators of pain, joint tissues were collected for histological and molecular analysis, and IBA-1, GFAP and CGRP were measured in spinal cords by immunohistochemistry .

Results: Animals fed obesogenic (high-fat or western) diets showed behavioral indicators of pain beginning at 12 weeks and persisting up to 40 weeks of diet consumption. Histological indicators of joint degeneration were not detected in the IVD or knee until 40 weeks on the experimental diets. Mice fed the obesogenic diets showed increased intradiscal expression of inflammatory cytokines and circulating levels of MCP-1 compared to control. Linear regression modeling demonstrated that age and diet were both significant predictors of most pain-related behavioral outcomes, but not histopathological joint degeneration.

Conclusion: Diet-induced obesity accelerates IVD degeneration and knee OA in mice; however, pain-related behaviors precede and are independent of histopathological structural damage. These findings contribute to understanding the source of obesity-related back pain and the contribution of structural IVD degeneration.

3.3 Introduction

Obesity – traditionally defined as a body mass index over 30 – is a worldwide epidemic. Obesity substantially increases the risk of developing metabolic, cardiovascular, neurological and musculoskeletal diseases¹, and with the prevalence nearly tripling over the last 30 years¹, it poses a large public health concern. Obesity decreases both life expectancy² and quality of life, and is associated with increased disability, mental illness and unemployment^{1.3}. A significant contributor to obesity-induced disability is low back pain (LBP) ^{4.5}, which is the single most common cause of long-term pain and disability worldwide⁶. Despite efforts to improve the clinical management of LBP, treatments are limited to symptomatic relief, often without treating the underlying cause of the pain⁷. This is largely due to an incomplete understanding of the tissues and pathways involved in the initiation and progression of LBP. While several tissues appear to be involved in LBP, including the paraspinal muscles, ligaments, and facet joints^{8–10}, degeneration of the fibrocartilaginous intervertebral disc (IVD) is believed to be the major contributor to pain in an approximately 40% of cases⁸.

Despite the clinical associations between LBP, IVD degeneration, and obesity, the underlying mechanisms and biological pathways responsible remain unknown. One contributing factor appears to be increased mechanical loading. Excess weight alters the mechanical load experienced by the IVD¹¹, a known regulator of IVD cellular function^{12,13}. Increased body weight is associated with indices of lumbar disc degeneration including disc space narrowing and decreased lumbar disc signal intensity detected by MRI^{14,15}. In articular cartilage, excess weight and altered mechanical loading has also been suggested to contribute to osteoarthritis (OA)¹⁶, a degenerative musculoskeletal disease with many similarities to IVD degeneration¹⁷. Of note, increased mechanical load alone does not account for the association between obesity and OA, as obese individuals also present more frequently with OA in non-weight bearing joints, such as the hand¹⁸.

In addition to increased mechanical load, metabolic abnormalities associated with obesity impact musculoskeletal health^{19,20}. Obesity is associated with chronic metabolic disorders including hypertension, diabetes mellitus and dyslipidemia, collectively known as metabolic syndrome²¹. In the context of OA, it is postulated that each component of metabolic syndrome may independently contribute to disease progression, as comprehensively reviewed by Zhuo et al.²² Specifically, alterations in the release of systemic factors (inflammatory cytokines, adipokines), nutrient exchange, advanced glycation end-products (AGEs) levels and glucose/lipid metabolism are believed to be major contributors to OA progression^{19,22}. Studies from multiple groups have demonstrated using mouse models that obesity induced by a high-fat diet accelerates the progression of both age- and surgically-induced knee OA²³⁻²⁷, accompanied by behavioral indicators of pain²³. Aside from its role in energy storage, adipose tissue is also a major endocrine organ and has been shown to secrete hormones termed adipokines (e.g. leptin, adiponectin, visfatin, resistin) and inflammatory cytokines (e.g. TNF- α , IL-6, TGF- β)²⁸. Studies investigating the role of adipokines have highlighted their importance in obesity-associated

pathologies. For example, leptin-deficient mice become obese yet they do not develop knee OA, suggesting leptin may play a key role in obesity-induced OA²⁹. In the IVD, exposure of NP cells to adipokines, such as leptin and resistin, promotes catabolic metabolism associated with increased expression of matrix remodeling enzymes such as MMP and ADAMTS genes^{30,31}. Adipokines also appear to play a role systemically as modulators of pain sensitivity³². In addition to back pain, obese individuals are more likely to develop chronic pain conditions such as fibromyalgia, headaches and abdominal pain³³. While the underlying mechanisms linking obesity and chronic pain remains unknown, it has been suggested that systemic immune and endocrine alterations play a role in the altered pain response³⁴. This systemic modulation of pain may contribute to LBP in addition to structural alterations and local inflammation within the IVD itself.

While there is extensive clinical evidence supporting the association between obesity, LBP and IVD degeneration^{35,36}, no studies have directly assessed whether diet-induced obesity accelerates IVD degeneration, back pain, or investigated biological mediators underlying this association. The current study was designed to investigate whether chronic consumption of a high-fat or high-fat, high-sugar western diet alters the progression of age-related IVD degeneration or back pain using the mouse as a model.

3.4 Methods

3.4.1 Mice and Diets

Wild-type, male, C57BL/6N (Charles River: Wilmington, MA, USA) mice were used. Mice were fed standard chow (Envigo 2018) after weaning and randomized at 10-weeks of age into one of three diet groups (n=9-16 mice/group; **Table 3.3, Supplementary Table** 1) based on previous reports of obesity and metabolic derangement in mice^{26,37}: high-fat diet (60% kcal fat, 21% kcal carbohydrate; Envigo TD.06414), western diet (45% kcal fat, 41% kcal carbohydrate; Envigo TD.10885), or standard chow (18% kcal fat, 58% kcal carbohydrate). Mice remained on the experimental diets until sacrifice at 5, 8 or 11.5 months-of-age (12-, 24-, 40-weeks on diet, respectively). Mice were housed in standard cages and maintained on a 12 hr light/dark cycle, with food and water consumed *ad libitum*; food consumption and body weight were measured weekly. All aspects of this study were conducted in accordance with the policies and guidelines set forth by the Canadian Council on Animal Care and were approved by the Animal Use Subcommittee of the University of Western Ontario (protocol 2017-154; **Appendix B**).

3.4.2 Characterization of pain-associated behavior

Behavioral analysis was conducted on mice following 12-, 24-, or 40-weeks on experimental or control diets. Behavioral studies were preceded by a two-week habituation to the neurobehavioral testing facility, and mice were habituated to all tests one week prior to data collection. On data collection days, animals were habituated to the testing room for 1 h before test start. To avoid confounding variables associated with the diurnal cycle, all behavioral assessments were conducted between 8 and 11 AM.

3.4.2.1 Stretch induced axial discomfort

Stretch-induced axial discomfort was measured using the tail suspension test and grip force during axial stretch, as described previously^{38–40}. For the tail suspension test, spontaneous reaction to gravity-induced stretch was assessed in mice suspended by the base of their tails

for 180 s. Two observers blinded to the experimental groups independently scored the duration of time spent by mice in immobility, full extension, rearing, or self-supporting using ANY-maze software (Stoelting Co.: Wood Dale, IL). Voluntary activity was quantified for 5 min immediately before (pre) and after (post) tail suspension using open field activity monitors (AccuScan Instruments, Omnitech Electronic: Columbus, OH), to quantify movement-evoked discomfort. The difference in total distance between the two open-field sessions (post - pre) was calculated for each mouse.

For the grip force assay, mice were positioned to grab onto a metal bar attached to a grip force meter (Stoelting Co.: Wood Dale, IL), and then gently pulled back by their tails to exert axial stretch. Tolerance was assessed by measuring the grip strength, in grams, for each mouse at the point of release averaged over 3 trials.

3.4.2.2 Hind limb sensitivity to mechanical and cold stimuli

Mechanical sensitivity was measured through application of calibrated Von Frey filaments (Stoelting Co.: Wood Dale, IL) to the plantar surface of the hind paw for 3 s or withdrawal. 50% withdrawal threshold was calculated using the Chaplan up-down method⁴¹. The stimulus intensity ranged from 0.07-6.0 g, beginning with a stimulus intensity of 1.4 g. Cold sensitivity was assessed by measuring the total time spent by mice in behavior evoked by evaporative cooling of acetone (flicking, stamping or licking of ventral surface of the paw) during the first 40 s following application of 50 μ L acetone to the ventral surface of the hind paw. The test was carried out twice for each paw, with at least 5 min recovery between each test. Times were then averaged between paws.

3.4.2.3 Spontaneous activity

Voluntary locomotor activity was assessed using open field activity monitors (AccuScan Instruments, Omnitech Electronic: Columbus, OH). Mice were placed into individual boxes and their activity was monitored over 2 h. This was repeated for 3 consecutive days and values were averaged for each mouse.

3.4.3 Micro-computed tomography (Micro-CT)

Forty-eight hours before sacrifice, μ CT imaging was performed using a cone-beam imaging system (eXplore SpeCZT scanner, GE Healthcare Biosciences: London, CAN). For imaging, mice were anesthetized using 2-3% inhaled isoflurane (CA2L9100, Baxter: Mississauga, CAN) infused with oxygen at a flow rate of 1.0 mL/min. To maintain sedation, a nose cone apparatus was used to administer 1.75% inhaled isoflurane for 20 min while scanning was performed. During a single 5 min rotation of the gantry, 900 Xray projections were acquired (peak voltage of 90 kVp, peak tube current of 40 mA, and integration time of 16 ms). A calibrating phantom composed of air, water, and cortical bone-mimicking epoxy (SB3; Gammex, Middleton WI, USA) was included in each scan. Data were reconstructed into 3D volumes with an isotropic voxel spacing of 50 µm and scaled into Hounsfield units (HU). Using MicroView software (GE Healthcare Biosciences) three signal-intensity thresholds (-200, -30, and 190 HU) were used to classify each voxel as adipose, lean, or skeletal tissue, respectively. Custom software was used to calculate tissue masses from assumed densities of 0.95 (adipose), 1.05 (lean), and 1.92 (skeletal) g/cm³, as previously reported 42 .

3.4.4 Histological analysis

Intact lumbar spine segments (L1-S1) and knees were isolated, fixed, decalcified and paraffin embedded, as previously described⁴³. Spines were sectioned sagittally, and knees were sectioned coronally at a thickness of 5 µm using a microtome (Leica Microsystems: Wetzlar, DEU). Lumbar spines were stained using a 0.1% Safranin-O/0.05% Fast Green. Knees were stained with either 0.1% Safranin-O/0.05% Fast Green or 0.04% Toluidine Blue. Sections were imaged on a Leica DM1000 microscope, with Leica Application Suite (Leica Microsystems: Wetzlar, DEU).

To evaluate IVD degeneration, spine sections were scored by 2 independent scorers using the modified Boos system⁴⁴. Knee joint health was assessed using the murine Osteoarthritis Research Society International (OARSI) histopathological scale⁴⁵. Articular surfaces of the medial femoral condyle (MFC), medial tibial plateau (MTP), lateral femoral condyle (LFC) and lateral tibial plateau (LTP), were scored by two blinded observers and averaged. For each knee joint surface, scores from 10 serial sections spanning 500 μ M of the joint were summed to represent OARSI score for each quadrant. Total scores from each of the four quadrants were then added together to generate whole joint OARSI score. Thickness of articular cartilage of the MTP and LTP was quantified using the OsteoMeasure7 Program (v.4.2.0.1, OsteoMetrics Inc., Decatur, GA, USA). Articular cartilage thickness within each quadrant was averaged using three serial sections spanning 150 μ M of the weight-bearing region of the knee. Using the same histomorphometry system, trabecular bone area was calculated by measuring the total surface area of the bone between the articular cartilage and growth plate and subtracting the area of bone marrow. Measurements were taken for both medial and lateral compartments of the joint and averaged from 3 serial sections.

3.4.5 Gene expression analysis

Thoracic IVDs (4-5 per mouse; 5-8 mice per diet/per timepoint) were isolated by microdissection, placed in TRIzol reagent (Thermo Fisher Canada: Mississauga, ON, CAN) and homogenized using a PRO250 tissue homogenizer (PRO Scientific: Oxford, CT, USA). RNA was extracted according to manufacturer's instructions, quantified using a NanoDrop 2000 spectrophotometer (Thermo Fisher Canada: Mississauga, ON, CAN), and 0.5 µg was reverse transcribed into complementary DNA (cDNA) (iScript; Bio-Rad Laboratories (Canada): Mississauga, ON, CAN). Gene expression was assessed by realtime PCR using a Bio-Rad CFX384 instrument. PCR analyses were run in triplicate using 120 ng of cDNA per reaction and 310 nM forward and reverse primers with 2x SsoFast EvaGreen Supermix (Bio-Rad Laboratories (Canada): Mississauga, ON, CAN) using optimized PCR parameters and primers (Table 3.4, Supplementary Table 2). Primers were designed and validated to have efficiency values between 90 and 120%. Transcript levels were calculated relative to a 6-point standard curve made from pooled cDNA generated from murine IVD explants treated with lipopolysaccharide for 4 days (50 mg/mL; Thermo Fisher Canada: Mississauga, ON, CAN).

3.4.6 Immunohistochemistry

The intact spinal cord was removed, dissected to separate the upper (L1-L2) and lower (L3-L6) segments of the lumbar enlargement, and fixed in 4% paraformaldehyde (PFA) for 24
h at 4°. Tissues were cryoprotected for 4 days in 10% sucrose and embedded in optimal cutting temperature compound (Tissue-Tek O.C.T; Sakura Finetek US: Torrance, CA, USA) and stored at -20°C. Tissues were sectioned on a cryostat (Leica Microsystems: Wetzlar, DEU) in the transverse plane at a thickness of 14 μ m, thaw mounted onto gelatin-coated slides, and stored at -80°C.

Slides were brought to room temperature, washed twice in PBS and blocked using 5% donkey serum, 0.1% Triton X-100 in PBS for 2h at room temperature. Sections were incubated overnight in a humidified chamber at 4°C in 5% donkey serum in PBS (with 0.1% Triton-X) containing primary antibodies directed against glial fibrillary acidic protein (GFAP) (1:500; G3893, Sigma-Aldrich: St. Louis, MO, USA), ionized calcium binding adaptor molecule 1 (IBA-1) (1:1000; AB-10341, Abcam: Cambridge, UK), or calcitonin gene-related peptide (CGRP) (1:750; BML-CA1137, Enzo Biochem: New York, NY, USA). Slides were rinsed 3 x 10 min in PBS-T (PBS + 0.01% Triton X-100) and then incubated for 45 min at room temperature with secondary antibodies diluted 1:500 in PBS: Alexa Fluor 488 conjugated donkey anti-mouse IgG for GFAP (A-21202, Thermofisher: Waltham, MA, USA); Alexa Fluor 594 conjugated donkey anti-rabbit IgG for Iba-1 (A-21207, Thermofisher: Waltham, MA, USA); or Alexa Fluor 488 donkey anti-sheep IgG for CGRP (A-11015, Thermofisher: Waltham, MA, USA). Slides were rinsed 3 x 10 min in PBS, dipped in deionized water, and cover slips mounted using Fluoroshield Mounting Medium with 4',6-diamidino-2-phenylindole to visualize nuclei (ab104139, Abcam: Cambridge, UK). Tissue sections were imaged using a Leica Microsystems DMI6000B fluorescence microscope and DFC360FX camera with Leica Advanced Application Suite software (Version 2.7.0-9329, Leica Microsystems: Wetzlar, DEU). A region of interest (ROI) was manually defined to contain lamellae 1-4 of the spinal cord dorsal horn using ImageJ software. The dorsal horn was differentiated from surrounding white mater based on brightfield images. Integrated density of fluorescence within the ROI was used to quantify astrocyte/microglia density, and CGRP-immunoreactivity.

3.4.7 Serum analysis by Multiplex Assay

At euthanasia, blood was obtained by cardiac puncture, coagulated for 30 min at room temperature, and centrifuged at 3 000 rpm for 10 min at 4°C to collect serum. Serum (n=5-6 mice per diet/per timepoint) was diluted 2-fold in DPBS and analyzed using the LuminexTM 200 system (Luminex, Austin, TX, USA) by Eve Technologies Corp. (Calgary, Alberta). Thirty-two markers were simultaneously measured in each serum sample using the MILLIPLEX Mouse Cytokine/Chemokine 32-plex kit (Millipore, St. Charles, MO, USA) according to the manufacturer's protocol. The multiplex assay quantified Eotaxin, G-CSF, GM-CSF, IFN γ , IL-1 α , IL-1 β , IL-2, IL-3, IL-4, IL-5, IL-6, IL-7, IL-9, IL-10, IL-12 (p40), IL-12 (p70), IL-13, IL-15, IL-17, IP-10, KC, LIF, LIX, MCP-1, M-CSF, MIG, MIP-1 α , MIP-1 β , MIP-2, RANTES, TNF α , and VEGF. The assay sensitivities range from 0.3 – 30.6 pg/mL for the 32-plex.

3.4.8 Statistical Analysis

For all assays except histopathological scoring of the joints, outcome measures for mice within each time point were compared between the different diet groups by one-way ANOVA with Tukey's multiple comparisons test. For histopathological analysis, within each timepoint scores for mice were compared between the different diet groups by nonparametric Kruskal-Wallis test with Dunn's multiple comparison test. P < 0.05 was considered significant. Statistical analysis was conducted using GraphPad Prism 8 (Graphpad Software: San Diego, CA, USA).

To assess the effect of diet, adiposity, knee OA and IVD degeneration on behavioral, molecular, and histological changes assessed, bivariate and multivariate linear regression models were used to identify which variables remained independently associated with the other outcomes. Bivariate and multivariate modelling was conducted using STATA 16 (StataCorp LLC: College Station, TX, USA).

3.5 Results

3.5.1 Weights and adiposity

As expected, following 12, 24- and 40-weeks mice fed the high-fat and western diets showed a significant increase in body mass and weight gain compared to age-matched chow fed controls (**Figure 3.1A**). Analysis of body composition by micro-CT (**Figure 3.1B**) demonstrated that the increase in adiposity in mice fed the experimental diets was accompanied by a significant decrease in the percentage of both lean and skeletal tissues at all time points (**Figure 3.1C**). This analysis also showed a significant increase in overall bone mineral density (BMD) in mice fed the high-fat diet at all timepoints, and at the 24- and 40-week timepoints for mice fed the western diet, compared to age-matched chow fed controls (**Figure 3.1C**).



Figure 3.1: Chronic consumption of high-fat and western diets increases adiposity and bone mineral density in C57BL/6N mice.

(A) At all timepoints, mice fed the high-fat and western diets showed significant increases in overall weight and in weight-gain from baseline compared to age-matched chow fed controls. (B) Representative reconstructed micro-CT images of mice following 40-weeks of experimental diets. Isotropic surface-rendering of skeletal tissue (indicated in white) is overlaid with a mid-coronal slice where lean tissue is indicated in red and adipose tissue is indicated in yellow. (C) Quantitative micro-CT analysis of whole-body composition showed a significant increase in adiposity and significant decreases in both percentage of lean and skeletal and in mice fed the high-fat and western diet mice compared to agematched chow fed controls at all time points. A significant increase was also seen in bone mineral density in mice following consumption of the high-fat and western diets at the 24and 40-week timepoints. n=9-16 mice per timepoint, per diet. Data are displayed as mean \pm 95% CI; data points for each mouse are graphed within each group. *P<0.05, **P<0.01, ***P<0.001, ****P<0.001 by 2-way ANOVA.

