

EXPERIMENTAL AND COMPUTATIONAL STUDIES OF ARTICULAR
CARTILAGE AT HIGH RESOLUTIONS

by

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*Dedicated to
To my mother, and my late father*

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Syeda Saira Batool

ABSTRACT

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Articular cartilage is a thin layer of connective tissue covering the ends of bones in diarthrodial joints to minimize friction and distribute loads. The degradation of articular cartilage is the hallmark of osteoarthritis (OA), a complex joint disease that ranks as the number one cause of disability in the adult population. Conceptually, cartilage can be divided into three sub-tissue zones (the superficial zone SZ, the transitional zone TZ, and the radial zone RZ) across its thin thickness (depth), where each zone has a set of unique depth-dependent properties. These uneven structural and compositional variations across its thin thickness imply that any diagnostic technique should ideally have high resolution in imaging; otherwise, volume averaging within an image pixel could obscure any possibility of early disease detection. In this dissertation, we used quantitative microscopic magnetic resonance imaging (μ MRI) combined with polarized light microscopy (PLM) to study the articular cartilage in rabbits and its potential use as an animal model for OA. The utilization of animals plays a vital role in OA research, which enables the examination of the degradation of OA before any procedures are used on humans in the clinic.

This dissertation has six chapters. Chapters 1 and 2 contain the introduction, the background, and the literature review. Chapter 3 is an experimental MRI and PLM study of rabbit cartilage at sub-10 μm resolution, which established quantitatively the baseline characteristics of healthy rabbit cartilage between μMRI and PLM. The results, which are useful for the future investigation of OA using the rabbit model, have been published in the *Journal of Orthopaedic Research* (Batool & Xia, 2020). Chapter 4 is an experimental study to characterize the structural variations at different anatomical locations of femoral cartilage in young rabbits (12-14 weeks old) using μMRI and PLM. Knowledge of location-specific structural differences in the collagen network over the joint surface can improve the understanding of local mechanobiology and provide insights into tissue engineering, and degradation repairs This study has been published in the journal *Cartilage* (Batool & Xia, 2022). Chapter 5 presents a computational study that used a mathematical model to describe the role of collagen fibril mechanics in articular cartilage under 1D axial compression. This study is being finalized for a peer-reviewed journal. The final Chapter 6 summarizes this dissertation.

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LIST OF ABBREVIATIONS

A	Anterior
AC	Articular cartilage
AS	Articular surface
C	Central
CBI	Cartilage Bone interface
CPMG	Carr-Purcell-Meiboom-Gill Sequence
CT	Computed Tomography
GBD	Global Burden of Disease
dGEMRIC	delayed gadolinium-enhanced MRI of cartilage.
2D	Two-Dimensional, 2-Dimensions
3D	Three-Dimensional
ECM	Extracellular matrix
EM	Electromagnetic
FCD	Fixed charge density
FTIRI	Fourier-transform infrared imaging
GAG	Glycosaminoglycan
Gd-DTPA ⁻²	Gadolinium diethylene triamine penta acetic acid
¹ H	Proton
MRI	Magnetic resonance imaging

LIST OF ABBREVIATIONS—Continued

MT	Magnetization transfer
MTR	Magnetization transfer ratio
MSME	Multi-slice multi-echo
μMRI	micro-MRI
NHANES	National Health and Nutrition Examination Survey
NIH	National Institutes of Health
NMR	Nuclear magnetic resonance
OA	Osteoarthritis
OARSI	Osteoarthritis Research Society International
P	Posterior
PG	Proteoglycan
PLM	Polarized light microscopy.
ROI	Region of interest
RZ	Radial zone
SZ	Superficial zone
TE	Echo time
TEc	Echo time of leading contrast segment
TSL	Spin lock period
T1	The spin-lattice (longitudinal) relaxation
T1ρ	The spin-lattice relaxation in the rotating frame

LIST OF ABBREVIATIONS—Continued

T2	The spin-spin (transverse) relaxation
TG	Trochlear groove
TR	Repetition time, Trochlear ridge
TZ	Transitional zone
UHF	Ultra-high-field scanners
σ	Stress
ϵ	Strain

LIST OF PUBLICATIONS

FIRST AUTHORED ARTICLES

1. **Batool, S.**, Mahar, R., Badar, F., Tetmeyer, A., & Xia, Y. (2020). Quantitative μ MRI and PLM Study of Rabbit Humeral and Femoral Head Cartilage at Sub-10 μ m Resolutions. *Journal of Orthopaedic Research*, 38(5), 1052–1062.
2. **Batool, S.**, Hammami, M., Mantebea, H., Badar, F., & Xia, Y. (2022). Location-Specific Study of Young Rabbit Femoral Cartilage by Quantitative μ MRI and Polarized Light Microscopy. *Cartilage*, 13(1), 19476035221085144.
3. **Batool, S.**, Roth, B., & Xia, Y. (2023). Depth-Dependent Strain Calculations for Anisotropic Fibril Structures in Articular Cartilage by Finite Difference Methods, to be submitted to a peer-reviewed journal.

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CO-AUTHORED ARTICLES

The author has also contributed to the following published work.

1. Mahar, R., Batool, S., Badar, F., & Xia, Y. (2018). Quantitative measurement of T2, T1 ρ , and T1 relaxation times in articular cartilage and cartilage-bone interface by SE and UTE imaging at microscopic resolution. *Journal of Magnetic Resonance* (San Diego, Calif.: 1997), 297, 76–85.
<https://doi.org/10.1016/j.jmr.2018.10.008>

2. Mantebea, H., Batool, S., Hammami, M., & Xia, Y. (2021). Structural Morphology of Rabbit Patella and Suprapatella Cartilage by Microscopic MRI and Polarized Light Microscopy. *CARTILAGE*, 13(2_suppl), 356S-366S.
3. Mantebea, H., Batool, S., Singh, A., Hammami, M., Badar, F., & Xia, Y. (2022). Structural differences between the immature and mature articular cartilage of rabbits by microscopic MRI and polarized light microscopy. *Journal of Anatomy*. 240(6):1141-1151. <https://doi.org/10.1111/joa.13620>.

CHAPTER ONE

INTRODUCTION

1.1 Introduction

Articular cartilage (AC) is a thin layer of avascular and aneural tissue on the articulating ends of long bones in joints (Marijnissen & Lafeber, 2003; Maroudas, 1968). The gradual degradation of articular cartilage is a hallmark of osteoarthritis (OA), which is a chronic disease and one of the leading causes of disability that affects millions of adults each year. The loss of articular cartilage weakens the effective distribution of mechanical load over the joint surface, which leads eventually to joint failure and the surgical need for joint replacement. Currently, no disease-modifying drug has been approved for medical treatment, and most treatment guidelines only target symptom management. MRI is a non-invasive and non-destructive imaging tool with excellent soft tissue contrast to study the musculoskeletal system, including articular cartilage. Currently, the early diagnosis of osteoarthritis using clinical MRI is not achievable. There is a strong and unmet need in the clinic to find reliable and sensitive MRI-based imaging biomarkers, which are capable of detecting early degenerative changes for early intervention and tissue repair. At least three features of arthritic disease prevent the successful detection of early osteoarthritis in the clinic by clinical MRI.

- 1) Articular cartilage is thin (~2 mm in large human joints) and exhibits complex depth-dependent variations in both molecular structure and composition throughout its thinness. Conceptually, along with its thin thickness, cartilage can be divided into three histological zones (the superficial zone SZ, the transitional zone TZ, and the

radial zone RZ). Each zone has a characteristic thickness, a set of macromolecular concentrations, a unique orientation of the collagen fibrils, and a specific chondrocyte arrangement. These uneven structural variations across its thin thickness signify that any diagnostic technique, including MRI, requires high resolution imaging; otherwise, volume averaging in each image pixel could obscure any likelihood of early detection.

- 2) The image resolution in current diagnostic MRI is insufficient and lacks sensitivity to detect early changes in the tissue's fine structure and delicate functions, which precede the development of OA as a clinical disease.
- 3) OA is a slow-onset disease that spreads over many years. Because the tissue is avascular and aneural, early degradation remains silent to patients. By the time pain and stiffness are felt, chances for successful interventions in the initial stages may have been missed.

To overcome the first two obstacles, higher spatial resolution is required in the MRI of cartilage. Microscopic MRI (μ MRI) is a high-resolution version of clinical MRI, which can have tens of microns in resolution and hence become potentially a useful tool to quantify microscopic details of cartilage degradation. To address the final obstacle, the use of animal models in OA research allows highly controlled and predicted outcomes. In a typical OA model, degradation can be initiated by a particular event, and the certainties in the timing and type of the disease initiation can enable careful studies of molecular and pathological changes during the subsequent degradation process (Bendele, 2001; Little & Zaki, 2012). With the step-by-step understanding of the degradation process from the early (the most vital period) to the late stage, an intervention at the early stages before the

point-of-no-return could be designed to ultimately save the joints in human. Although canines are likely the most studied OA model (Gregory et al., 2012; Paatsama, 1992; Pond & Nuki, 1973; Poole et al., 2010), the use of canines is being discouraged in recent years due to societal concerns and cost-effectiveness. Among other animal models (Bendele, 2001; Gregory et al., 2012; Lampropoulou-Adamidou et al., 2014; Little & Zaki, 2012; Poole et al., 2010), the use of rabbits is becoming popular because of its close phylogenetical and genetic resemblance to humans, as well as its sufficient thickness (~0.5 mm) for easier creation of local defects (better than small rodents ~0.1 mm), relative inexpensiveness, and less societal sensitivity (Batiste et al., 2004; Borrelli et al., 2009; Esteves et al., 2018).

We used quantitative protocols in μ MRI combined with polarized light microscopy (PLM) to study the articular cartilage in rabbits as its potential use in an OA model. We aimed to test quantitative MRI at sub-10 μ m resolution to elucidate topographical and zonal variations in the structural properties of rabbit articular cartilage. We used PLM as an independent imaging tool to validate the μ MRI results. In addition to the experimental studies, a mathematical modeling project was carried out in this research to highlight the role of pre-stressed collagen fiber in the biomechanical compressive response of articular cartilage and calculating zonal strains in the tissue.

1.2 Organization of the Dissertation

This dissertation is divided into six chapters. The background is provided in Chapter 2 (articular cartilage and its composition, osteoarthritis, animal models, MRI and PLM imaging, biomechanics, and computational modeling). The original results are presented in Chapters 3, 4, and 5, which are based on the three individual projects, two of

which have been accepted by a refereed journal (Chapters 3 and 4), and a third has been submitted for a journal publication (Chapter 5). The final Chapter 6 provides the summary of the dissertation research and guidance for future work.

CHAPTER TWO

BACKGROUND

2.1 Articular cartilage

Articular cartilage (AC) is a thin layer (~0.1–3 mm, depending on the joint and animal of choice) of connective tissue that covers the articular ends of bones in diarthrodial joints to provide cushioning, protection, and load distribution during locomotion (Malda et al., 2013; Xia et al., 2016). Articular cartilage is also known as hyaline cartilage, named after the Latin word *hyalinus*, due to its smooth or glass-like appearance (Jeffrey & Watt, 2003). Mature cartilage contains no nerves or blood vessels and depends on internally self-diffusion and externally repetitive loading to intake nutrients and remove wastes (Mankin, 1994). Structurally, AC contains a small number of living cells (chondrocytes) and an extensive extracellular matrix (ECM) (Fig 2.1). The major constituents of its ECM are water (70-80% by wet weight), collagen (15-22% by wet weight), and proteoglycans (PGs) (4-15% by wet weight) (Fig 2.2) (Buckwalter & Mankin, 1998a; Maroudas, 1968, 1975a, 1975c; Maroudas & Venn, 1977; Buckwalter et al., 2005).

2.1.1 Water

Water is the most abundant molecular component of the ECM in cartilage, which has a depth-dependent concentration ranging from about 85% in the surface tissue to about 60% near the cartilage-bone interface. Judging by its molecular environment, at least two pools of water exist in AC: “bound” (a smaller pool) and “unbound” (a larger pool). When located close to the macromolecules in ECM, mainly PG and collagen, the water molecules have restricted and anisotropic mobility and hence are said to be bound;

further away from macromolecules water molecules are unrestricted and have more isotropic mobility, and hence can be considered unbound or free (Maroudas et al., 1991; Xia et al., 2016). Although water is a fluid, water molecules play an important role in the biomechanical properties of articular cartilage, by providing hydraulic pressure against external loads and forces. In addition, the flow of water through the ECM of cartilage and across the articular surface facilitates the transportation of nutrients and the removal of wastes from the tissue (Buckwalter & Mankin, 1998b).

2.1.2 Proteoglycans

The most abundant macromolecules in adult cartilage are proteoglycan (PG), an aggrecan (Roughley & Mort, 2014; Sophia Fox et al., 2009) that consists mainly of a protein core covalently attached to negatively charged side chains of glycosaminoglycans (GAGs), chondroitin, and keratan sulfates. GAGs are macromolecular polysaccharides with a linear chain structure (Horkay, 2012). The core protein typically has a molecular weight of ~250 000 Da and can have more than 100 GAG molecules laterally bound to it in a branched, bottlebrush-like configuration. Because the GAG molecules are heavily sulfated, they carry negative charges and generate fixed charge density (FCD) through collagen-fibril interaction, which is the main source of cartilage swelling pressure (~0.05–0.35 MPa) and viscoelastic behavior (Maroudas, 1968, 1970; Maroudas & Bannan, 1981). In cartilage, the PG concentration is also known to vary continuously through the tissue depth (being least just below the surface) (Mittelstaedt & Xia, 2015; Xia, Zheng, et al., 2008; Yin et al., 2011). PGs are responsible for the hydration of the matrix, stabilization of collagen networks, and giving the tissue its ability to resist compression due to repulsion between the negative charges.

2.1.3 Collagen

Collagen fibers play a crucial role in the mechanical functionality of articular cartilage, via its unique depth-dependent structure and its interactions with the other ECM molecules, mainly water and proteoglycans (Basser et al., 1998). Collagen is a structural protein and the main solid component of cartilage's ECM, which contains several different types of collagens. The most abundant type of collagen in the ECM of adult cartilage is type II (Buckwalter & Mankin, 1998b), which is a fibrillar protein consisting of tropocollagen forming a triple helix configuration (Eyre, 1991) (Fig 2.1). The tropocollagen molecules are assembled into small prototypic fibrils with an axial repeat length D of ~65–67 nm that is called the D-period (Antipova & Orgel, 2010), which gives collagen its characteristic striated appearance in high-resolution electron microscopy images. Scanning electron microscopy (SEM) images have revealed that multiple collagen fibrils bundle into large collagen fibers. The thickness or diameter of collagen fibers has been reported to vary with the cartilage depth and lies in the range between 25 nm near the surface and 200 nm in deep cartilage (Gottardi et al., 2016).

Collagen type II, together with type IX and XI collagens, form a fibrillar meshwork through the depth of the tissue in a trizonal manner in mature cartilage, which is absent in immature or neonatal cartilage (Julkunen et al., 2009). A unique structural feature of the collagen fibers in mature articular cartilage is their depth-dependent orientation, which can be used to subdivide the noncalcified cartilage into three sub-tissue zones (Fig 2.1). The fibers are parallel to the surface in the superficial zone (SZ) (10–15% of overall thickness), without any predominant orientation in the transitional zone (TZ) (20–40%), and perpendicular to the surface in the radial zone (RZ) (60–70%).

The radial fibers anchor the cartilage to the subchondral bone (Aspden & Hukins, 1981; Benninghoff, 1925; Fox et al., 2009). Below RZ, there is a zone of mineralized calcified cartilage that attaches to the underlying subchondral bone. A mineral front, the *tidemark*, distinguishes between the non-calcified and calcified tissues.

The role of the collagen fibers' orientation has been predicted to be the primary factor modulating the crack morphology in cartilage under indentation loading (Moo et al., 2021). During growth and maturation, structural variations in the collagen network are observed to occur across tissue depth, from a monolayer in neonates to a mixed multizonal layer and finally the standard trizonal structure in mature tissue (Julkunen et al., 2010; Xia et al., 2003). It remains not clear that how these structural variations in the collagen fibril network contribute to the functional response of the tissue. A current understanding is that once the tri-zonal structure is established in mature tissue, the concentrations of macromolecules in ECM are responsible for the tissue's resistance to compressive stress (Guterl et al., 2010; Han et al., 2011), through osmotic swelling pressure and electrostatic repulsion between negatively charged PGs.

The PGs bind to the collagen fibers via their core proteins, with their GAG side chains shown to form bridges between adjacent collagen fibers (Lewis et al., 2010). Without the collagen framework, the PGs could dissociate into smaller molecules. At equilibrium, the ECM in healthy cartilage is known to be in a state of pre-stress required to balance the swelling pressure induced by negatively charged PGs, which is vital for tissue load-bearing strength and homeostasis. Any imbalance between the PG swelling pressure and the collagen fibers restraining stress could alter the normal mechanical function of the tissue and be a manifestation or initiation of degradation.

2.1.4 Chondrocytes

Chondrocytes are the only kind of living cells in AC (Buckwalter & Mankin, 1998b). They are the primary structural, functional, and metabolic units, responsible for the balance of the ECM catabolism and anabolism (Kuettner, 1992). A single cell or a small group of cells are surrounded by a pericellular matrix, which is embedded in the ECM. The cells are sparsely spaced (1%–2% of the total cartilage volume) in adult cartilage and vary in numbers, sizes (approximately $\sim 10\ \mu\text{m}$), and shapes (oval to circular) through the depth of tissue (Choi et al., 2007; Quinn et al., 2013). The depth-dependent alignment of the collagen network in the ECM has been suggested to modulate the cell shape differently in different zones. Chondrocyte metabolism is chiefly controlled through mechanical stimuli generated in the surrounding matrix and can influence the biochemical conditions of the tissue (Mow et al., 1999). During skeletal growth, chondrocytes significantly undergo growth and differentiation and are responsible for synthesizing and remodeling the ECM (mainly collagen type II and proteoglycan). The collagen is synthesized by the chondrocytes during skeletal growth, which virtually ceases in the adult articular cartilage. In adult human cartilage, collagen turnover is significantly slower (several hundred years) than proteoglycan turnover (1–5 years) (Heinemeier et al., 2016; Maroudas, 1975b).

2.2 Depth-dependent characteristics of articular cartilage

AC is a thin layer of complex tissue. It exhibits unique depth-dependent physical and biological properties in terms of the content and structure of its constituent cellular and matrix components. Collagen fibers form unique structural frameworks in a depth-dependent manner as described before (Fig 2.1). In addition to the depth-dependent

structure of the collagen network, the distributions of all molecular components (water, proteoglycan, and collagen) are also not homogenous, and all vary with depth from the articular surface towards the cartilage-bone interface (Mittelstaedt & Xia, 2015; Rieppo et al., 2009; Xia, Zheng, et al., 2008). For mature and healthy cartilage, the SZ has a lower concentration of PGs and a higher concentration of water, when compared to the RZ (Mittelstaedt & Xia, 2015; Xia et al., 2016). In general, a clearer understanding of the depth-dependent variations of the ECM macromolecular components, in conjunction with the structural variations, is crucial in determining cartilage health and the alterations that occur from degradation (OA).

2.3 OA

AC can develop degenerative changes due to aging, excessive usage, and trauma. According to Osteoarthritis Research Society International (OARSI, 2015),

Osteoarthritis is a disorder involving movable joints characterized by cell stress and extracellular matrix degradation initiated by micro-and macro-injury that activates maladaptive repair responses including pro-inflammatory pathways of innate immunity. The disease manifests first as a molecular derangement (abnormal joint tissue metabolism) followed by anatomic, and/or physiologic derangements (characterized by cartilage degradation, bone remodeling, osteophyte formation, joint inflammation, and loss of normal joint function), that can culminate in illness.

The prevalence of the number of people living with OA worldwide has increased significantly; it nearly doubled from 1990 to 2019, according to the Global Burden of Disease study (Institution for Health Metrics and Evaluation, 2023). In the report of the

National Health and Nutrition Examination Survey I (NHANES I), an estimated 12% of US adults ages 25–74 years have clinically diagnosed OA (NHANES, 2020). OA is the most common type of arthritis in terms of prevalent cases and ranked 17th highest of the 369 diseases and injuries included in the Global Burden of Disease study (Vos et al., 2020). OA is also considered a concern of global health and economic burdens in terms of increase healthcare costs, a decline in the workforce, and loss of wages due to disability. The overall economic burden associated with OA in the US is estimated at \$136.8 billion annually (OAA, 2023; United States Bone and Joint Initiative, 2023).

The limited healing potential of articular cartilage imposes challenges to the efficacy of repair strategies. Currently, there are no drugs or treatments that can effectively regenerate or restore the damaged cartilage without surgical intervention. Clinically, it is extremely challenging to slow down the progression of OA over time; a total joint replacement is the common result for millions of OA patients worldwide. Therefore, there is a consistent interest in finding non-invasive imaging biomarkers with sufficient sensitivity to cartilage health and early degradation, to plan timely and effective interventions and to design new cartilage repair strategies. In current clinical practice, being noninvasive and having exceptional soft tissue contrast, MRI has proven to be an exceptional tool in OA management (clinical diagnostics and laboratory research). Since articular cartilage is a thin layer of tissue, however, the current resolution in clinical MRI is insufficient to detect early degenerative changes in articular cartilage.

2.4 Rabbit as an OA model

The use of animal models plays a vital role in OA research, as degradation in animals can be initiated by a particular event and monitored with time. Although some

studies have claimed the ability of the rabbit's cartilage to heal and repair rapidly after a traumatic experience, this model is still popular for intermediate-sized animals in OA research as it provides sufficient size/thickness for easier creation of local defects. Recently the reports on the radiographic signs of OA, associated with age and weight in aging domestic rabbits, make rabbits a reasonable choice to study both traumatically induced and naturally occurring OA (Arzi et al., 2011).

2.4.1 Age

The age or maturity of the animal is an important factor to consider before planning any orthopaedical experiments and studies. In the case of New Zealand white male rabbits, the average timeline for physal closure is ~ 6 months with some exceptions, where physes does not close until 9-10 months (Bland & Ashhurst, 1996; Isaksson et al., 2010). There are vivid age-related differences in tissue response to naturally occurring or trauma-induced OA, e.g., the post-traumatic healing response is reported to be weaker in the aged rabbits (24 months old) than in younger rabbits (4 months). In general, cartilage repair, and response to trauma or surgical interventions were all found to be age-dependent (Buckwalter & Mankin, 1998a; Calandruccio & Gilmer, 1962; Frenkel et al., 2005; Wei et al., 1997). There is speculation that the immature rabbit cartilage has some potential for self-healing, but the underlying mechanism is poorly understood. In addition, significant structural differences in collagen fiber organization and glycosaminoglycan distribution have been reported to present in different locations of the immature rabbit knee cartilage (Batool et al., 2022; Heise & Toledo, 1993; Julkunen et al., 2010), which is possibly related to the natural growth and maturation process. Such age-related differences should be kept in mind when choosing

the animal model for OA studies; and in some cases, skeletally mature animals might be a better choice.

2.4.2 Topographical variations

Cartilage has several topographical variations (i.e., among different locations on a single joint surface) in the biochemical, physical, optical, and mechanical properties (Brama et al., 2000a; Briant et al., 2015; Lee et al., 2016; Risch et al., 2021; Xia, Moody, Alhadlaq, et al., 2002). These topographical variations can cause significant differences in the results of a study whether it is morphological or mechanical, physical or chemical, in vivo or in vitro, and clinical or preclinical.

Topographical variations in healthy cartilage of a joint are dependent on age, gender, differences in the mechanical demands between joints, and sometimes in the different sites over the same joint (Roth & Mow; 1980, Kempson, 1991, Julkunen et al., 2010, Brama et al., 2000a; Batool et al., 2022)). Such variations include but are not limited to tissue thickness, cell shapes, density, collagen fibril organization, fibril orientation, the concentration of macromolecules (collagen, GAG) and water, stress-strain response, and local stiffness. Some joints are more complex in terms of topographical variations than others e.g., the knee is more complex than the hip and shoulder joints. These variations can influence any imaging and data analysis, and optimization of various research protocols. The choice of sampling sites from any single joint surface can therefore significantly influence the outcomes of biochemical or biomechanical studies of articular cartilage. Studying these variations is important in improving the consistency and accuracy of outcomes of biochemical and biomechanical studies.

2.5 Biomechanics

Since cartilage is a load-bearing tissue, biomechanical testing of cartilage is essential in OA diagnostics and research. Usually, the compressive modulus of cartilage can be correlated with the stiffness of the tissue. When cartilage is compressed using unconfined, confined, or indentation experiments, the integrity of the tissue can be determined by its response to compression (Setton et al., 1999).

2.5.1 Young's modulus

Several biomechanical parameters in cartilage can be measured using stress relaxation and creep experiments, e.g., instantaneous compressive modulus, aggregate modulus, Poisson's ratio, etc. The instantaneous compressive modulus is measured during the loading phase of stress relaxation. The aggregate modulus is a measurement of the stiffness of the tissue at the equilibrium when all fluid flow has stopped. Poisson's ratio is the ratio of the change in radial length to the change in axial length of the tissue.

Usually, the modulus of cartilage can be correlated with the stiffness of the tissue. Young's modulus μ , is a ratio of stress and strain, given by

$$\mu = \frac{\sigma}{\epsilon} \quad (2.1)$$

where $\sigma = \frac{F}{A}$ is stress and $\epsilon = \frac{\Delta L}{L}$ is strain, F is the resultant force, A is the contact area of the compressor and tissue, ΔL is the net displacement, and L is the uncompressed tissue length or thickness. The experimental setup to measure the stiffness of tissue typically consists of a rigid frame, load sensor, and compression platen. A rigid frame houses the sample, a load sensor measures force, and a micrometer or computer-controlled actuator

controls the compression plate movement. Under action, the platen compresses the sample in small and successive steps, and the load sensor measures the resultant force. The force and displacement are both recorded, and the elastic modulus can be calculated.

2.5.2 Viscoelasticity

During joint loading (compression), there is an immediate increase in interstitial fluid pressure which leads to a fluid flow out of the ECM. The displacement of the fluids in cartilage is a function of time because the fluid cannot escape from the matrix instantaneously. When the compressive load is removed, interstitial fluid would flow back (Mow et al., 1999). Experimentally, it has been shown that interstitial water pressure contributes to more than 90% of the initial load-carrying capacity at the cartilage surface, thereby reducing the stress acting upon the solid matrix (Ateshian, 2009; Ateshian et al., 1994; Ateshian & Wang, 1995). Any material whose properties have time dependency are said to be viscoelastic material.

2.5.3 Creep and Stress relaxation

Two biomechanical tests are commonly used to estimate the viscoelastic properties of the tissue. Creep is a time-dependent deformation of a viscoelastic material under the application of a constant stress at a constant temperature. A step-like load is applied to the cartilage, and the time dependence of the resultant deformation (strain) is monitored (Fig 2.3a).

Stress relaxation refers to the measurement when a constant displacement (strain) is applied to the cartilage and the time dependence of the induced stress is monitored. As the tissue is compressed, the deformation results in an increased stress and leads to fluid displacement, after which relaxation occurs (Fig 2.3b).

2.6 Depth-dependent compressive stiffness of articular cartilage

Articular cartilage deforms in a depth-dependent manner when compressed. Biomechanical testing in combination with different imaging techniques allows better visualization and understanding of how the cartilage responds under compression (Alhadlaq & Xia, 2005; Wang et al., 2015). In mature healthy cartilage, the RZ exhibits a stronger resistance to compression than the SZ and TZ (Fig 2.4b). Using fluorescence-labeled chondrocyte nuclei as intrinsic markers, Schinagl et al. (1996) measured the intra-tissue deformation within the cartilage matrix and found that the compressive modulus increased significantly with depth from the articular surface to the cartilage-bone interface, ranging from 0.079 MPa in the SZ to 2.10 MPa in the deep RZ. It is known that the close association between proteoglycan aggregates, interwoven collagen fibrils, and interstitial fluid provides compressive resilience to cartilage through its depth. It has also been reported that an increase in the depth-dependent GAG concentration, related to the FCD, may alter the depth-dependent biomechanical behavior. When compressed, the negatively charged sites on aggrecan pushed closer together. This effect increases their mutual repulsive force and adds to the compressive stiffness of the cartilage with depth (Chen et al., 2001; Mattson et al., 2017; Pfeiffer et al., 2008; Wilson et al., 2007).

Although the GAG concentration, which is related to the FCD, has been suggested to modulate the compressive resilience of the cartilage, the role of collagen fibrils should not be ignored. The intrinsic swelling of cartilage is resisted by the collagen network which is in the state of pre-stress (tension) and may play a role in resisting compression. The role of collagen fibers in articular cartilage during loading has been studied extensively but the pre-stressed condition is mostly ignored. Under small

deformations, the fibrils are in tension, while at larger deformations, the forces within the fibrils become compressive (Römgens et al., 2013), eventually leading to buckling (which is a reversible process). At both smaller and larger deformation, therefore, the fibrils in cartilage will likely contribute differently to the compressive stiffness of the tissue.

2.7 Computational models

Although the behavior of articular cartilage under compression and changes in the cartilage structure and composition can be studied by using in vivo and in vitro imaging techniques (MRI, CT, various optical microscopies), these imaging techniques alone are unable to quantify the local tissue stresses, strains, fluid flow produced inside ECM and the independent roles of different cartilage constituents (e.g., collagen, PGs, fluid, and ions) in the loading response due to the complexity of articular cartilage. Therefore, computational models in conjunction with imaging techniques have been employed to determine the material properties of cartilage under different loading conditions.

