

**THE DISTRIBUTION AND FUNCTION OF ELASTIN AND
ELASTIC FIBRES IN THE CANINE CRUCIATE LIGAMENT
COMPLEX**

Thesis submitted in accordance with the requirements of the University of Liverpool for the
degree of Doctor in Philosophy

By

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ABSTRACT

Anterior cruciate ligament rupture (ACL) is a major source of morbidity in the dog, leading to severe osteoarthritis of the knee joint and marked lameness. Following rupture, the ACL will not heal and in the dog, ACL rupture is thought to be the end stage of degenerative ligament disease (non-contact ACL injury). The extracellular matrix (ECM) of CLs has been extensively studied but little is known of the role of elastic fibres in the physiology of the ECM, the mechanics of CL function and in CL degeneration. Elastic fibres include polymers of fibrillins (microfibrils), bundles of microfibrils (oxytalan fibres) and elastin fibres (bundles of microfibrils with an elastin core). The hypothesis of this thesis is that elastin has a mechanical and a biological role in the canine cruciate ligament complex. It is further hypothesised that the distribution and function of elastic fibres will vary between three breeds of dog with differing risk of ACL rupture are: the greyhound with a low risk, the beagle with a low-to-moderate risk and the Labrador retriever with a high risk.

The distribution of elastic fibres, fibrillins and cells was investigated throughout the CL complex using a combination of histochemical staining and immunofluorescence. CL microanatomy was studied using Nomarski differential interference microscopy. Elastin was measured biochemically and compared to histologic assessment of tissue architecture, elastic fibre staining and other biochemical parameters. The biological effect of elastin degradation products (EDPs) was assessed in an *in vitro* ACL cell culture model. A low risk dog breed to ACL rupture (greyhound) was used in all investigations and comparisons were made with other breeds with regard to cellular and elastic fibre anatomy.

Differences in cell morphology between breeds with differing risk of ACL rupture may reflect fundamental differences in CL physiology possibly through altered cell-to-cell communication. Cellular and matrix changes, considered degenerative, were seen throughout the CL complex and may reflect adaptation rather than degeneration in certain dog breeds such as the greyhound. Elastin content ranged from 5.9 to 19.4% of ligament dry weight. This was a far greater proportion of canine CLs than previously. Elastin fibres may have a mechanical role in bundle reorganization following ligament deformation. The distribution of fibrillins 1 and 2 was different from the pattern previously reported in tendon and may represent a fundamental difference between ligament and tendon. In the greyhound CL there was a significant proportional increase in oxytalan fibre staining with advancing CL degeneration. This response was seen also in the Labrador retriever and the beagle but the increase in oxytalan fibre staining was less marked with advancing degeneration. Therefore production of oxytalan fibres may reflect a healing response in damaged CL tissue in breeds at a low risk of ligament rupture. Fragments of elastin containing the VGVAPG motif affect canine ACL cells *in vitro* resulting in increased transcription of fibrillin 2 mRNA. Additionally, there was synergism with TGF- β 1 resulting in upregulation of the elastin laminin receptor 1, through which EDPs are transduced. EDPs may thus have a role in response to injury in the CL.

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Dyfal donc a dyr y garreg

(Tapping persistently breaks the stone)

AUTHOR'S DECLARATION

I declare that the work in this dissertation was carried out in accordance with the regulations of the University of Liverpool. The work is original except where indicated by reference in the text.

Any views expressed in this thesis are those of the author and in no way represent those of the University of Liverpool.

This thesis has not been presented to any other university for examination in the United Kingdom or overseas.