3.5.2 Behavioral indicators of axial discomfort

We first investigated whether mice fed the high-fat or western diet showed behavioral indicators of stretch-induced axial discomfort using three complimentary assays established as indicators of discogenic back pain in a mouse model of degeneration⁴⁰: behavior during tail suspension, changes in spontaneous activity after tail suspension, and tolerance to axial stretching in the grip force assay. In the tail suspension test, no significant differences were observed between diet groups at any time point investigated (**Figure 3.2A**). Similarly, changes in spontaneously activity induced by the tail suspension assay (locomotion pre versus post tail suspension) were not altered between experimental diet groups and age-matched chow fed controls (**Figure 3.2B**). Grip force during axial stretch was reduced in mice fed the high-fat diet compared to age-matched chow fed controls at 12- and 40-weeks and in mice fed the western diet at all time points compared to control (**Figure 3.2C**), suggesting decrease tolerance to axial stretch.

3.5.3 Behavioral indicators of mechanical and cold sensitivity

Mechanical and cold sensitivity were assessed in the hind paw using the Von Frey assay and by measuring the response of mice to the evaporative cooling of acetone, respectively. Mice fed the western diet showed a significant increase in mechanical sensitivity at the 24and 40-week time points compared to age-matched chow fed controls, while mice fed a high-fat diet showed a significant increase in sensitivity only at the 24-week timepoint compared to controls (**Figure 3.3A**). In contrast, no significant difference was observed in sensitivity to cold between mice in either experimental diet group compared to agematched chow fed controls at any time point (**Figure 3.3B**).





(A) During tail suspension, the duration of time spent by mice immobile, in full extension, rearing or self-supported was quantified. No significant differences were detected between diet groups at any of the time points assessed. (B) Stretch-evoked discomfort was assessed using the open field assay, in which the total distance covered in 5 min immediately before (pre) and after (post) the 3 min tail suspension assay was quantified. Obesity induced by the high-fat and western diets did not alter behavior of mice in open field compared to agematched chow fed controls at any of the time points assessed. However, a significant difference was seen between mice fed a high-fat and western diet at the 40-week timepoint. (C) Grip force during axial stretch was reduced in obese mice. Mice fed the high-fat diet showed a significant decrease in grip force at the 12- and 40-week time points compared to age-matched chow fed controls. Mice fed the western diet showed a significant decrease in grip force at the 12- and 40-week time points compared to age-matched chow fed controls. Mice fed the western diet showed a significant decrease in grip force compared to age-matched chow-fed controls at all time points. n=9-16 animals per timepoint, per diet. Data are plotted mean \pm 95% CI; data points for each mouse are graphed within each group. *P<0.05, **P<0.01, ***P<0.001 by 2-way ANOVA.

3.5.4 Spontaneous locomotion

Behavior and locomotion in open field was assessed for all mice over a 2 h period. Mice fed the western diet showed a significant decrease in total distance travelled following 12 and 24 weeks compared to age-matched chow fed controls. Mice fed the high-fat diet showed a significant reduction in locomotion following 24-weeks compared to age-matched controls (**Figure 3.3C**). The number of rearing events was also significantly decreased in mice fed the high-fat and western diets at the 40-week time point compared to age-matched chow fed controls (**Figure 3.3C**).

3.5.5 Assessment of IVD degeneration

The effects of the high-fat and western diets on IVD health were assessed using both histopathological evaluation and molecular analysis (**Figure 3.4**). On average, no overt differences were detected in the histological appearance of lumbar IVDs between mice fed either the high-fat or western diet for 12- and 24-weeks compared to age-matched chow fed controls (**Figure 3.4A**). Accordingly, histopathological scoring using the modified Boos system showed no significant differences in degeneration between the groups. However, when data was analyzed by individual spinal level, mice fed the western diet showed significantly lower scores than their high-fat or chow fed counterparts at the L6S1 spinal level at the 12-week timepoint (**Figure 3.4B**). Following 40 weeks on the experimental diets, an accumulation of hypertrophic cells surrounded by a glycosaminoglycan-rich pericellular matrix was consistently detected in the inner annulus fibrosus of mice fed both the high-fat and western diets, but not in age-matched chow fed control mice (**Figure 3.4A** – black arrows). Despite this observation, histopathological



Figure 3.3: Obesity induced by a high-fat and western diet increases sensitivity to mechanical stimulation and alters spontaneous locomotion.

(A) Mechanical Sensitivity of the hind paw was assessed by manual application of Von Frey filaments using the Chaplin up-down method. Mice fed the western diet showed a significant decrease in withdrawal threshold at the 24- and 40-week time points compared to age-matched chow fed controls, indicative of increased mechanical sensitivity. Mice fed a high-fat diet showed a significant decrease in withdrawal threshold at the 24-week timepoint compared to control. (B) Sensitivity to cold was assessed by measuring the time spent in behavior evoked by evaporative cooling of acetone (flicking, stamping or licking of ventral surface of the paw) during the first 40s following application of acetone to the ventral surface of the hind paw. No significant differences were seen between the diet groups at any timepoint. (C) Spontaneous locomotor activity was recorded over three 2 hr sessions and averaged. Mice fed the western diet showed a significant decrease in the total distance travelled at the 12- and 24-week timepoint compared to age-matched chow fed controls, while mice fed the high-fat diet showed a decrease at the 24-week timepoint. The number of rearing events was significantly decreased in mice fed the high-fat and western diets compared to controls at the 40-week timepoint. n=9-16 animals per timepoint, per diet. Data are plotted mean \pm 95% CI; data points for each mouse are graphed within each group. *P<0.05, **P<0.01, ***P<0.001, ****P<0.0001 by 2-way ANOVA.

scoring revealed no significant degeneration in the diet groups compared to chow fed controls at the 40-week time point (Figure 3.4B). Histopathological analysis was paired with qPCR to quantify expression of markers of inflammation, neural ingrowth, and matrix degrading enzymes in thoracic IVDs to further investigate molecular changes associated with diet-induced obesity. Mice fed the high-fat or western diets showed no significant differences in the expression of the genes investigated at the 12-week timepoint compared to age-matched chow fed controls (**Figure 3.4C**). Mice fed the high-fat diet showed no significant differences in gene expression compared to chow fed controls at the 24-week timepoint for any of the genes investigated. At the 40-week timepoint, mice fed the highfat diet showed increased expression of inflammatory mediators (11-6, Ptsg2), neurotrophins (Bdnf), as well as matrix degrading enzymes (Adamts5) compared to agematched chow fed controls (Figure 3.4C). Mice fed the western diet showed increased expression of the inflammatory mediators *Il-1b* and *Ptgs2* compared to chow fed controls at 24 weeks; however, no significant differences in gene expression were detected at the 40-week timepoint (**Figure 3.4C**).

3.5.6 Assessment of degenerative changes in the knee

Since diet-induced obesity leads to other arthropathies, such as knee OA²³, we investigated degenerative changes to the knee joint as a potential contributor to the pain-related behavioral outcomes we assessed. We focused this analysis on the 24- and 40-week timepoints where changes in behavioral measures were most consistently identified. No overt histological differences were detected in the knee joints of mice fed either the high-fat or western diet for 24 weeks compared to age-matched chow fed controls (**Figure**



Figure 3.4: Effect of diet-induced obesity on the intervertebral disc.

(A) Representative histological sections of the L6/S1 spinal level of the lumbar spine stained with safranin-o/fast green from mice fed control chow, high-fat, or western diet for 12, 24, or 40 weeks. The accumulation of hypertrophic cells was detected within the inner annulus fibrosus of mice fed the high-fat and western diets for 40 weeks (highlighted by black arrows). (B) Evaluation of the grade of histopathological IVD degeneration using the modified Boos scoring system showed no significant differences between mice fed the control chow, high-fat or western diets at the 24- and 40-week timepoints. At the 12-week timepoint, mice fed the western diet showed a significant decrease in the degenerative score compared to mice fed chow and high-fat diets. n=9-16 animals per timepoint, per diet. Data are analyzed by Kruskal-Wallis test. (C) SYBR-based qPCR of intact thoracic intervertebral discs showed no significant difference between mice fed a chow, high-fat or western diet at the 12-week timepoint for any genes investigated. At the 24-week timepoint, significant increases in *Il-1b* and *Ptgs2* expression were seen in mice fed the western diet compared to control. By 40-weeks, significant increases in Il-6, Ptgs2, Bdnf and Adamts5 expression were seen in mice fed the high-fat diet compared to control. n=5-8 animals per diet/per timepoint. Analyzed by one-way ANOVA. All data are plotted mean \pm 95% CI; data points for each mouse are graphed within each group. *P<0.05, ***P<0.001.

3.5A). Histopathological scoring using the OARSI system supported these observations (Figure 3.5B). At the 40-week timepoint, mice fed the western diet showed decreased proteoglycan staining in the medial femoral condyle (MFC) compared to age-matched controls (Figure 3.5A), resulting in significantly higher OARSI scores (Figure 3.5B) indicating early signs of osteoarthritis. However, no significant differences were seen in the other compartments of the knee joint or in the cumulative OARSI scores for the whole joint. In addition to histopathological analysis, cartilage thickness was measured by histomorphometry. Although no significant differences in cartilage thickness were detected in mice following 24 weeks on the experimental diets, mice fed the high-fat diet for 40 weeks showed a significant decrease in cartilage thickness specific to the lateral femoral condyle compared to age-matched chow fed controls (Figure 3.5C). Histomorphometry was similarly used to measure the relative surface area of trabecular bone between the articular cartilage and growth plate. Mice fed the western diet for 40 weeks showed a significant increase in subchondral trabecular bone compared to age-matched chow fed controls specifically in the medial tibial plateau (Figure 3.5D), suggesting sclerosis of the subchondral bone.

3.5.7 Analysis of sensory neuroplasticity within the lumbar spinal cord To assess neuroplastic changes associated with neuroinflammation and chronic pain, lumbar spinal cords from mice at the 40-week time point were assessed for markers of astrocytes (glial fibrillary acidic protein, GFAP), microglia (ionized calcium-binding adapter molecule 1, IBA-1), and nociceptive innervation (calcitonin gene-related peptide, CGRP; **Figure 3.6A**). Although multiple mice in both the high-fat and western diet groups showed increased GFAP and IBA-1 staining in the upper and lower lumbar spinal cord



Figure 3.5: Effect of diet-induced obesity on the knee joint.

(A) Representative histological coronal sections of the medial knee compartment stained with safranin-o/fast green from mice fed either control chow, high-fat, or western diet for 24- or 40- weeks. Images are oriented with the medial femoral condyle (MFC) located superiorly, and the medial tibial plateau (MTP) inferiorly. White arrows indicate cartilage damage, presenting as loss of proteoglycan staining and focal fibrillation of the cartilage, and white boxes indicate osteophyte formation. (B) Histopathological grading of the knee joints using the murine Osteoarthritis Research Society International (OARSI) scale corresponding to MTP, MFC, lateral tibial plateau (LTP), and lateral femoral condyle (LFC), combined to generate the summed score for the whole joint. Mice fed the western diet for 40-weeks showed a significant increase in the degenerative score in the MFC compared to those fed the control chow. However, no difference was seen in the whole joint score between any of the groups at either timepoint. (C) Average articular cartilage thickness. After 40 weeks on the high fat diet, mice presented with decreased articular cartilage thickness on the LTP. No other differences were seen in any other joint compartment. Data analyzed by Kruskal-Wallis test. (D) Percent trabecular bone in the medial and lateral subchondral compartments of the tibia. Mice enrolled on the western diet for 40 weeks exhibited significantly more trabecular bone in the medial compartment for the tibia. n=9-16 animals per diet/per timepoint. Analyzed by one-way ANOVA. All data are plotted mean \pm 95% CI, *P<0.05.

compared to the average values for chow fed controls, quantification of fluorescent intensity within the dorsal horn showed no significant difference between diet conditions for any of the proteins investigated (**Figure 3.6B**).

3.5.8 Circulating inflammatory factors

Luminex xMAP multiplex assays were used to quantify a panel of 32 cytokines, chemokines, and growth factors in the serum of experimental mice. Mice fed a western diet showed increased levels of interleukin (IL)-1 β , IL-6, IL-10, and tumor necrosis factor alpha (TNF α) at the 12- and 40-week timepoints in their serum, however, no significant differences were seen when compared to chow fed control due to variability between animals (**Table 3.1**). Despite this variability, at the 40-week timepoint a significant increase in circulating monocyte chemoattractant protein 1 (MCP-1) was observed in mice fed a western diet showed for 24-weeks also showed significantly lower levels of circulating eotaxin and IL-1 α compared to chow fed controls, but no other differences in any of the other cytokines investigated (**Table 3.5, Supplementary Table 3)**.

3.5.9 Linear regression analysis

To directly examine associations in the context of the observed variability within each experimental group in our study, linear regression modeling was conducted to determine whether diet, adiposity or age were predictors of behavioral, histological, or systemic outcomes (**Table 3.2**). Bivariate modelling demonstrated that both adiposity and age are independent predictors of multiple indicators of pain including stretch-induced axial discomfort (grip force), mechanical sensitivity (Von Frey assay) and spontaneous



Figure 3.6: Effect if diet-induced obesity on neuroplastic changes within the lumbar spinal cord.

(A) Representative images showing transverse sections of the dorsal horn of the lumbar spinal cord used for immunohistochemical analysis of neuroplastic changes associated with chronic pain. Tissues were isolated from mice following 40 weeks of experimental diet. Slides were stained for glial fibrillary acidic protein (GFAP), ionized calcium-binding adapter molecule 1 (IBA1) and calcitonin gene-related peptide (CGRP). Yellow boxes indicate the region of interest for high magnification images (second row) for GFAP and IBA-1 (B) The fluorescence intensity averaged across the region of interest (lamellae 1-4 of the dorsal horn) of the upper (L1-L2) and lower (L3-L6) lumbar spinal cords. Mice fed a high-fat or western diet for 40-weeks showed no significant differences in immunoreactivity for any of the proteins investigated. n = 6-8 animals/group. Individual data points of the same colour indicates the same animal. Data are plotted mean $\pm 95\%$ CI.

Table 3.1: Multiplex analysis of cytokines, chemokines and growth factors in serum.

	12 Week					24 \	Neek			40 Week		
	Chow	HF	Western		Chow	HF	Western		Chow	HF	Western	
	n=6	n=6	n=6		n=5	n=6	n=6		n=5	n=6	n=6	
Analyte	Mean (SD)	Mean (SD)	Mean (SD)	p- value	Mean (SD)	Mean (SD)	Mean (SD)	p- value	Mean (SD)	Mean (SD)	Mean (SD)	p- value
IL-1B	13.9 (17.5)	32.3 (58.6)	39.4 (43.8)	0.54	22.5 (26.9)	15.3 (15.7)	41.1 (89.3)	0.72	2.0 (1.8)	2.0 (1.4)	6.4 (6.7)	0.16
IL-6	17.2 (14.3)	20.9 (28.1)	47.6 (67.0)	0.43	25.7 (31.9)	28.7 (35.0)	13.6 (12.2)	0.63	2.7 (3.8)	4.1 (1.9)	20.0 (30.2)	0.24
IL-10	45.8 (51.4)	56.6 (85.1)	153 (261.2)	0.47	19.0 (24.1)	17.2 (18.7)	15.1 (18.7)	0.95	0.5 (1.0)	0.4 (0.9)	2.1 (2.9)	0.25
IP-10	129.4 (53.8)	122.0 (32.4)	139.5 (38.7)	0.78	90.9 (53.2)	105.9 (43.8)	63.6 (49.5)	0.34	106 (14.3)	113.7 (24.2)	144.7 (67.9)	0.32
KC	453.4 (323.3)	690.2 (843.0)	856.0 (1324)	0.75	527.2 (363.1)	452.0 (230.1)	240.7 (108.5)	0.17	191.1 (85.0)	355.3 (223.4)	258.8 (139.5)	0.28
MCP-1	53.7 (58.7)	51.6 (46.4)	78.3 (65.6)	0.68	54.3 (23.8)	68.1 (42.3)	165.2 (259.5)	0.45	13.6 (9.6)	63.8 (39.1)	135.3 (93.7) *#	0.02*
TNFa	28.4 (31.4)	37.8 (54.2)	41.9 (35.0)	0.85	23.9 (23.6)	16.0 (22.7)	53.4 (96.2)	0.54	1.0 (1.9)	2.2 (4.1)	3.0 (5.1)	0.63
VEGF	2.2 (1.87)	6.8 (14.2)	13.1 (27.8)	0.59	2.9 (4.2)	1.5 (2.5)	1,9 (2.1)	0.73	5.4 (8.4)	2.1 (3.1)	1.0 (0.5)	0.33
Values are Tukey's po	displayed in st-hoc test.	n pg/mL an P<0.05 is	d analyzed by signific ant	one-way AN	IOVA. * indica	ates signific	antly different	from chow o	liet. # is sign	ificantly diff	erent from HF	diet by

whether diet, adiposity or age were predictors of behavioral, histological, or systemic outcomes (Table 3.2). Bivariate modelling demonstrated that both adiposity and age are independent predictors of multiple indicators of pain including stretch-induced axial discomfort (grip force), mechanical sensitivity (Von Frey assay) and spontaneous locomotion (open field). Despite their association with behavioral alterations, neither adiposity nor age were associated with measures of joint degeneration or levels of most circulating factors assessed (**Table 3.2**). Exceptions to this were TNF α and MCP-1, which were associated with age (bivariate), and the western diet (multivariate), respectively (**Table 3.2**). When diet, adiposity and age were accounted for, age was found to be the most robust predictor of all outcomes measured (Table 3.2). Aside from age, the multivariate model also showed that diet and adiposity are covariates for grip force and rearing in open field, respectively (**Table 3.2**). To determine if histopathological scores for joint damage were independently associated with behavioral indicators of pain, bivariate and multivariate regression modeling was completed. In this study, no significant association was detected between histopathological scores for IVD degeneration and painrelated behaviors. A weak but significant association was detected between histopathological scores for knee OA and grip force, but not for any of the other behavioral metrics assessed (Table 3.6, Supplementary Table 4).

3.6 Discussion

Obesity is one of the largest modifiable risk factors for the development of both IVD degeneration and associated LBP^{35,36}, yet the biological mechanisms underlying this association are currently unknown. With the prevalence of obesity on the rise¹, it is necessary to address this problem and identify underlying pathogenic mechanisms. The

	Bivar	iate (r)		Multiv	variate (β, r²)	
	%	Timepoint	Diet (3)	%	Timepoint	Whole
	Adipose				Adipose (ß)	(β)	model (r²)
Parameter			Experimental	Western	1197		
			(Obesogenic)				
Behavioral							
Grip Force	-0.41***	-0.45***	-21.96**	-6.23	0.4	-0.841**	0.45***
Mechanical Sensitivity (Von Frey)	-0.42***	-0.21*	-0.35	-0.14	-0.0007	-0.006	0.26***
Cold sensitivity (Acetone)	0.04	0.01	-	-	-	-	0.08
Tail Suspension Rearing	0.01	0.11	-	-	-	-	0.02
Immobility	0.18	0.19	-	-	-	-	0.05
Self-support	-0.22*	-0.30**	1.73	-0.46	-0.71	-0.74*	0.10*
Stretch	0.06	0.10	-	-	-	-	0.07
Open Field							
Locomotion	-0.49***	-0.35***	-1226	-59.3	-92.7	-93.5***	0.33***
Rearing	-0.52***	-0.40***	-0.88	-7.41	-2.87*	-2.059***	0.39***
Histopathology							
IVD degeneration	0.065	0.04	-	-	-	-	0.03
Knee OA (Cumulative OARSI)	0.07	0.08	-	-	-	-	0.05
Neuroinflammation							
GFAP	0.23	-				-	0.30
IBA-1	0.23	-	-	-	-	-	0.06
CGRP	0 024	-		-	-	-	0.18
Upper spinal cord	0.021						0.10
GEAP	0.21	-	-	-	-	-	0.08
IBA-1	0.33	-	-	-	-	-	0.12
CGRP	0.31		-	-	-	-	0.18
0011	0.51						0.10
Systemic factors							
II -1B	0.04	0.21	-	-	-	-	0 12
II -6	0.06	0.22	-	-	-	-	0.09
II -10	0.03	0.21	-	-	-	-	0.14
IP-10	0.03	0.07	-	-	-	-	0.001
KC	0.002	0.25	-	-	-	-	0.08
MCP-1	0.23	0.04	-	-	-	-	0.18
TNFa	0.03	0.30*	-	-	-		0.16
VEGF	0.01	0.12	-	-	-	-	0.06

Table 3.2:	Impact	of	diet-induced	obesity	and	age	on	behavioral,	molecular,	and
histological	l changes	s.								