In the early models of articular cartilage, the tissue was considered isotropic, homogenous, and biphasic (Mow et al., 1980). The biphasic theory models the tissue as a mixture of two phases: a charged porous permeable solid phase (collagen-proteoglycan matrix) and an interstitial fluid phase. Later, a triphasic model (Lai et al., 1991) of articular cartilage was introduced as an extension of the biphasic theory, where negatively charged proteoglycans are modeled to be fixed to the solid matrix, and monovalent ions in the interstitial fluid are modeled as additional fluid phases. According to the triphasic model, the swelling of articular cartilage depends on its fixed charge density distribution, the stiffness of its collagen-proteoglycan matrix, and the ion

concentrations in the interstitial fluid. These models, however, lack the structural inhomogeneity and anisotropic properties of ECM and consider cartilage to be a homogenous, isotropic, and viscoelastic material, which probably does not allow the discrimination between PGs and collagen-dependent mechanical properties, which are required for a more accurate understanding of cartilage mechanics. To address this issue, several fibril-reinforced models (fibril-reinforced poroelastic, poroviscoelastic, hyperelastic) have been introduced to include the effect of fibril anisotropy in traditional biphasic and triphasic models (Korhonen et al., 2003; Pierce et al., 2013; Soulhat et al., 1999; Wilson et al., 2005). These models all treat cartilage as inhomogeneous and anisotropic tissue and include many aspects of the real tissue structure and composition.

2.8 Quantitative MRI in cartilage research

2.8.1 Introduction

MRI is a sensitive and noninvasive imaging tool that can produce detailed anatomical images of biological specimens due to its high sensitivity to the dynamic motion of water molecules. The brain, spinal cord, and a number of connective tissues (muscles, ligaments, tendons, and articular cartilage) can all be seen much more clearly in MRI than in regular x-rays and computed tomography (CT). The technique is currently widely in use in clinical and preclinical settings for disease detection, diagnosis, and treatment monitoring. MRI is also safer than CT, since it does not utilize damaging ionizing radiation.

The working principle of MRI is nuclear magnetic resonance (NMR), which was first discovered in 1930s by I.I. Rabi, who measured the magnetic properties of atomic nuclei. He found that under a strong magnetic field, the spin states of the atomic nucleus

would flip if an electromagnetic (EM) wave at a specific frequency was incident on it.

The resonance occurs when the incident oscillation frequency matches the intrinsic frequency of the nucleus, which depends on the strength of the static magnetic field (B_0), the chemical environment and the magnetic properties of the nucleus involved. This intrinsic precessional frequency is known as the Larmor frequency and given as

$$\omega = \gamma B_0 \quad (2.2)$$

where γ is the gyromagnetic ratio ($\text{rad sec}^{-1}\text{T}^{-1}$). Among the nuclei found in the human body that have a nonzero nuclear spin (e.g., ^1H , ^{13}C , ^{17}O , ^{19}F , ^{23}Na , ^{31}P), hydrogen is by far the most dominant nucleus of interest. For the ^1H nucleus (proton), the gyromagnetic ratio is 42.57 MHz T^{-1} .

2.8.2 NMR to MRI

In 1946, Felix Bloch and Edward Purcell demonstrated the NMR phenomenon in bulk matter, for which they shared the Nobel Prize in Physics in 1952 (Bloch, 1946, 1953; Purcell, 1946). In the 1950s, Erik Odeblad and Gunnar Lindstrom were the first to use NMR to study various biological tissues including articular cartilage (Odeblad and Lindstrom, 1955; Xia and Stilbs, 2016). For the next 20 years from the 1950s to the 1970s NMR was used exclusively in spectroscopy to analyze the structures of organic molecules. The earliest use of NMR for diagnostic purposes was likely in 1971 when R. Damadian showed that the NMR relaxation parameters (T_1 and T_2 relaxation) were significantly different in malignant tumor tissue when compared to normal tissue.

The era of MRI started in 1973 when Paul Lauterbur performed the first imaging experiment using the principle of NMR. Given the linear proportionality between the precessional frequency of the nuclei (e.g., protons in water) and the magnitude of the

external magnetic field (i.e., Eq (2.1)), a position-dependent magnetic field (i.e., a field gradient) can encode the physical space where the specimen resides. By setting up three field gradients in three orthogonal directions, the frequency of the nuclei (e.g., protons) can be spatially encoded in any 3D or 2D space in the form of an image. For this work, the extension of NMR to NMR imaging (or MRI), P. Lauterbur and P. Mansfield shared the Nobel Prize in Medicine in 2003.

2.8.3 Instrumentation

A complete MRI system consists of a magnet, which is used to generate the B_0 field. A radiofrequency (rf) coil can act both as a transmitter coil to deliver the energy to the specimen (a B_1 field) and as a receiver coil to receive the signal generated by the transverse magnetization (the free induction decay, FID). The use of three independent gradient coils can produce the linear field gradients with their magnitudes varying in the x, y, and z directions respectively. The MRI system also needs at least one computer for human-machine interface and a complex array of electronic components.

2.8.4 MRI contrast

MRI signal is sensitive to the motional dynamics of the nuclei in the system. Since the proton (^1H) is the most common and sensitive nucleus in biological systems, most MRI experiments rely on the water molecules in any water-containing specimens (biological or not) for its signal. There are several fundamental molecular mechanisms that are responsible for the unique characteristics of the MRI signals. We will discuss a few of them that are relevant to this dissertation, emphasizing their influences on the MRI signal.

In any practical (bulk) sample, the vast numbers of nuclear spins of protons precessing in a strong magnetic field (B_0) can be viewed/averaged as a macroscopic magnetization M vector, pointing to the $+z$ direction in the thermal equilibrium condition. When an oscillating rf field (B_1 field) is applied in the transverse plane under the resonant condition (i.e., at the Larmor frequency), the macroscopic magnetization will be tipped towards the transverse plane (x-y plane, as in Fig 2.5a). After M is tipped to the transverse plane and the B_1 field is turned off, the nuclear spins will relax back to the $+z$ direction. Since the M vector can be separated into two components, M_z in the longitudinal z-axis and M_{xy} in the transverse x-y plane, the relaxation of the nuclear spins can be examined in transverse and longitudinal directions (Fig 2.5a).

The time rate of change of the longitudinal component of the net magnetization (M_z) is given in the classical mechanical description as

$$\frac{dM_z}{dt} = -\frac{(M_z - M_0)k}{T_1} \quad (2.3)$$

where M_0 is the unperturbed macroscopic magnetization of the spin system, and T_1 is a longitudinal relaxation time, characterizing the spin-lattice interaction (Fig 2.5c).

Similarly, the time rate of change of the transverse component of the net magnetization (M_{xy}) is given as

$$\frac{dM_{xy}}{dt} = -\frac{(M_x i + M_y j)}{T_2} \quad (2.4)$$

where, M_x and M_y are the x and y components of the magnetization and T_2 is a transverse relaxation time, characterizing the spin-spin interaction (Fig 2.5b).

T_1 (the spin-lattice relaxation time) and T_2 (the spin-spin relaxation time) are the two most common types of MRI contrast in water-rich specimens especially in the biological and clinical environment. In addition, there is a spin-lattice relaxation time in the rotating frame, or $T_{1\rho}$, which is the relaxation of the macroscopic magnetization being spin locked in the transverse plane. The difference between their characteristics is closely related to the dominant frequencies of the molecular environment, which can be explained with the aid of nuclear spin dynamics. T_1 is sensitive to the highest frequencies of molecular motions in the Larmor frequency range (tens and hundreds of MHz); $T_{1\rho}$ is sensitive to the intermediate frequencies in the range of several or tens of kHz; and T_2 is sensitive to the lowest frequencies of molecular motions involving static or slowly fluctuating magnetic fields. These relaxation times have been explored extensively by incorporating different relaxation weighting mechanisms in the imaging pulse sequences.

2.8.5 Magnetization prepared pulse sequences in MRI

Practically, an MRI experiment is under the control of a pulse sequence, which is a list of time events on when and how the rf field and various gradient fields are applied and the data is acquired. In any imaging experiment, the influence of the spin relaxations (in particular, T_2 relaxation) exists in the final image, different for different imaging sequences. Consequently, a true nuclear spin density (i.e., a true map of water content in a biological specimen) is difficult to quantify. In our MRI work, we commonly use 2D imaging pulse sequences in the format of magnetization preparation (Xia, 1998). This format of imaging sequences contains two clearly separated time-segments, a leading contrast segment and a subsequent imaging segment. The leading contrast segment uses a variable timing table to impose an image contrast (either T_2 contrast or T_1 contrast);

while the final imaging segment keeps all timings constant – hence the contrast effect of this imaging segment can be normalized in the final image analysis.

To impose a T2 contrast, a spin-echo or CPMG (Carr-Purcell-Meiboom-Gill) sequence can be used in the leading contrast segment. For example, in the spin-echo format, a 90° rf pulse is applied to tip the net magnetization into the transverse plane. At a time $TE_c/2$, a 180° rf pulse flips the dephasing magnetization about the transverse plane, which results in the re-phase of the spins at the time TE_c (where TE_c is the echo time between the 90° rf pulse and the echo center). A final -90° rf pulse tips the refocused transverse magnetization back to the longitudinal direction (z-axis), to prepare it for imaging. The imaging segment typically starts with a slice selection sequence, which has a soft 90° rf pulse together with one gradient pulse. Subsequently, a set of 2D imaging gradients in the orthogonal plane can be applied to sample the signal over the 2D k space (raw data matrix). Two important parameters in MRI are the echo time (TE_i), which is the time duration between the center of the first 90° soft rf pulse and the center of the signal in acquisition (Fig 2.6b), and the repetition time (TR), which is the time duration at which the sequence repeats itself. Using several TE_c , an exponential fit can be applied to the peaks of the signal, which enables the quantitative calculation of the T2 relaxation time, as

$$M_{xy}(TE) = M_{xy}(0) e^{-\left(\frac{TE}{T_2}\right)} \quad (2.5)$$

where $M_{xy}(0)$ is the net signal, and TE is the variable time in the contrast segment.

To impose a T1 contrast, an inversion recovery sequence can be used in the contrast segment, in which the magnetization is inverted along its longitudinal direction

(to the -z direction) by a 180° rf pulse. After a time, delay τ_i (the inversion time), a 90° rf pulse is applied to tip the magnetization into the transverse plane for signal detection. Therefore, the signal intensity depends on the delay time between the 180° and 90° rf pulses (Fig 2.6a). This process can be repeated for various τ_i , which imposes different T1 contrasts. The calculation of T1 can be performed using the following equation,

$$M(\tau_i) = M_0 \left(1 - 2 e^{-\left(\frac{\tau_i}{T1}\right)}\right) \quad (2.6)$$

where M_0 is the initial magnitude of the signal right after the 180° pulse, $M(\tau)$ is the signal depending on τ_i . Several τ_i are needed to calculate the T1 relaxation time.

To impose a T1ρ contrast, a 90° rf pulse tips the magnetization into the transverse plane; immediately after, the magnetization is spin-locked in the transverse plane by a second rf pulse in the direction of the magnetization vector. Under the resonance condition, this second rf pulse (a spin-lock B1 pulse) is the only field in the rotating frame of reference, which acts to keep (i.e., spin-lock) the magnetization in the transverse plane. However, the magnitude of the magnetization was established under B_0 , which is often too big to be maintained by the spin-lock B_1 field, hence the magnetization will shrink its magnitude according the power of the spin-lock field. The time duration and power of the spin-lock B1 pulse is known as the spin lock time (TSL) (Fig 2.6c). When the power of the spin-lock field is zero, T1ρ relaxation is identical to T2 relaxation.

2.8.6 Quantification of relaxation times in MRI of articular cartilage

Due to its excellent soft tissue contrast and its ability to map the molecular environment of tissue, MRI has been increasingly used in the diagnosis of OA and other musculoskeletal disorders. However, the measurement sensitivity to *early* cartilage

degeneration, occurring before morphological changes, is still limited. In the last two decades, several quantitative MRI techniques for cartilage imaging, such as sodium MRI, T1 ρ and T2 relaxations, diffusion MR, and delayed gadolinium-enhanced MRI of cartilage (dGEMRIC) have been used to assess different aspects of the extracellular matrix of cartilage and as potential biomarkers for early degradation (Baum et al., 2013; David-Vaudey et al., 2004; Duarte et al., 2019; Hänninen et al., 2021; Lammentausta et al., 2006; Li et al., 2007; Garrick, 2016; Nissi et al., 2007; Shapiro et al., 2002; Xia, Zheng, et al., 2008). The quantifications of relaxation times are currently the most used protocols in clinical MRI for imaging cartilage and joints.

2.8.7 Quantitative T2 in cartilage imaging

T2 imaging in mature articular cartilage is both depth-dependent and orientation-dependent (Mahar et al., 2018; Xia, Moody, & Alhadlaq, 2002), which reflect the combination of water content, PG content, collagen content and collagen fiber orientation in the ECM. When the normal axis of the cartilage surface is placed at 0° to the external magnetic field B₀, articular cartilage appears bilaminar or tri laminar in high-resolution T2 weighted MRI images. Quantitative analysis of T2 (0°) in healthy cartilage shows that T2 values in the deep zone are significantly shorter than in the SZ and TZ. This is because the deep zone collagen fibers are densely packed so restricting the mobility of water protons. In addition, at orientations close to 0°, there is a strong influence from the nuclear dipolar interaction in deep cartilage, which shortens the measured T2 values (Bydder et al., 2007; Xia, 2000b). A depth-dependent T2 profile at 0° is bell-shaped, which implies a classical three-zone structure in cartilage and has been used to subdivide cartilage thickness into multiple histological zones (Xia et al., 2001a). At the magic angle

(55° to B_0), this laminar appearance vanishes and the cartilage appears homogenous with a maximum signal intensity (Xia, 2000b). This is due to the minimization of the nuclear dipole interaction. The disappearance of the cartilage laminae associated with the 55° angle is known as the magic angle effect in MRI of cartilage (Bydder et al., 2007; Xia, 2000b). Due to the geometric factor in the dipolar interaction, $(3\cos^2\theta - 1)$ (Xia, 1998), the radial zone of cartilage has stronger T2 anisotropy than TZ and SZ.

Quantitative mapping of T2 relaxation time is one of the most used techniques in MRI of cartilage due to its unique relationship with the structural integrity and arrangement of the collagen network as well as the water content in the tissue. T2 has also been reported to increase in spontaneous degeneration of articular cartilage and used as a vital imaging biomarker to study the progression of OA (Baum et al., 2013; David-Vaudey et al., 2004).

2.8.8 Quantitative T1 in cartilage imaging

In contrast to the depth-dependent anisotropy in T2 relaxation, T1 is known to be isotropic and has a small dependency on its tissue depth. In healthy cartilage, T1 is associated with slow motional interactions between the water molecules and highly constrained PGs (Xia, 2013). A common usage of T1 in MRI of cartilage is via the measurement of T1 before and after doping the tissue with a negatively charged contrast agent, gadolinium diethylene triamine penta acetic acid (Gd-DTPA^{-2}). Since Gd-DTPA^{-2} is repelled by the negatively charged GAGs and tends to deposit in the inverse proportion to the local GAG concentration in tissue (Bashir et al., 1996; Owman et al., 2008), different depositions of Gd-DTPA^{-2} can be used to determine the local GAG concentrations. This technique is known as delayed Gadolinium-Enhanced MRI of

cartilage (dGEMRIC). Since the contrast agent contains heavy metal ions, the use of this procedure in a patient requires caution.

2.8.9 Quantitative T1ρ in cartilage imaging

A subtle feature of T1ρ relaxation is its dependency on the strength of the spin-lock field. With a sufficiently high spin lock field (e.g., > 2 kHz), the influence of the dipolar interaction to spin relaxation can be minimized (Wang & Xia, 2013). When the spin-lock field is zero, T1ρ relaxation is identical to T2 relaxation, which shows strong anisotropy in cartilage MRI. With an increase in the spin-lock power, T1ρ becomes less anisotropic, less depth dependent, and has higher values than T2. The measurements of T1ρ probe molecular fluctuations in the kHz range and are sensitive to the interactions between motion-restricted water molecules and their local macromolecular environment. Some studies showed T1ρ imaging could differentiate between normal cartilage and early-stage OA (Li et al., 2007; Mlynárik et al., 1999) and be correlated with proteoglycan loss in vitro (Duvvuri et al., 2002), which provides an attractive alternative to map GAG loss, an early sign of cartilage degeneration.

2.8.10 Spatial resolution in MRI of cartilage

A major limiting factor in clinical MRI of articular cartilage in humans is its imaging pixel resolution, typically 300-500 μm for 3 Tesla whole-body systems, which does not provide sufficient resolution to determine the zonal characteristics of human cartilage (2-3 mm total thickness), nor the detection of early (i.e., small) degenerative changes topographically in joint degeneration. Ample work is in progress to improve the resolution and sensitivity of clinical MRI of OA.

In small animals, articular cartilage is much thinner than the tissue in large primates and humans (Malda et al., 2013), which requires even higher spatial resolution. In this regard, the use of high-resolution microscopic MRI (μ MRI) is preferred to precisely identify the anatomical features of the cartilage from small animals. The rabbit cartilage is $\sim 1/4$ times thinner than human cartilage and it has been shown that a resolution on the order of 10 μm is required to successfully resolve its depth-dependent structural features (Batool et al., 2020). The fact that μ MRI shares identical physics principles and engineering architectures with clinical MRI ensures a translational pathway between the animal model study of OA to clinical detection and intervention (Xia, 2013).

2.9 PLM imaging in cartilage research

The PLM is a gold standard in histology, which is capable of analyzing the anisotropy of a specimen's optical properties such as refraction and absorption (Mehta et al., 2013). In articular cartilage, due to the orientation and organization of the collagen fibers in the tissue, cartilage is birefringent, which exhibits two different indexes of refraction along perpendicular axes.

2.9.1 Working principle of PLM.

Light is an electromagnetic wave that has two orthogonally oscillating components: electric and magnetic components, both orthogonal to the direction of light propagation. Natural light is nonpolarized which can vibrate in all perpendicular planes to the direction of propagation, while polarized light vibrations are restricted in a specific plane (Fig 2.7). Polarized light microscopy makes use of polarizing filters to study the anisotropic properties of materials, which have distinct optical axes and hence whose

optical properties dependent upon the orientation of their axis to the incident light. When light enters a non-equivalent axis in an anisotropic material, it is refracted into two orthogonal components (ordinary and extraordinary waves), which travel at different speeds through the specimen and experience different refractive indices, a phenomenon known as birefringence or (bi-refraction).

Birefringence (b) can be quantified by the numerical difference between these two indices and is given by relation.

$$b = |n_e - n_o| \quad (2.7)$$

where n_e and n_o are the extraordinary (slow) and ordinary (fast) axis indexes of refraction, respectively. After exiting the specimen, the light components become out of phase, and interfere between the two components as they are re-united along the same optical path. This enables the study of anisotropic materials, based on their birefringent properties.

A common PLM system consists of multiple polarizers and wave plates to measure the sample birefringence without the need to rotate the sample (Oldenbourg & Mei, 1995). Microscopes can be outfitted with a mounted camera to capture images of a specimen and be connected to a computer to analyze the images, to calculate several parameter maps such as the collagen fiber orientation and organization, and pericellular matrix around chondrocytes.

2.9.2 Retardation and azimuthal measurements

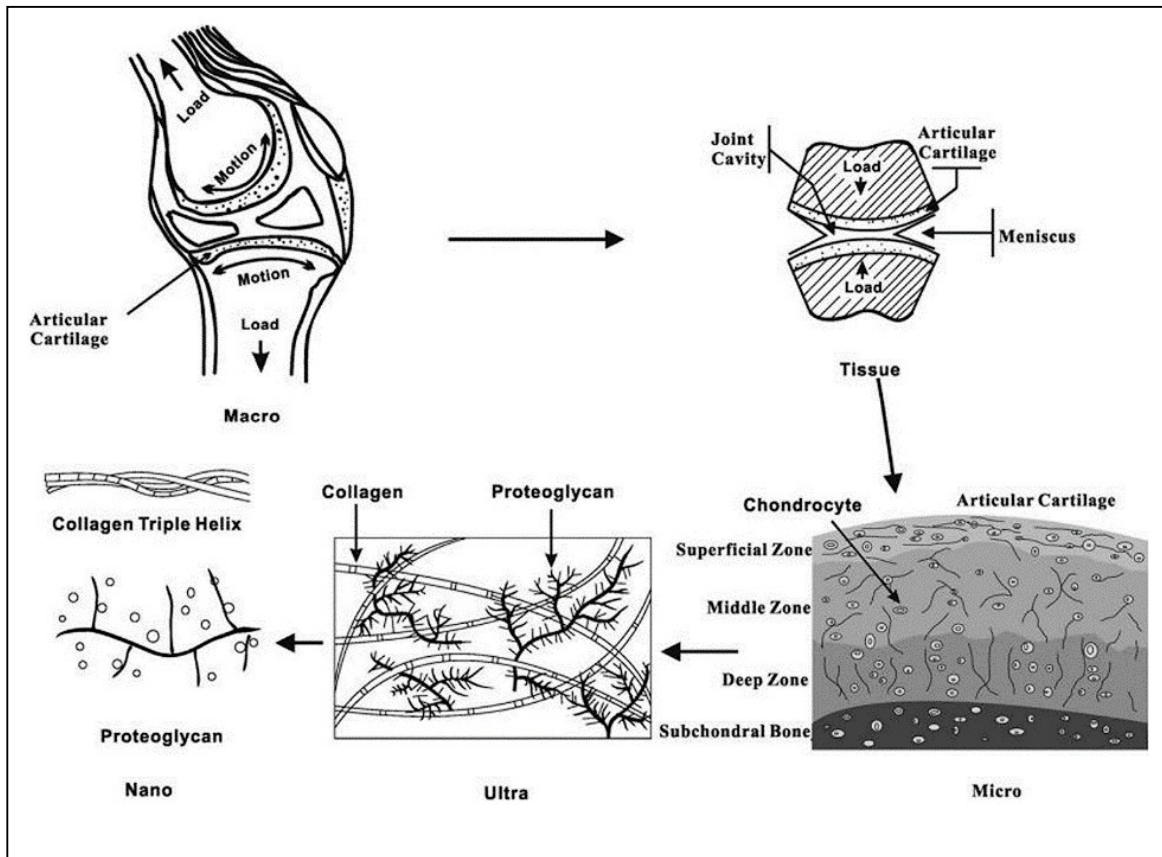
In articular cartilage, the birefringent properties of its ECM can be quantitatively examined by two parameters, retardation, and angle (azimuth), which correspond to the local fibril organization and the fibril orientation respectively (Xia et al., 2001a). The

retardation and azimuth calculations at high resolution in PLM show a depth-dependent anisotropic structure of the collagen network in cartilage. The retardation value is high in the SZ, low in the TZ, and high in the RZ related to the degree of collagen fiber organization and density in different zones. The azimuth calculation shows the collagen fibers are oriented parallel, randomly, and perpendicular to the articular surface in the SZ, TZ, and RZ, respectively (Lee & Xia, 2013). Mapping these two parameters has also permitted the detection of a disrupted collagen matrix due to the early onset of lesions and tissue deformation due to loading (Alhadlaq et al., 2004, 2007b).

Since the structural orientation and organization of the collagen matrix in cartilage are the cause of the magic angle effect and T2 anisotropy in MRI of cartilage, the quantitative correlation between MRI and PLM has enabled independent validation of the ultrastructural information in our comprehensive studies of cartilage and other connective tissues (Batool et al., 2020, 2022; Xia et al., 2001a). Chapter 3 and 4 results are based on the quantitative MRI relaxation data and PLM fibril structural data, which clearly show distinct features in tissue properties at different depths and at different anatomical locations.

Figure 2.1

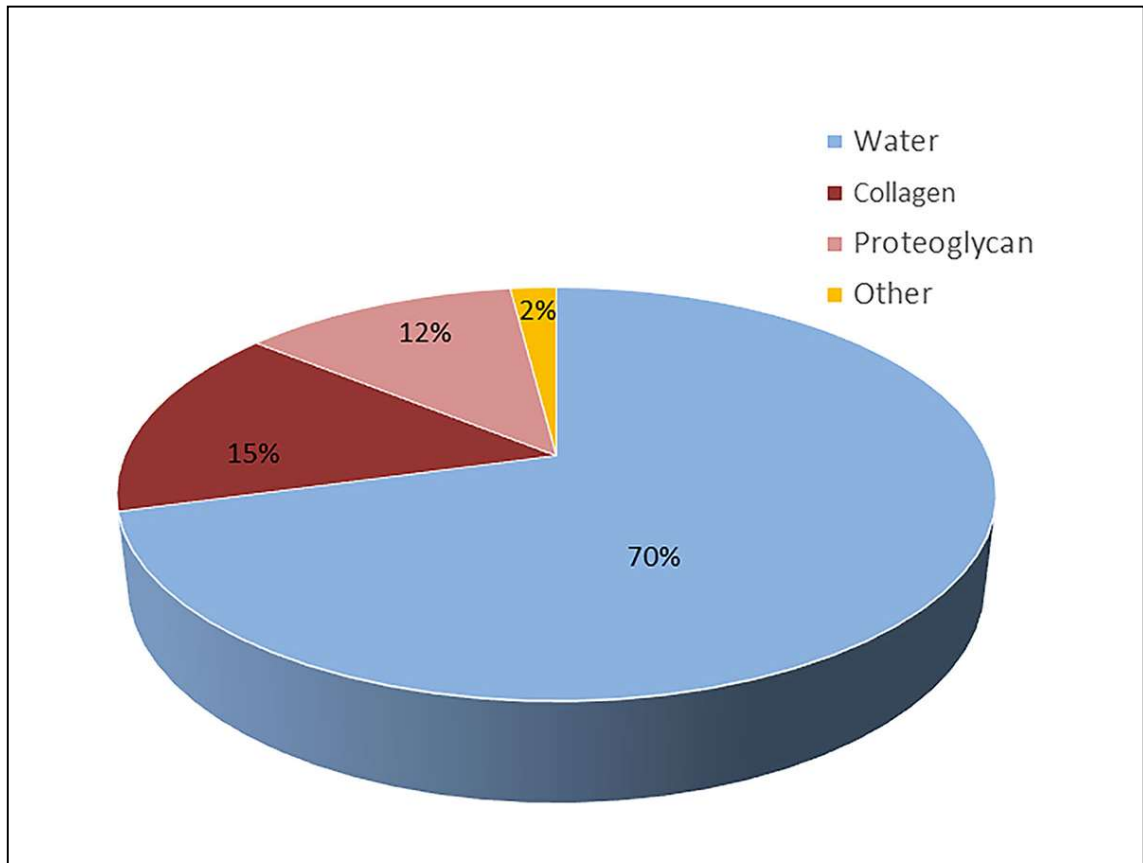
Articular cartilage structure from macro to micro scales



Note. Morphologically (micro-scale) articular cartilage mainly consists of chondrocytes (cells) and the surrounding extracellular matrix (ECM). There are three major components of ECM of articular cartilage, collagen (type II), proteoglycans (aggrecan), and water (not shown). Depth-dependently, cartilage can be divided into three sub-zones (superficial, middle, and deep zone). Reused with the permission of Ge, Z., Li, C., Heng, B., & Cao, G. (2012). *Functional biomaterials for cartilage regeneration. Journal of Biomedical Materials Research. Part A*, 100, Figure 1. Page 2527. Copyright by John Wiley and Sons.

Figure 2.2

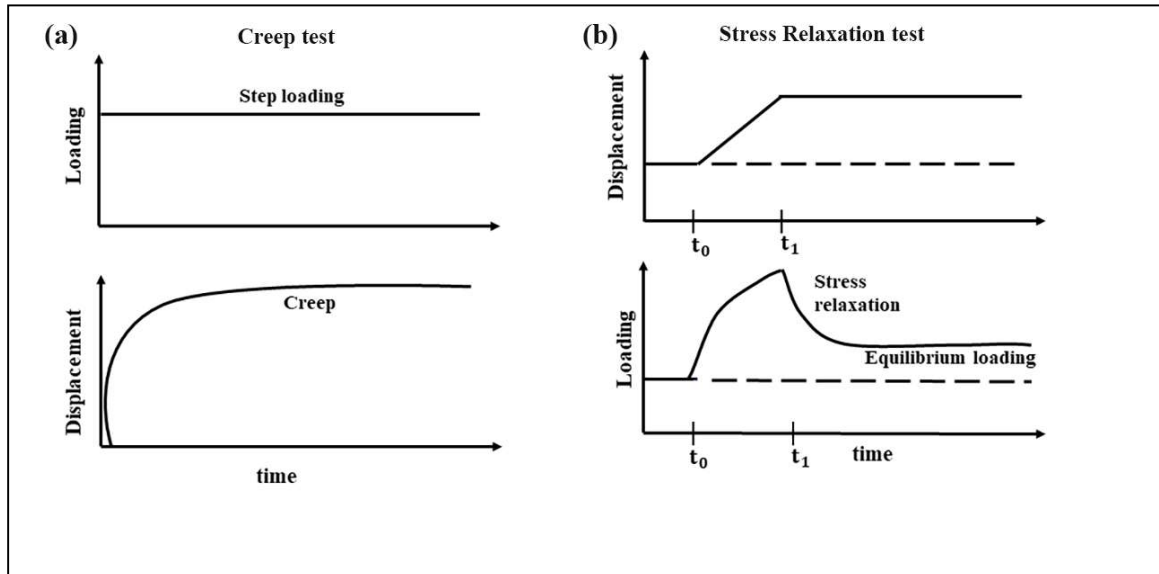
ECM Composition of articular cartilage



Note. Water (blue), constituting 65–80%, collagens (red) make up about 10–20% and proteoglycans (pink), constituting 10–15% of the wet weight of the AC, 1-3% other macromolecules. (yellow).