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LIST OF ABBREVIATIONS

ACL	Anterior cruciate ligament
ADAM	A disintegrin and metalloproteinase domain
ADAM-TS	ADAM with a thrombospondin motif
ANOVA	Analysis of variance
AP	Anterior-posterior
ASMA	Alpha smooth muscle actin
B2M	Beta-2 macroglobulin
cbEGF	Calcium binding epidermal growth factor
CCA	Congenital contractile arachnodactyly
CL	Cruciate ligament
CLSM	Confocal laser scanning microscopy
DAPI	4', 6-diamidino-2-phenylindole
cDNA	Cyclic deoxyribonucleic acid
Col1a2	Collagen type I ($\alpha 2$ chain) gene
Col2a1	Collagen type II ($\alpha 1$ chain) gene
Col3a1	Collagen type III ($\alpha 1$ chain) gene
CS	Chondroitin sulphate
CTSB	Cathepsin B
CTSK	Cathepsin K
DS	Dermatan sulphate
DMEM	Dulbecco's modified Eagle's Medium
DMMB	1, 9-dimethylmethylene blue
ECM	Extracellular matrix
EBP	Elastin binding protein
EDP	Elastin degradation peptides
EDTA	Ethylenediaminetetraacetic acid
EF	Elastin fibre
EGF	Epidermal growth factor
Eln	Elastin gene
ELR1	Elastin laminin receptor 1
EVH	Verhoeff's iodine-iron haematoxylin stain
Fbn1	Fibrillin 1 gene
Fbn2	Fibrillin 2 gene
FBS	Foetal bovine serum
FGF	Fibroblast growth factor
GAG	Glycosaminoglycan
GAPDH	Glyceraldehyde phosphate dehydrogenase
GH	Greyhound
H&E	Haematoxylin and eosin
HCl	Hydrochloric acid
IGF	Insulin-like growth factor
IL	Interleukin
KS	Keratin sulphate
LR	Labrador retriever
LTBP-1	Latent transforming growth factor β binding protein 1
M	Miller's stain
MAGP-1	Microfibril associated glycoprotein 1
MCL	Medial collateral ligament

MF	Microfibril
MFS	Marfan syndrome
M-MLV	Moloney - murine leukaemia virus
MMP	Matrix metalloproteinase
mRNA	Messenger ribonucleic acid
MS	Miller's score
mVS	Modified Vasseur score
MT-MMP	Membrane type-matrix metalloproteinase
NDIC	Nomarski differential interference contrast (microscopy)
NO	Nitric oxide
OA	Osteoarthritis
OF	Oxytalan fibre
OHPro	Hydroxyproline
PBS	Phosphate-buffered saline
PCL	Posterior cruciate ligament
PG	Proteoglycan
qPCR	Quantitative polymerase chain reaction
RA	Rheumatoid arthritis
RGD	Arg-Gly-Asp peptide sequence
RT-PCR	Reverse transcriptase polymerase chain reaction
SDFT	Superficial digital flexor tendon
sGAG	Sulphated glycosaminoglycans
SOX9	Sex determining region box 9 gene
TB	8-cysteine containing motif (thrombospondin motif)
TGF	Transforming growth factor
TIMP	Tissue inhibitor of metalloproteinase
TNF	Tumour necrosis factor
TPA	Tibial plateau angle
uPA	Urokinase plasminogen activator
Vcan	Versican gene

CHAPTER ONE: General introduction

CANINE CRUCIATE LIGAMENTS

1.1 Gross Anatomy

1.1.1 Overview of the knee

The knee (stifle) is a condylar synovial joint consisting of articulations between the femur and tibia and between the femur and patella. It is an extremely complex joint in both structure and function and ligaments are essential for the maintenance of these articulations (Tirgari and Vaughan, 1975b). The cruciate ligaments (CLs), anterior (ACL) and posterior (PCL), are the primary stabilisers of the knee and serve to limit anterior translation and rotation of the tibia (Arnoczky and Marshall, 1977, Moore and Read, 1996), (**Figure 1.1**). Within the knee joint, the ACL and PCL are in intimate contact, with the ACL twisted around the PCL. They are considered to function as a unit hence the term CL complex (Arnoczky and Marshall, 1977, Harari, 1993).