Note: Experimental/Obesogenic diets include the high-fat and western diet

current study was designed to investigate the longitudinal effects of diet-induced obesity on inflammation, IVD degeneration and pain using the mouse as a preclinical model. We show that obesity induced by both a high-fat and high-fat/high-sugar western diet led to behavioral indicators of pain in as little as 12-weeks, preceding gross structural changes to the IVD and knee joints. Following 40 weeks, changes in cellular morphology within the inner AF of the IVD were detected in mice fed both obesogenic diets compared to chowfed controls; however, these changes were not associated with increased histopathological degeneration. Chronic consumption of the high-fat diet was associated with increased expression of *Il-6, Ptgs2, Bdnf, Adamts-5,* and *Mmp-3* within the IVD and a decrease in articular cartilage thickness within the lateral tibial plateau. In contrast, chronic consumption of the western diet was associated with increased expression of *Il-1b* and *Ptgs2* within the IVD, histopathological features of early OA, subchondral bone sclerosis, and increased serum MCP-1 levels. These findings highlight the complex interplay between diet, adiposity, pain, inflammation, and joint health.

Rodent models have proven useful to study IVD and joint biology since they allow insight into biological processes that regulate tissue homeostasis and degeneration. Numerous models have been described in which IVD degeneration is induced through genetic manipulation, surgical disruption, chemical injection or aberrant mechanical loading⁴⁶. Although these studies contribute to understanding the cellular and molecular basis of IVD degeneration, few investigate the association with pain⁴⁷. Given the discordance between structural IVD degeneration and pain development in humans^{10,48,49}, and pre-clinical models⁵⁰, it is important to investigate both pain and structural IVD degeneration and their relation to one another. Although pain cannot be directly measured in animals, several indirect, quantitative behavioral assays have been developed to evaluate pain-like behaviors for a variety of different pain states^{51,52}. Many of these behavioral metrics are not specific to back pain and are used to assess joint, inflammatory, and neuropathic pain^{53,54}. Stretch-induced axial discomfort has however been established as a reliable measure of axial low back pain⁴⁰. In the current study, mice fed obesogenic diets showed significant impairments in grip force at all timepoints compared to control mice, suggesting axial discomfort. In contrast, no significant difference was detected in the tail suspension assay between mice fed obesogenic diets and controls. This potentially contradictory data may be influenced by the nature of the tail suspension assay, which was originally used to test depressive behavior in mice⁵⁵. In addition to musculoskeletal pathologies, obesity is also associated with depression³³. In fact, recent studies demonstrated that mice fed a highfat diet for 8 weeks showed increased time immobile in tail suspension, interpreted as a characteristic of depression-like behavior⁵⁶. This behavior is in contrast to both genetic and age-related mouse models of IVD degeneration and axial pain which show increased time rearing in tail suspension^{57,58}. This confounding factor may impact the outcome of tail suspension in the diet-induced obesity model, and consequently the interpretation of our findings.

In addition to indicators of stretch-induced axial discomfort, obese mice also displayed mechanical but not thermal (cold) hypersensitivity of the hind-paw. These alterations are consistent with results seen in a model of surgically induced IVD degeneration, where intradiscal injection of PBS induced mechanical hypersensitivity in rats⁵⁹. However, these results directly contrasted a genetic model of IVD degeneration, where SPARC-null mice show thermal but not mechanical hypersensitivity compared to WT control³⁹. While the

mechanisms underlying these differences are unclear, mechanical sensitivity is common in both inflammatory and neuropathic pain models⁶⁰, and different models of pain are likely to impact nociception through different pathogenic mechanisms⁶¹. Obesity also appeared to impact non-reflexive (spontaneous) pain behaviors including distanced travelled and rearing in open field. These measurements have been previously shown to be decreased in both inflammatory and neuropathic pain models⁶². Taken together, our findings indicate that mice fed a high-fat or western diet display pain-related behaviors starting at the 12week timepoint and persisting until the 40-week timepoint.

Previous studies established that high-fat diet-induced obesity accelerates OA progression in mice^{23,26}; however, pain-related behaviors were found to be independent of osteoarthritis severity²³. As IVD degeneration has been associated with many of the pain-related behaviors reported for OA³⁸, we characterized both IVD degeneration and OA-associated knee degeneration in our experimental mice to account for both as factors contributing to pain. Histopathological scoring of lumbar IVDs did not reveal significant degeneration caused by the high-fat or western diet at any timepoint. However, mice fed both obesogenic diets showed a consistent accumulation of hypertrophic-like cells in the inner annulus fibrosus at 40 weeks, suggesting early degenerative change. Moreover, increased expression of inflammatory mediators (Il-1b, Il-6, Ptgs2), matrix degrading enzymes (Adamts5) and neurotrophins (Bdnf) were detected in mice fed the high-fat and western diets. These inflammatory cytokines are known drivers of IVD pathogenesis associated with ECM degeneration and expression of neurogenic factors such as NGF and BDNF that can contribute to pain⁶³. Similarly, significant histopathological degeneration in the knee was only detected at the 40-week timepoint in mice fed the western diet. Linear regression modeling indicates that the behavioral indicators of pain assessed are independent of joint degeneration, except for the grip force assay for which knee OA was a significant predictor of impairment. Together these findings indicate that diet-induced obesity may potentially accelerate the progression of structural IVD degeneration and knee OA at the later timepoints; however, these changes are mild and are likely independent of most pain-related behaviors.

Findings from the current study suggesting that pain-related behaviors precede molecular and structural alterations in joint tissues raise questions related to the source of the pain observed. These findings are however consistent with the hypothesis that pain is multifactorial. Obesity is considered a state of chronic inflammation associated with increases in several circulating inflammatory cytokines such as TNFa and IL-6 in humans⁶⁴. Inflammation has been shown to contribute to peripheral and central sensitization and may lead to hyperexcitability of the nervous system and chronic pain 65 . Consistent with these findings, we demonstrate that consumption of the western diet led to increased levels of circulating MCP-1 in mice, a pro-algesic mediator that can increase primary afferent neuron activity^{66,67}. Obesity may also impact central pain processing; in mice consumption of a high-fat diet increases the activation of microglia⁶⁸ while exposure of cultured astrocytes to saturated fatty acids induces cytokine release and astrocyte inflammation⁶⁹. These neuroplastic changes can contribute to central sensitization through multiple mechanisms, including increased release of inflammatory factors contributing to modulation of synaptic activity⁶⁵. Although the averaged values of GFAP and IBA-1 expression in the spinal cord were not significantly different between diet groups in our study, multiple mice in both the high-fat and western diet groups showed increased activation of both microglia and astrocytes at the 40-week timepoint. These alterations to the nociceptive pathways at either the peripheral or central level may contribute to the pain response seen. While not assessed, obesogenic diets also lead to painful diabetic neuropathy, or nerve damage, which may also contribute to the pain phenotype observed⁷⁰. Given the diverse impacts of obesity, it could be postulated that obesity-related pain may arise either from hyperexcitability or damage to nociceptive pathways in conjunction with tissue damage, and these contributions may differ between animals and over time. A limitation of this study is the utility of behavioral tests used to assess pain in our model. While they serve as indicators of pain in other models, many assays used in the current study also have been used to test muscle strength⁷¹ and psychiatric disorders⁵⁵ which may be impacted by obesity^{33,72}. To further explore whether these behavioral responses are due to pain, future studies may wish to pharmacologically inhibit nociceptive pathways to confirm that behavioral differences are due to pain.

Although the average weight gain was similar between the two obesogenic diets evaluated in this study, important differences in outcomes were detected. Mice fed the western diet showed a more consistent pain response compared to control at all timepoints than mice fed the high-fat diet. Furthermore, at the 40-week timepoint knee osteoarthritis and increases in systemic inflammation were only seen in mice fed the western diet. These findings highlight the importance of dietary composition in the study of obesity. In the context of OA, dietary fatty acid and carbohydrate composition can significantly impact joint health^{24,73}. Diets high in saturated fatty acids or \Box -6 polyunsaturated fatty acids (PUFAs) induce more severe metabolic dysregulation and OA progression than diets enriched with \Box -3 PUFAs²⁴. Aside from fat, diets high in sucrose can also accelerate OA progression independent of weight gain⁷³. These findings may explain our results, as the composition of the western diet is higher in both saturated fats and sucrose than the high-fat diet. Dietary composition has also been shown to impact IVD health. Diets rich in advanced glycation end-products (AGE) precursors accelerate IVD degeneration in mice in parallel with insulin resistance⁷⁴.

A confounding factor in the interpretation of our findings was the substantial variability for many of the outcomes investigated, including substantial differences in weight gain between mice on both obesogenic diets. Despite controlling for genetics using an inbred strain, susceptibility to diet-induced obesity can be affected by social stress, microbiome composition and epigenetic mechanisms^{75–77}. Accounting for this variability, previous studies in mice demonstrated that cartilage damage induced by a high-fat diet is proportional to adiposity²³. In the current study, adiposity did not predict histopathological measures of joint degeneration; however, adiposity and age were significant predictors of behavioral indicators of pain. Highlighting the complexity of these models, increased adiposity was also not directly associated with increased circulating cytokine levels in our study. In humans, obese individuals can be classified as metabolically healthy (metabolically healthy obese (MHO)). MHO individuals are at less risk for developing obesity-related complications than metabolically abnormal obese individuals, including OA⁷⁸. The mechanisms underlying MHO are not well understood, however, genetic, epigenetic and environmental factors are thought to play a role^{79,80}. As adiposity does not directly correlate with systemic or neuroinflammation in the current model, it is important to investigate all aspects of metabolic syndrome (i.e. circulating lipids, glucose,

cytokines/adipokines, blood pressure) on both the musculoskeletal and nervous system in future models to determine their contribution to disease progression.

Taken together, this study highlights the complexity of the relationship between obesity, IVD degeneration and pain. Mice fed a high-fat or western diet showed pain-related behaviors that preceded structural joint degeneration in both the IVD and knee. The chronology of these findings may be of clinical importance, as pain may affect the progression of radiographic joint degeneration. While not directly investigated in IVD degeneration, knee pain has shown to be a predictor of accelerated radiographic OA through inflammation and reduced mobility⁸¹, which is also seen in IVD degeneration⁸². This raises the intriguing possibility that back pain may be both a consequence of structural IVD degeneration in the current model, and a contributor to it.

3.7 Supplementary Figures





Spontaneous Locomotion activity was recorded over three 2 hr sessions and averaged. (A) Mice fed the western diet showed a significant decrease in the average movement velocity at all timepoints compared to age-matched chow fed controls, while mice fed the high-fat diet showed a decrease at the 24-week timepoint. (B) The amount of time spent in the anxiety-inducing center area of the open field enclosure was decreased in mice fed the high-fat and western diets compared to controls but not significant at any timepoint. n=9-16 animals per timepoint, per diet. Data are plotted mean \pm 95% CI; data points for each mouse are graphed within each group. *P<0.05, **P<0.01, ***P<0.001, ****P<0.001 by one-way ANOVA.



Figure 3.8, Supplementary Figure 2: SYBR-based qPCR of thoracic IVDs (continued).

SYBR-based qPCR of intact thoracic intervertebral discs showed no significant difference between mice fed a chow, high-fat or western diet at the 12-week and 24-week timepoint for any genes investigated. At 40-weeks a significant increase was seen in *Mmp12* expression in mice fed a high-fat diet compared to chow control. n=5-8 animals per diet/per timepoint. Analyzed by one-way ANOVA. All data are plotted mean \pm 95% CI; data points for each mouse are graphed within each group. *P<0.05, ***P<0.001.

	Contr	ol diet	Obesogenic diets						
	Ch	wow	Hiç	gh fat	Western				
Product #	Envig	o 2018	Envigo	TD.06414	Envigo TD.10885				
Content	g	kcal%	% by	kcal%	% by	kcal%			
			weight		weight				
Protein	19.2	20	23.5	18.3	17.3	14.8			
Carbohydrate	67.3	70	27.3	21.4	47.5	40.6			
Fat	4.3	10	34.3	60.3	23.2	44.6			
kcal/g		3.85		5.1		4.7			
Ingredient	g/kg		g/kg		g/kg				
Casein	N/A		265		195				
L-Cystine	N/A		4		3				
Corn starch	N/A		0		55.46				
Maltodextrin	N/A		160		60				
Sucrose	N/A		90		340				
Cellulose	N/A		65.5		50				
Soybean oil	N/A		30		20				
Anhydrous Milkfat	N/A		0		61				
Cholesterol	N/A				1.5				
Lard	N/A		310		0				
Vitamin Mix	N/A		21		19				
Mineral Mix	N/A		48		43				
Dicalcium	N/A		3.4		0				
phosphate									
Choline bitartrate	N/A		3.0		3.0				
SFA (% of fat)	N/A		34.8		58.3				
MUFA (% of fat)	N/A		39.1		28.2				
PUFA (% of fat)	N/A		22.5		8.9				
ω6:ω3 ratio	N/A		13.0		7.2				

 Table 3.3, Supplementary Table 1: Experimental diet compositions.

NCBI Gene	Primer Sequence $5' \rightarrow 3'$
Name	
<i>Il6</i> Fwd	TCTCTGCAAGAGACTTCCATCCAGT
<i>Il6</i> Rev	AGTAGGGAAGGCCGTGGTTGTCA
<i>Il1b</i> Fwd	CCCTGCAGCTGGAGAGTGTGGA
<i>Il1b</i> Rev	TGTGCTCTGCTTGTGAGGTGCTG
Ptgs2 Fwd	GGCGCAGTTTATGTTGTCTGT
Ptgs2 Rev	CAAGACAGATCATAAGCGAGGA
Bdnf Fwd	TCATACTTCGGTTGCATGAAGG
Bdnf Rev	GACCTCTCGAACCTGCCC
Adamts5 Fwd	GGAGCGAGGCCATTTACAAC
Adamt5 Rev	GCGTAGACAAGGTAGCCCACTTT
Mmp3 Fwd	TTGTCCCGTTTCCATCTCTCTC
Mmp3 Rev	TTGGTGATGTCTCAGGTTCCAG
Adamts4 Fwd	GAGGAGGAGATCGTGTTTCCAG
Adamts4 Rev	CAAACCCTCTACCTGCACCC
Mmp12 Fwd	GCTTACCCCAAGCTGATTTCC
Mmp12 Rev	ATGTTTTGGTGACACGACGGA
Mmp13 Fwd	CTTCTTCTTGTTGAGCTGGAACTC
Mmp13 Rev	CTCTGTGGACCTCACTGTAGACT
Ngf Fwd	TGATCGGCGTACAGGCAGA
Ngf Rev	GCTGAAGTTTAGTCCAGTGGG
Vegf Fwd	CACTGGACCCTGGCTTTACT
Vegf Rev	GCAGTAGCTTCGCTGGTAGA
Annealing Temp	erature: 60°C

 Table 3.4, Supplementary Table 2: Real-time qPCR primer sequences.

 Table 3.5, Supplementary Table 3: Multiplex analysis of cytokines, chemokines and growth factors in serum (continued).