Figure 2.3

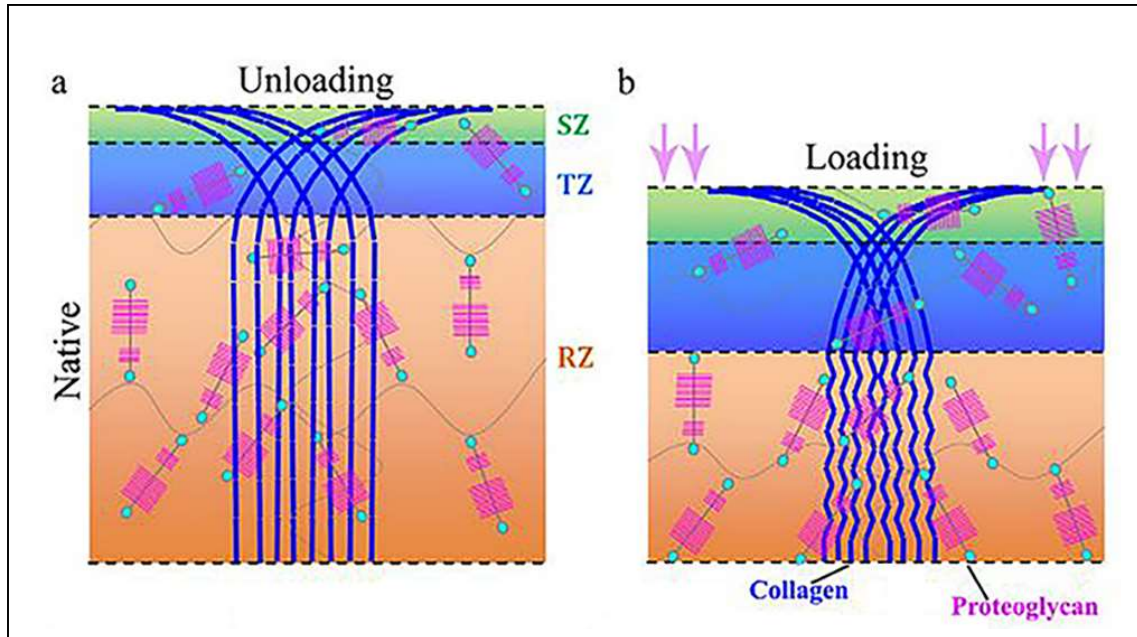
Schematic representation of stress relaxation and creep testing



Note. In creep testing (a) a constant stress or load (time-independent) is applied (step loading) and the resultant change in displacement with time is measured. In stress relaxation testing (b), the initial displacement/strain is ramped (t_0 to t_1) to a certain value and kept constant. The load or applied stress relaxation is measured from peak (t_1) to equilibrium.

Figure 2.4

The schematics of collagen fibril structure and proteoglycan distribution in unloaded and loaded cartilage.

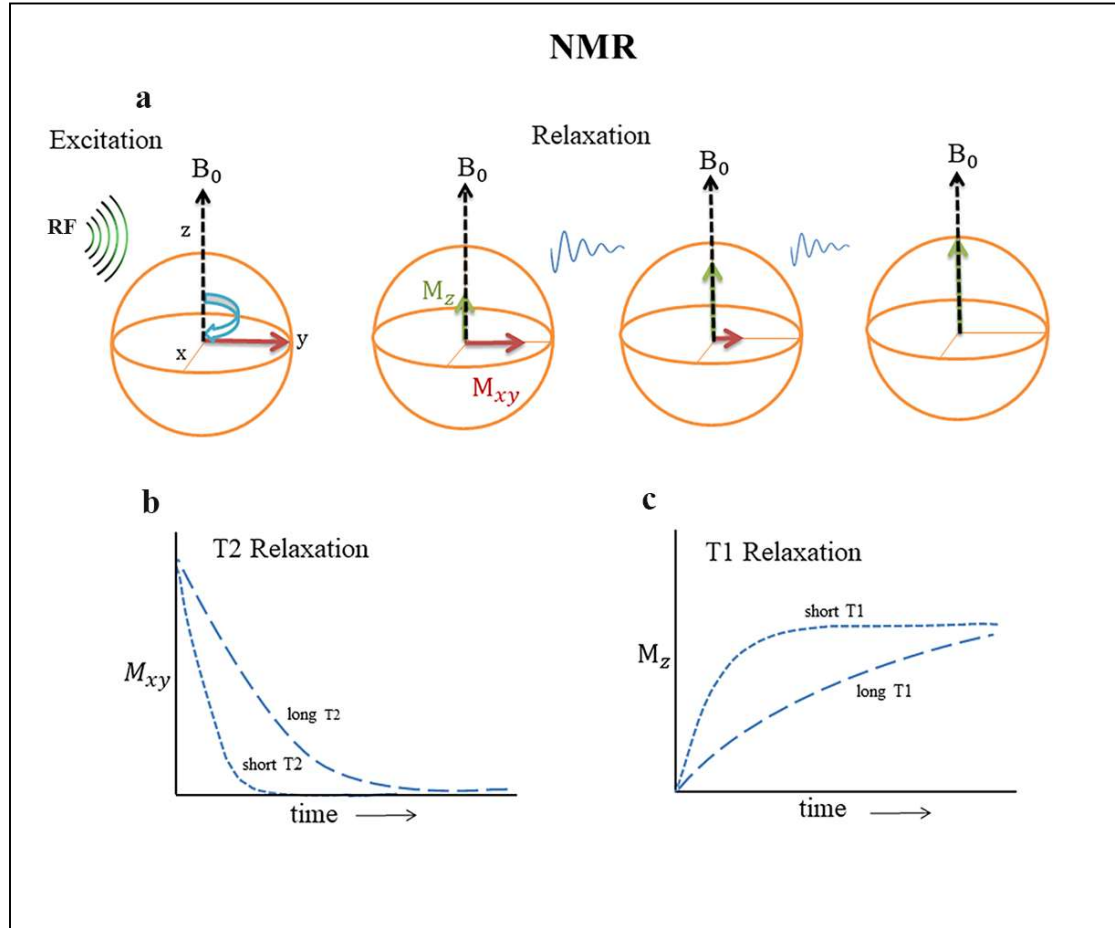


Note. Unloaded cartilage (a) with trizonal structure of collagen fibers and embedded PGs. Loaded cartilage (b), where collagen fibers in superficial and transitional zone deform easily but radial zone fibers resist compression and can have some type of localized zigzag patterns after loading. Reproduced with permission of Wang, N., Badar, F., & Xia, Y. (2015). MRI properties of a unique hypo-intense layer in degraded articular cartilage. *Physics in Medicine and Biology*, 60(22), 8709–8721.

Figure 5. Copyright by IOP Publishing, Ltd.

Figure 2.5

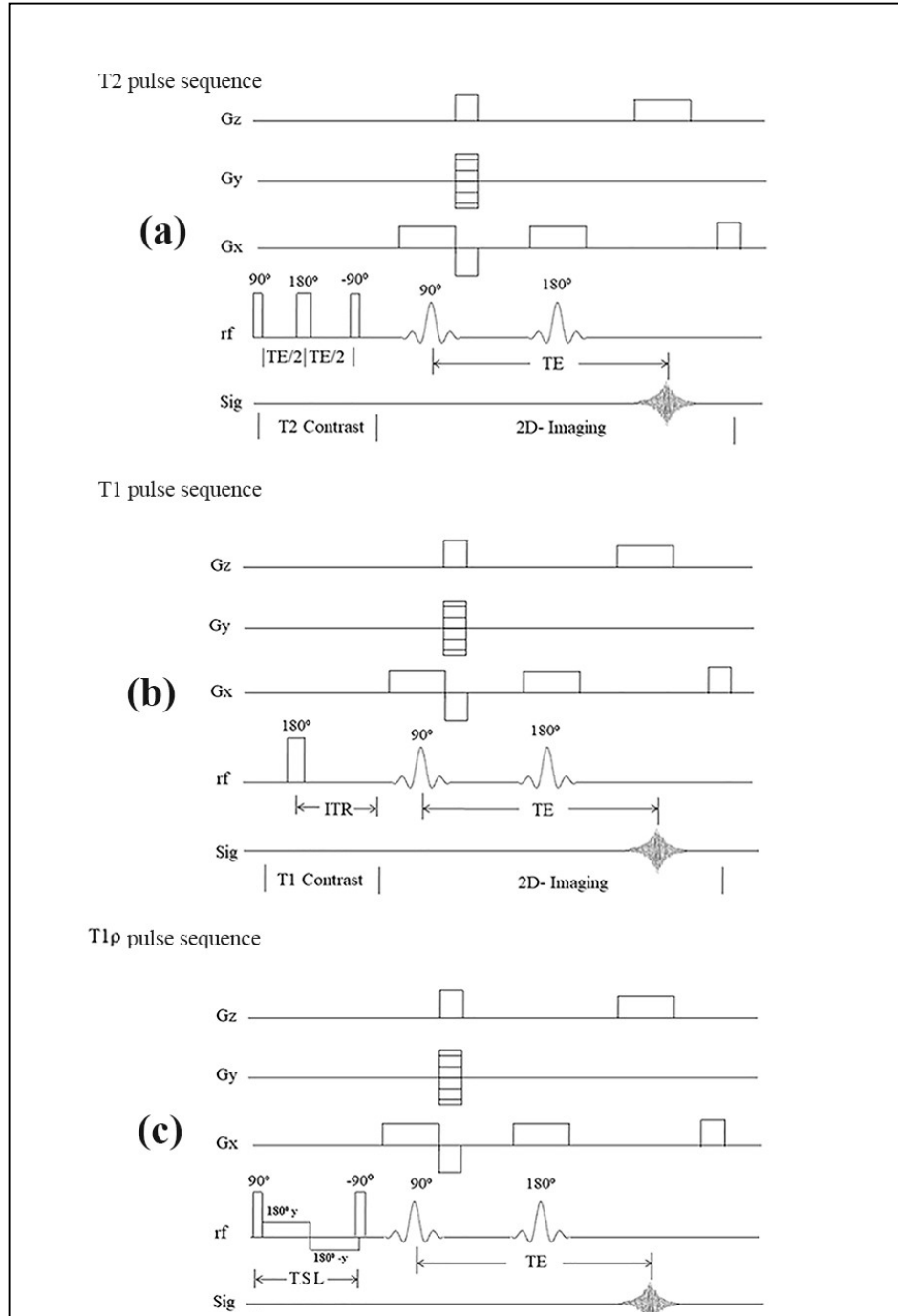
Schematics of working principle of nuclear magnetic resonance (NMR).



Note. (a) Excitation of protons in the magnetic field with radio frequency (RF) signal and subsequent relaxation in transverse (M_{xy}) and longitudinal direction (M_z). (b) Spin-spin relaxation (T2): time is taken for excited spinning protons to go out of phase with each other. (c) Spin-lattice relaxation (T1): time taken for the protons to realign with the external magnetic field (B_0).

Figure 2.6

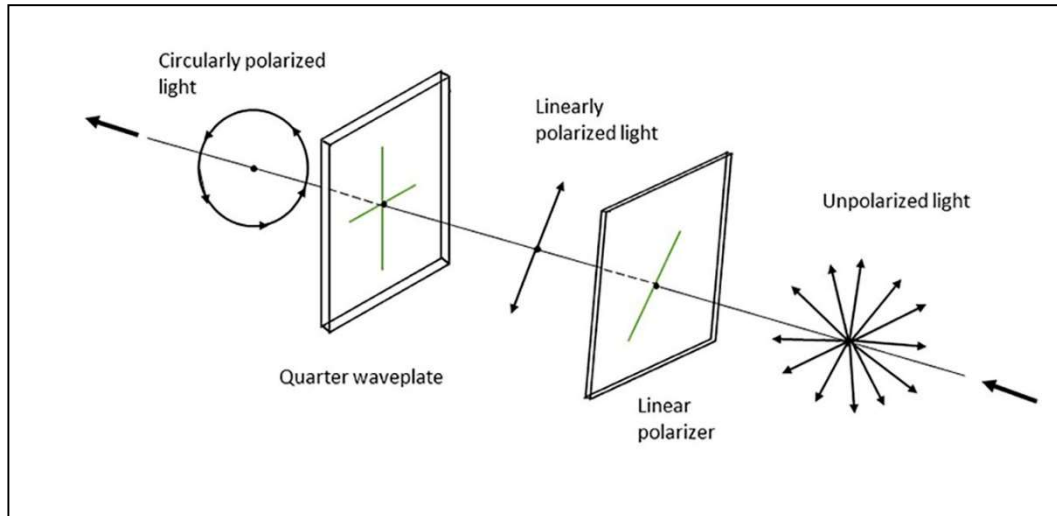
The graphical sketches show the SE-based and magnetization-prepared pulse sequences (T2, T1, T1 ρ).



Note. (a) quantitative T2, (b) quantitative T1 and (c) quantitative T1 ρ . (Gx- Slice selection gradient; Gy - Phase encoding gradient; Gz - read out gradient; TSL - spin lock period; TIR - inversion recovery time; TE - echo time). Originally published by Mahar, R., Batoool, S., Badar, F., & Xia, Y. (2018). *Quantitative measurement of T2, T1 ρ and T1 relaxation times in articular cartilage and cartilage-bone interface by SE and UTE imaging at microscopic resolution. Journal of Magnetic Resonance*, 297, 76–85, Figure 1. Reproduced with copyright permission.

Figure 2.7

Diagrammatic representations of linearly and circularly polarized light



Note. First, an unpolarized light beam enters from the right and propagates through first a linear polarizer oriented at 45° and then a quarter wave plate with its fast axis oriented vertically. The linear polarizer transmits fully linearly polarized light, while the quarter wave plate delays the horizontal component of this light by $\frac{1}{4}$ wave relative to the vertical component, producing left-handed circularly polarized light.

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CHAPTER THREE

QUANTITATIVE μ MRI AND PLM STUDY OF RABBIT CARTILAGE AT SUB-10 μ m RESOLUTIONS

3.1 Introduction

This chapter describes a high-resolution quantitative imaging study, which investigated the individual histological zones in the articular cartilage of rabbits. This study aimed to establish the baseline characteristic T2, T1 ρ , and T1 relaxation times in humeral and femoral cartilage in rabbits using quantitative protocols in microscopic MRI (μ MRI) and polarized light microscopy (PLM). To the best of our knowledge, this is the first correlated and quantitative MRI and PLM study of cartilage at sub-10 μ m resolution, which establishes the baseline characteristics of normal healthy cartilage. The quantitative MRI relaxation data and PLM fibril structural data collectively show distinct features in tissue properties at different depths of cartilage, different in individual histological zones. The establishment of these characteristic features of rabbit cartilage would provide a solid foundation for future utilization of the rabbit model in the OA investigation.

3.2 Materials and methods

3.2.1 Sample preparation

Five healthy rabbits (12-15 weeks New Zealand White, male) were used in this study. The animals were sacrificed for an unrelated biomedical study. Ethical approval for the use of spare tissues in our study was waived by the Institutional Animal Care and Use Committee (IACUC) because the specimens were the discarded tissue. The animals

were frozen immediately at -80 °C after necropsy and thawed at 4 °C before the specimen harvesting. Five intact (i.e., unopened) shoulder joints (the scapula and humeral heads) and three femoral heads from the hip joints were harvested; each was placed in a glass tube and imaged by MRI at low resolutions (70-82 μm pixel size). A cotton swab soaked with 154 mM protease-inhibitor (PI) saline solution was placed in the tube to avoid tissue drying. After each intact joint/bone was imaged, a diamond saw cut each humeral head into 5 slices (Fig 3.1a, b) and each femoral head into 4 slices (Fig 3.1c, d), where each slice was about 1.8 mm thick.

Two cartilage-bone specimens were selected from each humeral head (Fig 3.1a), which resulted in $n=10$ for the humeral specimens. Similarly, one cartilage-bone specimen was harvested from each femoral head (Fig 3.1c), which resulted in $n=3$ for the femoral specimens. Each cartilage-bone specimen (each about $1.8 \times 2 \times 2.5 \text{ mm}^3$) had the full thickness of articular cartilage still attached to the underlying bone. These specimens were bathed in the PI solution and imaged using the same MRI system at high resolution (9.75 μm pixel size).

3.2.2 MRI experimental

The MRI system was a Bruker AVANCE II console and a 7 Tesla/89 mm magnet (Bruker, Germany). The intact joints and bones were first imaged sagittally using the standard multi-slice multi-echo sequence, where each joint/bone had 8 or 10 slices and each slice had 10 intensity images. This resulted in 80 or 100 intensity images for each joint/bone. Each slice had a thickness of 1 mm with a gap of 0.5 mm (humeral) or 0.8 mm (femoral) between any two neighboring slices. The humeral joint imaging had an echo time (TE) of 6.64 ms, and a repetition time (TR) of 2.5 s. The 2D acquisition matrix

was 256×256 and the field-of-view (FOV) was $18 \times 18 \text{ mm}^2$, resulting in a 2D pixel size of $70.31 \text{ }\mu\text{m}$. The femoral head imaging had TE of 7 ms and TR of 2.20 s. The 2D acquisition matrix was 256×256 and FOV was $21 \times 21 \text{ mm}^2$, resulting in a 2D pixel size of $82.03 \text{ }\mu\text{m}$.

All cartilage specimens (10 humeral and 3 femoral) were imaged for quantitative T2, T1 ρ , and T1 at two orientations to the external magnetic field B_0 , 0° and 55° , which allowed the tissue properties to be influenced by the max and min dipolar interaction (Xia, 1997). All quantitative imaging used TR of 2 s, TE of 6.47 ms, and a slice thickness of 0.8 mm. The 2D matrix was 256×128 , which was reconstructed into 256×256 ; the FOV was $2.5 \times 2.5 \text{ mm}^2$, resulting in a 2D pixel size for all cartilage specimens at $9.75 \text{ }\mu\text{m}$.

To determine the depth dependent T2 anisotropy across the full depth of rabbit cartilage, quantitative T2 imaging was repeated at a series of angular orientations to B_0 for two humeral specimens. The selected orientations form a series of discrete sampling points along the geometrical factor $(3\cos^2\theta - 1)^2$ that is proportional to the non-zero dipolar Hamiltonian.

3.2.3 Quantitative μ MRI relaxation protocol

Quantitative T2, T1 ρ , and T1 imaging adapted the concept of magnetization preparation (Haase et al., 1993; Xia, 1997), which consisted of a head segment to prepare the magnetization and an imaging sequence where all timings were constant. The head segment for T2 was a CPMG sequence (Xia, 1997; Zheng & Xia, 2009a), where the echo times were 2, 8, 16, and 38 ms for both humeral and femoral specimens. The head segment for T1 ρ was a spin-lock sequence at a strength of 2000 Hz (N. Wang & Xia,

2011), where the spin-locking pulse lengths were 2, 8, 16, 38 ms for humeral and 2, 16, 40, and 60 ms for femoral. The head segment for T1 was an inversion-recovery sequence (Xia, 1997), where the inversion recovery times were 0, 0.8, 1.4, and 3.5 s for all specimens. Since only the timing of the head segment is altered during the experiments, the relaxation effect of the magnetization can be determined accurately.

3.2.4 Quantitative PLM protocol

After μ MRI, 7 humeral cartilage-bone specimens and 3 femoral cartilage-bone specimens were fixated in 10% neutral buffered formalin overnight at 4 °C. A number of cartilage-bone slabs (5-6 mm thick) were cut at the center of the humeral head (at the location of the μ MRI slices) across its width horizontally (hence containing both lateral and medial sides) and fixated. The fixated slabs and the cartilage specimens in μ MRI were sent to a histology service (Yale Pathology Tissue Services, CT) to section into 6 μ m thin sections using the paraffin method. 30 sections from the humeral cartilage specimens, 6 sections from the femoral cartilage specimens, and 5 sections from the whole-width humeral head slabs were imaged using PLM.

The PLM system consists of a digital setup (Cambridge Research & Instrumentation, MA) that is mounted on a polarized light microscope (Leica, Germany). The output of the CCD camera was processed (Oldenbourg & Mei, 1995) to generate two quantitative images: the optical retardation and the angular orientation. Both quantitative images describe different aspects of the collagen matrix in cartilage (Xia et al., 2001a). All cartilage specimens were imaged at the identical orientation under a 10x objective which yielded a pixel size of 1.0 μ m. All intact humeral heads were imaged under a 2.5x objective which yielded a pixel size of 4.0 μ m.

3.2.5 Image and data analysis

On each T2 map of the humeral and femoral bone, a three-pixel-wide region of interest (ROI) was selected, which covered the entire thickness of cartilage and extended beyond the cartilage. The ROI was chosen at an identical location on the joint surface from which the small cartilage-bone specimens would be harvested for μ MRI. From each ROI, one 1D profile along the cartilage depth was generated. On each quantitative 2D relaxation image, twenty parallel neighboring columns of data were selected and averaged to obtain one 1D profile. (Note that 20 consecutive columns in μ MRI at 9.75 μ m have approximately the same tissue width as 3 consecutive columns in the intact humeral cartilage resolution at 70.31 μ m.) On each quantitative PLM image, 195 parallel neighboring columns, which would have approximately the same width as the slice thickness in μ MRI, were averaged to produce one 1D profile.

Since the averaging from 2D ROI to 1D profiles in μ MRI and PLM occurs perpendicular to the tissue depth, the spatial resolutions of the 1D profiles along the tissue depth are still 70.31 μ m for MRI of the intact humeral, 82.03 μ m for MRI of the femoral head, 9.75 μ m for MRI of the cartilage-bone specimens, and 1 μ m for PLM of the intact joint/bone. All 2D images and 1D profiles, where each came from one independent imaging experiment, used identical scaling and settings without any additional adjustment. The post-acquisition analysis used the public-domain software ImageJ (NIH) and a commercial software KaleidaGraph (Reading, PA).

3.3 Results

3.3.1 MRI of Intact Joint and Small Cartilage-bone Specimens at Different Resolutions

Fig 3.2a and Fig 3.2c showed four intensity images for one intact humeral joint and one intact femoral head respectively, together with their quantitative T2 images. Note that the surface of the humeral cartilage was bordered by synovial fluid and other tissues since the shoulder joint was intact; in comparison, the surface of the femoral cartilage was bordered with air (which appeared black in the intensity images) since only the femoral head was imaged in the glass tube (due to the large size of the acetabulum of the hip). Fig 3.3b and Fig 3.3d showed the intensity and T2 images for one humeral specimen and one femoral specimen respectively. The high-resolution images of humeral and femoral cartilage appear similar. A thin black line was more visible near the femoral surface (Fig 3.2d) than the humeral surface (Fig 3.2b), indicating a lower T2 in the surface of the femoral cartilage.

3.3.2 Quantitative relaxation measurements at 9.75 μm resolution

Fig 3.4 shows the quantitative T2, T1 ρ , and, and T1 images and profiles in both humeral and femoral cartilage at 9.75 μm resolution, where the same tissue block was imaged twice using the identical protocol at two different orientations in B₀. These images and profiles reveal similarities and differences between humeral and femoral cartilage in rabbits, and between rabbit and canine humeral cartilage in the literature (Xia et al., 2001a, Lee & Xia, 2013). Between humeral (Fig 3.4a-3.4c) and femoral (Fig 3.4d-3.4f) cartilages in rabbits, the relaxation images and profiles appear similar at 55° to B₀, at which the dipolar interaction has the weakest influence to spin relaxation. This is

likely because both cartilages belong to the same type of ball-and-socket joint and hence are structured similarly. At 0° to B_0 (which has the strongest dipolar influence to spin relaxation), the shape of the T2 profile in femoral cartilage is well defined and narrower than humeral cartilage, the meaning of which at the fibril structure level needs further investigation.

Between the humeral cartilage in rabbits (Fig 3.4a-3.4c) and canines in our previous studies using the same protocols (Xia et al., 2001a; Xia, Moody, Alhadlaq, et al., 2002), the depth-dependent characteristics of these relaxation times in rabbit cartilage match broadly the corresponding characteristics in canine cartilage, where the bell-shaped curve in the T2 profiles (Fig 3.4a, 3.4d) implies a classical three-zone structure in cartilage, which can be used to sub-divide cartilage thickness into multiple histological zones (Lee & Xia, 2013; Xia et al., 2001a). One noticed that the T2 profile at 55° was not as uniform in values across the tissue depth, which should have an anatomic origin at the fibril level in the tissue. Using the established criteria (Lee & Xia, 2013; Xia et al., 2001a), the zonal division of cartilage thickness was carried out on all μ MRI images (Table 3.1). In both humeral and femoral cartilages, SZ is about 21 μ m thick, just over 2 pixels in μ MRI at 9.75 μ m/pixel. This thin thickness of SZ in rabbit cartilage puts a high-resolution requirement on μ MRI and any other imaging tools in the study of rabbit cartilage. A peculiar feature in the humeral data is that the L blocks were consistently thicker than the C blocks, which suggests a topographical variation of cartilage thickness over the joint surface (L and C blocks were two adjacent tissue blocks, about 1.8 mm apart on the joint surface). Table 3.2 summarizes the zonally averaged relaxation times, based on the zonal thickness data in Table 3.1. The biggest zonal variations are T2 at 0° ,

while the zonal data in T1 ρ and T1 have much less depth dependence and little orientational dependence.

Fig 3.5 compares the T2 profiles from the T2 images in Fig 3.4. Two features can be identified for humeral cartilage (Fig 3.5a). First, for the deeper part of humeral cartilage (about 75% of the total cartilage), T2 profiles at both resolutions match each other. Second, for the top 25% of the tissue, the low-resolution imaging missed the critical feature of the surface cartilage. The continuously increased T2 values towards the articular surface (the depth of 0) and beyond the articular surface (into the neighboring space) in Fig 3.5a demonstrates the consequence of low-resolution MRI of cartilage, which averaged cartilage with the surrounding synovial fluid – two tissues that have very different T2 relaxation times. Similarly, the low-resolution MRI of femoral cartilage (Fig 3.3b) missed out on the critical features of the surface cartilage (the top 25%), where the cartilage signal was reduced because of the averaging with air near the surface. In contrast, μ MRI can resolve the T2 characteristics of the surface tissue faithfully (Fig 3.5), which represent the superficial and transitional zones.

3.3.3 Depth-dependent T2 anisotropy

To study the depth-dependence of T2 anisotropy, two humeral specimens were each imaged at six different orientations to B_0 . Fig 3.5a showed one set of six T2 profiles from six independent T2 experiments, plotted together without additional scaling. These profiles demonstrated that T2 anisotropy in rabbit humeral cartilage is strongly depth-dependent and orientational-dependent, with the biggest anisotropy around 80 μ m to 160 μ m depths, representing the upper part of the radial zone. A close examination can be done by plotting T2 anisotropy at every tissue depth (every 9.75 μ m). Four such plots are

shown in Fig 3.5b. T2 anisotropy in rabbit humeral cartilage is prominent at 0 μm and stays equally prominent at 9.75 μm (not shown) – both pixels represent SZ cartilage. As the depth goes deeper, T2 anisotropy weakens and remains nearly identical within the depths of 29.25 μm to 48.75 μm , which signals TZ cartilage. However, T2 anisotropy remains noticeable even at its weakest (up to 40%), which reflects a residual fibril structure in TZ. From the depth of 68.25 μm , T2 anisotropy becomes stronger again, reaching a max at 351 μm (near the bottom of RZ). These pixel-by-pixel features in the depth-dependent T2 anisotropy precisely reflect the variations in the collagen organization across the entire thickness of cartilage (Xia, Moody, & Alhadlaq, 2002; Zheng et al., 2011; Zheng & Xia, 2009b).

3.3.4 Quantitative PLM results

The angle and retardation maps from the quantitative PLM represent the pixel-averaged fibril orientation and fibril organization of the collagen matrix in cartilage respectively. Fig 3.6 showed the retardation image of an intact humeral head. Several observations can be noted. First, there is a dark ‘line’ in the retardation image at ~ 50 μm below the articular surface, which represents the location of the least organized fibers in tissue (the center of TZ). Second, this dark line gradually vanishes towards both peripherals, which suggests the collagen architecture in peripheral cartilage is different from the central region. Thirdly, humeral cartilage has strong topographical variation in its thickness, from ~ 250 μm at the joint center to ~ 500 μm at the peripherals. Although this thickness variation is significant, it is not as complex as in some other joints such as the tibia (refer to Fig 1.10 in Chapter 1 of the book (Xia & Momot, 2016)). Finally, our cartilage specimens for μMRI and PLM were harvested from a slightly off-center

location in the humeral head, which explains the consistent thickness difference between our L specimens and our C specimens (Table 3.1). Fig 3.7 showed the quantitative PLM results from the same humeral specimen from which the μ MRI data were discussed (Fig 3.4). These PLM images from rabbit cartilage are very similar to the PLM images from canine cartilage ((Xia et al., 2001a). Briefly, the depth-dependent retardation (Fig 3.7a) had its lowest values about 50 μ m below the articular surface, which indicates the center of TZ; the high retardation values at the articular surface indicate the fine fibril organization at the articular surface while the high retardation values in the deep cartilage represent the fibril structure of RZ (closely and more densely packed). In the angle image (Fig 3.7b), the blue and red colors in this color scale showed $\sim 85^\circ$ orientational difference between the SZ and RZ fibrils. The tide mark in this rabbit tissue was not quite visible as in some mature cartilage (Xia et al., 2001a). Femoral cartilage was also analyzed by PLM, which yielded similar results as in humeral cartilage. Fig 3.7c showed one set of quantitative PLM profiles (from Fig 3.7a and 3.7 b), which were averaged over 6 tissue sections from one humeral specimen, together with the T2 profile from the same tissue block (Fig 3.4a). These profiles solidify the descriptions and conclusions for the angle and retardation images in the early part of this section.

3.3.5 μ MRI zones vs histological zones correlation

Using the established criteria (Lee & Xia, 2013; Xia et al., 2001a), the zonal division of cartilage thickness was carried out on all PLM images from 30 histological sections (from 5 humeral specimens). PLM zones from the humeral specimens were compared with the corresponding μ MRI zones from the same specimens (Table 3.3). Some disagreements can be noted for the thin zones between μ MRI and PLM, which is

likely caused by the difference in the imaging resolution (9.75 μm in μMRI , 1.0 μm in PLM). The correlations in the total tissue thickness and the radial zone are satisfactory. Similarly, the zonal division for femoral cartilage has also been carried out, where the μMRI and PLM results from the same femoral specimens were compared for their zonal thicknesses. In femoral cartilage, SZ was $\sim 7\text{-}9\%$, TZ $8\text{-}9\%$, and the radial zone was $\sim 82\%$ of the total cartilage.