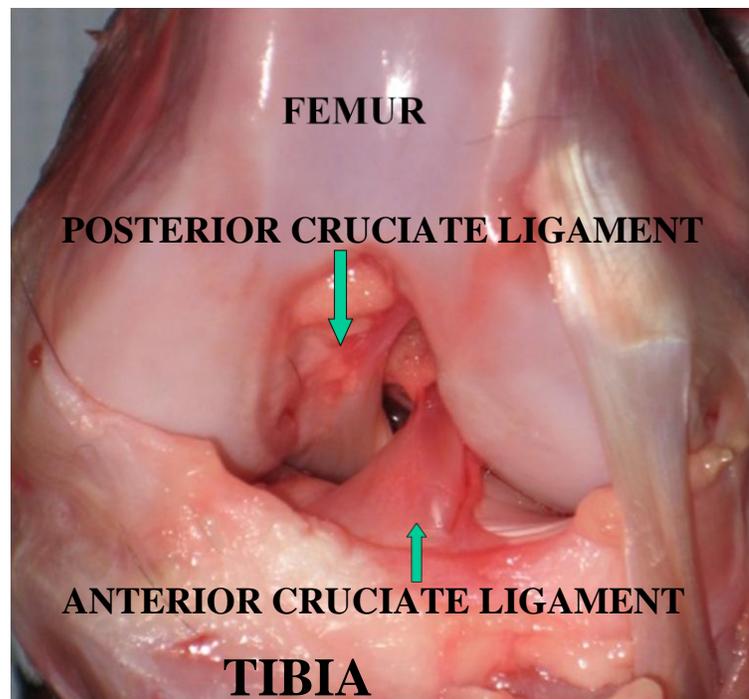


Figure 1.1: Cruciate ligament complex within the canine knee

1.1.2 ACL anatomy

The canine ACL runs anteriorly, medially, and distally in an outward spiral as it passes from the medial aspect of the lateral femoral condyle to the anterior intercondyloid area of the tibial plateau (Zahm, 1965, Haut and Little, 1969). Two separate bundles are observed *in-situ* in the dog, termed anteriomedial and posteriolateral bands. The anteriomedial subdivision is the longest, yet smaller component. It arises more proximally from the femur and inserts more anteriorly on the tibial attachment area, compared with the posteriolateral subdivision (Arnoczky and Marshall, 1977, Heffron and Campbell, 1978).

1.1.3 PCL anatomy

The PCL is a longer and wider structure than the ACL, narrowest at the midsection and fanning toward origin and insertion (Arnoczky and Marshall, 1977). It is subdivided into two bands, a larger anterior and smaller posterior, although these are often indistinct (Heffron and Campbell, 1978). They have reciprocating functions through flexion and extension with the anterior portion taut in flexion and loose in extension and the posterior portion taut in extension only (Arnoczky and Marshall, 1977, Harari, 1993).

1.1.4 Blood supply

The main supply to the central knee is the middle genicular artery penetrating the posterior joint capsule (Kobayashi et al., 2006). A well vascularised enveloping synovial tissue carries the vessels, with minimal supply from origin and insertion (Arnoczky et al., 1979). The core of the ACL is less well vascularised than the remainder (Tirgari, 1978, Vasseur et al., 1985, Narama et al., 1996). Anastomoses exist between endo- and epiligamentous networks (Kobayashi et al., 2006). The PCL may have a more substantial blood supply as there appears to be more epiligamentous vessels (Arnoczky et al., 1979). CLs may also gain nutrition from the synovial fluid (Kobayashi et al., 2006).

Although the ACL has been considered extra-articular due to the enveloping synovial epiligament, free passage of macromolecules from intra-articular synovial fluid to the substance of the ACL has been demonstrated (Kobayashi et al., 2006). As such free movement exists between the synovial fluid and ACL substance, a fall in intra-articular pressure, as may occur in osteoarthritis, is likely to adversely affect ACL blood flow.

1.1.5 Neurology

Innervation of the canine knee joint is from three major articular nerves arising from the saphenous, tibial and common peroneal nerve, although other nerves may contribute to a variable extent (O'Connor et al., 1982). The main trunk of the nerve bundles is found at the femoral origins of the CLs (Arnoczky, 1983). Nerves in the synovial envelope penetrate the CLs radially, and endoligamentous nerves course with the vessels, where their function is thought to be primarily associated with the regulation of blood flow and nociception (Kennedy et al., 1974). Mechanoreceptors within the substance may activate local reflex patterns to protect the ligament against tearing and warn against possible joint damage (Krauspe et al., 1992, Biedert et al., 1992).