	12 Week					24	Week					
	Chow	HF	Western		Chow	HF	Western		Chow	HF	Western	
	n=6	n=6	n=6		n=5	n=6	n=6		n=5	n=6	n=6	
Analyte	Mean (SD)	Mean (SD)	Mean (SD)	p- value	Mean (SD)	Mean (SD)	Mean (SD)	p- value	Mean (SD)	Mean (SD)	Mean (SD)	p- value
Eotaxin	2219	2657	3069	0.93	823.7	711.4	306.2	0.03*	777.5	818.8	722.2	0.74
	(3826)	(3647)	(3747)		(223.3)	(342.9)	(316.9) *#		(279.9)	(169.6)	(196.3)	
G-CSF	339.4	435.6	442.2	0.86	333.9	449.6	195.3	0.31	116.6	140.7	206.7	0.60
	(308.8)	(342.7)	(416)		(297)	(348.3)	(139.3)		(95.7)	(88.5)	(226)	
GM-CSF	78.9	58.6	67.4	0.98	79.4	55.8	61.8	0.88	33.2	21.1	59.0	0.66
	(101.8)	(83.3)	(132.6)		(92.0)	(78.9)	(64.6)		(44.4)	(55.6)	(102.8)	
IFNy	25.4	52.0	89.7	0.70	74.1	11.7	16.6	0.38	13.4	57.4	20.8	0.58
	(36.6)	(113.5)	(192.0)		(142.8)	(22.0)	(18.2)		(18.0)	(118.2)	(41.4)	
IL-1A	389.8	347.7	248.5	0.33	423.4	149	109.4	0.01**	329.6	399.7	514.6	0.74
	(152.8)	(90.7)	(221.2)		(253.8)	(79.4)	(47.5) *#		(182)	(153.6)	(631.4)	
IL-2	109.3	159.3	164.7	0.93	111.8	51.5	51.3	0.43	56.6	19.6	41.2	0.60
	(167.2)	(325.4)	(334.9)		(105.0)	(75.6)	(75.3)		(92.5)	(38.5)	(57.0)	
IL-3	7.9	9.4	13.9	0.79	3.9	5.9	4.8	0.87	0	0.3	1.3	0.49
	(8.8)	(10.7)	(23.6)		(4.2)	(6.5)	(7.1)		(0)	(0.6)	(3.0)	
IL-4	2.5	8.2	7.9	0.59	2.9	3.0	3.5	0.97	0.1	0.1	5.9	0.25
	(3.0)	(16.7)	(7.8)		(4.7)	(3.3)	(5.2)		(0.1)	(0.2)	(11.0)	
IL-5	19.4	37.3	23.4	0.58	27.1	33.7	22.2	0.80	6.5	6.4	15.1	0.26
	(16.9)	(44.9)	(17.5)		(33.2)	(37.9)	(14.3)		(3.6)	(5.9)	(15.2)	
IL-7	24.6	30.2	260.7	0.43	97.6	32.4	5.0	0.32	1.5	5.1	104.0	0.41
	(27.0)	(33.1)	(600.9)		(177.8)	(48.9)	(9.2)		(2.2)	(9.7)	(241.2)	
IL-9	143.7	139.7	378.0	0.57	235.6	48.5	105.8	0.29	243.6	82.6	128.9	0.42
	(145.9)	(151.3)	(724.7)		(318.7)	(36.6)	(139.1)		(334.6)	(88.5)	(112.0)	
IL-12	36.9	18.3	35.6	0.68	10.4	13.0	6.5	0.69	0(0)	122.9	0	0.40
(p40)	(48.8)	(22.2)	(43.9)		(14.6)	(15.1)	(9.5)			(291.5)	(0)	
IL-12	40.4	62.7	241	0.17	28.2	70.8	62.4	0.75	0 (0)	15.6	281.2	0.42
(p70)	(45.8)	(108.2)	(310.2)		(30.6)	(129.8)	(88.3)			(38.2)	(657.5)	
IL-13	137.6	133.5	210.2	0.84	231.7	80.3	87.0 (111)	0.43	62.9	55.9	107.3	0.62
	(141.2)	(147.7)	(383.2)		(356.4)	(74.3)			(65.5)	(32.4)	(146)	
IL-15	324.0	316.1	239.3	0.89	1231	269.5	58.6 (69)	0.34	3.0	140.4	983.7	0.42
	(357.3)	(357.3)	(294.7)		(2454)	(473.2)			(6.8)	(203.1)	(2191)	
IL-17	8.7	6.0	6.3	0.86	4.8	4.9	3.8	0.92	3.0	2.8	11.9	0.46
	(8.1)	(9.4)	(9.7)		(3.5)	(5.4)	(5.3)		(1.9)	(3.4)	(22.8)	
LIF	4.8	5.5	5.1	0.97	4.0	4.3	0.3	0.31	0	1.8	11.1	0.44
	(5.2)	(6.1)	(5.7)		(6.7)	(5.3)	(0.4)		(0)	(3.5)	(25.3)	
LIX	6937	8908	6397	0.61	9955	5235	4320	0.31	9699	10211	7224	0.24
	(4608)	(5462)	(3134)		(8707)	(5285)	(4376)		(1509)	(1959)	(4547)	
M-CSF	35.3	25.8	24.1	0.82	19.1	23.0	15.5	0.84	0.6	28.4	0.9	0.16
	(38.7)	(31.7)	(27.3)		(25.3)	(23.7)	(17.0)		(1.1)	(44.7)	(0.8)	
MIG	474.0	345.3	345.7	0.79	287.6	368.3	130.2	0.37	85.2	142.1	114.3	0.49
	(417.7)	(353.3)	(344.2)		(193.5)	(438.5)	(82.1)		(36.7)	(110.6)	(58.0)	
MIP-1A	249.6	333	587.6	0.69	417.8	178.9	191.2	0.28	261.4	109.2	257.8	0.58
	(373)	(539.3)	(1032)		(401)	(207.3)	(150.7)		(302.7)	(133.0)	(146.2)	
MIP-1B	126.5	104.7	144.1	0.92	217.2	97.1	123.7	0.62	88.3	100.7	210.6	0.55
	(161.5)	(126.1)	(204.2)		(328.9)	(124.6)	(130.1)		(116.3)	(155.2)	(286.9)	
MIP-2	236.8	195.8	256.3	0.64	127.4	149.8	111 (47.7)	0.40	168.8	198.9	161.7	0.33
	(113.5)	(93.1)	(123.7)		(21.6)	(62.6)			(21.4)	(69.6)	(15.0)	
RANTES	49.6	76.7	96.0	0.42	53.6	59.1	58.9	0.98	25.13	36.77	52.65	0.54
	(44.4)	(57.5)	(73.9)		(69.5)	(57.2)	(30.2)		(22.17)	(44.93)	(46.62)	
Values are dis	played in pg	/mL and an	alyzed by or	e-way ANO	VA. * indicate	essignifica	ntly different fr	om chow die	et. # is signific	antly differ	ent from HF	diet by
Tukey's post-h	octest. P<0).05 is signi	ificant									

	Bivaria	te (r)	Multivariate (β, r ²⁾						
Parameter	IVD	Knee OA	IVD degeneration	Knee OA (β)	Whole				
	degeneration		(β)		model (r ²)				
Behavioral									
Von Frey	0.002	0.206	-	-	0.053				
Grip Force	-0.015	-0.254*	-6.513	-0.371*	0.102*				
Acetone	0.135	0.038	-	-	0.021				
Tail									
suspension									
Rearing	0.089	0.042	-	-	0.004				
Self-Support	0.012	0.047	-	-	0.009				
Immobility	0.063	0.061	-	-	0.017				
Stretch	0.019	0.001	-	-	0.003				
Open Field									
Distance	0.090	0.109	-	-	0.022				
travelled									
Rest time	0.140	0.025	-	-	0.048				
Rearing	0.106	0.015	-	-	0.019				

 Table 3.6, Supplementary Table 4: Association between behavioral indicators of pain and histological joint damage.

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Chapter 4

Role of PPAR δ in the regulation of joint degeneration: context-specific roles in obesity and aging

Chapter 4 is authored by Kerr, G.J., To, B., Margalik, D., Stone, L.S., Beier, F., and Séguin, C.A., and is entitled Role of PPARδ in the regulation of joint degeneration: context-specific roles in obesity and aging. This study is currently unpublished but is in preparation for submission.

4 Role of PPAR δ in the regulation of joint degeneration: context-specific roles in obesity and aging

4.1 Co-authorship Statement

G.J. Kerr performed most of the experiments, contributed to study design and wrote the manuscript. B. To bred, genotyped, fed, and weighed the mice for the diet study (**Figure 4.1**). Additionally, B. To conducted the whole-body composition analysis (**Figure 4.1C-D**) and generated the reconstructed micro-CT image (**Figure 4.1E**). D. Margalik assisted with histology and scoring of the lumbar spine (**Figure 4.4**). B. To conducted all histological and histomorphometry analysis on the knee joints and G. J. Kerr assisted with histopathological scoring (**Figure 4.6**). Manuscript was written by G.J. Kerr with suggestions from Dr. C.A. Séguin.

4.2 Chapter Summary

Introduction: Despite being a major cause of disability worldwide, clinical treatment of intervertebral disc (IVD) degeneration and associated back pain is limited to symptomatic relief. While epidemiological studies have highlighted obesity as a major risk factor for the development of IVD degeneration and back pain, the biological mechanisms underlying this association are unknown. A potential mediator of obesity-associated IVD degeneration is the nuclear receptor peroxisome proliferator-activated receptor delta (PPAR δ), as its ligands are dysregulated in obesity and its deletion has previously been shown to play a protective role in articular cartilage in the context of osteoarthritis.

Methods: *Ex vivo*, intact mouse IVD explants were treated with a synthetic agonist of PPARδ (GW501516) and gene expression and glycosaminoglycan breakdown in the IVD were evaluated. *In vivo*, the role of PPARδ was assessed in an age- and obesity-induced models of IVD degeneration using male and female *Col2-Cre;Ppard*^{fl/fl} conditional knockout mice (*Ppard* KO). Following 50 weeks of experimental diet, mice were assessed for histological and molecular indicators of joint degeneration and behavioral indicators of pain.

Results: PPAR δ is expressed in the NP and AF and responds to PPAR δ agonism by increased expression of downstream target genes, indicating PPAR δ is functionally active in the IVD. Following 50 weeks of western diet, male mice showed indicators of IVD degeneration - changes not detected in female mice - and osteoarthritis induction in both sexes. *Ppard* KO mice showed protection from age-associated IVD degeneration but not from obesity-associated IVD degeneration or knee osteoarthritis. Pain-related behaviors were detected more consistently in female than male mice, despite a lack of

histopathological IVD degeneration. Bivariate and multivariate regression modeling indicates that sex and diet are robust predictors of many behavioral, histopathological, and molecular changes observed, while loss of PPAR δ is only a predictor of knee OA when sex and diet are accounted for. Regression analysis also reveals that alterations in pain-related behavior are independent of structural IVD degeneration.

Conclusion: While PPAR δ deletion may protect from age-associated IVD degeneration, it does not protect against obesity-associated IVD degeneration but may influence the pain associated with joint damage. This data demonstrates the complex, and context-dependent role of PPAR δ in the IVD and the need to evaluate multiple models of IVD degeneration for potential of clinical translation of novel therapeutic targets.

4.3 Introduction

Low back pain (LBP) is the single largest cause of disability worldwide, with a lifetime prevalence over 84% in Canada^{1,2}. Despite efforts to improve the clinical management of LBP, treatments are limited to symptomatic relief without treating the underlying cause of the pain³. While several tissues appear to be involved in LBP, including the paraspinal muscles, ligaments, and facet joints^{4–6}, degeneration of the fibrocartilaginous intervertebral disc (IVD) is believed to be the major contributor to pain in approximately 40% of cases⁴.

IVD degeneration has traditionally been considered a condition of "wear and tear"; however, this paradigm has shifted in recent years with the realization that IVD degeneration is an active process starting at the cellular level, ultimately leading to structural and functional failure^{7,8}. Although the etiology of IVD degeneration remains

unclear, it appears to be multifactorial, as epidemiological studies have highlighted several modifiable and non-modifiable risk factors^{7–10}. Of these, obesity is a major modifiable risk factor for the development of IVD degeneration and LBP. Obesity — defined as a body mass index (BMI) over 30 — substantially increases the risk of developing several metabolic, cardiovascular, neurological and musculoskeletal diseases¹¹. In the spine, increased body weight is associated with indices of lumbar disc degeneration including disc space narrowing and decreased lumbar disc signal intensity detected by MRI^{12,13}. Increased fat mass is also associated with increased LBP intensity and associated disability¹⁴. With the prevalence of obesity nearly tripling over the last 30 years¹¹, it poses a large public health concern.

Despite the clinical associations between obesity, IVD degeneration, and LBP, the biological mechanisms underlying this association remain elusive. While increased mechanical load is thought to underlie in part these associations¹⁵, systemic factors including the chronic low-grade inflammation associated with obesity are also likely involved¹⁶. Metabolic and inflammatory abnormalities associated with obesity contribute to the development of osteoarthritis (OA) ¹⁷, a degenerative musculoskeletal disease with many similarities to IVD degeneration¹⁸. Obesity is associated with chronic metabolic disorders including hypertension, diabetes mellitus and dyslipidemia, collectively known as metabolic syndrome¹⁹. In the context of OA, it is postulated that each component of metabolic syndrome may independently contribute to disease progression, as comprehensively reviewed by Zhuo et al.²⁰. Specifically, alterations in the release of systemic factors (inflammatory cytokines, adipokines), nutrient exchange, advanced glycation end-products (AGEs) levels and glucose/lipid metabolism are believed to be

major contributors to OA progression^{17,20}. Studies from multiple groups have demonstrated that obesity induced by a high-fat diet accelerates the progression of both age- and surgically-induced knee OA using mouse and rat models^{21–25}, accompanied by behavioral indicators of pain²¹.

While not fully explored in the context of musculoskeletal diseases, the peroxisomeproliferator activated receptor (PPAR) family are current therapeutic targets for the management of metabolic syndrome²⁶. The PPAR family of nuclear receptors consists of three isoforms (alpha, delta, gamma), that are activated by circulating fatty acid ligands²⁶, and play distinct roles in energy balance and lipid/ glucose metabolism²⁷. The two most studied isoforms, PPAR α and PPAR γ , are highly expressed in the liver and adipose, respectively, and agonists of these receptors are being used clinically for obesity-associated conditions including dyslipidemia and type II diabetes mellitus^{26,28,29}. Studies investigating the role of PPARy in cartilaginous tissues demonstrated that cartilage-specific deletion of PPAR γ leads to spontaneous OA development in mice³⁰. In contrast to PPAR α and PPAR γ , PPAR δ has broad expression patterns and affects glucose and lipid metabolism, cell differentiation, proliferation, apoptosis and immune regulation³¹. Using transgenic mouse models, studies have shown PPARS agonism improves insulin resistance, fat burning, and muscle endurance^{32–34}; however, PPAR δ agonism has also been shown to induce catabolic processes within articular cartilage and chondrocytes^{35,36}, suggesting tissue-specific or context dependent functions of PPARS. In explant and cell culture models, PPARS agonism leads to aggrecan and glycosaminoglycan degradation in cartilage through increased expression of Mmp-3 and Adamts-535,36, matrix remodelling enzymes implicated

in both OA and IVD degeneration^{37,38}. Moreover, mice with deletion of PPAR δ in articular cartilage are protected from cartilage damage in a post-traumatic model of OA³⁵.

Given that its ligands are dysregulated in obesity and the potential role it plays in articular cartilage, the present study aimed to investigate the role of PPAR δ in the IVD. We first investigated the effects of PPAR δ activation using an *ex vivo* organ culture system and validated that PPAR δ was functionally active within the IVD. We then examine the role of PPAR δ in the IVD *in vivo* using *Col2-Cre;Ppard*^{*fl/fl*} mice to delete PPAR δ within the NP and inner AF. We used this mouse model to examine the role of PPAR δ in both age-associated and obesity-associated IVD degeneration and pain, using both male and female mice to specifically investigate sex as a variable.

4.4 Methods

4.4.1 Mice and Ex Vivo organ culture

Ten-week-old, C57BL/6N mice (Charles River: Wilmington, MA, USA) were used to isolate lumbar and thoracic spinal segments. Following removal of surrounding ligaments and musculature, IVDs were isolated from the vertebral columns using a stereoscope. Intact IVDs were used in *ex vivo* culture experiments. NP and AF tissue were further microdissected (as reported previously³⁹) for gene expression analysis. For *ex vivo* organ culture, isolated IVDs (NP, AF, CEP) were equilibrated in organ culture media (DMEM low glucose (1g/L), 1% FBS, 1 % penicillin/streptomycin) overnight at 2% O₂. On the following day, explants were treated with PPARδ agonist GW501516 (Cedar Lane Labs: Burlington, ON, CAN) at concentrations of 0.01, 0.1, or 1 μ M in organ culture media for

PPARδ-KO mice were generated by breeding mice carrying the *Ppard* gene flanked by loxP sites (*Ppard*^{fl/fl}; B6.129S4-Ppard^{tm1Rev}/J, The Jackson Laboratories)⁴⁰ to mice carrying Cre recombinase under control of the type II collagen promotor (*Col2-Cre*)⁴¹, and genotyping was performed as previously reported⁴¹. Mice without cre-recombinase were used as controls (*Ppard*^{fl/fl}; *Col2-Cre*⁻ hereinafter referred to as WT) and cre-positive mice used as knockouts (*Ppard*^{fl/fl}; *Col2-Cre*⁺ hereinafter referred to as *Ppard* KO).

To determine the role of PPAR δ in the IVD, male and female mice were aged to 14 (n=9-13 per sex) or 25-months (n=4 male) of age and spinal tissues were harvested for histological and molecular analysis. To determine the role of PPAR δ in obesity-induced joint degeneration, (**Supplementary Figure 4.1**) mice were fed standard chow (Envigo 2018) until 10-weeks of age, when they were randomized into one of two diet groups (n = 9-13 mice per diet/ per genotype/ per sex;): standard chow (18% kcal fat, 58% kcal carbohydrate) or a high-fat, high-sugar western diet (45% kcal fat, 41% kcal carbohydrate; Envigo TD.10885) (**Supplementary Table 3.1**). Mice remained on the experimental diets until sacrifice at 60 weeks-of-age (50 weeks on diet). Mice were housed in standard cages and maintained on a 12 hr light/dark cycle, with food and water were consumed *ad libitum*; food consumption and body weight were measured weekly. All aspects of the study were conducted in accordance with the policies and guidelines set forth by the Canadian Council on Animal Care and were approved by the Animal Use Subcommittee of the University of Western Ontario (protocol 2017-154; **Appendix B**).

4.4.2 Dimethylmethylene blue assay

The dimethylmethylene blue dye binding assay was performed on digested tissue explants and media collected from IVD explant cultures. Following 4 days in culture, explants were digested for 3 days in papain digest (125 μ g/mL; Millepore Sigma Canada: Oakville, ON, CAN) at 60°C. The dimethylmethylene blue dye binding assay was performed as previously described⁴², and absorbance values were measured at a wavelength of 595 nm, with a reference wavelength of 655 nm on the Tecan Safire plate reader. GAG concentration in the media and tissue content were calculated based on a chondroitin sulfate standard curve, and GAG release into the media was normalized to the total GAG in each sample following papain digest.

4.4.3 Histological analysis

For *ex vivo* organ cultures, one lumbar IVD from each treatment group was fixed overnight in 4% PFA, decalcified for 24h in TBD-2 (Thermo Fisher Canada: Mississauga, ON, CAN) and paraffin embedded. For the *in vivo* experiments, intact lumbar spine segments (L1-S1) and knees were isolated, fixed, decalcified and paraffin embedded, as previously described⁴³. IVD explants and lumbar spines were sectioned sagittally, and knees were sectioned coronally at a thickness of 5 μ m using a microtome (Leica Microsystems: Wetzlar, DEU) and stained using 0.1% Safranin-O/0.05% Fast Green or 0.04% Toluidine Blue, using methods previously described^{43,44}.

To evaluate IVD degeneration, spine sections were scored by 2 independent scorers using the modified Boos system⁴⁵. Knee joint health was assessed using the murine Osteoarthritis

Research Society International (OARSI) histopathological scale⁴⁶. Articular surfaces of the medial femoral condyle (MFC), medial tibial plateau (MTP), lateral femoral condyle (LFC) and lateral tibial plateau (LTP), were scored by two blinded observers and averaged. For each joint surface, scores from 10 serial sections spanning 500 μ M of the joint were summed to represent OARSI score for each quadrant. Total scores from each of the four quadrants were then added together to generate whole joint OARSI score.

4.4.4 Gene expression analysis

were placed were placed in TRIzol reagent (Thermo Fisher Intact IVDs Canada: Mississauga, ON, CAN) (n=5-6 thoracic/lumbar IVDs per ex vivo treatment group; 4-5 thoracic IVDs per mouse from 5-8 mice per diet/per timepoint) and homogenized using a PRO250 tissue homogenizer (PRO Scientific: Oxford, CT, USA). To assess tissue-specific differences in gene expression, NP and AF tissue were further microdissected (as reported previously³⁹) from WT mice and placed in in TRIzol reagent (Thermo Fisher Canada: Mississauga, ON, CAN) and homogenized manually using a pestle or using a PRO250 tissue homogenizer (PRO Scientific: Oxford, CT, USA) for NP and AF samples, respectively. RNA was extracted according to manufacturer's instructions and quantified using a NanoDrop 2000 spectrophotometer (Thermo Fisher Canada: Mississauga, ON, CAN). For the ex vivo and in vivo studies 0.5 µg RNA was reverse transcribed into complementary DNA (cDNA) (iScript; Bio-Rad Laboratories (Canada): Mississauga, ON, CAN), while 0.15 µg RNA was reverse transcribed for the NP/AF comparison due to sample size limitations. Gene expression was assessed by real-time PCR using a Bio-Rad CFX384 instrument. PCR analyses were run

in triplicate using 120 ng of cDNA per reaction and 310 nM forward and reverse primers with 2x SsoFast EvaGreen Supermix (Bio-Rad Laboratories (Canada): Mississauga, ON, CAN) using optimized PCR parameters and primers at an annealing temperature of 60°C (**Supplementary Table 4.1**). Primers were designed and validated to have efficiency values between 90 and 120%. For the *ex vivo* study and NP/AF comparison, transcript levels were calculated using $\Delta\Delta$ Ct, with data normalized for input based on Ribosomal protein S29 (*Rps29*) and expressed relative to DMSO treated controls or AF samples, respectively. For the *in vivo* studies, transcript levels were calculated relative to a 6-point standard curve made from pooled cDNA generated from murine IVD explants treated with lipopolysaccharide for 4 days (50 mg/mL; Thermo Fisher Canada: Mississauga, ON, CAN).