3.4 Discussion

Among many MRI parameters that could be used to study cartilage degradation, T2, T1 ρ , and T1 relaxation times are currently the most commonly used parameters in clinical MRI of OA (Xia, 2013). These parameters are sensitive to molecular motions and their interactions with the tissue surroundings. This study has successfully quantified T2, T1 ρ , and T1 in both humeral and femoral cartilage in rabbits at 9.75 μm resolution. To the best of our knowledge, these μMRI results are the first quantitative characterization of the entire depth of humeral and femoral cartilage in rabbits at a sub-10 μm resolution. The complementary use of quantitative PLM in this study further enhances the μMRI results. Given our experience in canine studies (Alhadlaq et al., 2004; Lee et al., 2016; Lee & Xia, 2013; Xia, 1997; Xia, Alhadlaq, et al., 2008), the depth-dependent features of T2, T1 ρ , and T1 in rabbit cartilage were found to be like those in canine cartilage. This similarity implies a classical three-zone structure in rabbit cartilage and provides possibilities for doing comparative analyses between canine studies and leporine studies.

3.4.1 Challenges in studying rabbit cartilage – age, thickness, topographical variations.

There are at least three challenges in using the rabbit model for OA research. First, the growth plates in rabbit joints do not close until 6-9 months, beyond which those

animals can be considered skeletally mature (Bland & Ashhurst, 1996; Burr, 2002; Isaksson et al., 2010; Turunen et al., 2011). The animals in this study (also in many studies in the literature) were 3-4 months old, hence skeletally immature. However, the same studies (Kavanagh & Ashhurst, 1999; Turunen et al., 2011) also noted that the collagen matrix of articular cartilage in mature rabbits appeared to be “similar” to that at 12–14 weeks. In contrast to this similar conclusion between immature and mature rabbits, some previous μ MRI and PLM studies of canine humeral cartilage have found vastly different characteristics between young and mature canines (Xia et al., 2003; Xia, Moody, Alhadlaq, et al., 2002). In any case, the maturity of rabbit tissues should be noted for any study that creates an injury or intervention in the tissue and waits for a time for its development, as well as for any comparative analyses of the rabbit data in the OA literature.

Second, rabbit cartilage is thin, a few hundred microns in thickness in the largest joints. This thin thickness possesses challenges for several types of studies, including diagnostic imaging and surgery. In this imaging study, based on the birefringent information at the 1.0 μ m optical resolution, we determined the zonal thicknesses for the central part of humeral cartilage, where SZ and TZ were just over 30 μ m each. These findings demonstrate that high resolution (e.g., approximately 10 μ m) is required to analyze zonal differences in the rabbit cartilage. One subtle but important reason to use two resolutions to study rabbit cartilage in this MRI study comes from the imaging scaling in the MRI of cartilage (Xia, 2007). In this study, for a nominal thickness of 300-400 μ m in rabbit cartilage, there were approximately 4-5 pixels in our low-resolution MRI of intact bone heads (70-82 μ m pixel size) and 35 pixels in our μ MRI of tissue

blocks (9.75 μm pixel size). In contrast, although human knee and hip cartilage are much thicker, clinical MRI also has a bigger pixel size, which commonly resolves human cartilage in 3-5 pixels. Consequently, our low-resolution MRI of rabbit cartilage mimics almost exactly the tissue averaging situation in clinical MRI of human cartilage (Xia, 2007). Hence our experimental methodology and approach can be adapted to perform many OA projects using the rabbit model.

Thirdly, most of the tissue characteristics in cartilage change topographically over a joint surface, likely due to the consequence of load-bearing and motional patterns for a particular joint and a particular animal. Even for a ball-and-socket joint such as the humeral head in rabbits (which has considerably simpler variations than a knee joint), the thickness of cartilage ranges from about 250 μm at the center to twice the thickness at the periphery (Fig 3.6). These topographical variations exist on any joint surface (Xia, 2000a; Xia et al., 2003; Xia, Moody, Alhadlaq, et al., 2002). In this study, slightly off-center harvesting of the cartilage-bone specimens from the humeral head has led to noticeable variations in the quantitative measurement of the tissue properties by both μMRI and PLM (Fig 3.4-3.7). In μMRI results, for example, we can attribute the non-uniformity of the T2 depth profile at 55° (Fig 3.4) to be the results of these off-center specimens, where the collagen organization was not entirely perpendicular to each other between the superficial and radial fibrils, which was also confirmed by the PLM results (Fig 3.7c). These topographical variations can make the interpretation of the individual study difficult and the comparison among different studies in the literature challenging.

3.4.2 Limitations of study

This study has several experimental limitations. First, five animals were used in this study, from which five intact shoulder joints and three femoral heads from the hip joints were harvested. These animals belong to a total of over 20 nearly identical rabbits that we obtained from the same source and studied in our lab in different projects. We can confirm that the μ MRI and histology features from the tissues of other animals were highly consistent with the presentation in this study. Second, quantitative imaging can be influenced by the topographical variation of cartilage, i.e., the surface site where the specimen is harvested and studied. This variation could be significant for joints such as the knee. In humeral and femoral heads, by contrast, our experience indicates that the topographical variation has a much smaller influence on cartilage properties and hence the limited sampling sites in this study can represent most of the joint surface (apart from the peripherals). Third, this study demonstrates that the depth-dependent characteristics of the T2, T1 ρ , and T1 relaxation times in humeral and femoral cartilage in rabbits are similar to those in canine cartilage (Xia et al., 2001a; Xia, 2013). This similarity implies a classical three-zone structure in the rabbit cartilage and the possibility of doing comparative analyses between canine studies and leporine studies.

Table 3.1a

Averaged zonal thicknesses based on the T2-0° profiles of humeral cartilage (n = 10 specimens)

Specimen	Superficial Zone (μm)	Transitional Zone (μm)	Radial Zone (μm)	Total Tissue (μm)
1L	19.50	48.75	234.25	302.25
1C	19.50	48.75	224.25	292.50
2L	19.50	58.50	292.50	380.25
2C	29.25	48.75	234.25	302.25
3L	19.50	48.75	380.25	458.25
3C	19.50	48.75	360.75	429.00
4L	19.50	58.50	380.25	458.25
4C	19.50	48.75	302.25	370.50
5L	19.50	48.75	399.75	468.00
5C	29.25	58.50	370.50	458.25
Mean ± σ	21.45 ± 1.3	51.67 ± 1.6	317.90 ± 21.8	391.95 ± 22.5
%	6	13	81	100

Note. σ represents the standard error. % is the relative percent thickness of each zone

based on the total thickness.

Table 3.1b

Averaged zonal thicknesses based on the T2-0° profiles of femoral cartilage (n = 3 specimens)

Specimen	Superficial Zone (μm)	Transitional Zone (μm)	Radial Zone (μm)	Total Tissue (μm)
1C	19.50	29.25	253.50	302.25
2C	29.25	48.75	282.75	360.75
3C	29.25	39.00	292.50	360.75
Mean ± σ	26 ± 3.3	39 ± 5.6	276.25 ± 11.7	341.25 ± 19.5
%	8	11	81	100

Note. σ represents the standard error. % is the relative percent thickness of each zone

based on the total thickness.

Table 3.2a

Averaged orientation dependent (0°, 55°) zonal T2, T1ρ, and T1 relaxation times based on the zone division of humeral cartilage (n = 10 specimens).

	T2 (ms)		T1ρ (ms)		T1 (s)	
	0°	55°	0°	55°	0°	55°
SZ	20.46 ± 1.55	51.31 ± 3.87	86.32 ± 2.92	88.82 ± 3.97	1.25 ± 0.02	1.32 ± 0.02
TZ	37.96 ± 1.66	58.99 ± 2.75	92.49 ± 3.12	93.60 ± 3.52	1.21 ± 0.02	1.21 ± 0.01
RZ-I	12.04 ± 0.58	62.94 ± 2.91	90.84 ± 2.67	103.63 ± 2.67	1.26 ± 0.03	1.27 ± 0.02
RZ-II	7.24 ± 0.61	37.55 ± 1.93	53.97 ± 2.27	65.67 ± 1.85	0.95 ± 0.03	1.01 ± 0.02

Note. Data are in mean ± standard error. RZ-I and RZ-II are the top and bottom equal halves of the radial zone.

Table 3.2b

Averaged orientation dependent (0°, 55°) zonal T2, T1ρ, and T1 relaxation times based on the zone division of femoral cartilage (n = 3 specimens)

	T2 (ms)		T1ρ (ms)		T1 (s)	
	0°	55°	0°	55°	0°	55°
SZ	17.16 ± 3.59	48.10 ± 7.47	68.34 ± 3.88	76.01 ± 5.77	1.18 ± 0.09	1.29 ± 0.08
TZ	33.30 ± 3.73	51.15 ± 1.82	76.30 ± 3.23	79.94 ± 4.72	1.21 ± 0.01	1.26 ± 0.04
RZ-I	10.90 ± 0.65	46.19 ± 3.60	82.32 ± 9.06	89.58 ± 1.06	1.23 ± 0.05	1.27 ± 0.03
RZ-II	4.30 ± 2.10	25.89 ± 2.50	35.63 ± 4.22	51.25 ± 2.19	0.75 ± 0.07	0.90 ± 0.06

Note. Data are in mean ± standard error. RZ-I and RZ-II are the top and bottom equal halves of the radial zone.

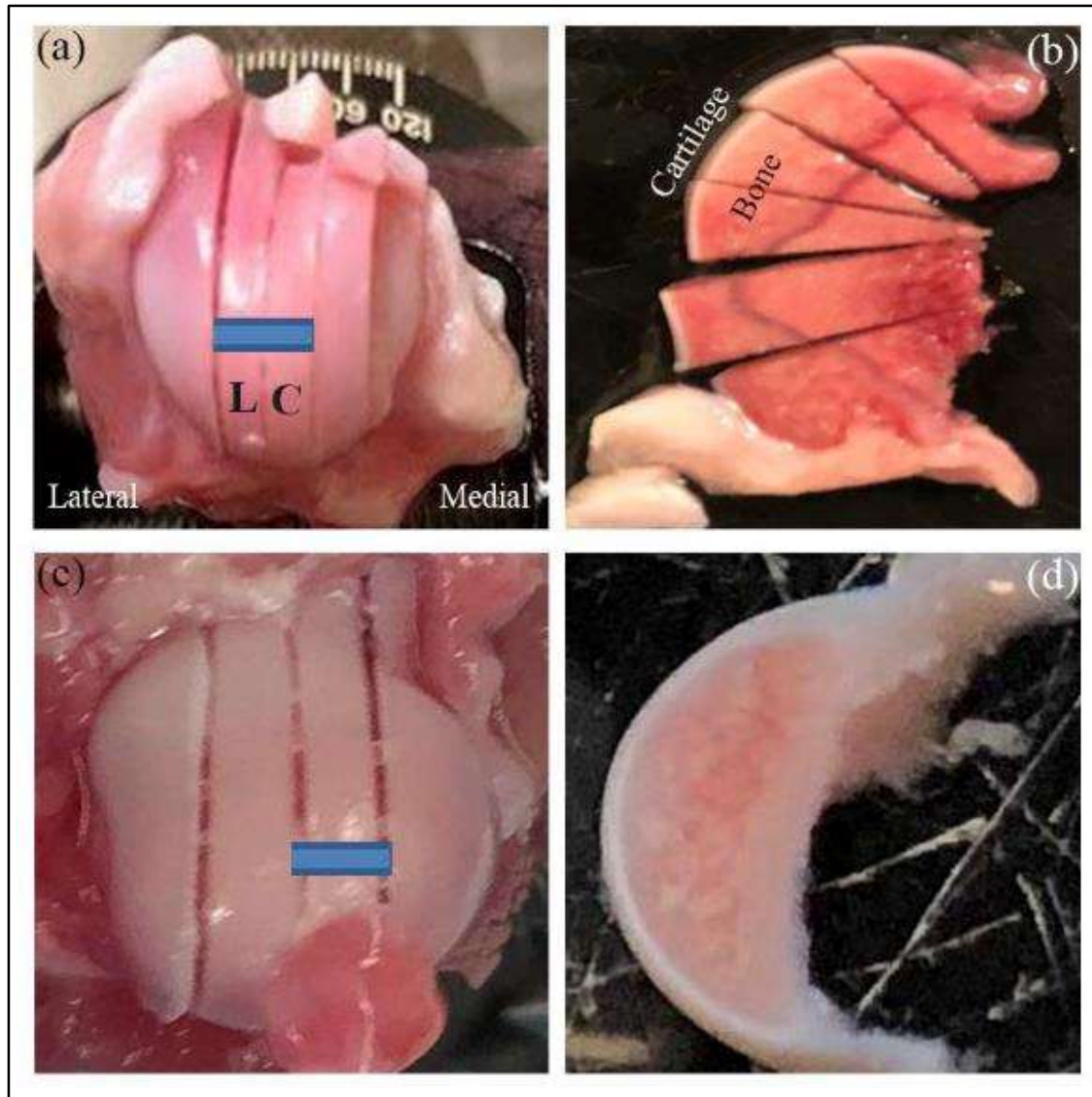
Table 3.3*Correlations of μ MRI vs PLM zonal thickness (μ m)*

	μMRI Thickness	PLM Thickness	t-probability
SZ	21.45 $\pm$ 1.95	29.09 $\pm$ 2.36	0.04
TZ	52.65 $\pm$ 2.38	31.69 $\pm$ 1.24	<0.001
RZ	366.60 $\pm$ 16.82	379.68 $\pm$ 15.76	0.58
Total Tissue	440.70 $\pm$ 18.13	440.46 $\pm$ 16.82	0.95

Note. Correlations of the averaged zonal thickness in humeral cartilage between μ MRI (n = 5 cartilage specimens) and PLM (n = 30 histological sections from 5 μ MRI specimens at 6 sections per specimen). Data are in mean $\pm$ standard error. The t-probabilities were calculated using Student t-test analysis.

Figure 3.1

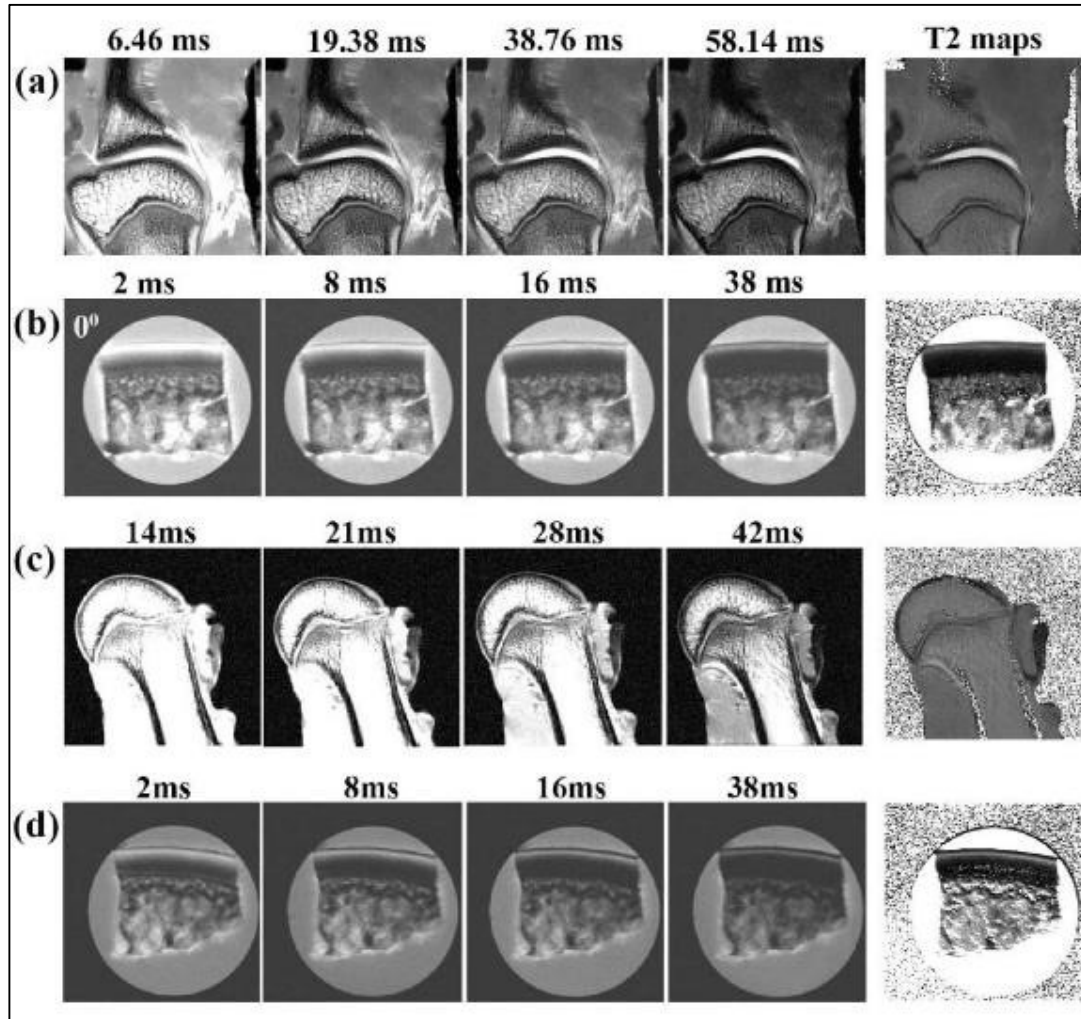
Rabbit humeral and femoral head and histological sections



Note. The top left and right corners are the greater and lesser tubercles respectively (a). After an MRI of the intact shoulder joint, four cuts were made over each intact head, as shown in the photo. The two peripheral slabs were not used in specimen imaging because of their irregularity. The second and third slabs (each about 1.8 mm thick) were selected for the quantitative μ MRI and PLM. The rectangular shape marks the approximate location of the imaging slice in the high-resolution imaging. (b) The side view of a central slab cut into small specimen blocks, through the humeral head. (c) A femoral head, where a central slab is shown in (d).

Figure 3.2

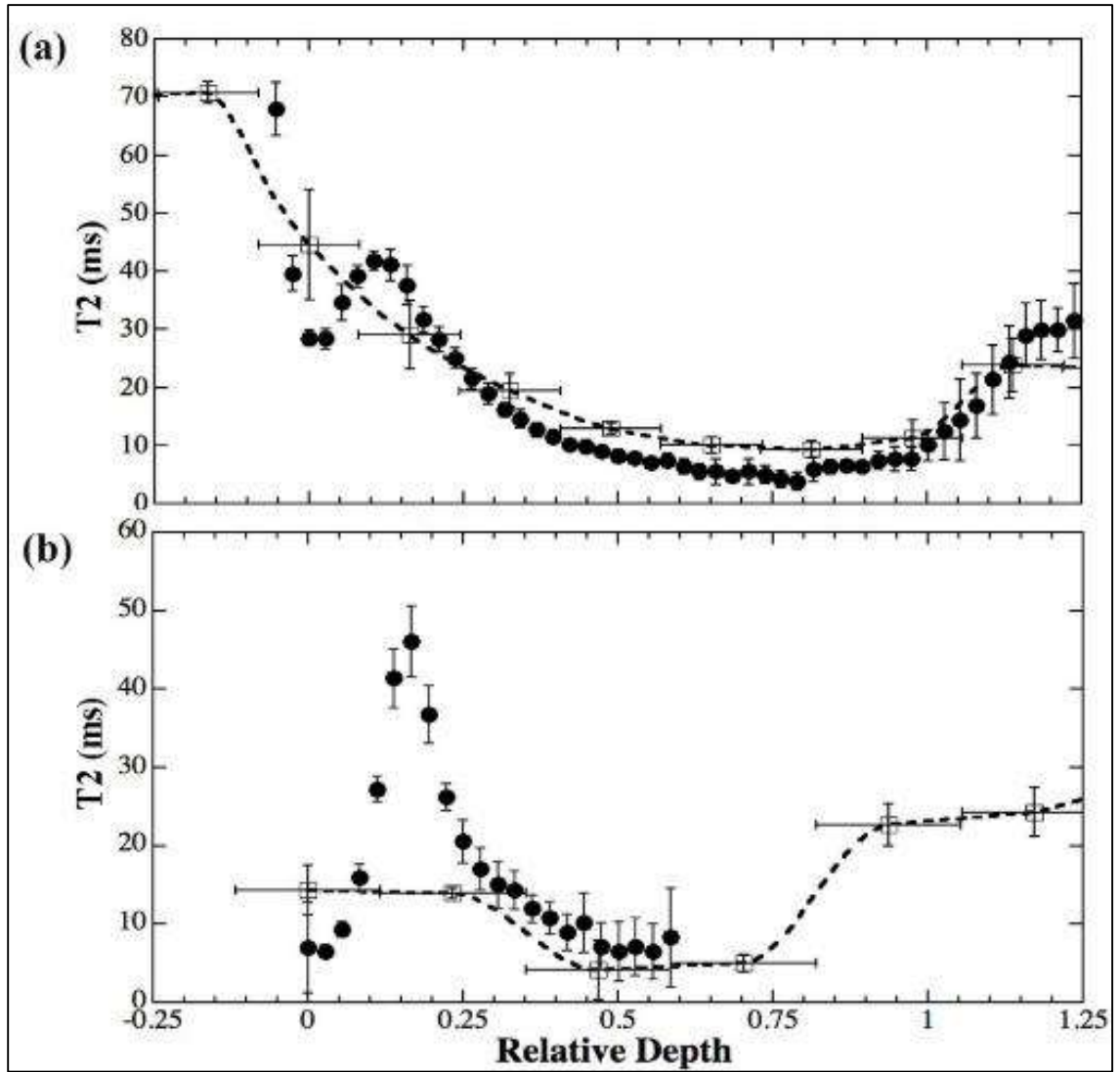
T2-weighted intensity images and T2 maps of an intact shoulder, the femoral head (80 $\mu\text{m}/\text{pixel}$) and small cartilage bone plugs (9.75 $\mu\text{m}/\text{pixel}$)



Note. An intact shoulder joint (a) and a humeral cartilage specimen (b) at different echo times, at the pixel resolution of 70.31 μm and 9.75 μm respectively. (c) and (d) show similar images from a femoral head, at the pixel resolutions of 82.03 μm and 9.75 μm respectively. The quantitative T2 images from each series of ten T2-weighted intensity images were shown on the most-right images. Since the direction of the magnetic field, B_0 was vertically up in the figure, the orientation of the cartilage specimens was 0° to B_0 (Fig 3.2b, 3.2d), which was like the surface orientation of the tissue when the intact joint was imaged in the magnet in sagittal view (Fig 3.2a, 3.2c). All T2 images were displayed with the same up/low limits (0-100 ms).

Figure 3.3

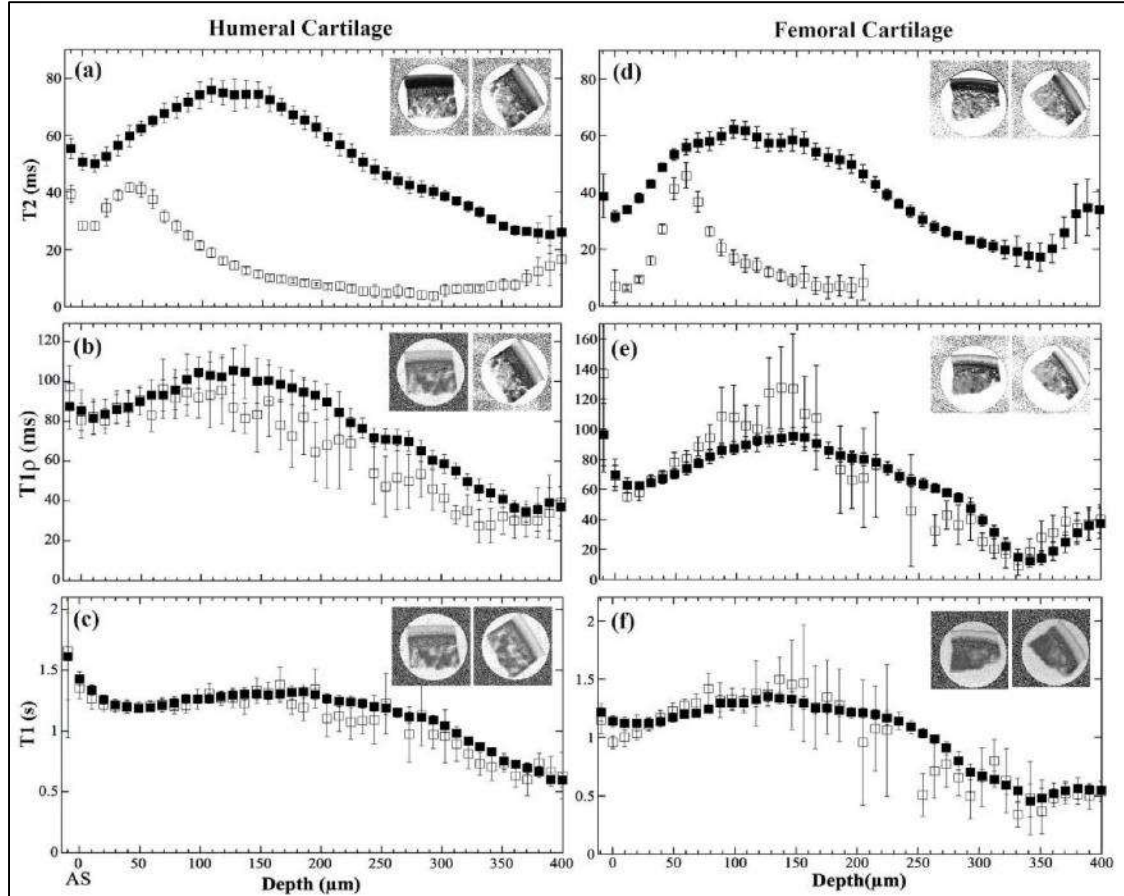
The depth dependent T2 profiles from the T2 images at low ($\sim 80 \mu\text{m}/\text{pixel}$) and high ($9.75 \mu\text{m}/\text{pixel}$) resolution



Note:(a) humeral cartilage at 70.31 $\mu\text{m}/\text{pixel}$ (open squares) and 9.75 $\mu\text{m}/\text{pixel}$ (solid dots) and (b) femoral cartilage at 82.03 $\mu\text{m}/\text{pixel}$ (open squares) and 9.75 $\mu\text{m}/\text{pixel}$ (solid dots), respectively. The depth scale is relative, with 0 marking the articular surface and 1 marking the cartilage-bone interface. The dashed lines fit through the T2 profiles in the low-resolution MRI. The vertical error bars represent the standard deviation of the data, and the horizontal error bars in the low-resolution mark the pixel sizes. Note that in (b), μMRI T2 (0°) are shown till the tissue depth of 0.6, after which the image intensities become very low and the calculated T2s become unreliable.

Figure 3.4

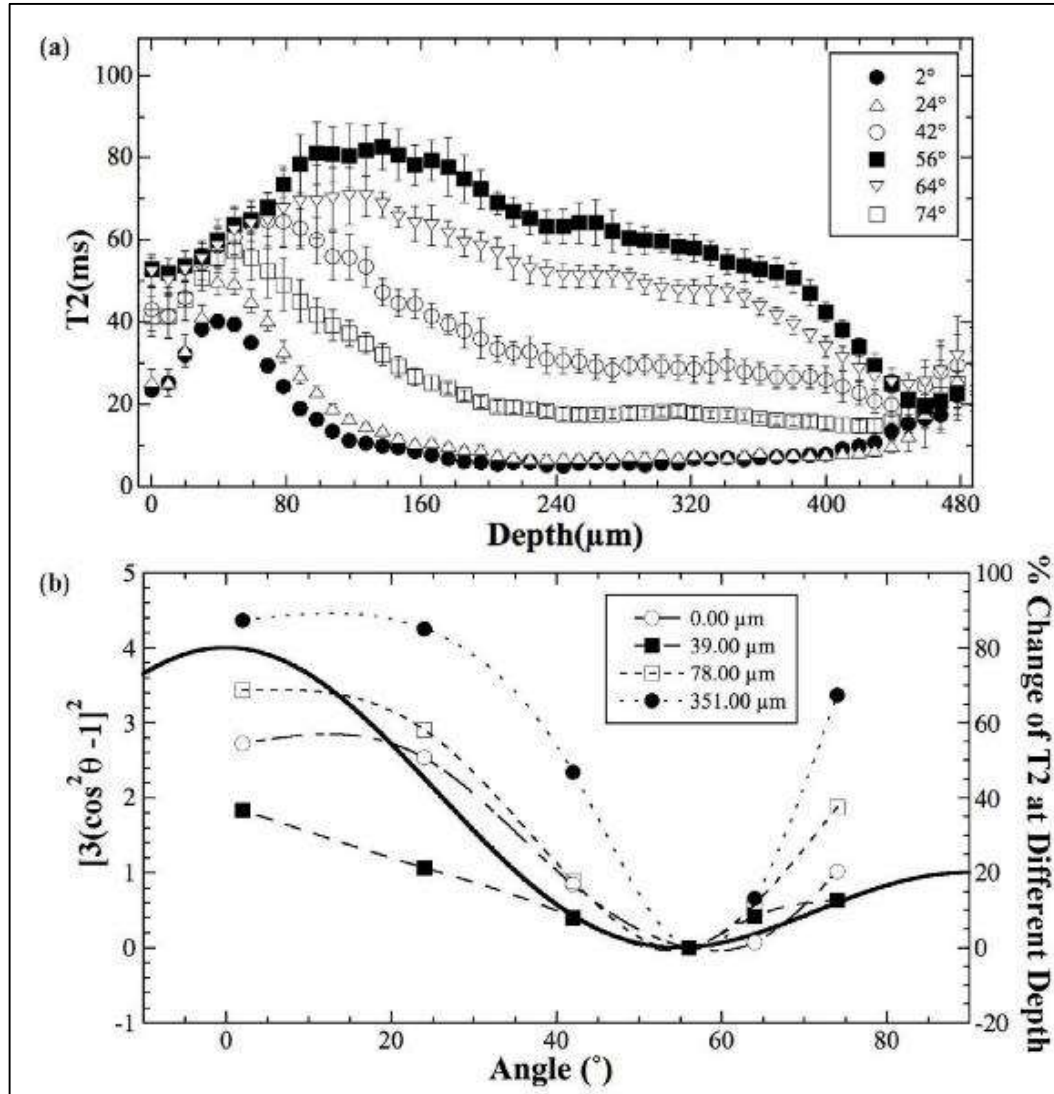
The quantitative T2, T1ρ, and T1 images and their respective 1 D profiles at 9.75 μm/pixel for humeral and femoral cartilage



Note. (a)-(c) humeral cartilage, and (d)-(f) femoral cartilage. The solid dots are the profiles at the 55° specimen orientation, while the open squares are the profiles at the 0° specimen orientation to the external magnetic field B_0 .