1.1.6 Ultrastructural anatomy

1.1.6.1 Fascicular subdivision

In man and dogs, CLs have been described as containing twisted collagenous fascicles and fibre bundles that are subdivided into fascicles, subfascicular units, fibres, and fibrils (Arnoczky, 1983, Yahia and Drouin, 1989). A simpler subdivision has been proposed with only two divisions of bundles and fascicles with a fascicular membrane that does not concentrically bound the fascicle (Clark and Sidles, 1990). Cells in the CL substance are predominantly fibroblasts which may vary from a spindle to rounded-shape (Arnoczky, 1983). These are isolated or in short rows of 2-3 cells and tightly applied to collagen bundles.

Both CLs are covered in a fold of synovial membrane, incompletely bisecting the joint in the sagittal plane (Arnoczky and Marshall, 1977). The synovial envelope of the ACL originates at the intercondylar notch and extends to the anterior tibia where it communicates with a fold of the distal joint capsule (Alm and Stromberg, 1974). The PCL is ensheathed by two folds of synovial membrane (Alm and Stromberg, 1974). These enveloping membranes are termed the epiligament. It has been divided to an intima and subintima: the intima comprising a single layer of synoviocytes and the subintima mainly areolar tissue containing small vessels, fibroblasts and some adipocytes (Heffron and Campbell, 1978, Vasseur et al., 1985). The synovial epiligament is far more cellular than the CL substance (Arnoczky and Marshall, 1977). The epiligament is an incomplete barrier and there is ready contact with synovial fluid (Kobayashi et al., 2006, Tang et al., 2009).

Each ligament component consists of multiple elliptical fascicles (Heffron and Campbell, 1978). The peripheral subunits follow a spiral path of waviness around the fascicle axis (Yahia and Drouin, 1989, Amis and Dawkins, 1991, Kennedy et al., 1974). Each subfascicle contains bundles of collagen fibres, but these are not orientated isometrically during knee motion (Amis and Dawkins, 1991). Each change in knee position recruits fibres differently (Butler et al., 1992). It has been postulated that individual fibres change their length by straightening their crimp, a period banding of collagen observed under crossed polarizing filters (Amis and Dawkins, 1991, Yahia and Drouin, 1989, Boorman et al., 2006).

The collagen fibres are formed by fibrils, themselves an organisation of repeated collagen subunits. Fibrils have a uniform crimp parallel to the long axis of the fibril: the internal fibrils are almost straight and those on the periphery undergo maximal crimp (Alm et al., 1974, Hayashi et al., 2003a). The collagen fibrils of the PCL are larger (Brunnberg, 1989).

Organisation of the fascicles and smaller units has not been fully resolved. The difficulty in separating individual fascicles in the gross specimen implies a firm structural association (Frank, 2004). In the annulus fibrosus of the human intervertebral disc, it has been demonstrated that structural cohesion between the fascicles is by complex linking elements which control movement (Pezowicz et al., 2005). These are thought to contain elastin (Smith and Fazzalari, 2006, Yu et al., 2007) and may have a role in fascicle realignment following deformation (Szirmai, 1970). A lateral supporting structure for collagen fibres within bands of the rabbit straight patellar ligament has been suggested as a mechanism for protecting against ligament fibre damage (Boorman et al., 2006).