4.4.5 Micro-computed tomography (Micro-CT)

At sacrifice, mice in the diet study were immediately imaged by micro-CT (Locus Ultra, GE Healthcare Biosciences). Whole-body scans were acquired using 1000 projection images obtained over a single 16 s rotation (80 kv, 55 mA tube current, 16 ms exposure). A calibrating phantom composed of air, water and cortical bone-mimicking epoxy (SB3; Gammex, Middleton WI, USA) was scanned together with each of the animals. Data were reconstructed into 3D volumes with an isotropic voxel spacing of 154 µm and scaled into Hounsfield units (HU). Using MicroView software (GE Healthcare Biosciences) three signal-intensity thresholds (-200, -30, and 190 HU) were used to classify each voxel as adipose, lean, or skeletal tissue, respectively. Custom software was used to calculate tissue

masses from assumed densities of 0.95 (adipose), 1.05 (lean), and 1.92 (skeletal) g/cm³. Volumetric measures were averaged over voxel volume⁴⁷.

4.4.6 Behavioral measures of pain

In the week prior to endpoint, mice in the diet study were assessed for behavioral indicators of pain. All mice were moved from conventional housing to the neurobehavioral testing facility two weeks prior to testing and habituated to all behavioral tests one week prior to data collection. On data collection days, animals were habituated to the testing room for 1 h before test start. To avoid confounding variables associated with the diurnal cycle, all behavioral testing was completed between 8 and 11 AM.

4.4.6.1 Stretch-induced axial discomfort

Stretch-induced axial discomfort was measured using the grip force assay, as described previously^{48–50}. For the grip force assay mice grabbed onto a metal bar attached to a grip force meter, and gently pulled back by their tails to exert axial stretch. Tolerance was assessed by measuring the grip strength (Grip Force Meter, Stoelting Co., Wood Dale, IL), in grams, for each mouse at the point of release.

4.4.6.2 Hind limb sensitivity to cold and mechanical stimuli

Hind paw sensitivity to cold was assessed by measuring the time spent in behavior evoked by evaporative cooling of acetone (flicking, stamping or licking of ventral surface of the paw) during the first 40 s following application of 50 μ L acetone to the ventral surface of the hind paw. Mechanical sensitivity of the hindlimb was measured through application of calibrated Von Frey filaments (Stoetling Co., Wood Dale, IL) to the plantar surface of the hind paw for 3 s or withdrawal. 50% withdrawal threshold was calculated using the Chaplan up-down method⁵¹, with a starting stimulus intensity of 1.4 g.

4.4.6.3 Spontaneous activity

Spontaneous locomotor activity was assessed using open field activity monitors (AccuScan Instruments, Omnitech Electronic, Columbus, OH). Mice were placed into individual boxes and their activity was monitored over 30 min.

4.4.6.4 Rotarod

Locomotor capacity was assessed with rotarod (30 mm diameter; Omnitech Electronics Inc: Columbus, OH, USA) that accelerated (0-30 rpm) for the first 60s and maintained at a maximal speed for an additional 240s. The experimental endpoint occurred when the animal fell from the rotating rod, detected by an infrared sensor. Animals that fell off once within the first 30 s were placed back onto the rod, and video recordings were used to confirm latency times.

4.4.7 Pharmacological manipulation of pain

To assess whether behaviors indicative of stretch-induced axial discomfort (grip force) and mechanical sensitivity (Von Frey) were caused by pain, assays were repeated on all mice following administration of a weak opioid (Tramadol hydrochloride, 50 mg/kg, intraperitoneal (i.p.); Medisca: Richmond, BC, CAN). Testing was conducted 20 min and 45 min after tramadol injection for the Von Frey and grip force assay, respectively.

4.4.7.1 Tail flick test

To assess the anti-nociceptive effects of tramadol, the latency of tail flick response was measured prior to injection, 20 min and 60 min after injection. Tail flick latency is the amount of time between applying heat to the tip of the tail through submersion in a hot water bath (set to 52°C), and voluntary tail withdrawal. The cutoff time was 12 s to prevent tissue injury and the antinociceptive response to tramadol was expressed as a percentage of the maximal possible effect (%MPE), as previously reported⁵².

4.4.7.2 Spontaneous locomotion

To assess whether tramadol had a sedative effect, spontaneous activity in open field was measured over 10 min, 35 min after tramadol injection.

4.4.8 Statistical analysis

For all assays except histopathological scoring of the joints, groups of male and female mice were compared between the different diet groups and genotypes by two-way ANOVA with Tukey's multiple comparisons test. For histopathological scores, within each timepoint mice were compared between the different diet groups by non-parametric Mann-Whitney U test. P < 0.05 was considered significant. Statistical analysis was conducted using GraphPad Prism 8 (Graphpad Software: San Diego, CA, USA).

To assess the effect of sex, genotype, and diet on behavioral, molecular, and histological changes, bivariate and multivariate linear regression models were used to identify which variables remained independently associated with all other outcomes. Bivariate and

multivariate modelling was conducted using STATA 16 (StataCorp LLC: College Station, TX, USA).

4.5 Results

4.5.1 IVD cells express PPARδ and respond to pharmacological activation

To determine whether PPAR δ was expressed and functional, intact IVD explants from 10week-old WT mice were treated with increasing concentrations of the synthetic PPAR δ agonist GW501516 for 4 or 7 days (Figure 4.1A). After both 4 (Figure 4.1B) and 7 days (Figure 4.1C) of treatment, expression of known downstream PPAR δ effector genes including Pyruvate Dehydrogenase Kinase 4 (Pdk4) and Angiopoietin-Like 4 (Angptl4) were significantly increased with agonist treatment. To assess the relative expression of PPAR8 in distinct IVD compartments, expression of Ppard was assessed NP and AF tissues isolated by microdissection from 10-week-old WT mice. NP cells showed significantly lower expression of *Ppard* than AF cells (Figure 4.1D). Genes involved in metabolism and oxidative stress altered by PPAR δ agonism in chondrocytes were also evaluated³⁶. No significant difference in the expression levels of ATP-binding cassette transporter (Abca1), carnitine palmitoyltransferase 1A (Cpt1a), Glutathione S-transferase A4 (Gsta4), or Insulin Induced Gene 1 (Insig1) were detected in the IVD following agonist treatment (Figure 4.11, Supplementary Figure 2). After 7 days, treatment with 0.01 µM GW501516 significantly upregulated the expression of Thioredoxin-interacting protein (*Txnip*) in IVD explants (Figure 4.11, Supplementary Figure 2B).



Figure 4.1: PPAR δ is expressed in the IVD and responds to pharmacological agonism. (A) Representative histological sections of IVD explants cultured for 4 or 7 days in media supplemented with DMSO (vehicle control) or the PPAR δ -specific agonist GW501516 (0.01 µM, 0.1 µM, or 1 µM) stained with Safranin-O/Fast Green. (B & C) SYBR-based real-time qPCR analysis of IVD explants treated with GW501516 for 4 or 7 days show significant increases in the expression of known PPAR δ targets *Pdk4* and *Angptl4* (n=5-6 biological replicates). (D) SYBR-based real-time qPCR analysis of *Ppard* expression in NP and AF tissues isolated by microdissection. The expression of *Ppard* in the AF is significantly higher than expression in the NP (n=10-11). Data are presented as mean ± 95% CI. *P<0.05, **P<0.01, ***P<0.001, ***P<0.001 by One-way ANOVA with Tukey's test compared to DMSO control for organ culture and by Welch's t-test for comparison of AF and NP tissue samples.

4.5.2 Effect of PPARδ activation on proteoglycan levels in the IVD

To determine the effects of PPAR δ activation on glycosaminoglycan (GAG) levels within intact IVDs, explants were treated with a vehicle control (DMSO), 0.01 μ M, 0.1 μ M, or 1 μ M GW501516 for 4 days (**Figure 4.12, Supplemental Figure 3**). While no gross morphological changes were observed, an apparent loss of proteoglycan staining was observed in explants cultured in the presence of 1 μ M GW501516 compared to DMSO control (**Figure 4.12, Supplementary Figure 3A**). To confirm these observations, levels of GAG were quantified in both the tissue explants and culture media using the dimethylmethylene blue dye binding (DMMB) assay. PPAR δ agonism was associated with a slight but not statistically significant increase in the release of GAG into the media (**Figure 4.12, Supplementary Figure 3B**). SYBR-based qPCR analysis was used to assess the expression of extracellular matrix genes and genes involved in matrix degradation. No significant differences were seen in the expression of matrix degrading enzymes (**Figure 4.12, Supplementary Figure 3C**) or extracellular matrix components (**Figure 4.12, Supplementary Figure 3D**).

4.5.3 Deletion of PPARδ may protect against age associated IVD degeneration

We next assessed the effect of PPARδ deletion on age associated IVD degeneration. A conditional knockout mouse was generated using the Col2-Cre that our lab has demonstrated is expressed throughout the nucleus pulposus, inner two thirds of the AF and cartilage endplate, in addition to in addition to cartilaginous tissues, similar to other Col2-

Cre transgenic mouse lines^{53,54}. WT and *Ppard* KO mice were aged to either 14 or 25 months-of-age and IVD health was assessed through histological analysis. At the 14-month timepoint, no discernable differences were detected in IVD health between WT and *Ppard* KO mice (**Figure 4.2A**). Accordingly, histopathological scoring showed no difference between WT and *Ppard* KO mice at any individual spinal level or in the average histopathological score in both male and female mice (**Figure 4.2B**). At the 25-month timepoint, male WT mice demonstrated advanced IVD degeneration, including a loss of proteoglycan content in the NP, accumulation of fibrotic matrix in the NP, and the loss of a distinct boundary between the NP and AF. In contrast, *Ppard* KO mice show more consistent maintenance of overall IVD structure and preservation of the NP cellularity and proteoglycan content (**Figure 4.2C**). While histopathological scoring indicated that *Ppard* KO animals had less degeneration on average than WT mice, the differences were not significant (**Figure 4.2D**).

4.5.4 Wild-type and *Ppard* KO mice gain weight on the western diet To investigate whether PPARδ would play a similar role in obesity-induced IVD degeneration, at 10 weeks of age male and female WT and *Ppard* KO mice were randomized into groups fed either a high-fat/high-sugar western diet or chow control for 50 weeks (**Figure 4.10, Supplemental Figure 1**). As expected, following 50 weeks on the experimental diet both male and female mice fed the western diet showed a significant increase in body mass (**Figure 4.3A**) and weight gain (**Figure 4.3B**) compared to sex and genotype-matched chow fed controls. Analysis of whole-body composition by micro-CT demonstrated that for both WT and *Ppard* KO, male and female mice fed the western diet



Figure 4.2: Loss of Ppard alters age associated IVD degeneration.

Representative images of mid-sagittal sections of lumbar IVDs stained with Safranin-O/Fast Green from 14-month old or 25-month-old wild-type (WT) and PPAR\delta knockout (*Ppard* KO) mice paired with evaluation of the grade of histopathological degeneration using the modified Boos scoring system. (A) At 14 months-of-age, no differences in was detected in the histological appearance of IVD tissues between WT and PPAR^δ KO mice. (B) At the 14-month timepoint, no differences in histopathological scores of IVD degeneration were seen between genotypes at either individual lumbar spinal levels or in the average degeneration score calculated for all lumbar IVDs. (C) At 25 months-of-age, male *Ppard* KO mice lacked histopathological signs of advanced IVD degeneration seen in WT mice, including a loss of proteoglycan content and accumulation of fibrotic matrix in the NP and the loss of a distinct boundary between the NP and AF. (D) At the 25-month timepoint, male Ppard KO mice showed lower degenerative scores at individual lumbar levels compared to WT; however, these changes did not result in significant differences in histopathological scores. Data are plotted mean ± SEM and analyzed by Mann-Whitney Utest (n=9-13 for 14-month timepoint; n=3-5 for 25-month timepoint). *P<0.05, ***P<0.001.





(A) Following 50 weeks on the western diet, WT and *Ppard* KO mice showed a significant increase in body weight compared to chow-fed control. (**B**) Mice fed the western diet showed increased weight gain from baseline compared to chow-fed controls, regardless of sex or genotype. (C) Representative reconstructed micro-CT images of mice following 50-weeks of experimental diets. Isotropic surface-rendering of skeletal tissue (indicated in white) is overlaid with a mid-coronal slice where lean tissue is indicated in red and adipose tissue is indicated in yellow. Micro-CT based analysis of whole-body composition demonstrated a significant increase in (D) adipose tissue mass in all groups of mice fed the western diet compared to chow-fed controls; however, no differences were detected in (E) bone mineral density (BMD). n=9-13 animals per genotype, sex, and diet. Data are plotted as mean \pm 95% CI. **P*<0.05, ***P*<0.01, ****P*<0.001, *****P*<0.001 by 2-way ANOVA with Tukey's test.

showed an increase in adipose tissue compared to chow controls (**Figure 4.3D**). No significant difference was seen in bone mineral density between diet groups in either WT or *Ppard* KO mice (**Figure 4.3E**).

4.5.5 Deletion of PPARδ does not protect against obesity related IVD degeneration

The effects of the western diet on IVD health was assessed through histopathological evaluation (**Figure 4.4**) and molecular analysis (**Figure 4.5**). In males, mice fed the western diet showed indicators of IVD degeneration not evident in chow control in both genotypes (**Figure 4.4A**). Histopathological changes included abnormal matrix deposition within the NP and a loss of clear demarcation between the NP and AF. Although no overt differences were detected between WT and *Ppard* KO mice, the western diet induced more consistent degeneration in male *Ppard* KO mice than in WT mice. Accordingly, histopathological scoring using the modified Boos scoring system showed significant increases in degeneration in the L6S1 spinal level of male *Ppard* KO mice fed a western diet compared to their chow-fed controls, but not in WT mice (**Figure 4.4B**). In females, no overt differences were detected in the histological appearance of the lumbar IVDs between the diet groups or genotypes (**Figure 4.4C**). As such, histopathological scoring did not show any differences in degeneration of lumbar IVDs between diet groups in female WT or *Ppard* KO mice (**Figure 4.4D**).

To further investigate the role of PPARδ and changes associated with diet-induced obesity, SYBR-based qPCR was used to quantify the expression of extracellular matrix components (**Figure 4.5A**), matrix degrading enzymes (**Figure 4.5B**), and markers of inflammation and neural in-growth (**Figure 4.5C**) in thoracic IVDs. Despite no changes in IVD





Representative mid-sagittal sections of lumbar IVDs stained with Safranin-O/Fast Green from male (A) and female (C) wild-type (WT) and PPAR δ knockout (*Ppard* KO) mice fed either a chow control or western diet for 50 weeks. Male mice fed a western diet showed a loss of the demarcation between the NP and AF compared to chow control. This was observed in both WT and *Ppard* KO male mice, to a greater extent in *Ppard* KO animals. In female mice, no observable differences were seen between diet group or genotype. Evaluation of the grade of histopathological IVD degeneration using the modified Boos scoring system showed that (C) male *Ppard* KO mice fed a western diet showed significantly greater degeneration than their chow-fed controls at the L6/S1 spinal level. (D) No significant differences in histopathological scoring were seen between diets in female mice. n=9-13 mice per sex, genotype, and diet. Data is presented as mean \pm 95% CI. **P*<0.05 by Mann-Whitney U test. histopathology, at 14 months-of-age male *Ppard* KO mice demonstrated increased expression of *Acan, Mmp3, Mmp13*, and *Il-1b* compared to WT mice (chow diet controls). Following 50 weeks of western diet, male WT mice showed a significant increase in the expression of *Acan* and *Il-1b* compared to chow fed controls, while male *Ppard* KO mice showed a significant increase in *Il-1b* expression compared to chow fed controls (**Figure 4.5A & 4.5C**). Male *Ppard* KO mice in the western diet group also showed a significant increase in *Il-1b* expression compared to male WT mice in the western diet group (**Figure 4.5C**). Similar patterns of gene expression were not detected in female mice. No differences in IVD gene expression were detected between female WT and *Ppard* KO mice. In female mice, changes associated with western-diet induced obesity were limited to down-regulation of *Col1a1* expression in WT mice compared to chow fed controls (**Figure 4.5A**).

4.5.6 Deletion of PPARδ does not protect against obesity-related knee OA

Diet-induced obesity contributes to other arthropathies, such as knee OA²⁹. Moreover, loss of *Ppard* protects from surgically induced posttraumatic OA³⁵. We therefore examined the role of PPARδ in obesity-induced OA. In male mice, chronic consumption of the western diet in both WT and *Ppard* KO mice led to increased erosion of the articular cartilage and osteophyte formation at the joint margins compared to sex-matched chow fed controls (**Figure 4.6A**). This corresponded to a significant increase in histopathological scores (**Figure 4.6B**). A similar trend was observed in female mice (**Figure 4.6C**). While female WT mice had more degeneration on average when fed the western diet compared to chow fed controls, this difference was not significant. Consumption of the western diet induced



SYBR-based qPCR analysis was conducted to assess the expression of (A) extracellular matrix genes, (B) matrix remodeling genes, and (C) proinflammatory and neurotrophic factors in IVDs of wild-type (WT) and PPAR δ knockout (*Ppard* KO) mice fed either a chow control or western diet for 50 weeks. Male *Ppard* KO mice showed increased expression of *Acan*, *Mmp3*, *Mmp13*, and *Il-1b* compared with WT mice. No significant differences were detected in female *Ppard* KO mice compared to WT controls. WT male mice fed the western diet showed significantly increased expression of *Acan* and *Il-1b*, while male *Ppard* KO mice fed a western diet only showed significant increases in *Il-*1b compared to genotype-matched chow-fed controls. Female WT mice fed the western diet showed significant decreases in *Col1a1* expression compared to chow-fed controls. n=8 animals per sex, genotype, and diet. Data are plotted mean \pm 95% CI. **P*<0.05, ***P*<0.01, ****P*<0.001 by 2-way ANOVA with Tukey's test.



Figure 4.6: Effect of diet-induced obesity and PPARô deletion on knee joint health. (A & C) Representative coronal sections of the medial knee compartment stained with toluidine blue male and female wild-type (WT) and PPARô knockout (*Ppard* KO) mice fed either a chow control or western diet for 50 weeks. Images are oriented with the medial femoral condyle (MFC) located superiorly, and the medial tibial plateau (MTP) inferiorly. Western diet-induced obesity was associated with indicators of osteoarthritis in all groups of mice assessed compared to chow-fed controls, marked by erosion of articular cartilage (arrows) and osteophyte formation (outlined in white dashed box). (B & D) Quantification of joint degeneration using the murine OARSI histopathological scoring system showed significant increases in joint degeneration in mice fed the western diet compared to chow fed controls for male WT and *Ppard* KO mice as well as female *Ppard* KO mice. n=4-8 animals per sex, genotype, and diet. Data are plotted mean \pm 95% CI. **P*<0.05, ***P*<0.01 by 2-way ANOVA with Tukey's test.

a significant increase in histopathological scores in female *Ppard* KO mice compared to chow fed controls (**Figure 4.6D**).