Figure 3.5

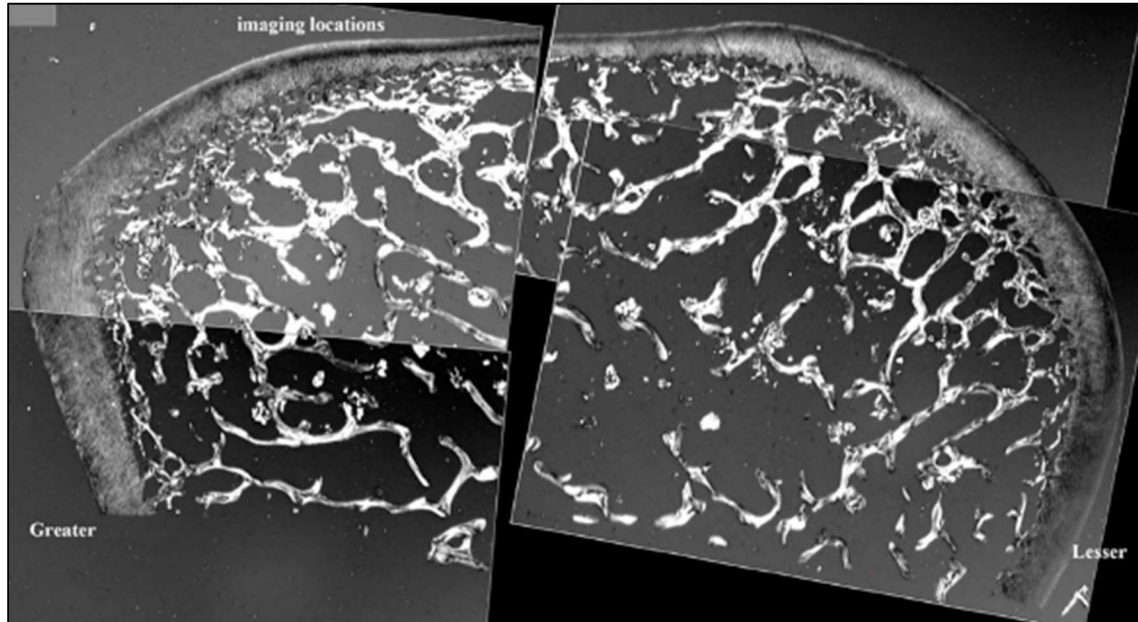
Depth-dependent T2 profiles and the plots of T2 anisotropy of humeral cartilage.



Note. T2 profiles at six orientations to B_0 , which follow the geometric factor in the dipolar interaction, $(3\cos^2 \theta - 1)^2$, that manipulates T2 relaxation in MRI. (b) The plots of T2 anisotropy (% change in T2 to the 55° data, as a function of the specimen orientation) at four tissue depths (at a $9.75 \mu\text{m}$ tissue depth step): $0 \mu\text{m}$ represents the first layer of the superficial zone; $39 \mu\text{m}$ is inside the transitional zone; $78 \mu\text{m}$ marks the boundary between the transitional zone and radial zone, and $351 \mu\text{m}$ locates deep in the radial zone. The solid line plots the $(3\cos^2 \theta - 1)^2$ in the dipolar interaction that dominates the T2 characteristics in the articular cartilage.

Figure 3.6

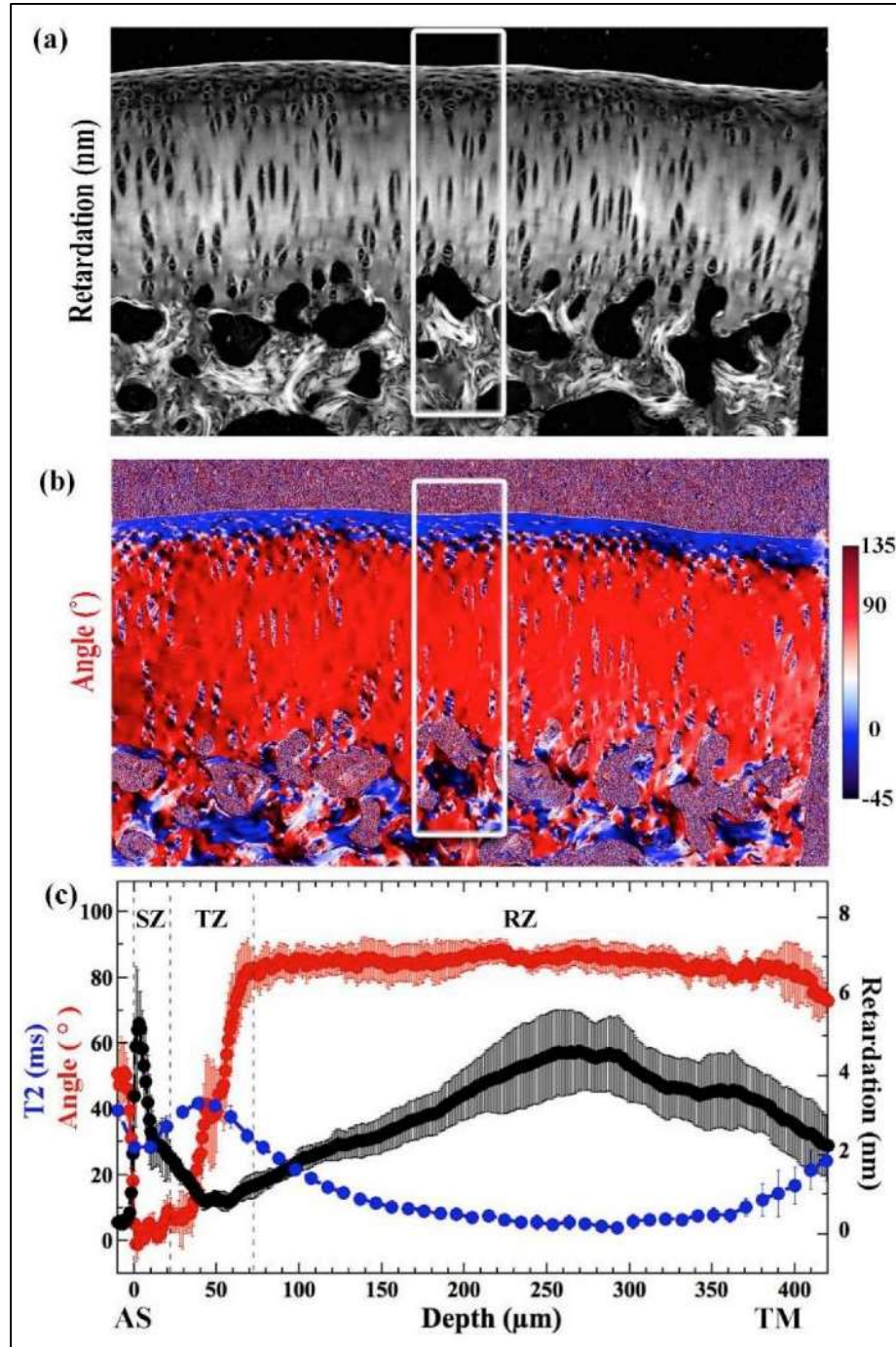
The retardation images from an intact humeral head of the shoulder joint, imaged by polarized light microscopy with a 2.5x objective (4 μm pixel resolution)



Note. The histological slice of the intact humeral head was imaged in four parts to cover the entire humeral head with a 2.5x objective (4 μm pixel resolution); the four quantitative retardation images were scaled using identical limits (0 – 10 nm); the images were pieced together in photoshop. The humeral head shows a strong curvature, flat in the central area and rounded towards both peripherals. (Some of the curvatures could also be influenced by the histological process after the underlying bone becomes soft in decalcification.) In the retardation images, large and small values indicate that the collagen fibrils are ordered, which could include the information for the fibril diameters and fibril packing density.

Figure 3.7

2D retardation (a) and angle (b) images in PLM, at $1\mu\text{m}$ pixel resolution of one humeral cartilage specimens



Note. In the angle images, a typical arch structure of the collagen matrix should have a nominal 90° orientational difference between the superficial and radial fibers, hence represented by the blue and red colors. The rectangular boxes in (a) and (b) show the location where the depth-dependent profiles were calculated and shown in (c). Also, in (c) is the T2 profile from μ MRI of the same specimen. Note that the oval shapes in the retardation image are the locations of the chondrocytes that used to be present in cartilage. Since no protocol was used in histology to preserve the cells, the cellular components were washed out, which yielded black color in the retardation images (i.e., background values). In contrast, the same empty cellular spaces are often filled with random numbers in the angle images, which is due to the imaging algorithm that determines two quantitative numbers at each pixel location). These inaccurate assigned angles and retardation values in the cellular cavities contribute to and exaggerate the size of the error bars in the PLM profiles. The vertical dashed lines mark the locations of the three histological zones. AS and CBI in the tissue depth mark the articular surface and cartilage-bone interface respectively.

CHAPTER FOUR

LOCATION-SPECIFIC STUDY OF YOUNG RABBIT FEMORAL CARTILAGE BY T2 μ MRI AND POLARIZED LIGHT MICROSCOPY

4.1 Introduction

In this study, we characterized the structural variations at different anatomical locations of femoral cartilage in young skeletally immature rabbits (12-14 weeks old). Quantitative MRI relaxation data and PLM fibril data revealed collaboratively distinct topographical variations in both cartilage thickness and collagen organization in the juvenile joint. This study aimed to better understand the structural specifics in skeletally immature tissue in the rabbit model, related to its potential in the use of osteoarthritis studies, and the remodeling of the collagen fibril network during growth and maturation at different anatomical locations over the same joint surface. Since any joint has specific 3D shapes/contours and a unique set of biomechanical and biochemical environments, any joint degradation at its early stages must have complex variations over joint surfaces, and knowledge of normal topographical variations will be useful to understand the degradation process dependent upon the location of the initiation event.

4.2 Materials and methods

4.2.1 MRI of intact joint

Four knee joints from 12 to 14 weeks old, healthy rabbits (White New Zealand, male) were obtained from a nearby higher-education institution. The animals were sacrificed for an unrelated approved biomedical study ((PROTO202000161). Ethical approval for the use of spare tissues in our study was waived by the Institutional Animal

Care and Use Committee (IACUC). The animals were frozen immediately at -80 °C after necropsy and thawed at 4 °C before specimen harvesting. Each intact joint was placed in a glass tube with a cotton swab soaked with physiological saline to avoid dehydration and imaged at room temperature using a Bruker AVANCE III HD NMR spectrometer equipped with a 7 Tesla/89 mm magnet and 25 mm RF resonator (Bruker, Germany).

All four intact knees were imaged in sagittal and coronal views using the standard multi-slice multi-echo sequence, where each imaging experiment had 8-10 slices and each slice had 6-8 intensity images. Each slice had a thickness of 0.8 mm with a gap of 1mm between any two neighboring slices. The imaging echo time (TE) was 8.23 ms, and the repetition time (TR) was 3 s. The FOV was 22×22 mm² and the 2D acquisition matrix was 256×128, which was later reconstructed to 256×256, resulting in a pixel size of 86 μm.

4.2.2 μMRI of cartilage-bone specimens

After imaging the intact joints, two knees were opened (Fig 4.1a) to harvest the cartilage-bone specimens from the femoral medial condyles. Each condyle was cut into 3 slices (Fig 4.1e), which were ~1.6-1.8 mm in thickness. Three individual cartilage-bone specimens were harvested from three separate locations on the same (middle) condyle slice (Fig 4.1f), which were named anterior (A), central (C), and posterior (P), respectively. Each cartilage-bone specimen was rectangular, approximately 1.8×2×2.5 mm³ in size, and still had a full thickness of articular cartilage attached to the underlying bone. All cartilage-bone specimens were equilibrated in saline with 154 mM protease inhibitors. The total number of cartilage-bone specimens for μMRI was six (two for each location).

Each of the cartilage-bone specimens was imaged independently five times using identical imaging parameters, except for the physical orientations of the specimen to the external magnetic field B_0 , ranging from 0° to 125° (where 0° is defined as the normal axis of cartilage surface being in parallel with B_0) with the use of a home-built rotating device (Xia, Moody, & Alhadlaq, 2002). A magnetization-prepared 2D spin-echo sequence was used for the quantitative T2 imaging of all cartilage-bone specimens, with TR of 2 s, TE of 6.47 ms, and a slice thickness of 0.8 mm. The location of the slices throughout each series of imaging experiments was identified to be the same during the project. The 2D matrix was 256×128 , which was reconstructed into 256×256 ; the FOV was $2.5 \times 2.5 \text{ mm}^2$, resulting in a pixel size of $9.75 \text{ }\mu\text{m}$ for all cartilage specimens (Batoool et al., 2020; Xia, 1997). The echo times of the leading contrast segment (TEc) had four increments: 2, 16, 22, and 46 ms, for the central (C) and posterior (P) specimens and 2, 16, 46, and 60 ms for the anterior (A) specimens.

4.2.3 PLM specimens and protocol

All six cartilage-bone specimens after the non-invasive μMRI were processed for histology. In addition, two remaining intact joints were processed for whole-joint histology, where the full-length and width slabs (4-5 mm thick) were cut in transverse planes (across tibiofemoral and patellofemoral regions, as in Fig 4.1b and 4.1.c) and along the middle of the medial condyle (including all 3 μMRI sites, as in Fig 4.1f). All histology specimens were fixated in 10% neutral buffered formalin overnight at room temperature. All fixated specimens were sent to a histology service (Yale Pathology Tissue Services, CT), which cut the specimens at specified locations into $6\text{-}\mu\text{m}$ thin sections using the paraffin embedding method. Three sections from each cartilage-bone

specimen (the μ MRI samples) and three sections from each longitudinal and lateral slab were imaged using a digital PLM system (Leica, Germany & Cambridge Research & instruments, MA). Each imaging acquisition in the digital PLM system generated two quantitative images: the optical retardation (in units of nm) and the angular orientation (in units of $^\circ$). The angle image represents the voxel-averaged orientations of the collagen fibrils, whereas the retardation image represents the voxel-averaged organization of the collagen fibrils in tissue (Xia et al., 2001a).

Each cartilage-bone specimen (A, C, P) had at least three full-thickness histological sections, which were imaged individually at identical orientations under 10x and 40x objectives. This yielded a pixel size of 1 μ m and 0.25 μ m respectively. The 10x images were used for quantitative retardation and azimuthal profiles and the 40x images were used to observe the territorial matrix surrounding the individual chondrocytes at different depths. Histological sections of the intact femoral condyles were imaged under 2.5x and 5x objectives, which yielded pixel sizes of 4 μ m and 2 μ m, respectively. These low-resolution optical images were used to study the heterogeneity of collagen fibril organization across the intact femoral condyles in orthogonal directions.

4.2.4 Image and data analysis

Quantitative maps of T2 relaxation were calculated by a single exponential fitting of the data on a pixel-by-pixel basis for all specimens. Each cartilage-bone specimen was imaged in μ MRI five times at five different orientations to B_0 , which resulted in 20 independent proton images and 5 calculated T2 maps for each specimen. On each quantitative 2D T2 image, one rectangular region of interest (ROI) with 15 parallel neighboring columns was selected manually and averaged into one 1D depth-dependent

T2 profile, which enabled the observation of the depth-dependent T2 anisotropy in the tissue and the comparison among profiles from different imaging experiments. Since the column-averaging occurred perpendicular to the tissue depth, the pixel resolution in the T2 profiles was still 9.75 μm along the tissue depth. The width and location of ROI were the same for all independent imaging experiments of each sample, using the image analysis software ImageJ. All 2D images and 1D profiles, where each came from one independent imaging experiment, used identical scaling and settings without any additional adjustment.

On each quantitative PLM image of cartilage specimens obtained at 10x objective, 146 parallel neighboring columns, which would have approximately the same width as the ROI thickness in μMRI , were averaged to produce one pair of 1D quantitative retardation and angle profiles across all tissue zones. The 40x images enabled the visualization of the shape and orientations of chondrocytes concerning the surrounding territorial and interterritorial matrix at different depths (histological zones) among all histological sections from three locations. The post-acquisition analysis used the public-domain software ImageJ (NIH) and commercial software KaleidaGraph (v4, Reading, PA).

4.3 Results

4.3.1 MRI of intact knee joint

Before opening the knee joints, the intact knees were imaged by μMRI at the pixel resolution of 86 μm . Fig 4.2 shows the sagittal and coronal views of T2-weighted intensity images respectively, together with their respective quantitative T2 images. The epiphyseal cartilage (growth plate) appears brighter and homogenous in intensity; in contrast, signal

variations can be seen over the thickness of articular cartilage, especially in the femur, even at the low 86 μm resolution. The appearance of articular cartilage in T2-weighted MRI images was highly sensitive to its orientation to the external magnetic field B_0 (vertically up). Because of its curved articular surface, the femoral condyle provides a convenient model to study the orientation effect of cartilage T2 relaxation to B_0 (i.e., the magic angle effect), if the quality of cartilage is the same at every location. If T2-sensitive structural variations exist as expected in the juvenile joint during development, the interpretation of imaging results needs additional considerations. In our imaging results, among the three topographical locations on the central medial femur, the central location (C) was about at the magic angle ($\sim 54.74^\circ$) to B_0 (sagittal view), where the influence of the dipolar interaction to the spin-spin relaxation was minimized; consequently, the cartilage in the central region appeared thicker than at the two other locations, anterior (A) and posterior (P). Away from the magic angle location, the femoral cartilage appeared bilaminar with a high-intensity surface layer and a low-intensity deep layer. Due to the pixel size, signal variations for thinner regions (A and P) can be influenced by volume averaging.

4.3.2 PLM images of intact femur joint at 4 μm resolution

Fig 4.3 is the quantitative retardation images of the whole joint by PLM, at the three cross-sectional surfaces shown previously in Fig 4.1f, Fig 4.1b, and Fig 4.1c. (Multiple retardation images were acquired individually in PLM and pieced together without any additional intensity adjustment.) Fig 4.3a is the retardation image of a longitudinal section of a femoral medial condyle at 4 μm pixel resolution (from the tissue block as in Fig 4.1f), which showed that femoral cartilage was thinner at peripheries and thicker in the central region. Three topographical variations were determined from this

image. First, the central cartilage was about 600 μm in thickness, twice the thickness when compared with the anterior and posterior cartilage ($\sim 300 \mu\text{m}$). This PLM observation is consistent with the MRI results of knee joints in Fig 4.2. Second, the values of retardation were the lowest at the anterior site and highest at the central site, where most of the deep central cartilage had high retardation (i.e., high coherence in fibril organization) and most of the anterior site had very low retardation (i.e., low coherence in fibril organization). Finally, the lowest values of retardation at any topographical location occurred consistently at approximately 35 μm to 40 μm below the articular surface, regardless of the total thickness of the cartilage at this location.

Fig 4.3b and 4.3c are the retardation images of two cross-sections taken transversely or orthogonally from the anterior (patellofemoral) locations of the femur (which opposes the patellar cartilage) and the central (tibiofemoral) location of the femur (which opposes the tibial cartilage) as shown in Fig 4.3a (as well as shown previously in Fig 4.1b and 4.1c). Two observations can be made for these cross-sections. First, there is little difference in cartilage thickness across most of these transverse/orthogonal sections (except at the peripheral edges). Second, the femoral cartilage (Fig 4.3b) from the central cross-section seemed to have more variations in the structural patterns across the tissue depth. For example, this occurred between the cartilage at the trochlear ridges (TR) to the cartilage in the trochlear groove (TG). For these reasons, one ROI was selected from the central location of the medial femoral head (shown in Fig 4.3c), and two ROIs were selected from the anterior cartilage opposing the patellar cartilage (shown in Fig 4.3b).

4.3.3 MRI of site-specific T2 anisotropy results

Fig 4.4 shows the T2 anisotropy results at the higher spatial resolution of $9.75\ \mu\text{m}$ in μMRI , from three specimens acquired from the anterior, central, and posterior regions of the femoral medial condyle, shown previously in Fig 4.1f and Fig 4.3a. (Note that each specimen was imaged five times between the orientations of 0° and 120° . Since these images appeared similar visually, only two images for each location were shown in the figure. The most insightful information on T2 anisotropy among each set of five images can be obtained from the depth-dependent profiles in Fig 4.4, which contain all orientations at all locations.) The images and anisotropy profiles of cartilage from the central (C) location have familiar features that assemble the well-known T2 anisotropy under the influence of the dipolar interaction (Xia, 1998, 2000b). Based on the data at 55° , the cartilage thickness was $\sim 500\ \mu\text{m}$ at the central location. When the specimen was oriented at 0° , where the dipolar interaction is the strongest, most of the deep tissue had little signal in the spin-echo-based imaging sequences. This caused the truncation of the T2 (0°) profile to a depth of $\sim 200\ \mu\text{m}$, beyond which the signal became too low to be reliably used to calculate T2 (the echo time in the imaging sequence is 6.47 ms). Compared to the central cartilage, the posterior cartilage was about 50% thinner, at $\sim 350\ \mu\text{m}$. The T2 (0°) profile followed approximately the bell-shaped curve as seen in the central cartilage, but the peak was not as sharp and occurred deep in the percentage thickness. The rest of the anisotropy profiles did not follow strictly the T2 anisotropy as observed in the central cartilage. This implies that the overall orientational structure of the collagen fibers in the posterior cartilage might be slightly different from the central cartilage. But it largely followed the usual three-zone structure as seen in mature cartilage. The anterior cartilage had the most curvature and its T2 anisotropy profiles had the most complicated variations. Two ROIs

(A1, A2) were selected for the anterior specimen, where A1 is located at the anatomical location of the trochlear ridge (TR) and A2 is in the proximity of the trochlear groove (TG). The T2 anisotropy profiles from the A1 location lost all recognizable features as shown in the central cartilage, which implies its collagen matrix did not have any of the usual three-zone structure.

4.3.4 Quantitative PLM images of cartilage-bone specimens

Fig 4.5 shows the quantitative PLM results at 1 μm pixel resolution from the cartilage specimens at three topographical locations on the medial femur, whose μMRI data was presented in Fig 4.4. In the quantitative retardation images, the small black ‘holes’ inside the articular cartilage are the locations where the cells and cell clusters used to reside (the histology procedure did not preserve the cellular details). In these angle images, the blue and red colors indicated a 90° difference between the fibril orientations; the same 90° difference also existed between the black and white colors, which had a 45° shift in the orientation to the blue and red combination.

Several features can be identified among the specimens obtained from three different anatomical locations of the joint. First, the PLM images and profiles for cartilage from the central location showed that the tissue had the arched three-zone fibril architecture, where the surface fibers and the radial fibers had a 90° difference in their angular orientations (represented by the blue and red colors respectively in the image). This observation agrees with the quantitative μMRI data discussed previously in Fig 4.4, and this agreement between the μMRI and PLM data is also consistent with the observations from many mature cartilage tissues from larger animals such as canines (Xia et al., 2001a). With this understanding, one can easily see that the collagen fibers in the

posterior cartilage also have the arched three-zone architecture, except that the tissue is thinner.

Second, as we have observed in the μ MRI T2 anisotropy data (Fig 4.4c), cartilage from the anterior location has a fibril architecture that is far from the arched three-zone structure. This μ MRI observation can be confirmed by the high-resolution PLM data in Fig 4.5. Comparing the 2D images between the anterior location and the central location, it is easy to recognize the continued variation of fibril structure even within the short length of the anterior cartilage block (~ 1.5 mm). From the orientations of the black oval-shaped cell voids in the retardation images, which follow the local orientations of the collagen fibers in the territorial matrix of the tissue, one recognizes the transition from the right end (towards the G side) of this specimen, which still has some similarity to the arched three-zone structure, and to the left end (towards TR side) of this specimen, where most of the deep cells become parallel with the articular surface, representing the parallel architecture of collagen fibers at this location. Comparing the quantitative 1D retardation profiles of all locations (Fig 4.5c), it is evident that the overall retardation values are lower for anterior than central and posterior specimens (weak mutual organization).

Third, among the three locations, the central cartilage was the thickest (550 μ m), about 50% thicker than cartilage at both anterior and posterior locations (250 μ m-300 μ m). Although different in their total thickness, both central and posterior cartilage had approximately the same thickness in their superficial zones (10-15 μ m), and an identical tissue retardance minimum (~ 30 -35 μ m below the articular surface, marked by the arrow in Fig 4.5.c).

4.3.5 Comparison between μ MRI and PLM

Fig 4.6 compares the T2 (0°) profiles from μ MRI and the retardation profiles from PLM, for the four topographical locations on the femoral condyle. In these quantitative profiles, the location of the minimum retardation indicates the location of the least ordered collagen fibrils in the tissue. By comparison, the location of the maximum T2 relaxation at 0° indicates the location of the least-ordered water molecules in the tissue. Among the four locations, only the central cartilage (Fig 4.6c) has a matched correlation between the T2 peak and the retardation minimum, which is consistent with the previous observations in mature cartilage (Xia, Moody, Alhadlaq, et al., 2002). At the A2 location (Fig 4.6b) and in posterior cartilage (Fig 4.6d), the T2 peaks are less well-defined and the peak correlation between μ MRI and PLM is lacking. Cartilage at the A1 location (Fig 4.6a) does not have any recognizable feature that can be used to interpret the arched three-zone fibril structure.

4.3.6 Territorial matrix

Fig 4.7 shows the high-resolution quantitative angle and retardation images (40x objective, 0.25 μ m/pixel) of the cellular clusters at different depths, from the same specimens used before in the 10x imaging (Fig 4.5). Morphologically, one can use quantitative retardation images to view the cellular structure, from which one can easily determine the zonal orientation of the cells and cellular clusters. In the SZ of all three sample locations, all cells had a spindle shape and were oriented in parallel with the articular surface. In addition, the territorial and interterritorial fibers ran parallel to the articular surface (i.e., having the blue color in the angle images). In the central and posterior cartilage, the cellular clusters in TZ were more circular, which implied the lack of a dominant orientation. Furthermore, the cellular clusters in RZ are all oriented

vertically – perpendicular to the articular surface, which can also be seen from its red color in the angle images. In contrast to the central and posterior cartilage, the cellular structures in RZ at the anterior location oriented approximately with the articular surface, which gives the anterior cartilage its lack of arched zonal structure. A higher cellular density can also be noted in the anterior cartilage – in particular, the anterior clusters were fatter than the central and posterior clusters, which implies larger aspect ratios of the oval shapes (aspect ratio = width/length).

Quantitatively, higher retardation values can be found in the proximity of cells and cellular clusters, which indicates the presence of a cocoon-shaped fibril structure surrounding each cell or each group of cells. The same cocoon-shaped fibril structure was also seen from the quantitative angle images when they were enlarged further, in which the angle values change in a cyclic manner, i.e., the same color appears on the opposite side of the same cell or cellular cluster. Since a collagen fiber has neither head nor tail, the same angle value appears on the opposite side of any circular shape, which should have two 90° orientational differences, between the blue and red colors, and between the white and black colors. For the central and posterior cartilage, the background colors in the angle images, i.e., the orientations of the territorial matrix in the tissue, had the expected 90° transition between the surface and deep cartilage, indicating an arched three-zone structure. In contrast, the anterior cartilage (A1) had the same blueish color for all regions of the tissue, indicating an absence of a three-zone structure.

4.4 Discussion

To the best of our knowledge, this is the first quantitative study of the topographical features of the femoral cartilage in a rabbit model by μ MRI and PLM at microscopic resolutions. Compared with the use of larger animals, rabbits are small, inexpensive, and have more societal acceptance. Since rabbit cartilage is thin, any depth-dependent examination requires high resolution in imaging. In recent years, the use of rabbit models in biomedical research has been gradually increasing (Batoool et al., 2020; Esteves et al., 2018; Frenkel et al., 2005; Julkunen et al., 2009; Sah et al., 1997; Wakitani et al., 1998). Still many features of the rabbit tissues, even in healthy control animals, have not been well characterized, which complicates the understanding of degenerative characteristics in tissue.

4.4.1 Topographical structural variations in juvenile joints

A better understanding of the developmental changes from juvenile tissue to mature tissue is critically important since younger populations can develop osteoarthritis several years after traumatic injuries. Studies of the juvenile canine humeral head and ovine distal metacarpus (vanTurnhout et al., 2010; Xia et al., 2003) have revealed several site-specific collagen structures ranging from the arched three-zone structure to complex multi-zone architecture, which can be seen on the same joint surface. Such site-specific collagen structures are absent in mature animals (Xia, Moody, Alhadlaq, et al., 2002). These experimental observations support the notions that there are specific imaging features that are unique in juvenile joints, and that not all locations on the same juvenile joint develop into the three-zone structure simultaneously. A probable explanation for these developmental differences at different locations on the same joint surface could be

either variable growth rates, a pattern of the bone epiphysis, or exposure to different mechanical loading conditions at different anatomical sites of the same joint (Brama et al., 2000b; Heinemeier et al., 2016; Hudelmaier et al., 2001; Julkunen et al., 2010; van Turnhout et al., 2010; Peters et al., 2018; Thonar & Sweet, 1981; Wong et al., 2009).

This study showed that for 3-month-old juvenile rabbits, the cartilage at the central and posterior locations along the femoral medial condyles had already developed into the classic three-zone architecture, which is commonly seen in mature joints. In contrast, the cartilage at the anterior location, the trochlear ridge (TR) of the patellofemoral region of the same joint, had no zonal architecture. This topographical difference could have two causes, developmental and mechanical. The developmental cause is associated with the maturity of the cartilage, where distinct differences between the juvenile and mature cartilage in large animals have been noticed (Xia et al., 2003; Xia, Moody, Alhadlaq, et al., 2002). The mechanical cause is associated with the differences in the force patterns between the anterior location and the central and posterior locations. While the central and posterior cartilage is subject to large compressive loading, the anterior cartilage is subject to constant tangential motion with the patellar cartilage. Hence, the collagen fibers may adapt to the local mechanical environment.

Since MRI T2 anisotropy is extremely sensitive to the dynamic environment of water molecules in biological tissue, it is expected that the fibril architecture and structural variations of the collagen network can be detected non-invasively and non-destructively by MRI, if a sufficient resolution can be achieved (as in μ MRI). At the same time, this study also employed quantitative PLM protocols at the higher optical

resolution, which are uniquely sensitive to the mutual organization and overall orientation of the collagen matrix based on different physical principles. It has been shown previously that the retardance minimum below the articular surface is closely associated with the geometric center of the transitional zone, which is represented by a maximum in quantitative T2 anisotropy of adult animals. The correlation between the two quantitative protocols can define the zonal architectures of the collagen network, from the classic three-zone architecture to more complicated variations in juvenile tissues.

4.4.2. Intervention at an early stage

Understanding the differences between the juvenile and mature cartilage associated with the developmental stages and mechanical environments in individual animals (and humans) can contribute to a better intervention of the cartilage degradation process at an early stage. The collagen network in cartilage is responsible for the structural stability of the tissue, which remodels continuously during growth and adolescence periods. After skeletal maturation, cartilage maintains a stable structure for the rest of the lifespan with only minor adaptations (Heinemeier et al., 2016). Studying the site and age-dependent mechanisms in tissue remodeling provides an opportunity to intervene/repair cartilage optimally at different ages when the damage and degradation occur, which could prevent or postpone advanced stages of osteoarthritis. For example, sports-related injuries, common in young adults, can develop into osteoarthritis within a brief time after the injury. Failure in cartilage repair could happen if the repair tissue does not fit the specific age and topographical dependent features of the damage location.

The development of arcade-like structures with age is an important and interesting phenomenon, the understanding of which would be beneficial to the design of superior

materials and procedures as it would be more stable and better able to effectively handle the loading environment. Therapeutic improvements could potentially be implemented by identifying and investigating the localized physiological and mechanical factors involved at different sites on the same joint. For example, osteochondral allograft transplantations have been used to treat focal lesions in young and active patients. Considering the topographical variation in the fibril architecture can be helpful in dealing with the site morbidity and improve the integration, if the repair grafts can be designed to better match with neighboring/donor cartilage structurally.(Reddy et al., 2007).

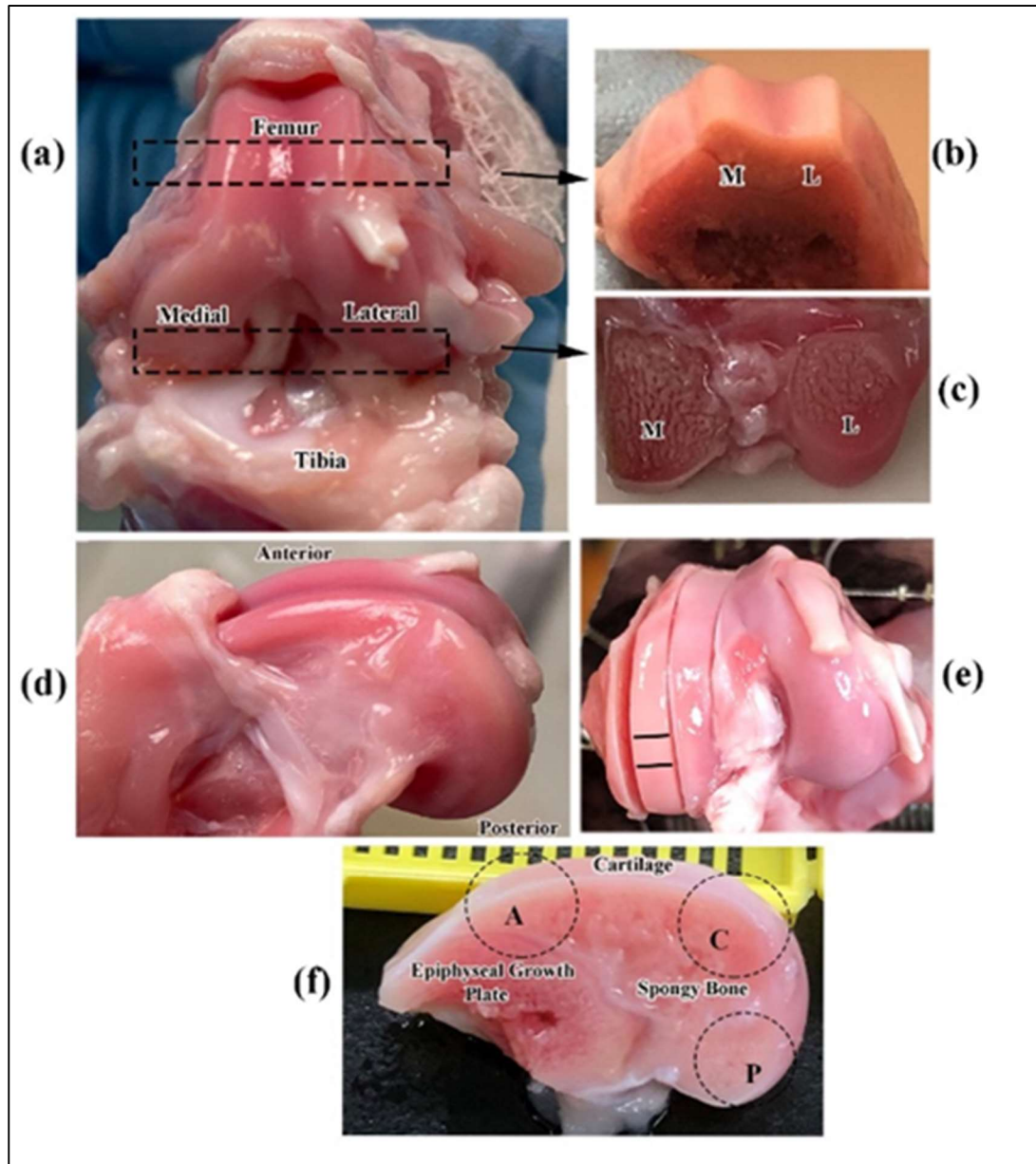
4.4.3. Experimental notes

Structural changes in collagen fibril orientation are closely linked to compositional changes (PG and collagen content) in connective tissue. Due to technical limitations, we did not perform any histochemical analysis to determine any compositional difference in the cartilage at different sites. In addition, since the development of the collagen network is a gradual process, we recognize the potential need for more time points (i.e., different-aged animals) to fully understand the developmental variations in the tissue architecture. The number of μ MRI sampling locations was three, which were 4-6 mm apart from each other along the same small joint. It should also be noted that any topographical variations are highly dependent on the plane/location of histological cutting and the type of joint. Finally, any sample with curved surfaces (e.g., the anterior specimen in Fig 4) cannot be set in any single experiment with the entire surface at a particular orientation. This would also be true for any intact joint in human MRI. Any deviation from the nominal orientation would affect the T2 anisotropy, mainly via the geometric factor in the dipolar spin Hamiltonian (Xia,

1998). Note that in physics, anisotropy is a descriptive term, which refers to the characteristic of a measurement that exhibits different values when measured along with different directions. In MRI, the direction in the sample space is defined by the polarizing magnetic field B_0 . There are various methods to analyze the characteristic of T2 anisotropy. The first and original method analyzes the depth-dependent profiles directly (Xia, 1998) (as Fig 4c of this study). One can also combine these depth-dependent profiles into 2D or pseudo-3D anisotropy maps, either without image interpolation (Xia, Moody, & Alhadlaq, 2002) or with image interpolation (Hänninen et al., 2021). Furthermore, one can calculate the percentage differences in these depth-dependent profiles (Hänninen et al., 2021). In addition, if T2 anisotropy has been measured at many different orientations in the 0° - 360° angular space, one can calculate the relative contents of two orthogonal fibril directions on a pixel-by-pixel basis (Xia, Moody, & Alhadlaq, 2002). In this study, our goal is to characterize the structural variations at different anatomical locations of femoral cartilage in young rabbits. The direct comparisons among the μ MRI and PLM profiles (Fig 6) are sufficient to clearly show the nature of these location-specific anisotropies quantitatively.

Figure 4.1

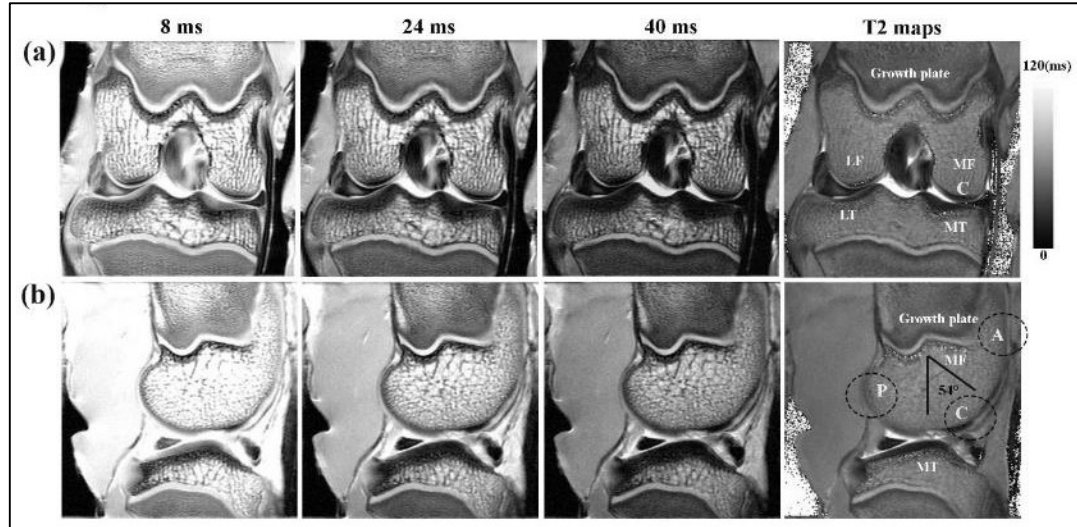
The right knee of a 3-month-old, rabbit, and its histological sections.



Note. (a) Both medial and lateral condyles of a femur are shown. (b) The cross-sectional slice at the patellofemoral location of the femur covers both medial and lateral condyles before its split. (c) The cross-sectional slice at the tibiofemoral location of the femur covers both condyles. (d) Side view of an un-sectioned femur medial condyle. (e) The medial condyle is sectioned into three ~1.6-1.8 mm thick cartilage-bone slices longitudinally. The central slice, which runs up to the unsplit part of the femur, is further cut into small rectangular cartilage-bone specimens ($\sim 1.8 \times 2 \times 2.5 \text{ mm}^3$) for high-resolution μMRI and PLM. Two thin lines mark the location of the central specimen on the central slice. (f) Side view of a central slice of medial condyle with three designated locations (A, C, and P) for μMRI specimens.

Figure 4.2

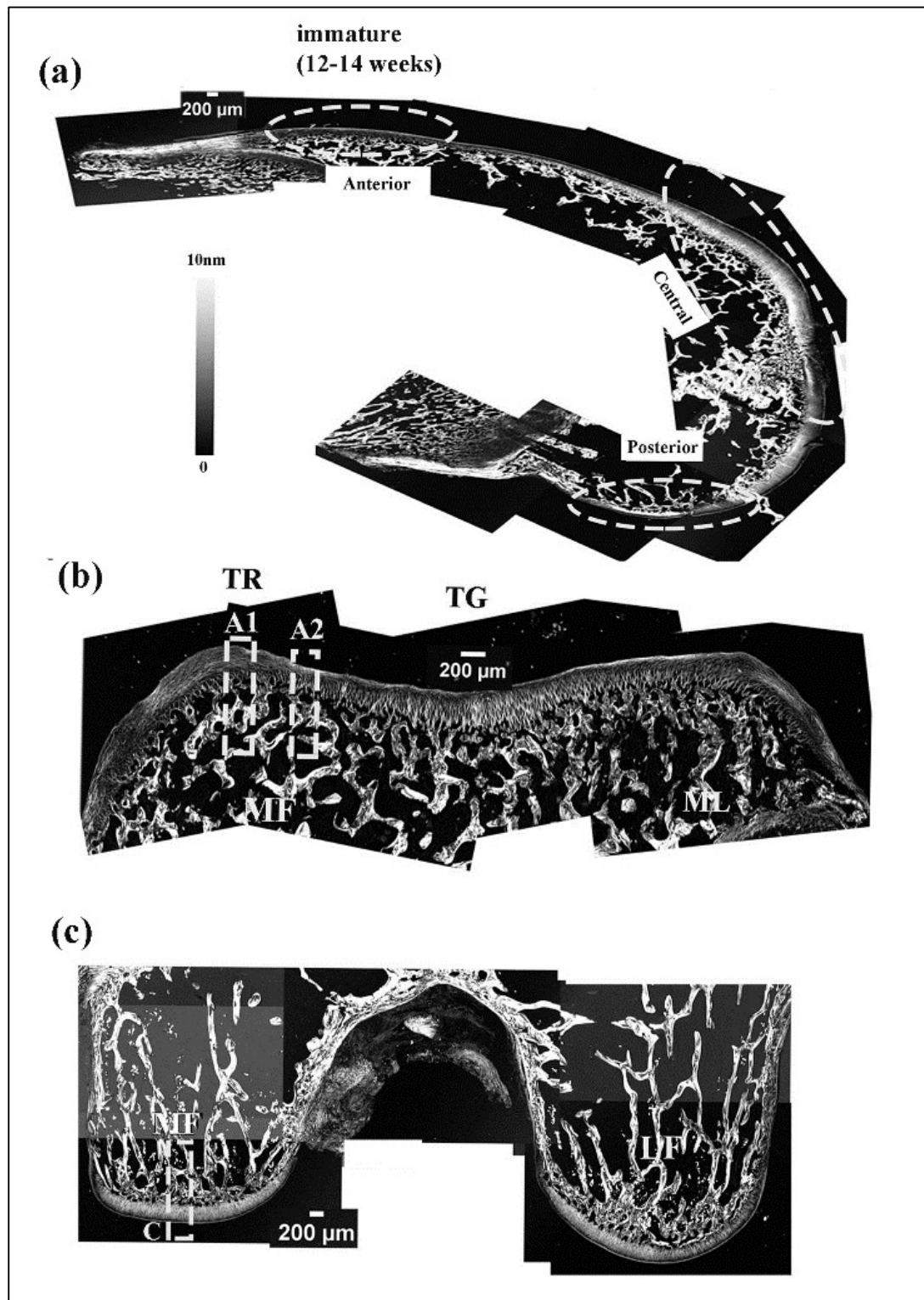
Coronal and sagittal T2-weighted intensity images and corresponding T2 maps



Note. Each set of T2-weighted intensity images had 10 individual images, each at a different echo time. These intensity images were used to calculate one quantitative T2 image. Three designated locations, labeled as central (C), posterior (P), and anterior (A) on the medial condyle, were selected for μ MRI T2 experiments. All images are displayed using the same up/low limits on a greyscale. (MT: medial tibia; MF: medial femur).

Figure 4.3

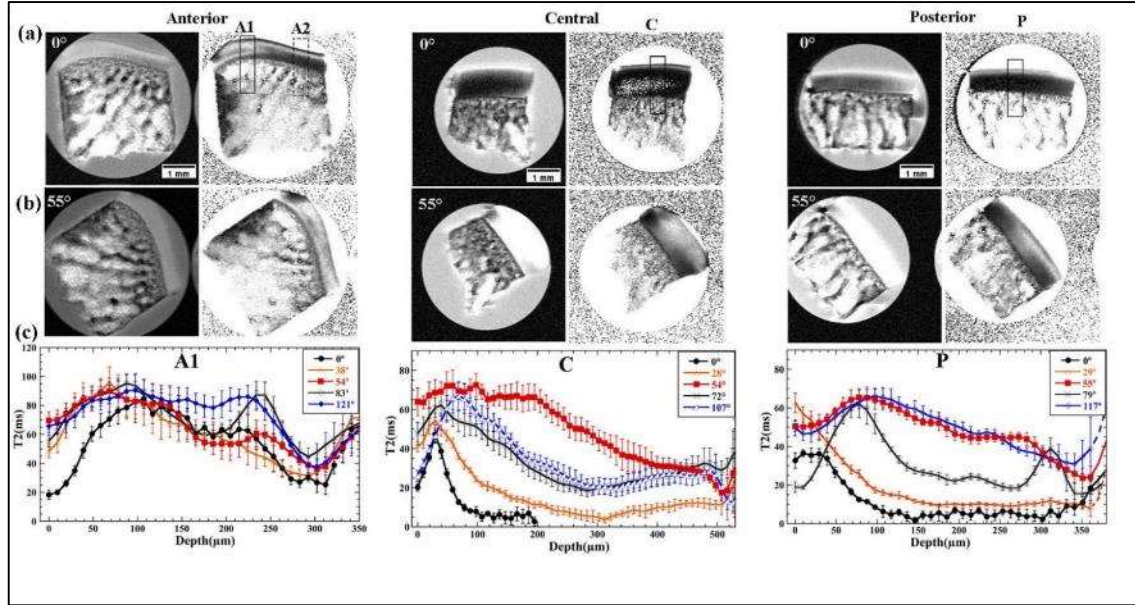
The quantitative retardation images.



Note. (a) The longitudinal slice from a femoral medial condyle (4 $\mu\text{m}/\text{pixel}$) displayed with identical intensity limits (0–10 nm). Three white elliptical shapes show the locations where the small specimens were harvested for μMRI . (b) The retardation images (2 $\mu\text{m}/\text{pixel}$) from a tissue slice that was harvested transversely to the slice in (a), approximately at the anterior location. (c) The retardation images (4 $\mu\text{m}/\text{pixel}$) from a tissue slice that harvested transversely to the slice in (a), approximately at the central location where the femur has two distinct medial and lateral condyles. The transverse planes on these slices are approximately at the same imaging planes as in μMRI . Three dashed rectangular boxes (A1, A2, and C) in b and c are the ROIs selected on the cartilage-bone specimens for quantitative imaging. (MF: medial femoral; ML: medial-lateral; TR: trochlear ridge; TG: trochlear groove).

Figure 4.4

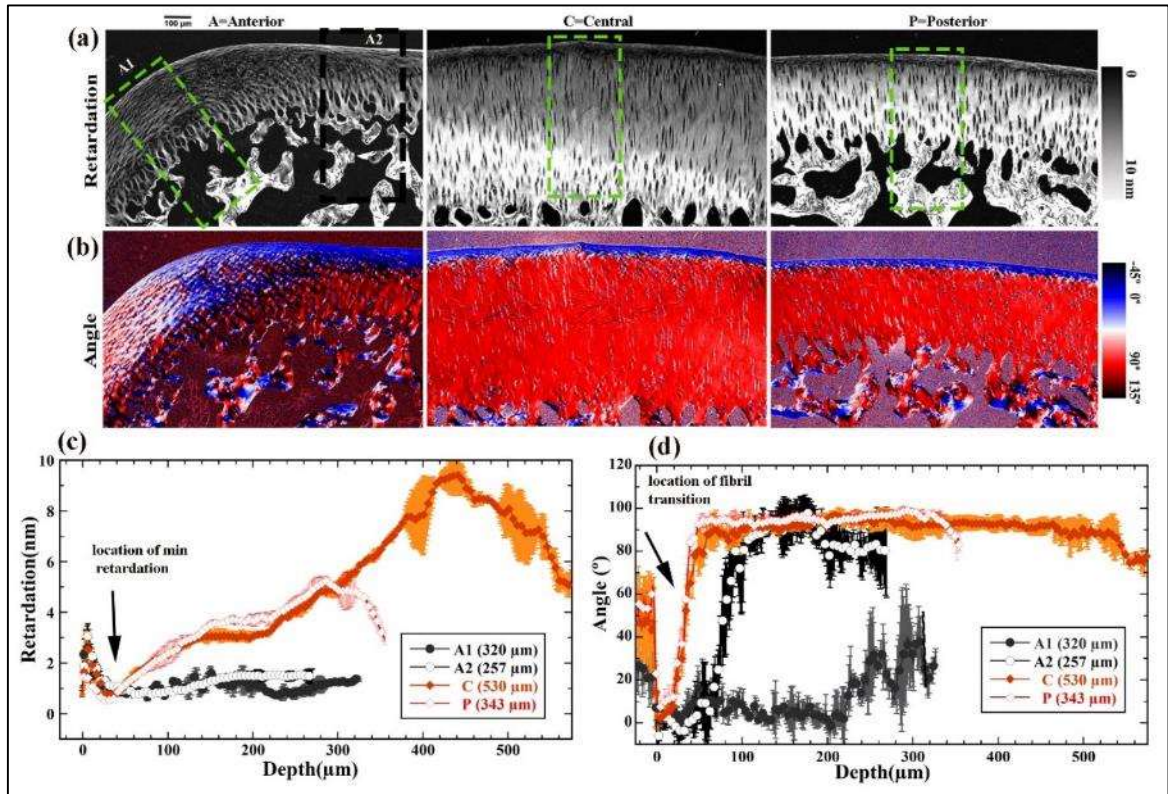
2D μ MRI intensity images (left) and corresponding T2 images (right) of cartilage-bone specimens (A, C, P) at 9.75 μ m/pixel at (a) 0° and (b) 55° and (c) 1D profiles at 5 different orientations.



Note. Each of the three samples was imaged under identical parameters except its orientations in B_0 . Since the anterior specimen had the most curved surface, two ROIs (A1, A2) were selected, where the A1 location is closer to the trochlear ridge (TR) and A2 is towards the trochlear groove (TG). (c) Quantitative T2 profiles from cartilage in ROIs (A1, C, and P), where the standard deviations of the averaged T2 data over the width of the ROI are represented by the vertical error bars.

Figure 4.5

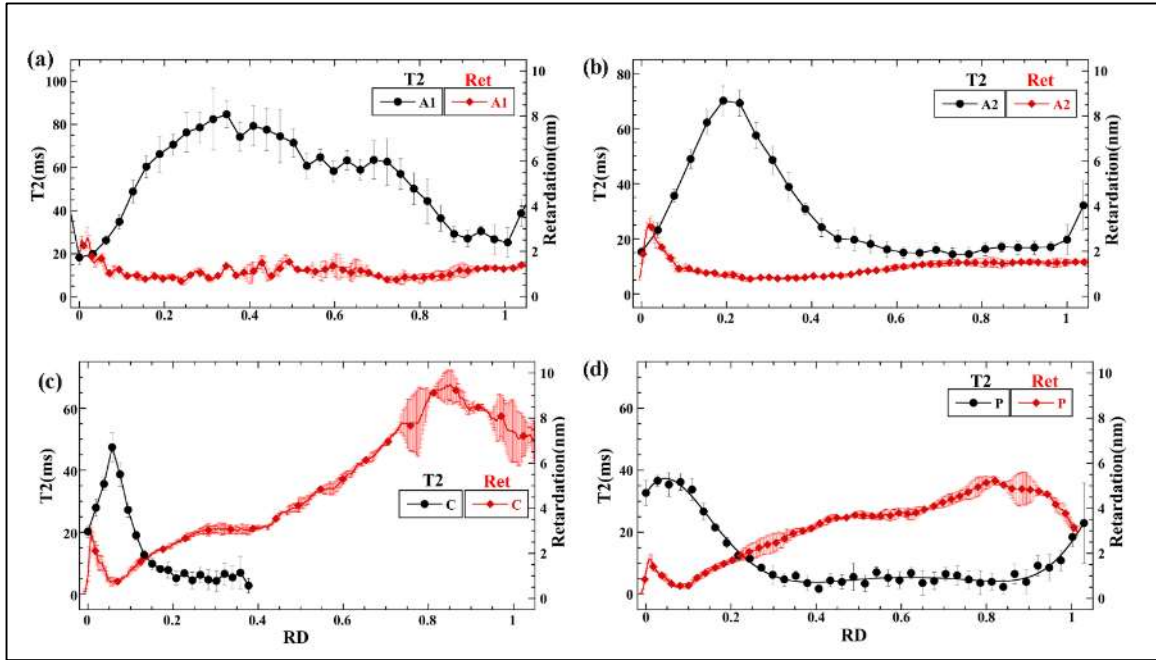
2D quantitative PLM images and 1D quantitative profiles (retardation and angle) at 1- μm pixel resolution



Note. The rectangular boxes in (a) show the locations where the depth-dependent profiles for both retardation and angle were calculated and shown as the profiles in (c) and (d). The quantitative profiles for each location are the mean values (retardation and angle) obtained by averaging three histological sections from each specimen. The vertical error bars are the standard deviations in the averaging. The profiles are plotted for every 5th data point, but the standard deviations are shown for all data points. In the angle images, the blue and red colors defined a 90° orientational difference in collagen fibers. The white color represents 45° in the angle, which can be seen in large regions of the deep tissue in A1 (trochlear ridge). The thickness of the cartilage for each location is listed next to the figure legends in a black rectangular box.

Figure 4.6

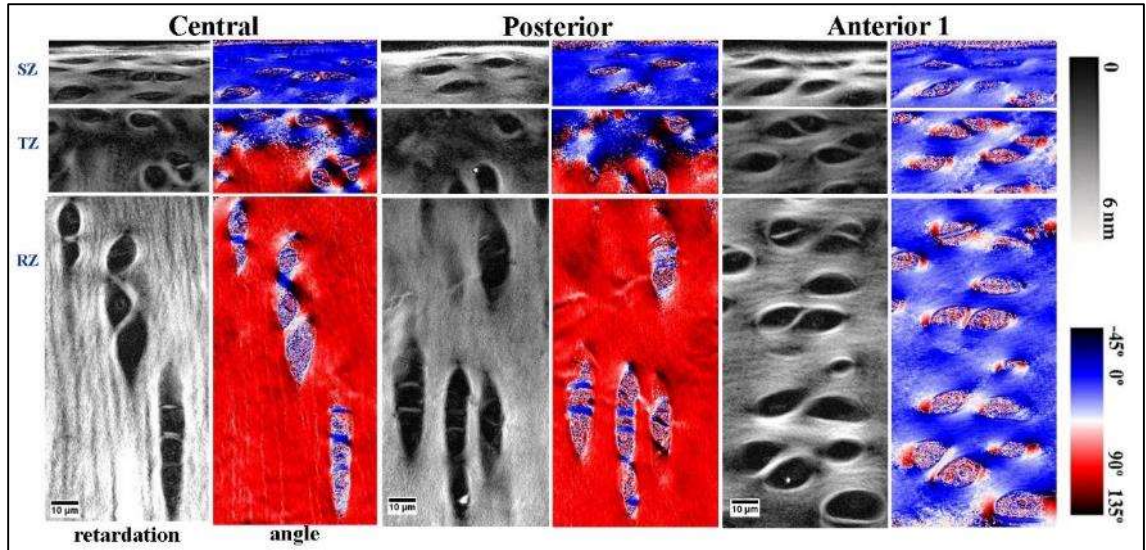
Comparison between the quantitative T2 profiles at 0° and the quantitative retardation profiles, at four different locations in femoral cartilage



Note. 0 in the relative depth marks the articular surface; 1 in the relative depth marks the cartilage-bone interface (CBI).

Figure 4.7

Quantitative PLM images of the three locations of tissue at 0.25 μm pixel resolution at different histological zones.



Note. (a) SZ (superficial zone) images of all three locations at 5% of their relative thickness, (b) TZ (transitional zone) images of all three locations at 10% of their relative thickness, and (c) (RZ) (radial zone) images at 40% of their relative thickness.

CHAPTER FIVE

DEPTH-DEPENDENT STRAIN CALCULATIONS FOR ANISOTROPIC FIBRIL STRUCTURES IN ARTICULAR CARTILAGE

5.1 Introduction

The mechanisms giving cartilage its mechanical resilience involve complex compositional and structural variations across the tissue thickness (depth). The major constituents of its ECM are interstitial fluid (water and ions such as Na^+ , Cl^- , etc.), and macromolecules such as proteoglycans (PGs) and collagens (Sophia Fox et al., 2009). PGs in cartilage are considered unstructured; in contrast, collagen in mature healthy cartilage forms a network with a depth-dependent tri-zonal structure (Fig 5.1b, left). In addition to the depth-dependent structure of the collagen network, the distributions of molecular components (water, proteoglycan, and collagen) are also not homogenous, and all vary with depth from the articular surface towards the cartilage-bone interface (Mittelstaedt & Xia, 2015; Rieppo et al., 2009; Xia, Zheng, et al., 2008). The depth-dependent variations of the molecular components, in conjunction with the macro and microstructural variations, may control its unique depth-dependent and direction-dependent mechanical deformation response to external loading (Chahine et al., 2004; Chen et al., 2001; Korhonen et al., 2008).

In this study, we used a simple 1D mixture model to study the mechanical deformation response of articular cartilage under 1D axial compression (confined, equilibrium). The model included the structural inhomogeneity of collagen fibers and inhomogeneous distribution of FCD to calculate intra-tissue strains and displacements in

the tissue. This model highlighted the role of collagen fibril mechanics in conjunction with the osmotic swelling pressure of articular cartilage. The 1D model was computational and solved in MATLAB using finite difference methods (Provided in Appendix B). The fibril stiffness was introduced as an anisotropic modulus that varies with fibril orientation through depth. We assumed that collagen fibers were stiffer to stretch parallel to their length (in the RZ) than perpendicular to them (in the SZ). The intrinsic structural inhomogeneity of collagen fibers and inhomogeneous distribution of FCD caused an anisotropic pre-strain in the tissue and increased the tissue resistance against axial loading, specifically in the radial zone. The simulation results (intra-tissue strains and displacements) were applied to different scenarios (mature, immature, healthy, and degraded cartilage) and qualitatively explain many experimental findings. In particular, we examined the relative importance of collagen fibril intrinsic stiffness towards osmotic swelling contributing to the response of cartilage to compression. We hypothesized that the mechanical compressive properties of cartilage were not only dependent upon FCD but also the structured pre-strained collagen fibril network can counterbalance the mechanical compression. The knowledge of the structure and mechanical properties of ECM surrounding the cells is vital not only for understanding but also designing novel engineering scaffolds and tissues that possess normal mechanical functionality.