4.5.7 Behavioral indicators of stretch-induced axial discomfort

To investigate the association between obesity, back pain, and the potential role of PPAR δ , we first investigated whether mice fed the western diet showed behavioral indicators of stretch-induced axial discomfort. Decreased grip force during axial stretch has been previously established as an indicator of discogenic back pain in a mouse model of IVD degeneration³⁸. Loss of PPAR^δ did not alter grip force in either male or female mice (Figure 4.7A). In male WT and *Ppard* KO, mice fed the western diet showed slight decreases in grip force during axial stretch, but these differences were not statistically different from chow fed control. In female WT and Ppard KO, mice fed the western diet showed a significant reduction grip force during axial stretch, indicative of axial discomfort (Figure 4.7A). To determine whether these behaviors were due to pain, the grip force assay was repeated on all mice following administration of a dose of the weak opioid tramadol (50 mg/kg I.P.). For all mice, the anti-nociceptive properties of the drug at this dosage were confirmed with the tail-flick assay (Figure 4.13, Supplementary Figure 4) and potential sedative effects of the drug were evaluated by measuring voluntary open field locomotion (Figure 4.14, Supplementary Figure 5). In male mice, tramadol treatment did not significantly alter grip force in any group (Figure 4.7B). In WT and *Ppard KO* female mice, tramadol treatment did not significantly affect grip force, however, the baseline significant difference associated with western diet-induced obesity in *Ppard* KO mice was no longer evident (Figure 4.7C).



Figure 4.7: Diet-induced obesity reduces grip strength in female mice.

(A) Stretch-induced axial discomfort was assessed in all groups of chow (control) and western diet fed mice. Loss of PPAR δ did not alter grip force in either male or female mice. Female wild-type (WT) and PPAR δ knockout (*Ppard* KO) mice fed the western diet showed reduced resistive grip force during axial stretch compared to chow-fed controls. To determine whether this was associated with pain, the grip force assay was repeated on all mice after receiving a dose of the weak opioid tramadol (50 mg/kg I.P.). (B) In male mice, tramadol did not significantly affect grip force in any group. (C) In female WT mice, tramadol did not significantly alter grip force in mice on both diets and did not prevent the significant difference associated with western diet-induced obesity. In female Ppard KO mice, tramadol did not significantly improve grip force in either diet group; however, the significant difference associated with western diet-induced obesity was no longer evident. Graphs on the left show the percent of baseline, which represents the grip force following tramadol treatment normalized to the pre-treatment (baseline) values for each animal. Graphs on the right show the average grip fore (g) for mice in each group in the absence and following administration of tramadol. n=9-13 mice per sex, genotype, and diet. Data are plotted mean \pm 95% CI. **P<0.01 between diets in each treatment group; †indicates significant (P < 0.05) tramadol-induced differences in grip force for mice fed the chow (black) or western (red) diet by 2-way ANOVA with Tukey's test.

4.5.8 Behavioral indicators of cold and mechanical sensitivity

Cold sensitivity was assessed by measuring the response to evaporative cooling of acetone on the hind-paw. Cold sensitivity was not affected by loss of PPAR δ or by western dietinduced obesity in either male or female mice (**Figure 4.8A**). Mechanical sensitivity was then measured in the hind paw using the Von Frey assay. Loss of PPAR δ did not alter mechanical sensitivity in either male or female mice (**Figure 4.8A**). In males and female, WT mice fed the western diet showed a significant decrease in tolerance to mechanical stimulation compared to sex-matched chow fed controls. In both male and female *Ppard* KO mice, no difference in mechanical sensitivity was seen in mice fed the western diet compared to chow fed controls (**Figure 4.8B**). To assess whether these behavioral differences to mechanical stimulation were due to pain, the Von Frey assay was repeated on all mice following administration of a dose of the weak opioid tramadol (50 mg/kg I.P.). In male (**Figure 4.8C**) and female (**Figure 4.8D**) mice, tramadol significantly decreased mechanical sensitivity (increased withdrawal threshold) and prevented the baseline differences detected in mechanical sensitivity caused by western diet-induced obesity.

4.5.9 Behavioral indicators of physical function and spontaneous locomotion

Locomotor capacity was first assessed using an accelerating rotarod assay where the outcome measure was time to fall. Loss of PPARδ did not alter locomotor capacity in the rotarod assay in either male or female mice (**Figure 4.9A**). Moreover, western diet-induced obesity did not affect performance of male mice in the rotarod assay; no significant difference was seen in time to fall in either WT or *Ppard* KO mice compared to chow fed controls (**Figure 4.9A**). In females however, western diet-induced obesity was associated



Figure 4.8: Diet-induced obesity increases sensitivity to mechanical stimuli in wild-type mice.

(A) Sensitivity to cold was assessed by measuring the time spent in behavior evoked by evaporative cooling of acetone following application to the ventral surface of the hind paw. No differences were detected between any of the groups. (B) Mechanical sensitivity of the hind paw was assessed by manual application of Von Frey Filaments using the Chaplin updown method. Male and female wild-type (WT) mice fed the western diet showed increased mechanical sensitivity compared to chow-fed controls, indicated by a decreased tactile response threshold. Mechanical hypersensitivity was not detected following consumption of the western diet in PPAR^δ knockout (*Ppard* KO) mice. To determine whether mechanical sensitivity was associated with pain, the assay was repeated on all mice after receiving a dose of the weak opioid tramadol (50 mg/kg I.P.). In (C) male and (D) female mice, administration of tramadol significantly decreased mechanical sensitivity; withdrawal thresholds in mice fed the western diet were not different from those of chow-fed controls. Graphs on the left show the percent of baseline, which represents the withdrawal thresholds following tramadol treatment normalized to the pre-treatment (baseline) values for each animal. Graphs on the right show the average grip fore (g) for mice in each group in the absence and following administration of tramadol. n=9-13 animals per sex, genotype, and diet. Data are plotted mean $\pm 95\%$ CI. *P<0.05 between diets in each treatment group; †indicates significant (P < 0.05) tramadol-induced differences in withdrawal threshold for mice fed the chow (black) or western (red) diet by 2-way ANOVA with Tukey's test.



Figure 4.9: Diet-induced obesity affects physical function and locomotion.

(A) Locomotor capacity was assessed using an accelerating rotarod assay. No significant differences were seen in male mice between diet groups or genotypes. In female mice, obesity induced by the western diet was associated with decreased motor function in both wild-type (WT) and PPAR δ knockout (*Ppard* KO) mice compared to chow-fed controls. (B-D) Spontaneous activity was measured over 30 min in open field. No differences in spontaneous activity were observed between WT and *Ppard* KO mice. Compared to chow-fed controls, WT male mice fed the western diet showed reduction in distance and velocity of locomotion as well as vertical activity (rearing); changes not observed in *Ppard* KO mice. Compared to chow-fed controls, female WT and *Ppard* KO mice fed the western diet showed reduction in distance and velocity of locomotion as well as vertical activity of locomotion as well as vertical activity (rearing); changes not observed in *Ppard* KO mice. Compared to chow-fed controls, female WT and *Ppard* KO mice fed the western diet showed reduction in distance and velocity of locomotion as well as vertical activity (rearing); n=9-13 animals per sex, genotype, and diet. Data are plotted mean \pm 95% CI. **P*<0.05, ***P*<0.01, ****P*<0.001 by 2-way ANOVA with Tukey's test.

with a significantly decreased time to fall compared to chow fed controls, for both WT and *Ppard* KO mice (**Figure 4.9A**).

Spontaneous locomotion in open field was also assessed for all mice over a 30 min period. Loss of PPAR δ did not alter total locomotion, velocity of movement or rearing in either male or female mice (**Figure 4.9B-D**). Male WT mice fed the western diet showed a significant decrease in total distance travelled (**Figure 4.9B**), average velocity (**Figure 4.9C**), and the number of rearing events (**Figure 4.9D**) compared to sex-matched chow fed controls. However, these differences were not observed in male *Ppard* KO mice. In females, both WT and *Ppard* KO mice fed the western diet showed significant decreases in the total distance travelled, average velocity, and the number of rearing events compared to sex-matched chow fed controls (**Figure 4.9B-D**). These data suggest that diet-induced obesity impairs physical function in female, and to a lesser degree, in male mice.

4.5.10 Statistical modeling

To account for the large amount of variability within the data and outcome measures assessed in the diet study, linear regression modeling was conducted to assess whether sex, genotype or diet were predictors of the behavioral, histological, and molecular differences observed (**Table 4.1**). Bivariate modeling demonstrated that the western diet alone was a significant predictor of all behavioral alterations except for cold sensitivity, as well as histopathological degeneration of the knee and alterations in extracellular matrix gene expression in the IVD (**Table 4.1**). Sex was found to be an independent predictor of rotarod performance, spontaneous locomotion, histopathological changes to the IVD, and expression of inflammatory and matrix degrading enzymes (*Mmp3, Mmp13, Adamts4, Il-1b, Bdnf*). When sex, genotype and diet were adjusted for using multivariate analysis,

	Bivariate (r)			1 1	Multivariate (β, r ²)			
Parameter	Sex	Genotype	Diet		Sex (β)	Genotype (β)	Diet (β)	Whole Model (r²)
Behavioral								
Von Frey	0.03	0.10	-0.28**		0.03	0.13	-0.35**	0.09*
Grip force	0.09	0.17	-0.57***		2.35	5.36 (p=0.057)	-18.44***	0.36***
Acetone	0.02	0.017	0.10		-	-	-	0.01
Rotarod	0.22*	0.152	-0.33**		10.22*	6.93	-15.47***	0.18***
Open Tield	0 24**	0.04	0 50***		4450 40***	154.00	4025 00***	0 27***
Total Distance	-0.31	-0.04	-0.52		6.20	-154.69	-1930.00	0.37
Rearing	-0.09	-0.19	-0.51		-0.39	-15.85 (p=0.06)	-40.92	0.30
Velocity	0.29**	0.12	-0.48***		1.70***	-0.71	-2.74***	0.34***
Histopathology								
L6S1 Boos	0.323**	0.17	0.21 (p=0.056)		1.67**	0.82	1.08*	0.17**
Average Boos	0.05	0.19	0.18		-	-	-	0.07
Cumulative OARSI score	-0.22	0.28	0.53***		-9.97*	9.83*	19.53***	0.40***
Gene expression								
Col1a1	0.09	0.11	0.06		-	-	-	0.02
Col2a1	0.08	0.23	0.27*		3.51	13.39	16.12*	0.13*
Acan	0.22	0.18	0.34**		16.09	13.09	27.00**	0.19**
Mmp3	0.89***	0.08	0.12		62.14***	1.60	6.03	0.81***
Mmp13	-0.84***	0.14	0.06		-1.38***	0.31**	0.15	0.75***
Adamts4	-0.85***	0.03	0.03		-118.19***	3.75	8.29	0.72***
ll-1b	0.65***	0.11	0.16		4.44***	0.49	0.99	0.45***
11-6	0.13	0.07	0.17		-	-	-	0.06
Bdnf	-0.42 ***	0.03	0.23		-48.38***	0.07	27.11*	0.23**

Table 4.1: Impact of sex, genotype, and diet on behavioral, molecular and histological changes.
similar trends were seen for all the outcomes measured, with sex and diet being covariates for many behavioral and histological outcomes. Focusing on histological outcomes, sex, diet, and genotype were all significant predictors of histopathological knee OA, while sex and diet were significant predictors of IVD degeneration (**Table 4.1**). In addition to looking at the association between sex, diet, and genotype on behavioral outcomes, we wanted to determine whether histopathological joint damage was predictive of pain-related behavior, independent of sex, diet, and genotype. To do so we constructed another bivariate and multivariate linear regression model (**Table 4.3**, **Supplementary Table 2**), and this analysis indicated that knee OA was associated with alterations to grip force and differences in spontaneous locomotion; however, IVD degeneration was not significantly associated with any of these behavioral outcomes.

4.6 Discussion

Previous research suggests that the nuclear receptor PPARδ plays a catabolic role in articular cartilage, and that genetic deletion protects against both post-traumatic and ageassociated OA^{35,55}. Our study represents the first examination of the role of PPARδ in the IVD and the role of PPARδ in an obesity-induced model of IVD degeneration and osteoarthritis. The present study shows that PPARδ is expressed and functionally active in the IVD. We evaluated the effect of PPARδ inactivation on age- and obesity-associated IVD degeneration *in vivo* using *Col2-Cre;Ppard*^{fl/fl} mice to delete PPARδ within the NP and inner AF. With age, *Ppard* KO mice showed considerably less IVD degeneration than WT controls, yet in a western diet-induced obesity model, *Ppard* KO mice showed a higher proportion of moderate-severe IVD degeneration compared to WT animals. Moreover, loss of PPARδ in joint tissues appeared to modulate behavioral indicators of pain and physical function. Taken together, these finding illustrate a complex, and likely context-dependent role of PPAR δ in the IVD and articular cartilage.

A member of the nuclear receptor superfamily, PPAR δ acts as a transcription factor whose ligands are fatty acids and their metabolites, ultimately altering transcription of downstream target genes^{33,56}. Previous studies have highlighted common downstream targets of PPAR δ in chondrocytes as well as other tissues including *Pdk4* and *Angptl4*^{36,57}, which we showed were induced in IVD explants treated with the PPAR δ agonist GW501516. These findings indicate that PPAR δ is functionally active in the IVD. In articular cartilage explants, PPAR δ agonism induces proteoglycan degradation, mediated through upregulation of matrix degrading enzymes including *Mmp3* and *Adamts5*³⁵. Although, IVD explants treated with the PPAR δ agonist showed a loss of safranin-o staining within the NP, no significant differences were seen in the expression of any matrix degrading enzymes investigated. These findings may be impacted by the use of intact IVDs in organ culture and potential differential effects of PPAR^δ agonism in its distinct tissue types. Anatomically, the IVD is a heterogeneous structure composed of three distinct, yet interdependent tissues: the gelatinous nucleus pulposus (NP), the fibrocartilaginous annulus fibrosus (AF), and cartilage endplates. In adults, the major cell type of the NP are chondrocyte-like NP cells, which secrete a matrix very similar to that of articular cartilage, in contrast to AF cells which are phenotypically more similar to fibroblasts⁵⁸. Gene expression analysis in the current study demonstrated that *Ppard* expression is significantly higher in the AF than in the NP, highlighting the need to evaluate the effect of PPAR δ activation in each tissue type independently.

Previous work by our group demonstrated that deletion of PPAR δ in articular cartilage protects against both surgically-induced³⁵, and age-associated knee OA⁵⁵. Consistent with these findings, deletion of PPAR δ within the IVD appears to protect from age associated IVD degeneration, largely within the NP. In our current study, analysis of whole IVD gene expression at 14 months-of-age (a timepoint not associated with histopathological change) showed increased expression of aggrecan as well as genes involved in matrix degradation (*Mmp3*, *Mmp13*) and inflammation (*Il-1b*) in male *Ppard* KO mice. While these results suggest early degenerative changes, inhibitors of IL-1 β signaling and matrix metalloproteinases significantly impact the effects of these proteins^{59,60}, and their expression levels were not quantified and may counteract this increased expression. Further studies are however required to examine the underlying mechanisms behind this protective effect, specifically focused on characterizing molecular changes within IVD subcompartments in *Ppard* KO mice over time.

Given the protective effect of PPAR δ deletion in age-associated IVD degeneration, we wanted to determine whether this phenotype would be recapitulated in a model of diet-induced obesity. Since PPAR δ ligands are dysregulated in obesity³³ we postulated that its deletion would protect against both obesity-induced IVD degeneration and osteoarthritis. In contrast to our hypothesis, our findings demonstrated that western diet-induced obesity accelerated IVD degeneration in male *Ppard* KO mice. Moreover, a greater proportion of male *Ppard* KO mice fed the western diet showed severe degeneration than did male WT mice fed the western diet, although these differences were not significant. Like in the IVD, western diet-induced obesity accelerated OA progression in both WT and *Ppard* KO mice. Although histopathological scores of OA severity were not different between *Ppard* KO

and WT mice on the western diet, statistical modeling identified genotype as a significant predictor of knee joint damage when sex and diet were accounted for. Taken together, these findings underscore a context dependent role for PPAR δ in joint degeneration.

While the exact role of PPAR δ is not known in the IVD, PPAR δ activation regulates the transcription of many genes involved with lipid metabolism in chondrocytes, including enzymes associated with fatty acid oxidation^{35,36}. These studies postulated that in a posttraumatic model of OA, PPAR δ -mediated cartilage degeneration may result from increased production of reactive oxygen species (ROS) as a consequence of increased fatty acid oxidation³⁵. While this increase in ROS production may be detrimental to cartilage or the IVD under physiological conditions, PPARδ-mediated lipid catabolism may be essential in pathological conditions such as obesity. Studies investigating the role of diet-induced obesity determined that free fatty acids induce lipotoxicity in chondrocytes leading to cell death⁶¹. As a major regulator of lipid metabolism, PPARδ may directly contribute to the prevention of lipotoxicity. Supporting this, studies in human skeletal muscle cells demonstrated that PPAR^δ activation prevents lipoxicity, through reductions in fatty-acid induced inflammation and endoplasmic reticulum stress⁶². In addition to the direct effects of PPAR δ deletion, alterations in PPAR α and PPAR γ signaling should also be considered, as share common ligands with PPARS and therefore may contribute to obesity-induced changes in joint tissues⁶³. Together, these putative mechanisms may contribute to the context-specific role demonstrated for PPAR δ in joint degeneration.

Although there are many similarities in the initiation and progression of IVD degeneration and OA (as reviewed in¹⁸), our data highlights important differences. Consistent with our previous study in wild-type mice⁶⁴, animals fed a western diet often showed moderate to

severe knee OA compared to chow control, while alterations to the IVD were subtle and histopathological degeneration was localized to a single spinal level (L6-S1). In the knee, metabolic alterations associated with obesity are postulated to be the major contributor to joint degeneration. Supporting this, leptin deficient mice become obese yet do not develop knee OA, suggesting that systemic factors may play a key role in obesity-induced OA^{65} . In the IVD, the relative contributions of mechanical and metabolic factors to obesity-induced degeneration remain unknown. While speculative, the data from the current study suggests that biomechanical factors may play a more substantial role in obesity-associated IVD degeneration. In this study, obesity-induced histopathological IVD degeneration was localized to the lowest lumbar spinal level (L6-S1). In humans, accelerated IVD degeneration at the lumbosacral joint (L5S1) is attributed to increased mechanical loading at this site⁶⁶. If systemic factors were the major contributor to obesity-associated IVD degeneration in our model, we would expect to see a more homogenous pattern of IVD degeneration at multiple spinal levels; observations not seen in the current study. However, the analysis of IVD degeneration was limited to a single timepoint in the current study and it may be that generalized IVD degeneration may follow a more chronic exposure to the obesogenic diet. It should also be noted that in addition to altered mechanical loading, the size and geometry of IVDs differ based on spinal levels⁶⁷, which may impact the diffusion of nutrients and systemic molecules into the IVD thereby impacting levels of systemic metabolic effectors. Future studies should explore the relative contributions of mechanical and metabolic factors in the development of obesity-induced IVD degeneration and back pain.

Despite the utility of using rodent models to study IVD biology, few studies also assess the clinical problem associated with IVD degeneration - pain⁶⁸. In humans and animals, there is often a discord between radiographic IVD degeneration and pain^{69,70}. For example, two animals may have equally degenerative IVDs, yet have different pain-related behaviors⁷⁰. Although pain cannot be directly measured in animals, several indirect, quantitative behavioral assays have been developed to evaluate pain-like behaviors for distinct pain states^{71,72}. While many behavioral metrics are non-specific and are used to assess OA pain⁷³, stretch-induced axial discomfort is a validated measure of axial low back pain⁵⁰. In the current model, western diet induced obesity was associated with stretch-induced axial discomfort in female mice fed the western diet, measured using the modified grip force assay. To determine whether these differences were truly a measure of pain, mice were treated with Tramadol, a centrally acting opioid analgesic previously shown to improve grip strength and mechanical hypersensitivity in other pain models^{74–76}. However, in the current study, obesity-associated differences in grip force were not altered by tramadol. Since grip force is also used to assess muscle strength⁷⁷, and neuromuscular deficits have been suggested as an initiating factor in obesity-induced OA²¹, we postulate that obesityinduced neuromuscular deficits may contribute to the behavioral differences observed and should be further explored in the context of IVD degeneration and back pain.