5.2 Materials and methods

The fundamental assumptions used in this study are described in Fig 5.1a and 5.1b. Cartilage is assumed to be compressible and permeable (water flows in and out). Initially, the unstretched and unloaded tissue (stress $\tau = 0$) has cross-sectional area A ,

length L , and FCD concentration C_c (Fig. 5.1a, left). We model the collagen as a series of linear springs with intrinsic stiffness denoted by a mechanical modulus (μ). Due to osmotic swelling pressure caused by FCD, the tissue swells causing a pre-strain ε_{pre} to the collagen fibers and the effective concentration of FCD drops from its unstrained value C_c (assuming the number of molecules remains the same but the volume changes) as tissue is diluted by additional water, giving a strain-dependent concentration.

$$C = \frac{C_c}{(1 + \varepsilon_{pre})} \quad (5.1)$$

During the pre-strain condition of the collagen fibers (without any load applied), let u_{pre} be the distance it displaces in the radial direction, so strain $= \varepsilon_{pre} = \frac{u_{pre}}{L}$ (Fig 5.1a, middle). When tissue is loaded under uniaxial compression ($\tau \neq 0$), let u be the distance it displaces from the initial unloaded state. The resultant strain will be $\varepsilon = \frac{u}{L}$ (Fig 5.1a, right). The observed strain, e , is the net difference between displacements (u_{pre} and u) over the pre-compression length ($L + u_{pre}$). Tissue with length $L = 1$ mm under uniaxial load (confined compression) is used to model intra-tissue strains and displacements at equilibrium, when the osmotic swelling stress in cartilage is balanced by the elastic stress exerted by a pre-stressed collagen network (Basser et al., 1998). In that case, the total stress (τ) in the tissue can be represented as the osmotic stress (τ_{FCD}) minus the elastic stress of the collagen network (τ_{CN})(Basser et al., 1998),

$$\tau = \tau_{FCD} - \tau_{CN} \quad (5.2)$$

$$\tau = CkT - \mu\varepsilon_{pre} \quad (5.3)$$

For simplicity, a 1D analysis of cartilage occurs in confined compression. The total normal stress (perpendicular to the articular surface) can be modeled as Eq. (5.1), where, $\tau_{FCD} = CkT$ is an osmotic term that depends on the concentration of FCD (C), the Boltzmann constant k , and the absolute temperature T , and $\tau_{CN} = \mu\varepsilon_{pre}$ is the linear elastic stress exerted by the collagen fibers.

When pre-stressed tissue is under zero external stress ($\tau = 0$), Eq. (5.3) can be written as

$$\frac{C_c kT}{\mu} = \varepsilon_{pre}(1 + \varepsilon_{pre}) \quad (5.4)$$

Solving Eq. (5.4) quadratically gives the pre-strain as

$$\varepsilon_{pre} = \frac{-1 + \sqrt{1 + \frac{4C_c kT}{\mu}}}{2} \quad (5.5)$$

When tissue is stressed under uniaxial compression ($\tau \neq 0$), the observed strain, e , is

$$e = \frac{u_{pre} - u}{(L + u_{pre})}, \quad (5.6)$$

$$e = \frac{\varepsilon_{pre} - \varepsilon}{(1 + \varepsilon_{pre})}. \quad (5.7)$$

From Eq. (5.2), total stress can be written as

$$\tau = \frac{C_c}{(1 + \varepsilon)} kT - \mu\varepsilon \quad (5.8)$$

Rearranging Eq. (5.8) and using Eq. (5.4) gives

$$\frac{\tau}{\mu} = \frac{\varepsilon_{pre}(1 + \varepsilon_{pre})}{(1 + \varepsilon)} - \varepsilon. \quad (5.9)$$

Solving Eq. (5.9) for ε using the quadratic equation gives

$$\varepsilon = \frac{-\left(1 + \frac{\tau}{\mu}\right) + \sqrt{\left(1 + \frac{\tau}{\mu}\right)^2 + 4(\varepsilon_{pre}^2 + \varepsilon_{pre})}}{2} \quad (5.10)$$

The observed strain can be calculated using Eqs. (5.5), (5.7), and (5.10)

$$e = \left(\frac{\varepsilon_{pre}}{1 + \varepsilon_{pre}} \right) + \left\{ \frac{\left(1 + \frac{\tau}{\mu}\right) - \left(\sqrt{\left(1 - \frac{\tau}{\mu}\right)^2 + 4(\varepsilon_{pre}^2 + \varepsilon_{pre})} \right)}{2(1 + \varepsilon_{pre})} \right\} . \quad (5.11)$$

Using approximation $\tau \ll \mu$, Eq. (5.11) can be written as,

$$\tau = \mu(1 + 2\varepsilon_{pre})e , \quad (5.12)$$

so, the effective modulus, μ_{eff} , is

$$\mu_{eff} = \mu(1 + 2\varepsilon_{pre}) \quad (5.13)$$

During compression and under small strains, the effective modulus of the tissue is governed by the contributions from its intrinsic stiffness μ and the ratio $\frac{C_c k T}{\mu}$. The fibril pre-strain ε_{pre} — which depends on the FCD (C_c) and the intrinsic stiffness of collagen network (μ), both of which vary with depth (Fig 5.1b, middle and right) — can also be non-uniform and depth-dependent and depends on variations of these two parameters (Eq. (5.5)). The observed strain (e) under applied load can be calculated using Eq. (5.11) considering uniaxial compression (confined, constant load, equilibrium) and will be non-uniform along the depth.

5.2.1 Depth-wise variation in fiber orientation

Non-calcified, healthy, mature cartilage is commonly subdivided into three structural zones (SZ, TZ, RZ), where the lower/deeper end of the radial zone is anchored to the subchondral bone. The collagen fibers in different zones have preferred orientations, as shown in Fig 5.1b (left). The tissue is divided into three zones based on the depth-dependent angle profile θ of the collagen fibers (Xia et al. 2001)

$$\theta(y) = \frac{\pi}{4} \left[\tanh\left(\frac{y - m1}{m2}\right) + 1 \right]. \quad (5.14)$$

To represent a typical tri-zonal architecture of mature cartilage of depth, $L = 1$ mm, we take $m1 = 0.3$ mm and $m2 = 0.09$ mm, where θ is the angle between the fiber direction and the plane of the articular surface ($\theta \sim 0$ in SZ and $\theta \sim \pi/2$ in RZ), y refers to the depth of tissue with zero at the articular surface, $m1$ refers the center of TZ or the location of most random fibers, and $m2$ determines the rate at which fibers are changing their orientation. The thickness of each zone can be determined using Eq. (5.13) (Xia et al. 2001).

5.2.2 Depth-wise variation in FCD and μ

The depth-wise increase in FCD (Fig 5.1b) is incorporated into the model using

$$C_c(y) = C_0 \exp\left(\frac{y}{r}\right) \quad (5.15)$$

where C_0 is the concentration of FCD near-surface ($y = 0$) and r is the rate at which concentration varies with the depth. The typical value of FCD ranges from 50–250 mM for human cartilage (Chen et al., 2001b). The calculation of FCD used in the simulations is based on a depth-dependent GAG concentration (Mittelstaedt & Xia, 2015; Xia, Zheng, et al., 2008) for mature canine cartilage, assuming 2 moles of negative charge per mole of chondroitin sulfate (one sulfate and one carboxylate), which has a molecular weight of 502.5 g/mole. These values (159–398 mM) lie within the range reported in the literature, as shown in Fig 5.1b (right). The concentration of FCD, which depends on PG concentration, is not only depth-dependent but also varies among species, types, and locations in the joint and it has also been reported to change during growth and

maturation (Julkunen et al., 2009). Therefore, the choice of C_0 and r would be sample-specific to fit the simulation results.

Another assumption in our model was that the intrinsic stiffness of the collagen network (denoted by modulus μ) in articular cartilage is anisotropic and depth-dependent (it varies with fiber orientation along its depth) as shown in Fig 5.1b (middle). We assume the collagen fibers are stiffer parallel to their length (in the RZ) than perpendicular to it (in the SZ). The stress-strain relationship in anisotropic tissue has been derived previously (Wijesinghe & Roth, 2019). For the 1D case, the collagen network modulus can be written as

$$\mu(y) = \mu_0 + \mu_1 \sin^4 \theta \quad (5.16)$$

where μ_0 is the modulus of the collagen network near the surface in the SZ ($\theta = 0$) and μ_1 is the additional contribution to the modulus in the RZ ($\theta = \pi/2$). Since there were no independent values reported in the literature for collagen network intrinsic stiffness, the values we have used (0.4 MPa for SZ, 0.6 MPa for RZ) are comparable to the values of solid matrix stiffness of immature bovine cartilage from carpometacarpal joints (Wang et al. 2002). The corresponding μ_0 and μ_1 are

$$\mu_0 = 0.4 \times 10^6 \quad (\text{Pa}) \quad (5.17)$$

$$\mu_1 = 0.08 \times 10^6 \exp\left(\frac{y}{0.00099}\right) \quad (\text{Pa}) \quad (5.18)$$

In general, when an external force deforms an elastic body, the resistance to deformation is called its stiffness. It could be a function of material properties, material orientation, geometric dimensions, loading directions, type of constraint, and choice of spatial region, where loads and constraints would be applied. In the case of cartilage,

when loaded or stretched in tension, resistance to tensile deformations is offered by the intrinsic stiffness of the collagen fibers (Wijesinghe & Roth, 2019) which we assume in our model is anisotropic and increases with depth. The pre-strain ε_{pre} , which depends on the FCD (C_c) and the intrinsic stiffness of collagen network μ (Eq. (5.5)), both of which varied with depth (Eqs. (5.15) and (5.16)), would be non-uniform and depth dependent. Eq. (5.11) implies that the observed stress-strain relationship is nonlinear when the cartilage is compressed, even if the collagen stress-strain relationship is linear (defined in the text under Eq. (5.2)).

To appreciate the relative influence of collagen and proteoglycan, assume $\varepsilon_{pre} < 1$ and $\tau \ll \mu$. The effective modulus can be obtained by simplifying Eqs. (5.11) and (5.12)

$$\mu_{eff} = \mu + 2C_c kT \quad (5.19)$$

which is the sum of a term relating to the intrinsic stiffness of the collagen network and a term relating to the contribution of FCD (twice the osmotic term). As the stress becomes large, the observed strain e in Eq. (5.11) becomes independent of stress. If τ becomes larger than $\mu\varepsilon_{pre}(1 + \varepsilon_{pre})$, the collagen network alone cannot support compressive stress and our model no longer holds (loss of pre-stretch corresponds to $\varepsilon = 0$).

5.3 Results

5.3.1 Nonlinear strain (e vs τ relationship)

Fig 5.2 shows the e versus τ relation for $\varepsilon_{pre} = 1$. For small, applied stresses, the stress vs strain relation is linear, but with increased stress, it tends to be nonlinear even

though the elastic stress-strain relationship was linear in the model. If $\tau \gg 2\mu$, the collagen loses pre-stretch ($\varepsilon = 0$).

5.3.2 Effect of variations of FCD and μ on strain

The observed strain (Eq. (5.11)) was simulated in Fig 5.3, for the given ranges of FCD and μ as two independent parameters. When both FCD and μ are small the predicted strains are large (a desirable trend for SZ of compressed cartilage), while for larger FCD and μ the strains appear smaller (RZ trend).

5.3.3 Depth-dependent strain

The simulated tissue strain is obtained by introducing depth-dependent parameters FCD and μ (Eq. (5.15) and Eq. (5.16)) considering mature healthy cartilage with $L = 1$ mm, $m1 = 0.3$ mm and $m2 = 0.09$ mm. The FCD near the surface ($C_0 = 159$ mM) was obtained from experimental studies (Mittelstaedt and Xia 2015; Xia, Zheng, and Bidthanapally 2008), and varied with depth ($r = 0.9$ mm in Eq. (5.15)). The values of FCD used in this simulations are based upon the depth-varying GAG concentration (159 mM at the surface to 398 mM at radial zone) reported for mature canine humeral cartilage obtained from different experimental studies (Mittelstaedt and Xia 2015; Xia, Zheng, and Bidthanapally 2008; Zheng et al. 2009) (μ MRI, μ CT, and biochemical); all of them provided similar values and are strongly correlated. For preliminary analysis, the intrinsic modulus of the collagen fibers near-surface was 0.4 MPa, which increases to 0.6 MPa in deep regions as described above (Eqs. (5.16-5.18)). The simulated intra-tissue strain decreases non-homogenously through the depth ($\tau = 0.1$ MPa) as shown in Fig 5. Positive stress or observed strain corresponds to compression in our model. The tissue strain is highest near the articular surface and

decreases (~50%) from the SZ to RZ with the middle ‘kink’ occurring in the transitional zone at a depth between 0.2–0.4 mm.

5.3.4 Force vs intra-tissue displacement relation.

The strains averaged over the thickness of each zone (as determined by Eq. (13)) are represented as zonal strains. The intra-tissue displacement for each zone of tissue was obtained by integrating the depth-varying observed strain ‘ e ’ along the depth and then averaging the displacement results over the thickness of each zone. In general, the RZ has the least mean intra-tissue strain and displacement compared to the SZ and TZ, consistent with what is observed experimentally (Chen et al., 2001; Kahn et al., 2015).

Fig 5.5 represents the relationship between applied force (stress) and simulated intra-tissue displacements for three zones (SZ, TZ, RZ) of a thin rectangular sample tissue, compressed (surface-to-surface) by increasing stress (τ) in discrete steps (stress relaxation). The applied stress (τ) can be converted into force (F) as $F = \tau \times A$, where A is the surface area ($A = \text{thickness} \times \text{width}$) in contact with the loading platen (0.25 mm^2). The thickness of tissue is assumed to be $\sim 120 \text{ }\mu\text{m}$, the width is $\sim 2 \text{ mm}$, and it is compressed with a thin glass platen (thickness $\sim 100 \text{ }\mu\text{m}$). The thickness and width are in the x and z directions and y refers to the depth of tissue i.e., direction perpendicular to cartilage surface. These simulated zonal relationships between force and displacement agree with the known mechanical response of cartilage under compression and are comparable to the experimental measurements of the intra-tissue displacements in mature canine humeral articular cartilage (Kahn et al., 2015) (linear model, 10–20% error). The total depth of tissue used in the simulation is 0.50 mm , compared to a cartilage depth of 0.55 mm mentioned in the experimental study (Chen et al., 2001; Kahn et al., 2015). The

rest of the parameters are the same as described above to simulate depth-dependent strain for mature tri-zonal cartilage.

5.3.5 Effective modulus for cartilage zones (stress vs strain)

Fig 5.6a presents the stress vs strain relation for three zones for the selected parameters (FCD, μ , $m1$, $m2$) as described earlier (Fig 5.4, Fig 5.5). The mean strain appears to increase linearly with increasing stress for all three zones. The slope of linear fit represents the effective modulus of each zone. The modulus increases with depth, from ~ 1 MPa in SZ to 1.6 MPa in RZ.

With the adjustment of μ (at the deep region, RZ) in the range between 1–1.5 MPa instead of 0.6 MPa, the simulation produced similar depth-dependent intra-tissue displacements and intra-tissue strains when compared with the data reported for human femoral head cartilage (Chen et al., 2001) ($\tau = 0.01 - 0.4$ MPa). The value of μ was chosen by trial and error. The FCD used in the simulation is the same as mentioned before ($C_0 = 159$ mM, $r = 0.9$ mm), since the FCD range reported in the study by Chen et al. (2001) lies in the range reported by several other studies (Mittelstaedt & Xia, 2015; Xia, Zheng, et al., 2008; Zheng et al., 2009).

The loss of PG can cause a loss in FCD which can be attained through different enzymatic digestion (e.g., trypsin) and is the characteristic feature of degraded tissue. As a result, tissue can experience an abnormal increase in intra-tissue strains and displacements for a given otherwise-normal stress. Fig 5.6b presents the stress vs zonal strain relation for three zones of PG depleted or degraded tissue, with μ the same as normal (0.4 – 0.6 MPa), assuming no change in intrinsic properties of collagen fibers. The maximum depth-wise loss in FCD was introduced in the model using Eq. (5.15) and

adjustment of C ($C_0 = 1.59$ mM) which results in a decrease of pre-strain and a considerable increase in observable strain e for all three zones. The rest of the parameters are the same unless otherwise stated. The effective modulus (linear fit of stress vs zonal strain) decreases ~ 50 % for each zone as compared to normal native tissue (Fig 5.6a) and is consistent with experimental findings; selective enzymatic digestion of PGs can cause a significant drop in elastic compressive modulus of articular cartilage (Broom & Silyn-Roberts, 1990; Lyyra et al., 1999).

5.3.6 Strains between homogeneous and inhomogeneous tissues

Fig 5.7 presents several intra-tissue strains simulated for a given stress ($\tau = 0.1$ MPa) under different depth-varying conditions, including FCD and μ , between homogenous (e.g., neonatal cartilage) and inhomogeneous (e.g., healthy, mature) cartilage, $L = 1$ mm. For comparison, a constant FCD ($C \sim C_0 = 159$ mM, $r \rightarrow \infty$) and a constant intrinsic modulus of collagen network ($\mu \sim \mu_0 = 0.4$ MPa, $m1 = 0$, $m2 \rightarrow \infty$) were used for the homogenous case (no depth variations) keeping the rest of parameters the same as before. This condition of constant intrinsic modulus of the collagen network can be attained in neonatal cartilage (Julkunen et al., 2009) or nasal cartilage (Xia et al. 2012) with a uniform fibril organization where most fibers align parallel to an external surface i.e., $\theta = 0$. The inhomogeneous case of both FCD and μ used the same parameters as to simulate depth-dependent strain in Fig 5.4. By comparison, the intra-tissue strain in articular cartilage decreases significantly with depth in the middle and deep zones when it has non-homogenous distributions of both FCD and μ (mature tissue).

5.4 Discussion

Articular cartilage has unique depth-dependent structural and material anisotropy. A better understanding of anisotropy in biological tissues is important for predicting material properties and designing novel engineering structures. Considering the tissue structural anisotropy, the effect of the force loading direction can be explored through the examination of its structural and material characteristics in the loading direction, related mostly to the anisotropic structure of the collagen network in cartilage.

The role of collagen fibers in articular cartilage under compression has been studied extensively (Alhadlaq et al., 2007b; Kääb et al., 1998; Moger et al., 2009; Wilson et al., 2007) and depth-dependent deformation and fibril reorientation responses have been reported (Alhadlaq et al., 2007a; Chen et al., 2001; Kääb et al., 1998). In mature healthy cartilage, the RZ exhibits stronger resistance to compression than the SZ and TZ. Various theoretical models (Ateshian et al., 2004; Julkunen et al., 2013; Schwartz et al., 1994; Smith et al., 2016) (e.g.; isotropic elastic, isotropic biphasic or triphasic, fibril-reinforced poroelastic, etc.) have been developed to explain the depth-dependent compressive response of articular cartilage.

In this study, we have presented the simulation results for 1D loading (normal to the articular surface) only. The model uses the structural anisotropy of the fibril network to explain the depth-dependent mechanical compressive properties of the tissue. We intend to explore the role of the zonal structure of pre-stressed collagen fibers in the loading response of cartilage and provide some insights to design advanced anisotropic fibril-reinforced models of cartilage. We can examine the nonlinear strain response of articular cartilage. Our model emphasizes the structural arrangement of the collagen network that

influences the compressive nature of cartilage. A critical assumption in our model is that collagen fibers are stiffer along their longitudinal axis (RZ) than perpendicular to their axis (SZ), as illustrated in Fig 5.1c. This assumption is introduced into our modeling as a fibril modulus that varies with fiber orientation and depth (Fig 5.1b middle). When combined with a depth-wise increase in proteoglycan content (represented by the FCD; Fig 5.1b right), the collagen fibers in RZ appear to be more resistant (minimal observed strain) to uniaxial compression than SZ and TZ (maximum observed strain). Although the exact role of fibrillar mechanics in the collagen network is not clear, an increase in the fibrillar pre-strain may be associated with increasing fibrillar D-period (nanoscale structure) with depth (Inamdar et al., 2019), which in turn may be controlled by the known depth-wise variation in proteoglycan content and subsequent swelling pressure.

The fibril pre-strain in our model depends on the intrinsic stiffness μ of collagen fibers (anisotropic; higher longitudinal than transverse) and FCD. Since FCD varies with cartilage depth in addition to fibril orientation, our simulated fibril pre-strain was higher in RZ (FCD and μ higher) than SZ (FCD and μ lower). The effective modulus for SZ qualitatively agrees with experimental results (Kahn et al., 2015), while the RZ modulus (1.6 MPa) is significantly smaller than what the experimental study found (~6 MPa). We hypothesize that this difference arises because of the choice of elastic stiffness (μ) for RZ fibers (it could be higher than 0.6 MPa) and not the FCD since the FCD can be measured more easily and accurately. This hypothesis is supported by Chen et al. (Chen et al., 2001), who highlight the role of other factors (unknown) besides FCD at different depths of tissue to explain the marked increase in confined compression modulus of human femoral head articular cartilage in the deep radial zone (RZ).

When this pre-stressed structure is compressed perpendicular to the articular surface, the simulated intra-tissue displacement and observed strain decreased with depth below the articular surface (minimal observed strains appear for RZ). The simulation results are in agreement with several macro-scale experimental findings of higher compressive modulus in RZ than SZ (Kahn et al., 2015; Schinagl et al., 1996; C. Wang et al., 2002; C. Wang et al., 2003) in articular cartilage, except for some which reported the opposite trend (McLeod et al., 2013). We speculate that the model prediction of higher compressive modulus in RZ (smaller strains, strain hardening) can be explained if we assume that the intrinsic stiffness of collagen fibrils reaches a maximum when tissue is mature (closure of growth plate). If tissue is immature (growth plate open), we hypothesize that the intrinsic stiffness of collagen fibers in RZ could be smaller than in mature tissue even if both have radial fibers in the RZ. A recent study has quantified the structural differences between the immature and mature articular cartilage of rabbits by using microscopic MRI and polarized light microscopy (Mantebea et al., 2022) and we speculate that in the case of cartilage, the intrinsic stiffness of collagen fibers arises not only because of their orientation but also because of the presence of prototypic fibrils that can fuse or interlace into thicker fibrils or cross-linking, particularly in the RZ (Depalle et al., 2015; Gottardi et al., 2016; McLeod et al., 2013), which increase as tissue matures. This inter-fibril organization stabilizes the collagen assembly by the intramolecular and intermolecular linkages. This hypothesis can be tested by comparison of compressive properties of cartilage at different stages of growth and maturation, which involves structural adaptations including collagen fiber architecture.

A polarized Raman spectroscopy study combined with the nanoindentation technique (Bergholt et al., 2016) reported a marked increase in the elastic modulus in the deeper part of cartilage tissue and no correlation between composition and local mechanical modulus, which emphasized the contribution of ECM microstructural anisotropy to the tissue oppressiveness. We suggest that the local mechanical modulus of the anisotropic articular cartilage should not only depend upon the concentration of macromolecules (PG, collagen) but also the intrinsic structural properties of the extracellular matrix (ECM microstructure), including cross-linking, a shift in fiber orientation, mutual organization, etc. of the collagen fibers, which all influence fibril pre-strain and consequently the compressive stiffness.

In native mature cartilage, fibrils are pre-strained by the proteoglycans almost to their maximum limit, while this pre-strain is lost upon enzymatic digestion (e.g., trypsin, chondroitinase ABC) (Inamdar et al., 2019). This loss can lead to a significant decrease in tissue compressive modulus (Hunziker et al., 1997), but it has also been reported that PG depletion should not affect some intrinsic material properties of collagen fibers (e.g. cohesive strength, orientation, fiber-to-fiber interaction, length, width, etc.) which depend upon the structural linkages within the fibril meshwork itself (Hunziker et al., 1997) and could be a maturation-related phenomenon (Stammers et al., 2020). So, when simulating the observed strains in PG-depleted cartilage (Fig 5.6b), we assumed that loss of PG should not change the intrinsic stiffness of collagen fibers. The loss in fibril pre-strain in our case is due to the loss of FCD (PG) not the μ .

Our model can be used to estimate the mechanical response of both mature and immature articular cartilage if the selection of parameters is made accordingly. The actual

developmental changes of articular cartilage in animal growth involve complex mechanical and biochemical changes dependent upon compositional and structural changes of the extracellular matrix macromolecules including PGs and collagen fibers. We speculate that the intra-tissue strains and displacements during uniaxial compressive loading could be higher in the deep zone of immature or neonatal tissue, which is devoid of radial fibers and possesses approximately a non-structural organization with most fibers parallel to the articular surface. So, structurally a homogeneous tissue (neonatal) with no change in intrinsic stiffness of collagen fibers (0.4 MPa) through depth and with no change or increase in FCD to depth (Fig 5.7) will have a smaller compressive modulus than structurally inhomogeneous tissue (mature), which is consistent with Gannon et al.'s (2015) findings. In general, it is useful to study the local strains in the tissue as they may regulate cell function and metabolism (Torzilli et al., 2006).

5.4.1 Assumptions and Limitations

Applied stress in our model is global stress and the slope of a linear fit to stress vs strain does not represent a real quantification of zonal modulus but gives qualitative information about the local modulus based on the contribution of FCD and collagen fibril pre-stress. In addition, the relationship between the stress and the mean zonal strain appears linear within the model limits (smaller strains) and our choice of parameters. If a nonlinear relationship exists between stress and strain in the tissue, the predicted zonal modulus will change.

The compositional and ultrastructural features of cartilage lead to inhomogeneity and anisotropy in mechanical properties; sometimes it is difficult to distinguish the independent effect of one rather than the other. The functional properties of cartilage

depend on its non-homogeneous extracellular matrix molecular composition and structure. Besides normal depth-wise variations in composition and structure, the proteoglycan and collagen vary significantly in content during growth and maturation along with strong structural adaptations. Currently, the clear answer to the question of what component of collagen fibers determines the intrinsic stiffness (resistance to pre-stretch in case of osmotic swelling i.e., in pre-strain condition) and how it varies along with the depth is not clear. In general, the intrinsic properties of the collagen network can depend on its collagen content, type, size/diameters of collagen fibers, crosslinking, fiber-to-fiber interaction, fiber orientation, and strength of macromolecular interactions modulated by the collagen network. It should ideally be independent of GAG/FCD concentration but can be affected by the interactions involving the collagen network and PG together.

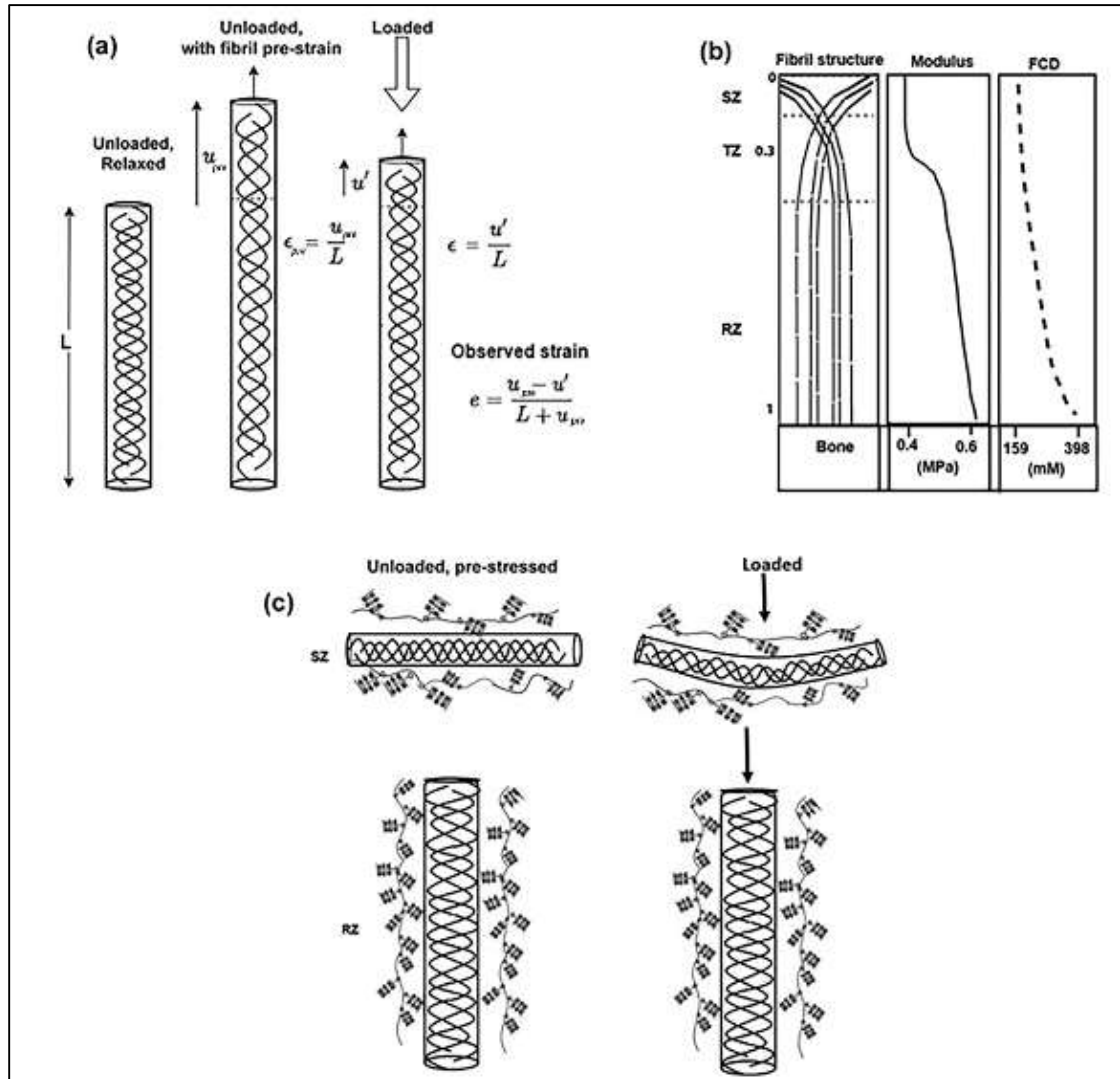
It has also been reported that articular cartilage obtained from different joints of the same animal can be different in its composition and similarly, the cartilage obtained from weight-bearing and non-weight-bearing regions of the same joint can offer different stiffness under the same loading conditions. Therefore, for the theoretical analysis and comparisons of the compressive behavior under the same loading conditions, the choice of parameters including collagen network stiffness depends upon the sample and its conditions under consideration, which requires sophisticated and careful measurements of biochemical and biomechanical gradients.

Modern models are complex and require several parameters to describe the overall mechanical behavior. With a great number of tissue components and material parameters, the models become prone to uncertainties caused by the assumptions, for

example, in the interactions between the model components. Although the model may be plausible in theory, these uncertainties need to be addressed and the model implementation and function should be verified before the model can be applied.

Figure 5.1

Schematics of the modeling



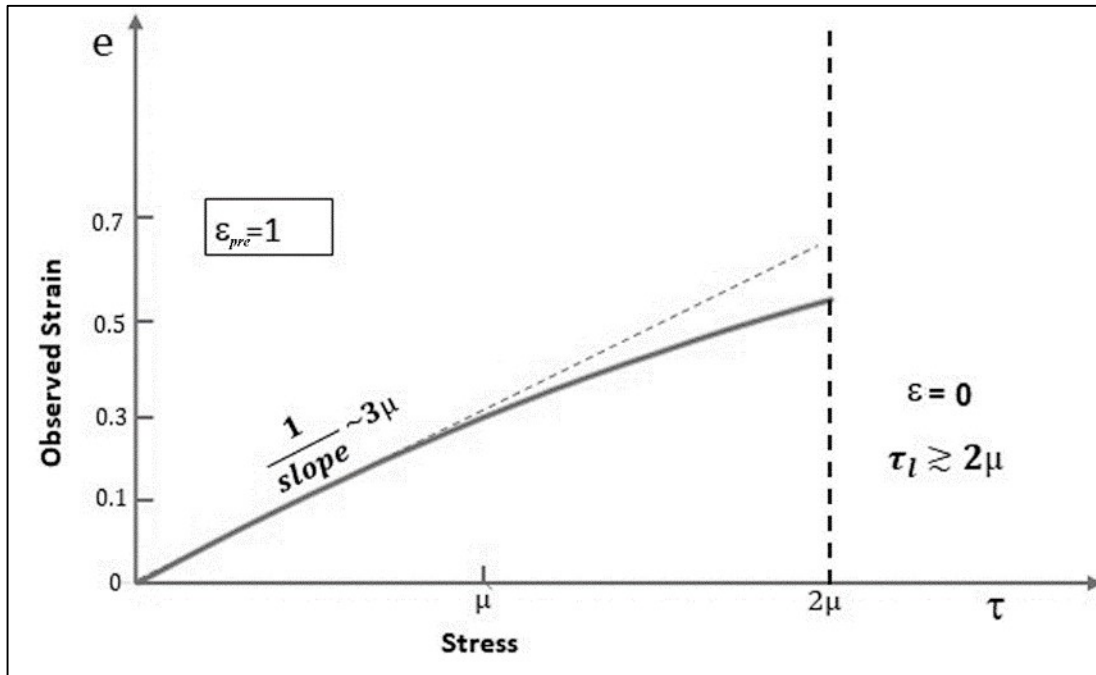
Note. Consider a 1D tissue of length L (embedded with collagen fibrils), without any pre-strain (Relaxed, zero pre-strain condition). Then an unloaded tissue with 1D pre-strain ε_{pre} , which is due to the internal FCD that causes 1D displacement u_{pre} . When the pre-stressed tissue is under loading (i.e., a confined compression), the 1D tissue length changes from u_{pre} to u' , where the subsequent strain is represented by ε . The ε represents the observed strain because of the difference between two strains (ε_{pre} and ε).

(b) The depth-dependent variations in articular cartilage, where the collagen fibers have a tri-zonal organization, the 1D depth-dependent modulus (μ) of the collagen fibers, and the depth-dependent FCD that represents the concentration changes of proteoglycans. The values shown in the modulus and FCD are used in the simulation.

(c) The schematics of collagen fibers (loaded and un-loaded) in SZ (parallel and thinner in diameters $\sim 50 \mu\text{m}$) and RZ (perpendicular and thicker in diameters $\sim 100\text{-}200 \mu\text{m}$). The PGs entangle around the fibers (lesser in content in SZ than RZ) and cause the fibril pre-strain which depends on both FCD and μ . Fibers in SZ are easier to compress than the fibers in RZ under uniaxial loading (the third panel). No significant structural changes happen to the radial zone collagen fibers under loading.

Figure 5.2

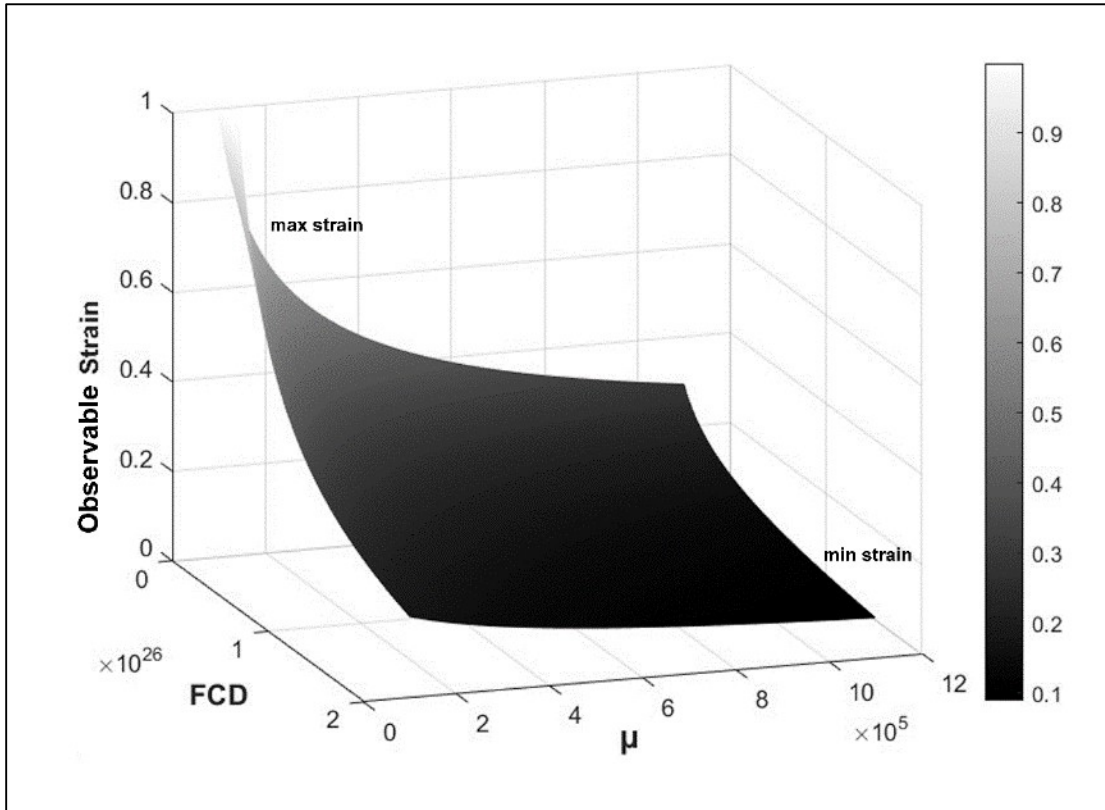
The relationship between applied stress τ and observed strain e for an assumed pre-strain $\varepsilon_{pre} = 1$.



Note. Originally the collagen pre-stress (μ) was assumed linear, and the stress-strain relationship appears nonlinear when τ becomes larger than μ . At smaller stress, the effective modulus ($1/\text{slope}$) is 3μ ; the collagen fibers lose pre-strain ($\varepsilon = 0$), and can no longer support compressive load if the applied stress becomes greater than 2μ .

Figure 5.3

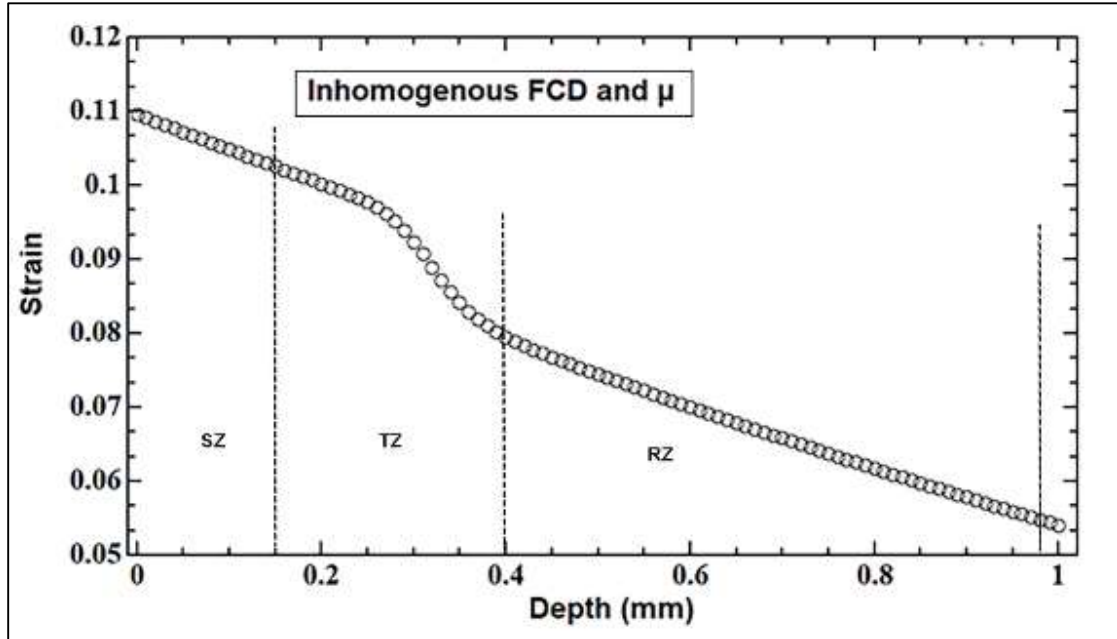
Effect of varying FCD and μ on simulated strains for a given stress ($\tau = 0.2$ MPa)



Note. When both FCD and μ are towards the smaller end, the strains are larger (a known trend for SZ of cartilage under compression), while for larger FCD and μ , the strains appear smaller (RZ trend).

Figure 5.4

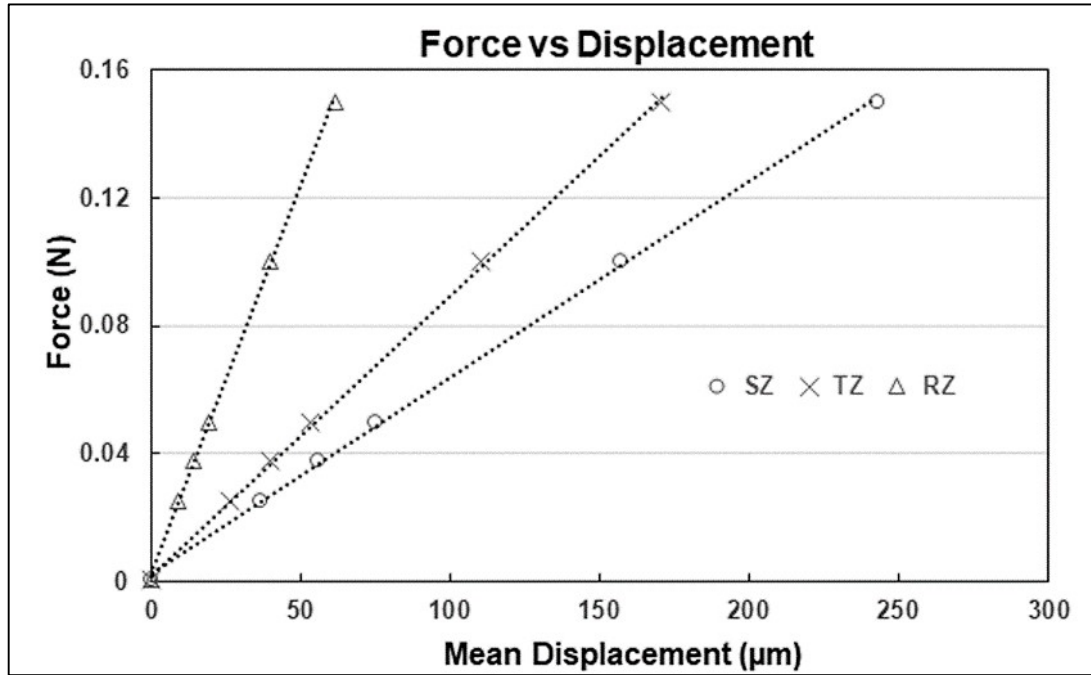
The simulated observed strain e over depth when both FCD and μ are-varying with depth.



Note. The tri-zonal tissue has a length $L = 1$ mm ($m1 = 0.3$ mm, $m2 = 0.09$ mm), and under the applied stress $\tau = 0.1$ MPa. The tissue strain is the highest at the articular surface and decreases from the surface zone (SZ) to the deep radial zone (RZ).

Figure 5.5

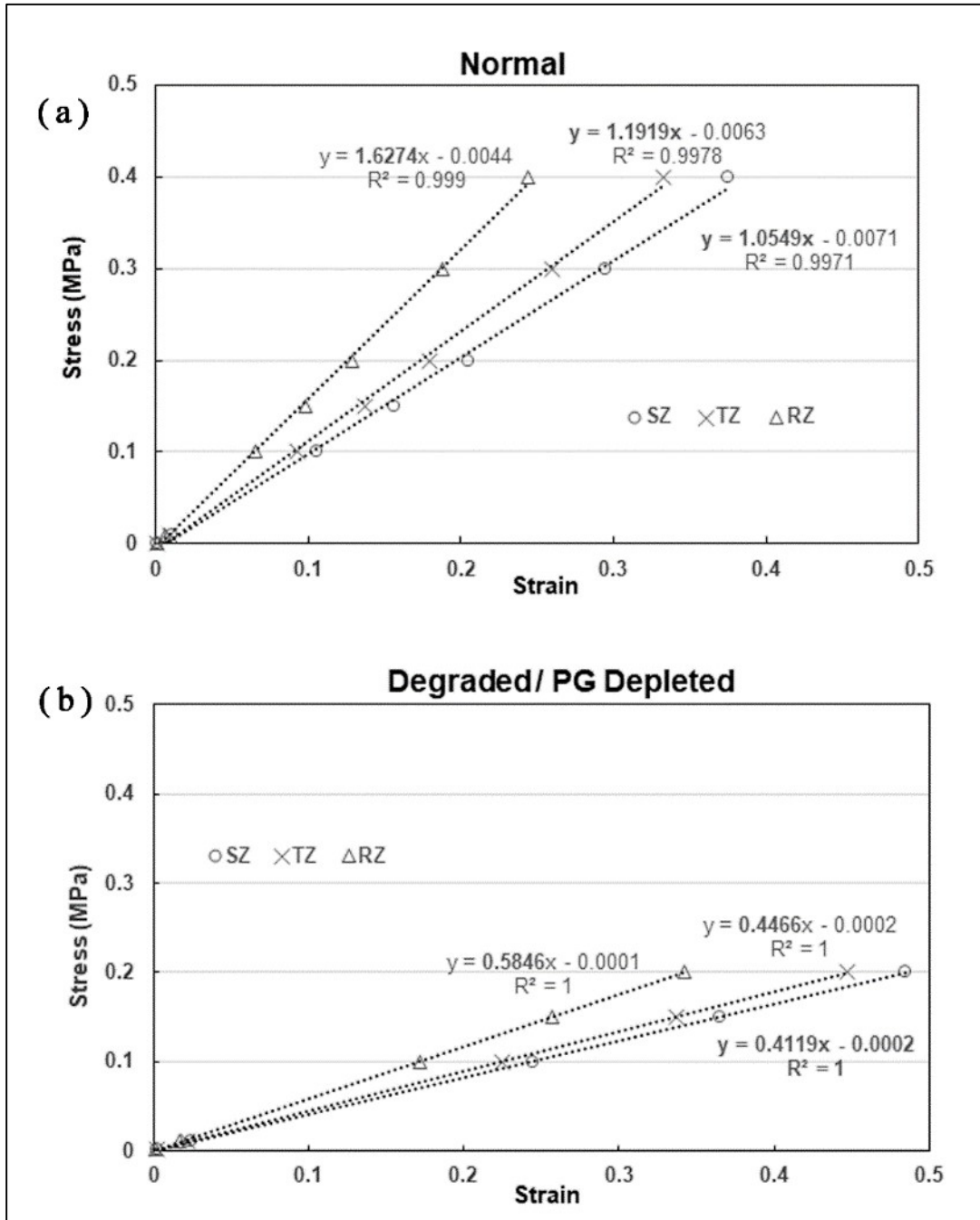
The equilibrium force vs intra-tissue displacement plots for three pre-defined zones (SZ, TZ, RZ).



Note. The intra-tissue displacements increased linearly with increasing force/stress for all three zones but the increase in intra-tissue displacements for SZ and TZ is significantly higher as compared to RZ. The zonal strains (SZ, TZ, and RZ) in cartilage are obtained by averaging the depth-dependent strain values for the respective zones defined by Eq. (5.13), where SZ is 15% of the total thickness, TZ is 25% of the total thickness, and RZ is the $\sim 60\%$ of the total thickness. The total depth of tissue used in the simulation is 0.5 mm.

Figure 5.6

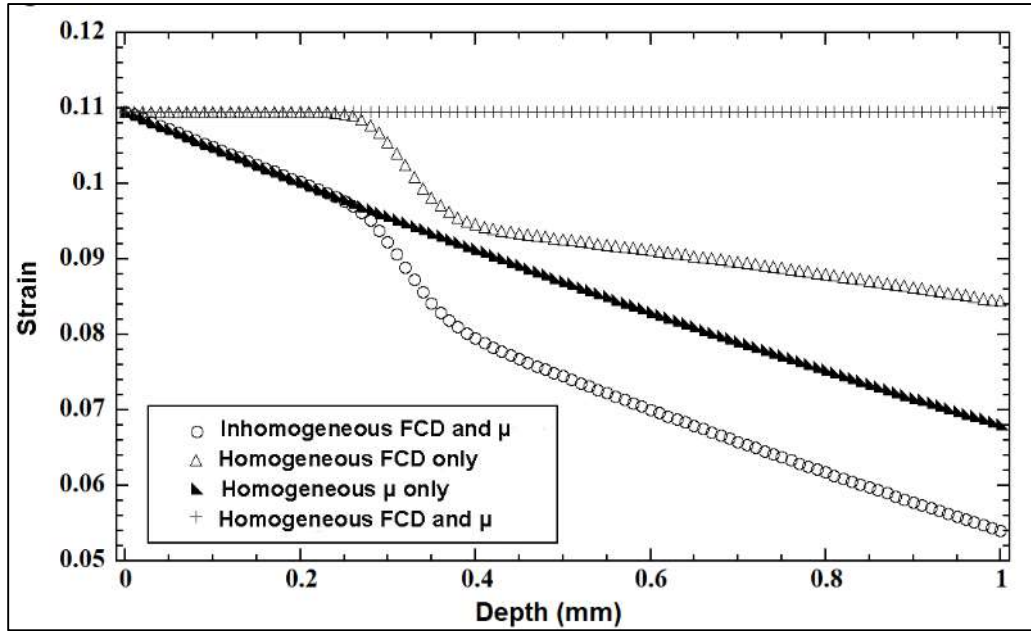
The effective compressive modulus, μ_{eff} , in three zones in a simulated 'mature healthy' and 'mature degraded cartilage'



Note. The slope of linear fit using stress-strain relationship represents the μ_{eff} . (a) For the SZ the modulus is ~ 1 MPa (consistent with several experimental studies) and increases to ~ 1.6 MPa (smaller than what is reported by other studies) in the deep region (RZ). The discrepancy in RZ modulus was attributed to the choice of elastic stiffness (μ) for RZ fibers (which could be higher than 0.6 MPa). (b) Loss of PG or FCD results in a decrease in pre-strain ε_{pre} , an increase in the intra-tissue strains ($\sim 50\%$) for the given stress, and consequently a decrease in effective modulus (slope of linear fit) for each zone (60% for SZ, 62% for TZ, and 64% for RZ). In other words, the amount of force required to compress the tissue is smaller than the normal tissue. The depth-wise loss of FCD was introduced in the model by adjusting SZ FCD (C_0) from 159 mM to 1.59 mM ($\sim 99\%$ decrease), and in RZ from 398 mM to 79.6 mM, ($\sim 80\%$ decrease), while the μ (0.4 MPa-0.6 MPa) is the same as used for the normal case.

Figure 5.7

The simulated strains for the given stress ($\tau = 0.1$ MPa) at different parameter conditions (FCD, μ , homogenous and non-homogenous) as the function of the tissue depth



Note. A constant FCD (159 mM) and constant intrinsic modulus of collagen fibers (0.4 MPa) were used for the homogenous tissue. Homogenous distributions of both FCD and μ along the depth generated constant strain values. When non-homogenous distributions of both FCD and μ were added to the model, the intra-tissue strain decreases significantly with increasing depth.

CHAPTER SIX

SUMMARY AND FUTURE DIRECTIONS

6.1 Summary

This dissertation contains three individual projects; two of them have been published in peer-reviewed journals and the third one will be submitted to a journal.

In the first project, we successfully established the baseline characteristics of the rabbits' humeral and femoral cartilage, using quantitative MRI relaxation times (T_2 , $T_{1\rho}$, T_1) and quantitative PLM measures (retardation, angle). To the best of our knowledge, these μ MRI results are the first quantitative characterization of the entire depth of humeral and femoral cartilage in rabbits at a sub-10 μm resolution; the highest MRI resolution in cartilage research. Comparing the sub-10 μm μ MRI results with low-resolution MRI results (70–82 μm) demonstrates the importance of having the sufficient resolution in MRI for the zonal assessment of articular cartilage. The quantitative PLM imaging (retardation, angle) at 1.0 μm and 4.0 μm pixel resolution further verify and enhance the high-resolution MRI results, which provides the anatomic knowledge that can be correlated with non-invasive MRI results. This project provides the baseline data for any future use of the rabbit model in the bench-top investigation of OA.

In the second project, we studied the topographical variations among different locations on the medial femoral condyle of juvenile rabbits. Distinctly different characteristics of tissue properties especially the arrangement of collagen fibers and chondrocytes that were found in the cartilage at different locations on the femoral condyle, likely related to the structural variations of different matrices in cartilage. Such

structural variations were not present in mature rabbit articular cartilage (>1 year). Such variations have a direct impact on the quantitative measurements of T2 relaxation times and PLM measurements (angle, orientation). Our high-resolution imaging findings in the location-specific differences in knee joint's collagen fiber organization can help to better understand juvenile tissue diagnostic imaging, developmental biology, localized mechanobiology, tissue engineering, and design of more successful repair strategies.

In the third project, a mathematical model was used to calculate the anisotropic collagen fibril mechanics in articular cartilage under 1D axial compression (perpendicular-to-surface direction). Under the assumption that collagen fibers are stiffer parallel to their length than perpendicular to it when combined with a depth-varying concentration of FCD, the model successfully predicted the typical depth-dependent trends of zone-specific strains and displacements (both decreasing with depth). In general, the model supports the hypothesis that the depth-dependent compressive response of articular is not only dependent upon the concentration of proteoglycans but also the anisotropic mechanical properties of the pre-strained collagen fibril network. A better understanding of anisotropy in biological tissues is important for predicting material properties and designing novel engineering structures.

6.2 Future directions

To study the depth-dependent characteristics of rabbit cartilage, sub-10 μm pixel resolution is sufficient and necessary in quantitative MRI. One of the limitations of using a magnetization-prepared spin-echo sequence to achieve sufficient SNR at high resolution is the long scan time. We plan to introduce compressed sensing techniques combined with our currently used pulse sequences to accelerate the acquisition process of

T2, T1 ρ , and T1 quantification on cartilage imaging without sacrificing accuracy. Recent new approaches (Geethanath et al., 2013; Madelin et al., 2012; Pandit et al., 2016) with compressed sensing (CS) have a reasonable potential to provide accurate and fast quantitative cartilage assessment and provide a way to make quantitative imaging techniques more accessible for clinical applications.

We have identified distinct variations in terms of collagen fibril structure and chondrocyte arrangement in juvenile rabbit femur cartilage. In the future, we intend to use this topographical knowledge to design location-specific biochemical, and biomechanical studies on rabbit femur cartilage to obtain more insights into their microenvironment and their role in development and biomechanics. We are interested to know why some sites (usually central in joint) attain trizonal architecture earlier than others (peripheral) and which factors are responsible for converting mono-zonal homogenous fibril architecture to mature trizonal architecture. New studies can include mapping the metabolic activity of chondrocytes and biochemical compositional differences present on the joint surface. We would further like to explore whether the difference in the structural development of collagen fiber at different locations of joints during growth and maturation can have any implication in case of tissue degradation and can this knowledge be useful in improving cartilage repair. Our location specific T2 anisotropy and PLM retardation and orientation data can be correlated with biomechanical behavior (modulus, stiffness, etc.) of structurally different locations.

We intend to use our mathematical model in conjunction with biomechanical testing on cartilage, to explore the role of structural anisotropy in tissue loading response and to separate the structure-composition effect. When compressed, tissue strains are

predicted to be higher in the deep zone in the absence of radial fibers (no trizonal architecture). This prediction can be tested using mechanical compression tests on sites with and without zonal arrangement and measuring the zonal strains.

APPENDIX A

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Fig 2.1

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Fig 2.4

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Fig 2.6



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Quantitative measurement of T2, T1p and T1 relaxation times in articular cartilage and cartilage-bone interface by SE and UTE imaging at microscopic resolution
Author: Rohit Mahar, Syeda Batool, Farid Badar, Yang Xia
Publication: Journal of Magnetic Resonance
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APPENDIX B

MATLAB CODE USED FOR THE STRAIN CALCULATIONS IN CHAPTER FIVE

%% Analysis of intra tissue strains with varying fiber orientation and FCD.

% Defining parameter

```
Ny=101; % number of nodes
D=0.001; %length of tissue 1 mm or 0.001 m
dy=D/(Ny-1);
for j = 1: Ny %y= depth.
    y(j)=(j-1) *dy;
end
t=zeros (Ny);
E=zeros (Ny);
tau=0.1*106; % applied stress
K=1.380649*10-23; % Boltzmann constant j/k
T=293; % 293 K, Room Temperature
% Define angle
for j=1: Ny
    t(j) = pi/4*(tanh((y(j)-(.0003))/.00009) +1.0);
    % t(j) =0;
end
% Depth dependent intrinsic stiffness of collagen fibers
for j=1: Ny
    mu0 = (0.4*106);
    mu1 = (0.08*106) *exp(y(j)/0.00099);
    mu(j) = mu0+mu1*sin(t(j)). ^4;
    % mu(j) = mu0; % for homogenous case
end
% Depth dependent concentration of FCD/GAG
```

```

C0=9.57*10^25;      % Surface GAG 40 mg/ml (wet weight), converted in to 'no
                    % of particles/m3'

for j = 1: Ny
    C(j)=C0*exp(y(j)/.0009)      %r = 0.0009.
    % C(j)=C0;                  % for homogenous case
end

% Pre-strain
for j = 1: Ny
    E(j)=0.5*(-1+sqrt(1+(4*C(j)*(K)*(T)/mu(j))));
end

% Calculation of intra-tissue strain e
for j = 1: Ny
    e(j)=(E(j)/(1+E(j))) +(0.5/(1+E(j))) *(1+(tau/mu(j))-sqrt((1-
    (tau/mu(j)).^2+4*E(j)*(1+E(j))));
end

%Calculation of displacements
for j=Ny: -1:2
    u(Ny)=0;
    u(j-1) =u(j)-dy.*e(j);
end

%figure;
%plot(y,e,'-*');
%title('Intra-Tissue Strain ')
%xlabel('Displacement (m)')
%ylabel('Strain')
%figure;
%plot(y,u)
%title('Intra-Tissue Displacement')
%xlabel('Displacement (m)')
%ylabel('Intra-Tissue Displacement(m)')

```

%%% Fig 5.3 (Phase diagram)

% Define parameters

$\tau = 0.2 \times 10^6$;

$K = 1.380649 \times 10^{-23}$;

$T = 293$;

$C = 0.2 \times 10^{23} : 2 \times 10^{23} : 2 \times 10^{26}$;

$\mu = 0.1 \times 10^6 : 0.001 \times 10^6 : 1.099 \times 10^6$;

for i=1: length(C)

for j =1: length(mu)

$E(i,j) = 0.5 * (1 + \sqrt{1 + ((4 * C(i) * K * T) ./ \mu(j))})$;

$e(i,j) = (E(i,j) ./ (1 + E(i,j))) + (0.5 ./ (1 + E(i,j))) * (1 + (\tau ./ \mu(j)) - \sqrt{(1 - (\tau ./ \mu(j))^2 + 4 * E(i,j) * (1 + E(i,j))})$;

end

end

figure;

mesh(C,mu,e);

xlabel('FCD')

ylabel('μ')

zlabel('Strain')

%%%%%%%% END %%%%%%%%%

%% Fig 5.3 (Phase diagram)

% Define parameters

$\tau = 0.2 \times 10^6$;

$K = 1.380649 \times 10^{-23}$;

$T = 293$;

$C = 0.2 \times 10^{23} : 2 \times 10^{23} : 2 \times 10^{26}$;

$\mu = 0.1 \times 10^6 : 0.001 \times 10^6 : 1.099 \times 10^6$;

for i=1: length(C)

 for j =1: length(mu)

$E(i,j) = 0.5 * (1 + \sqrt{1 + ((4 * C(i) * K * T) ./ \mu(j))})$;

$e(i,j) = (E(i,j) ./ (1 + E(i,j))) + (0.5 ./ (1 + E(i,j))) * (1 + (\tau ./ \mu(j)) - \sqrt{(1 - (\tau ./ \mu(j)).^2 + 4 * E(i,j) * (1 + E(i,j)))})$;

 end

end

figure;

mesh(C,mu,e);

xlabel('FCD')

ylabel('μ')

zlabel('Strain')

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