Obesity induced by the western diet in WT mice was also associated with mechanical, but not thermal hypersensitivity of the hind paw; findings that were consistent with previous work from our group⁶⁴. Mechanical sensitivity was inhibited by tramadol, indicating that these behavioral differences were likely as a result of pain. Interestingly, *Ppard* KO mice did not show obesity-induced mechanical sensitivity, despite having increased prevalence of IVD degeneration and knee OA. While the mechanisms underlying this phenotype in *Ppard* KO mice are not understood, previous work done by our group has demonstrated that subcutaneous injection of a PPAR δ antagonist reduced pain-related behaviors in a surgically-induced model of OA, despite not improving structural joint degeneration⁷⁸. Together, these findings demonstrate that western diet-induced obesity induces pain related behavior in mice, and intriguingly, that loss of PPAR δ in joint tissues may modulate this outcome.

Our study also demonstrated that obesity-induced pain behaviors were more consistent in female compared to male mice. These findings are supported by both clinical and basic research, which show that the prevalence of chronic pain is higher in females than males⁷⁹. While several underlying mechanisms have been suggested, including cognitive and sociocultural differences, differences in gonadal sex hormones also play a biological role in pain⁷⁹. Several studies establish the pro-nociceptive role of estrogen that is mediated through its receptors, estrogen receptor α (ER α) and estrogen receptor β (ER β)⁸⁰. In normal mice and in a model of inflammatory pain, global deletion of ERa or ERB eliminated underlying sex differences seen in mechanical sensitivity⁸⁰. In addition to the sexual dimorphism observed in pain-related behaviors, sex differences were also seen in the progression of structural IVD degeneration induced by the western diet. Male mice fed the western diet showed accelerated IVD degeneration compared to chow controls; a result not observed in female mice. While the underlying mechanism are currently unknown, previous studies have illustrated that female mice fed a high-fat diet have an increased capacity for adipocyte enlargement, resulting in reduced adipose tissue inflammation, fat deposition in non-adipose tissues, and better insulin sensitivity compared to male mice⁸¹.

These sex-dependent metabolic phenotypes may modulate the systemic effects of obesity and thereby alter the progression of structural IVD degeneration. Furthermore, sex may also impact PPAR δ signaling. Estrogen receptors (ER α , ER β) can inhibit PPAR α and PPAR γ signaling through competitive binding of hormone response elements and reduced availability of coactivators^{82,83}. Although not examined to date, a similar interaction may exist between estrogen receptors and PPAR δ , potentially impacting the results of this study.

Taken together, this study highlights the complex, and context-dependent role of PPARδ within the IVD and joint tissues. Our analysis demonstrates that deletion of PPARδ in cartilaginous tissues plays a protective role in age-associated IVD degeneration yet does not protect against obesity-associated IVD degeneration or osteoarthritis. These findings underscore potential differences in the pathogenesis of joint degeneration and associated pain based on the underlying cause. As many promising basic science discoveries fail to enter clinical use⁸⁴, these differences should be further explored in the context of IVD degeneration and applied to clinical trials and basic research.

4.7 Supplementary Figures



Figure 4.10, Supplementary Figure 1: Schematic overview of PPARδ diet study design.



Figure 4.11, Supplementary Figure 2: Effect of PPARδ activation on metabolic gene expression in IVD explant cultures.

SYBR-based real-time qPCR analysis of gene expression in IVD explants incubated for (A) 4 days or (B) 7 days with increasing concentrations of the PPAR δ agonist GW501516. Expression of *Abca1*, *Cpt1a*, *Gsta4*, and *Insig1* were not altered by PPAR δ agonist treatment at either time point. After 7 days of treatment, a significant increase in the expression of *Txnip* was induced by 0.01 µM agonist treatment. Values are represented as the mean \pm SEM. (n= 3-5 independent trials). **P*<0.05 compared to DMSO control by one-way ANOVA with Tukey's test.



Figure 4.12, Supplementary Figure 3: Effect of PPARδ agonism on extracellular matrix in IVD explant cultures.

(A) Representative sections of IVD explants treated for 4 days with DMSO or 1 μ M PPAR δ agonist GW501516 stained with Safranin-O/Fast Green. Explants treated with GW501516 show less intense safranin-o staining within the NP, suggesting proteoglycan loss. (B) The amount of sulfated glycosaminoglycan (s-GAG) released into the media and remaining within IVD explants was quantified using the dimethylmethylene blue assay. Data is presented as mean \pm 95% CI (n= 3 independent trials, P < 0.05 versus DMSO). SYBR-based qPCR analysis of (C) matrix remodeling genes and (D) extracellular matrix genes in IVD explants after 4 days incubation with GW501516 or vehicle control (DMSO). No significant difference in gene expression was detected for any of the genes investigated. Expression was normalized to the expression of the house-keeping gene *Rps29* and expressed relative to DMSO control. Data is presented as mean \pm 95% CI (n= 5-6 independent trials, P < 0.05 versus DMSO).





To determine the anti-nociceptive response of tramadol, the latency of the tail flick response was assessed prior to (baseline) and after IP injection of tramadol (50 mg/kg). (A) The tail flick response was elicited by applying radiant heat to the tip of the tail and the withdrawal time recorded. A cutoff time of 12 seconds was used to avoid tissue injury. (B) The response to thermal stimulation was plotted as the percentage of maximal possible effect (%MPE) which is calculated as $[(T_1 - T_0)/(T_2-T_0)] \times 100$. T₀ and T₁ were the tail flick latency times before and after administration of tramadol and T₂ is the cutoff time. No significant differences were seen between any condition. n=9-13 animals per sex, genotype, and diet. Data are plotted mean $\pm 95\%$ CI.



Figure 4.14, Supplementary Figure 5: Tramadol induced a slight sedative effect in male mice.

Spontaneous locomotion was measured over 10 min to assess the sedative effects of tramadol. (A) In male mice, tramadol (50 mg/kg I.P.) had a slight sedative effect leading to reduced spontaneous movement in both diet conditions, a change detected to a greater extent in chow-fed mice compared to mice fed the western diet. (B) In female mice, average spontaneous locomotion increased with the addition of tramadol. Graphs on the left show the percent of baseline, which represents the distance travelled following tramadol treatment normalized to the pre-treatment (baseline) values for each animal. Graphs on the right show the distance travelled for mice in each group in the absence and following administration of tramadol. n=9-13 animals per sex, genotype, and diet. Data are plotted mean \pm 95% CI. ***P*<0.01 between diets in each treatment group; †indicates significant (*P*<0.05) tramadol-induced differences in distance travelled for mice fed the chow (black) or western (red) diet by 2-way ANOVA with Tukey's test.

	Primer Sequence $5' \rightarrow 3'$
Pdk4 Fwd	CCGCTTAGTGAACACTCCTTC
Pdk4 Rev	TGACCAGCGTGTCTACAAACT
Angptl4 Fwd	CATCCTGGGACGAGATGAACT
Anptl4 Rev	TGACAAGCGTTACCACAGGC
Cidea Fwd	TGGAAAAGGGACAGAAATGG
Cidea Rev	TCTCGTACATCGTGGCTTTG
Ppard Fwd	TCCATCGTCAACAAAGACGGG
Ppard Rev	ACTTGGGCTCAATGATGTCAC
Collal Fwd	CTGGCGGTTCAGGTCCAAT
Collal Rev	TCCAGGCAATCCAGGAGC
Col2a1 Fwd	GCACATCTGGTTTGGAGAGACC
Col2a1 Rev	TAGCGGTGTTGGGAGCCA
Acan Fwd	CTGGGATCTACCGCTGTGAAG
Acan Rev	GTGTGGAAATAGCTCTGTAGTGGAA
Mmp3 Fwd	TTGTCCCGTTTCCATCTCTCTC
Mmp3 Rev	TTGGTGATGTCTCAGGTTCCAG
Mmp13 Fwd	CTTCTTCTTGTTGAGCTGGAACTC
Mmp13 Rev	CTCTGTGGACCTCACTGTAGACT
Adamts4 Fwd	GAGGAGGAGATCGTGTTTCCAG
Adamts4 Rev	CAAACCCTCTACCTGCACCC
Il-1b Fwd	CCCTGCAGCTGGAGAGTGTGGA
Il-1b Rev	TGTGCTCTGCTTGTGAGGTGCTG
Il6 Fwd	TCTCTGCAAGAGACTTCCATCCAGT
Il6 Rev	AGTAGGGAAGGCCGTGGTTGTCA
Bdnf Fwd	TCATACTTCGGTTGCATGAAGG
Bdnf Rev	GACCTCTCGAACCTGCCC
Adamts5 Fwd	GGAGCGAGGCCATTTACAAC
Adamts5 Rev	GCGTAGACAAGGTAGCCCACTTT
Abcal Fwd	GCTACCCACCCTACGAACAA
Abcal Rev	GGAGTTGGATAACGGAAGCA
Cpt1a Fwd	TCAATCGGACCCTAGACACC
Cpt1a Rev	TGGTAGGAGAGCAGCACCTT
Gsta4 Fwd	GCTGGAGTGGAGTTTGAGGA
Gsta4 Rev	TGTGTCAGCATCATCCCATC
Insig1 Fwd	GACGAGGTGATAGCCACCAT
Insig1 Rev	TGGCCCATTCTCTTGAAC
Txnip Fwd	GGTCTCAGCAGTGCAAACAG
Txnip Rev	AGCTCGAAGCCGAACTTGTA

 Table 4.2, Supplementary Table 1: Realtime qPCR primer sequences

	Bivariate (r)		Multivariate (β, r ²⁾		
Parameter	IVD	Knee OA	IVD	Knee OA (β)	Whole
	degeneration		degeneration (β)		model
Behavioral					
Von Frey	0.08	0.25	-	-	0.07
Grip Force	-0.16	-0.56***	-0.64	-0.45***	0.32***
Acetone	0.06	0.01	-	-	0.002
Rotarod	0.10	0.20	-	-	0.04
Open field					
Total Distance	-0.13	-0.41**	-403.8	-44.0**	0.19*
Rearing	0.12	0.33*	-	-	0.13
Velocity	0.09	-0.49***	0.06	-0.08***	0.24**

Table 4.3, Supplementary Table 2. Association between behavioral indicators of pain and histological joint damage.

4.8 References

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Chapter 5

5 General Discussion

5.1 Overview

The overall objective of my thesis was to assess how mechanical loading and obesity impact intervertebral (IVD) physiology. Despite being established risk factors for the development of both IVD degeneration and related back pain, the biological mechanisms by which these factors contribute to disease initiation and progression remain elusive. To address this, we utilized mouse models to understand the biological impact of both mechanical loading and obesity on the IVD. In order to contextualize our findings within systemic changes to musculoskeletal health, we also investigated the impact of these risk factors on the knee joint, as there are similarities between the IVD degeneration and knee osteoarthritis (OA) in terms of both risk factors and pathological processes¹.

We first investigated the impact of mechanical loading, specifically whole-body vibration (WBV) on IVD health in **Chapter 2**. This study built upon previous research from our lab which demonstrated that whole-body vibration (WBV) induces IVD degeneration and knee osteoarthritis (OA) in as little as 4 weeks in a mouse model, using vibrational parameters that model those used clinically^{2,3}. Previous studies were however limited to the analysis of only young, male mice of a single outbred stock (CD1) and do not accurately recapitulate the diverse human population currently using WBV platforms. To determine whether these findings would be consistent in another mouse model, I exposed inbred C57BL/6 mice to the same vibrational parameters as the previous studies^{2,3} for either 4 or 8 weeks. Analysis of these results revealed that the same vibrational parameters previously shown to induce histopathological IVD degeneration and knee OA in CD1 mice did not affect joint health

in C57BL/6 mice. Furthermore, molecular changes seen at the transcriptional level in the IVD were not observed in the C57BL/6 mice. From this initial study, we concluded that C57BL/6 mice were resistant to WBV-induced joint degeneration, at least at the specific vibrational parameters used.

Next, I investigated the impact of diet-induced obesity on IVD degeneration and associated pain in Chapter 3. Despite the established clinical association between obesity, IVD degeneration and low back pain^{4,5}, there have been no studies to directly investigate the impact of diet-induced obesity on IVD degeneration and pain. To address this, I characterized the longitudinal impact of diet-induced obesity on pain-related behavior and structural IVD degeneration in a mouse model. In this study, male wild-type C57BL/6N mice were randomized into one of three diet groups (chow control, high-fat, or a highfat/high-sugar western diet) and fed the experimental diets for 12, 24, 40 or weeks. At endpoint, all mice were assessed for behavioral indicators of pain as well as histological IVD degeneration. We demonstrated that diet-induced obesity led to pain-related behaviors in as little as 12 weeks, with behavioral differences persisting until the 40-week timepoint. Histopathological analysis showed that diet-induced obesity did not affect IVD health until the 40-week timepoint, where early degenerative changes were observed. At the molecular level, mice fed the western diet showed increased expression of *Il-1b* and *Ptgs2* in the IVD at the 24-week timepoint, while mice fed a high-fat diet showed increased expression of Il-6, Ptgs2, Bdnf, and Adamt5 at the 40-week timepoint. Since behavioral indicators of pain preceded structural and molecular alterations to the IVD, we then investigated other potential mechanisms to explain the pain phenotype observed. We quantified markers of neuroinflammation and systemic inflammation, as they have been previously shown to

impact pain sensitivity^{6,7}. While no significant differences were found in the levels of most markers investigated, mice fed the high-fat and western diet tended to show elevated levels of inflammatory cytokines and neuroinflammation, which could impact the pain phenotype seen.

Lastly, we investigated the role of peroxisome proliferator-activated receptor delta (PPAR δ) in the IVD and its role in age- and diet-associated IVD degeneration in **Chapter** 4. Since previous studies have demonstrate that PPAR δ activation has a deleterious role in other cartilaginous joints⁸, we sought to determine its role in the IVD. Intact IVDs were isolated and treated in organ culture with the synthetic PPARδ agonist GW501516. Gene expression analysis showed upregulation of known PPAR δ target genes (*Pdk4* and Angptl4)⁹, indicating that PPAR δ was functional and responsive in IVD tissues. Similar to articular cartilage⁸, PPAR δ agonism appeared to decrease proteoglycan staining in the NP; however, no differences were seen in the expression of matrix degrading enzymes. PPAR δ expression was then measured in different compartments of the IVD, demonstrating a 2fold greater expression of *Ppard* in the AF compared to the NP. To assess the role of PPAR δ in the IVD, we characterized mice with cartilage-specific deletion of PPAR δ in age-associated IVD degeneration. At 14 months-of-age loss of *Ppard* did not alter the gross histological features of the IVD; however, gene expression analysis showed an upregulation of matrix metalloproteinases (*Mmp3*, *Mmp13*) and *Il-1b* which are typically associated with accelerated degeneration^{10,11}. At 25 months-of-age, male *Ppard KO* mice appeared to be protected from age-associated degeneration observed in WT mice; however, differences in histopathological scores were not significant.

Given that PPAR δ ligands are dysregulated in obesity¹², and the potential protective role of PPAR δ inactivation in age-associated IVD degeneration, we then investigated the effects of PPAR δ inactivation in a model of western diet-induced obesity. After 50 weeks on the western diet, male *Ppard* KO mice showed significantly greater IVD degeneration than chow-fed controls. Moreover, a greater proportion of *Ppard* KO mice on the western diet showed moderate-severe IVD degeneration than WT mice on the western diet, although average histopathological scores were not significantly different between the two groups. These findings suggest that, contrary to our hypothesis, loss of *Ppard* may accelerate obesity-induced IVD degeneration. Of note, neither PPAR δ inactivation nor western dietinduced obesity impacted IVD degeneration in female mice, indicating important sexspecific effects. To determine whether IVD degeneration was associated with pain, we carried out a series of behavioral assays and demonstrated that western diet induces pain related behavior in both male and female mice, with a more consistent response seen in females. Interestingly, despite significant increases in joint degeneration, male *Ppard* KO mice fed the western diet did not show pain-related behaviors when compared to chow-fed control. While these findings need to be further explored, they suggest that PPAR δ may contribute to painful joint degeneration.

5.2 Contributions and significance of findings

Back pain is an extremely prevalent, debilitating condition that has a significant impact at both on the individual and societal level^{13–15}. The aim of this thesis was to answer several key questions surrounding the role of two environmental risk factors associated with IVD degeneration and associated pain using the mouse as a preclinical model. Our studies highlight several consistent findings which are important considerations in future clinical

and pre-clinical research. First, our studies underscore that the relationship between environmental risk factors (e.g. mechanical loading, obesity), IVD degeneration, and back pain is complex. While these risk factors may drive IVD degeneration and pain in certain circumstances, their effects on IVD health is influenced by interactions with nonmodifiable risk factors such as age, genetic background, and sex (i.e. effects of WBV on CD-1 versus C57BL/6 mice, obesity-induced IVD degeneration in male versus female mice). Second, although clinical or histological features of IVD degeneration may be similar at endpoint, disease pathogenesis may be driven by entirely different mechanism based on the degeneration-inducing catalyst (i.e. PPAR δ regulates age but not obesityinduced IVD degeneration). Lastly, our studies highlight the importance of taking an integrated approach (assessing both musculoskeletal and nervous tissues) when investigating the relationship of these risk factors with joint degeneration and pain. Mechanical loading and obesity impact other tissues including articular joints, connective tissue, and the nervous system, which may interact with, and contribute to both structural IVD degeneration and pain (Figure 5.1). Aside from these general themes, each chapter independently contributes important findings illustrating the role of mechanical loading and obesity in the initiation and progression of IVD degeneration and associated pain.

In **Chapter 2**, we present work in which we show that C57BL/6 mice are resistant to WBVinduced joint damage using vibrational parameters that have been shown to induce joint damage in CD1 mice, and model those used clinically. While this data was negative, it highlights the need for more comprehensive evaluations of the direct impact of WBV on joint health in the human population. Vibrational parameters such as duration, frequency, and acceleration, along with the direct load experienced by the joints may determine whether WBV therapy is therapeutic or detrimental.

Chapters 3 & 4 present novel data characterizing the impact of diet-induced obesity on IVD degeneration and pain over time in a mouse model. Despite being one of the largest modifiable risk factors for the development of IVD degeneration and back pain^{4,5}, no studies have directly assessed whether diet-induced obesity accelerates IVD degeneration, back pain, or investigated biological mediators underlying this association. Our studies show that diet-induced obesity leads to behavioral indicators of pain starting at the 12-week timepoint, yet histological indicators of IVD degeneration are not seen until the 40-week timepoint, increasing in severity following 50 weeks. The chronology of these findings raises the possibility that IVD degeneration may not be the primary causative factor in some cases of obesity-associated LBP. While speculative, it could be postulated that systemic inflammation caused by obesity directly alters the neural excitability, leading to a pain response in absence of overt joint tissue damage. This systemic inflammation, coupled with inactivity because of pain, may accelerate the IVD degeneration in this model. Aside from obesity-related inflammation, dyslipidemia, hyperglycemia, and hypertension may also impact musculoskeletal and nervous tissues and contribute to the findings observed (Figure 5.1). While the etiology remains elusive, our findings demonstrate that diet-induced obesity contributes to both structural IVD degeneration and pain in a mouse model. These findings lay the groundwork for future studies investigating potential biological mechanisms underlying these associations.



Figure 5.1: Schematic illustrating the proposed contributions of mechanical loading and obesity to IVD degeneration and associated pain.

This thesis highlights the complex interaction between environmental risk factors (i.e. mechanical loading and obesity) and non-modifiable factors (i.e. age, genetic background, sex) on IVD degeneration and associated pain. In addition to affecting the IVD directly, these factors can also impact other tissues (i.e. other joints, muscles, bone, nervous tissues) which may contribute to both structural IVD degeneration and associated pain. Solid lines indicate direct assessments made in this thesis, and dashed lines indicate potential contributions to IVD degeneration and pain not directly investigated in this thesis.

In **Chapter 4**, we investigate the nuclear receptor PPAR δ as a potential mediator of ageand obesity-associated IVD degeneration. Our findings demonstrated that similar its role in articular cartilage, deletion of PPAR δ may protect against age-associated IVD degeneration¹⁶. However, deletion of PPARδ did not protect from obesity-induced IVD degeneration or knee OA. In fact, on the western diet a higher proportion of *Ppard* KO mice showed moderate/severe IVD degeneration than did WT mice. These findings indicate that PPAR δ plays a context dependent role in the IVD (Figure 5.2) and highlight that the etiology of IVD degeneration is likely multifactorial and providing evidence that disc degeneration should be treated as a group of diseases with separate pathogenesis but a common radiological/histological endpoint. In addition to evaluating the impact of PPAR δ on the IVD health, this study also illustrates the importance of assessing the correlation between structural IVD degeneration and pain, and sex-related differences. Obesity induced by the western diet accelerated IVD degeneration in male mice but did not significantly impact IVD health in female mice. In contrast, female mice showed more consistent pain-related behaviors on the western diet than male mice, despite having no significant IVD degeneration.

5.3 Limitations of research

As with other MSK tissues, human IVD tissue is difficult to obtain, especially healthy tissues or those corresponding to the early stages of disease. As such, animal models are often used to understand the pathophysiology of IVD degeneration and associated back pain¹⁷. Rodents are the most frequently used small animal model based on their ease of use, low cost, and the ability to manipulate their genome¹⁷. Despite their widespread use, some limitations associated with the use of mouse models include the persistence of notochordal



Figure 5.2: Schematic overview of proposed context-dependent role of PPAR δ in the IVD.

Our findings highlight the complexities of nuclear receptor signaling in the IVD and other cartilaginous tissues. While PPAR δ deletion was protective in a model of age-associated IVD degeneration, it did not appear to play a protective role in a model of obesity-related IVD degeneration. Furthermore, a higher proportion of *Ppard* KO mice showed more severe IVD degeneration on the western diet than WT mice, suggesting that in the context of obesity loss of PPAR δ may accelerate IVD degeneration.

cells in the NP into adulthood, and altered mechanical loading as reviewed recently¹⁸. Although there is an obvious size difference between human and mouse IVDs, and the limitations mentioned above need to be acknowledged, both species have similar geometry, biomechanics and IVD extracellular composition ^{19–21}.

In Chapter 2, we show that C57BL/6 mice are resistant to WBV-induced joint degeneration, in contrast to previous studies by our group demonstrating the deleterious effects of WBV in CD-1 mice^{3,22}. However, these experiments were performed with only a single protocol of WBV and little is known regarding what parameters of WBV (i.e. frequency, amplitude, duration of exposure) may be beneficial or detrimental to joint health. Previous studies have reported that vibration magnitude is attenuated differently in different stains of mice²³, and that the resonant frequency range also differs between strains of mice²³, raising the question about what loads are being experienced by the joint tissues themselves. While mechanical loading is a key regulator of IVD cell homeostasis, cells can only respond to local mechanical forces, and can have differential responses dependent on both the cell type and loading parameters applied^{24–27}. If the local mechanical load experienced by the joint tissues differed between CD1 and C57BL/6 mice, this could explain the differential response. Additional factors including baseline bone properties or differences in posture have also been shown to impact WBV transmission through the body and could be further explored to better understand the differences observed between the strains of mice^{28,29}.

In **Chapter 3**, we establish the effect of diet-induced obesity on IVD degeneration and pain. Designed as a proof-of-concept, this study had several limitations. First, only male mice were used. Studies in humans have shown that females have a higher prevalence of

spinal disorders and chronic pain conditions than males^{30,31}. Second, this study was limited to only a single formulation of the high-fat and western diets, limiting the generalizability of our results. While not investigated in the context of the IVD, recent studies examining the effects of diet-induced obesity on knee OA show that dietary composition is an important determinant of OA progression in obesity. Diets high in saturated fatty acids or ω -6 polyunsaturated fatty acids (PUFAs) induce more severe metabolic dysregulation and OA progression than diets enriched with ω -3 PUFAs³². Aside from fat, diets high in sucrose also accelerate OA progression independent of weight gain³³. A third limitation is the use of behavioral assays to assess pain. While many of the behaviors examined are altered in experimental pain models, most can be impacted by additional biological and psychosocial factors^{34,35}. In systemic conditions such as obesity, the impact of co-morbidities on behavioral outcomes should be considered. For example, grip force during axial stretch and tail suspension assays have been shown to measure of discogenic back pain³⁶, but are also used to assess muscle strength and the effects of anti-depressants, respectively^{37,38}. Moreover, other joints should also be assessed, as obesity has been shown to globally impact joint health which may contribute to the pain-related behavior observed³⁹. Finally, it is becoming increasingly accepted that IVD degeneration is not just a disease of the IVD, but also the adjacent structures including the muscle, ligaments, vertebrae and facet joints¹. Understanding the direct effects of obesity on these structures, and their relation to IVD degeneration and pain, is an important consideration for future studies.

In **Chapter 4**, we established the effect of PPAR δ deletion in the IVD using both age and obesity-induced models of degeneration. One limitation of this study was the Cre-driver used. Using fluorescent reporter mice, our group has shown that *Col2-Cre* targets the NP,

inner 2/3 of the AF, and the CEP. However, in addition to effecting recombination in the IVD, the *Col2-Cre* is also active in cartilaginous tissues of all joints as well as synovial fibroblasts and some osteoblasts^{40,41}. Given that many of the behavioral indicators of pain would be impacted by degeneration associated changes in other joints⁴², use of the *Col2-Cre* impairs our ability to draw conclusions about the relative contribution of IVD changes and the precise role of PPAR δ to the pain-related phenotype observed. A second limitation common to all studies is the ability to assess tissue-specific changes within the IVD. In all three studies, gene expression analysis was conducted on intact IVDs. Given that the IVD is a heterogenous structure, composed of distinct tissues with diverse cellular phenotypes in each tissue⁴³, the tissue-specific effects of each intervention were evaluated.

In **Chapter 4** we used an *ex vivo* IVD organ culture system to evaluate the effect of PPARδ activation. Organ culture has been suggested as a physiologically relevant model in which cells remain in their endogenous microenvironment, while *in vitro* conditions are modulated¹⁷. One limitation of the organ culture system relates to the heterogeneity of the IVD⁴³ and the inability to differentiate the role of PPARδ activation within each cell type (NP, AF, and CEP). Although monolayer culture would allow for cell type specific experimentation, expansion of primary NP and AF cells in monolayer culture has been shown to alter cellular processes and phenotype¹⁷.

5.4 Future Directions

The findings of this thesis can be further expanded on in several ways, some of which address the limitations detailed above. To better understand the effects of vibrational loading on the IVD it would be important to quantify the transmission of vibration within joint tissues *in vivo* during WBV. One potential method to quantify site-specific vibrational

amplitude in live mice is through non-invasive x-ray imaging of mice carrying zirconium dioxide fiducial marker beads embedded around the joint tissues (i.e. spine and knee joint) during WBV. Similar protocols have been previously used to assess the effects of cyclic loading on strain in the porcine anterior cruciate ligament⁴⁴. This study would allow us to assess the effects of posture, soft tissue dampening and resonance effects on the transmission of vibration to specific anatomical locations (*i.e.* spine, knee). Site-specific findings could then be correlated to the biological effects of WBV characterized in each respective tissue. Future work can also follow up on putative biological pathways underlying this association using transcriptomics and proteomic approaches.

To further characterize the role of diet-induced obesity in IVD degeneration and associated pain, future studies should take a more integrated approach given the systemic effects of obesity on the body. In addition to investigating structural joint degeneration within the IVD, the effects of obesity should be directly assessed in surrounding spinal tissues including muscles, ligaments, and the facet joints to determine if their dysfunction contributes to IVD degeneration and/or axial pain. In the context of pain, the impact of obesity should be assessed longitudinally through a repeated measures design, where animals undergo behavioral testing prior to randomization and at regular intervals throughout the time course of the study (e.g. every 4 weeks). These findings would allow us to determine the point at which pain-related behaviors initiated and whether they persisted in each animal. These findings could then be correlated with non-invasive, *in vivo* assessments of joint degeneration such as T2-weighted MRI, which has been shown to detect degenerative changes in both the IVD and articular cartilage of the knee in rodent models^{45,46}. Pathological changes within the peripheral and central nervous system should

be more thoroughly examined to determine potential mechanisms underlying the pain response associated with diet-induced obesity. Examples include quantification of markers of peripheral nerve injury (ATF3) in the dorsal root ganglia (DRG) and the production of immune and glial mediators (e.g. TNF α , IL-1 β , BDNF) in the spinal cord^{47,48}.

To more robustly characterize the cellular pathways responsible for the association between obesity and IVD degeneration, a systems biology approach using multiple -omics technologies (i.e. epigenomics, transcriptomics, proteomics, metabolomics) could be used to independently assess isolated NP and AF tissues in this model. To specifically address the heterogeneity of cell types within the IVD, single-cell RNA sequencing could also be used. Finally, future studies could also be designed to investigate potential biological pathways or dietary constituents that are associated with accelerated IVD degeneration and pain.

Characterization of the role of PPAR δ in age- and obesity-associated IVD degeneration was limited by the transgenic model used. To address the off-target effects associated with use of the *Col2-Cre*, future studies could be conducted using the *Noto-Cre* mouse strain, which our group has been shown to be specific for notochord-derived cells of the NP⁴⁹. To address tissue specific-differences, AF-specific Cre-drivers such as *Gdf5-Cre* or *Scx-Cre* may also be used, although these are also associated with off-target effects in other MSK tissues^{50,51}. In addition to using a systems biology approach (as detailed above) to assess the effects of PPAR δ agonism in the NP and AF, the interplay between different nuclear receptors should also be investigated. There is substantial overlap between nuclear receptor signaling⁵², and given the similarity between the ligands and DNA binding domains between the PPAR subtypes⁵³, the consequence of deleting PPAR δ on PPAR α and PPAR γ
signaling should be investigated. To do so, chromatin immunoprecipitation with subsequent DNA sequencing (ChIP seq) could interrogate alternate binding of PPAR α and PPAR γ when PPAR δ is no longer present.

Taken together, the results of this thesis highlight the complex nature of mechanical loading and obesity on IVD health. Furthermore, these studies highlight the importance of assessing for pain in conjunction with structural IVD degeneration as they are not always mutually exclusive. Importantly, for effective clinical translation, potential therapeutic targets should be evaluated in multiple different preclinical models of IVD degeneration that consider factors such as genetics, sex, and age.

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Appendices

APPENDIX A: PERMISSION TO REUSE MATERIAL

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	C578L/6 mice are resistant to joint degeneration induced by whole-body vibration					
Osteoartivia	Author: G.J. Kerr, M.R. McCann, J.K. Branch, A. Ratneswaran, M.A. Pest, D.W. Holdsworth, F. Beier, S.J. Dixon, C.A. Séguin					
(Publication: Osteoarthritis and Cartilage					
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APPENDIX B: ANIMAL PROTOCOL NOTICE OF APPROVAL



AUP Number: 2017-154 PI Name: Seguin, Cheryle AUP Title: Mouse Models to Characterize Intervertebral Disc Development and Disc Disease Approval Date: 11/01/2017

Official Notice of Animal Care Committee (ACC) Approval:

Your new Animal Use Protocol (AUP) 2017-154:1: entitled "Mouse Models to Characterize Intervertebral Disc Development and Disc Disease " has been APPROVED by the Animal Care Committee of the University Council on Animal Care. This approval, although valid for up to four years, is subject to annual Protocol Renewal.

Prior to commencing animal work, please review your AUP with your research team to ensure full understanding by everyone listed within this AUP.

As per your declaration within this approved AUP, you are obligated to ensure that:

1) Animals used in this research project will be cared for in alignment with: a) Western's Senate MAPPs 7.12, 7.10, and 7.15

http://www.uwo.ca/univsec/policies_procedures/research.html

b) University Council on Animal Care Policies and related Animal Care Committee procedures

http://uwo.ca/research/services/animalethics/animal care and use policies.htm

2) As per UCAC's Animal Use Protocols Policy,

a) this AUP accurately represents intended animal use;

b) external approvals associated with this AUP, including permits and scientific/departmental peer approvals, are complete and accurate;

- c) any divergence from this AUP will not be undertaken until the related Protocol Modification is approved by the ACC; and
- d) AUP form submissions Annual Protocol Renewals and Full AUP Renewals will be submitted and attended to within timeframes outlined by the ACC. e) http://uwo.ca/research/services/animalethics/animal_use_protocols.html

3) As per MAPP 7.10 all individuals listed within this AUP as having any hands-on animal contact will

- a) be made familiar with and have direct access to this AUP; b) complete all required CCAC mandatory training (training@uwo.ca); and
- c) be overseen by me to ensure appropriate care and use of animals.

4) As per MAPP 7.15,

a) Practice will align with approved AUP elements; b) Unrestricted access to all animal areas will be given to ACVS Veterinarians and ACC Leaders;

c) UCAC policies and related ACC procedures will be followed, including but not limited to:

- i) Research Animal Procurement
- ii) Animal Care and Use Records iii) Sick Animal Response
- iv) Continuing Care Visits

5) As per institutional OH&S policies, all individuals listed within this AUP who will be using or potentially exposed to

hazardous materials will have completed in advance the appropriate institutional OH&S training, facility-level training, and reviewed related (M)SDS Sheets, http://www.uwo.ca/hr/learning/required/index.html

Submitted by: Copeman, Laura on behalf of the Animal Care Committee University Council on Animal Care

CURRICULUM VITAE

Geoffrey Kerr

Post-secondary Education and Degrees

2015-Present	PhD in Physiology and Pharmacology (Collaborative Program in Musculoskeletal Health) The University of Western Ontario (London, Ontario) Schulich School of Medicine & Dentistry, Bone & Joint Institute Supervisor: Dr. Cheryle Séguin, PhD
2011-2015	Bachelor of Medical Science (BMSc. 2015) Honors Double Major in Physiology & Pharmacology The University of Western Ontario (London, Ontario) Schulich School of Medicine & Dentistry
Publications	
2017	Geoffrey J. Kerr , Matthew R. McCann, Jennifer K. Branch, Michael A. Pest, Anusha Ratneswaran, David W. Holdsworth, Frank Beier, S. Jeffrey Dixon, and Cheryle A. Séguin. C57BL/6 mice are resistant to whole-body vibration-induced joint degeneration. Osteoarthritis and Cartilage. 2017 Mar; 25(3):421-425.
2017	Geoffrey J. Kerr *, Matthew A. Veras* (co-first author), Min Kyu Mark Kim, Cheryle A. Séguin. Decoding the intervertebral disc: unravelling the complexities and cell phenotypes and pathways associated with degeneration and mechanotransduction. Semin Cell Dev Biol. 2017 Feb; 62:94-103.
Fellowships a	nd Awards
2019 2019	Mogenson Trust Award (Hari and Gudrun Sharma Award) Canadian Institute of Health Research National Poster Competition, Silver
2019 2019 2018	Medal Canadian Institute of Health Research Travel Award SixSense Case Competition – First Place Canadian Bone and Joint Young Investigators Forum – Top Scientific Abstract

2017	Canadian Connective Tissue Conference – Oral Presentation Award
2017	Natural Sciences and Engineering Research Council of Canada – Post-
	graduate scholarship
2017	Transdisciplinary Bone & Joint Training Award
2017	Ontario Graduate Scholarship (Declined)
2016	Physiology and Pharmacology Research Day – Poster Award
2016	Canadian Connective Tissue Conference – Poster Award
2015	Transdisciplinary Bone & Joint Training Award

Teaching

2019-2020	PP 3000 – Physiology & Pharmacology Laboratory, UWO
2016-2019	Physiology 2130 – Introduction to Human Physiology, UWO
2015-2016	Physiology 3130Y – Physiology Laboratory, UWO

Supervision

2019-2020	1-4 th Year Honors Student
2018-2019	1-4 th Year Honors Student
2017-2018	1-4 th Year Honors Student
2016-2017	1-4 th Year Honors Student

Presentations at Scientific Meetings

Oral Presentations

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Poster Presentations

2019	Orthopedic Research Society – PSRS Spine Research Symposium
	"Investigating the relationship between diet-induced obesity, intervertebral
	disc degeneration and back pain." (Skytop, Pennsylvania, USA)

2019	Canadian Institute of Health Research National Poster Presentation
	"Investigating the relationship between diet-induced obesity, intervertebral
	disc degeneration and back pain." (Winnipeg, Manitoba)
2019	London Health Research Day
	"Investigating the relationship between diet-induced obesity, intervertebral
	disc degeneration and back pain." (London, Ontario)
2019	Canadian Pain Society Scientific meeting
	"Investigating the relationship between diet-induced obesity, intervertebral
	disc degeneration and back pain." (Toronto, Ontario)
2019	Gordon Research Conference in Cartilage Biology and Pathology
	"Investigating the relationship between diet-induced obesity, intervertebral
	disc degeneration and back pain." (Galveston, Texas, USA)
2019	Gordon Research Seminars in Cartilage Biology and Pathology
	"Investigating the relationship between diet-induced obesity, intervertebral
	disc degeneration and back pain." (Galveston, Texas, USA)
2018	Physiology & Pharmacology Research Day
	"Investigating the role of peroxisome proliferator-activated receptor delta
	in intervertebral disc health and degeneration." (London, Ontario)
2018	Canadian Bone and Joint Conference
	"Investigating the role of peroxisome proliferator-activated receptor delta
	in intervertebral disc health and degeneration." (London, Ontario)
2017	Physiology & Pharmacology Research Day
	"Male mice are more susceptible to whole-body vibration induced joint
	damage than age-matched female mice." (London, Ontario)
2017	Gordon Research Conference Cartilage Biology and Pathology
	"Male mice are more susceptible to whole-body vibration induced joint
	damage than age-matched female mice." (Lucca, Italy)
2017	Gordon Research Seminars Cartilage Biology and Pathology
	"Male mice are more susceptible to whole-body vibration induced joint
	damage than age-matched female mice." (Lucca, Italy)
2016	Physiology & Pharmacology Research Day
	"Genetic strain-specific differences in the effect of whole-body vibration
	on joint health." (London, Ontario)
2016	Canadian Connective Tissue Conference
	"Genetic strain-specific differences in the effect of whole-body vibration
	on joint health." (Hamilton, Ontario)
2016	Canadian Bone and Joint Conference
	"Genetic strain-specific differences in the effect of whole-body vibration
	on joint health." (London, Ontario)
2016	London health Research Day

"Genetic strain-specific differences in the effect of whole-body vibration on joint health." (London, Ontario)

2015 **Physiology & Pharmacology Research day** "The effect of repeated exposure to whole-body vibration on intervertebral disc and joint health." (London, Ontario)