1.1.6.2 Cellular organisation

The cells in ligament have an important role in maintaining the extracellular matrix (ECM) and controlling responses to altered mechanical load and injury (Frank, 2004). The ECM determines the mechanical properties of a ligament. The mechanical environment of the cell has been shown to influence cell morphology in other normal connective tissues such as tendon and cartilage (Giori et al., 1993, Matyas et al., 1994, Ralphs et al., 1998). Cells in tensile load in the absence of significant compressive load have cytoplasmic processes which are frequently long and extend parallel perpendicularly or transversely to the collagen fibres (Bruehlmann et al., 2002, Lo et al., 2002a). The detection of gap junctions in association with these cell connections suggests the potential to coordinate cellular and metabolic responses

throughout the tissue through cell-to-cell communication (Bruehlmann et al., 2002, McNeilly et al., 1996, Lo et al., 2002b). This elaborate three-dimensional structure has been termed the cell matrix, and has been described in tendon (Ralphs et al., 1998), meniscus (Hellio Le Graverand et al., 2001a) as well as in CL (Lo et al., 2002b). The cell matrix is dynamic, and changes have been noted in healing (Lo et al., 2002b), injury and degenerative joint disease (Hellio Le Graverand et al., 2001d).

This cellular matrix has a number of possible functions:

a) Organisation of collagen

Classically the ECM has been thought of the scaffold in which cells proliferate. However, the cellular matrix may itself act as the framework for ECM (Lo et al., 2002a). The processes allow detection of collagen at some distance from the cell (Birk and Zycband, 1994). In developing tissue, the cell number remains constant as the ECM is deposited in an orderly manner around each cell (Lo et al., 2002a). Importantly, cell orientation often precedes matrix deposition and alignment, particularly apparent in the intervertebral disc (Hayes et al., 1999).

b) Sensing the mechanical environment

The cellular network has the potential to connect the entire length of a ligament and may be critical in maintaining and remodeling the ECM directly influencing its mechanics (Lo et al., 2002a). Cells are thought to respond to mechanical environment *in vivo* (Banes et al., 1995). *In vitro*, gap junction inhibitors affected calcium wave propagation and inhibited collagen type I synthesis when tendon fibroblasts grown in monolayer culture were subjected to a mechanical load (Waggett et al., 2006). This implies that gap junction inhibitors can affect cellular response to mechanical stimulus, suggesting, at least in part, gap junctions regulate this response. *Ex-vivo* studies in tendon fibroblasts have shown an altered DNA and collagen synthesis pattern when gap junctions were inhibited (Banes et al., 1999). Taken together, these results suggest gap junctions and cellular processes are important in sensing and responding to the mechanical environment.

c) Injury

The effect of injury on connective tissue ECM has been studied, but little is known of changes to the cellular matrix (Birch et al., 1998, Hellio Le Graverand et al., 2001b, Hellio Le Graverand et al., 2001c). In the meniscus, the cell matrix in the periphery was disrupted in

response to either apoptosis or cellular proliferation resulting in cells retracting their cellular processes (Hellio Le Graverand et al., 2001b, Hellio Le Graverand et al., 2001d). This retraction resulted in isolated islands of cells with changes to their phenotype. Cells in these clusters further retracted the cytoplasmic processes and formed three types of morphological phenotype: stellate, round or mixed. With time the clusters of round cells become more prominent and increase in size possibly through proliferation. These changes resulted in changes to cell-cell and cell-matrix interactions (Hellio Le Graverand et al., 2001d). A change in the surrounding matrix was also reported, with increased matrix metalloproteinase (MMP)-13 and collagen type II degradation products localised to meniscal tears (Hellio Le Graverand et al., 2000). The failure or disruption of the cellular network may be the initiating event leading to disruption of the ECM (Lo et al., 2002a).

d) Healing

The healing of the ovine medial collateral ligament (MCL, normal functional healing) and the ACL (non-functional healing) have been compared (Lo et al., 2001, Lo et al., 2002b). At three months, both ACL and MCL showed prominent discontinuities in the scar formation (area devoid of nuclei). Scars in the MCL they were filled with cellular processes and were connexin 43 positive, whereas in the ACL they were devoid of cells, processes and gap junctions. Although these discontinuities may be a secondary change in the ACL, they are likely to disrupt chemical and electrical signaling between cells in the scar and uninjured tissue from communicating effectively. Thus a coordinated response to the mechanical environment may not be possible further compromising ligament integrity. Differences in the expression of α and β integrin subunits have also been noted in the MCL compared to the ACL (Schreck et al., 1995). Thus differences in the healing capacity of the two ligament types may be the result of complex changes in the relationships between cells and between cells and the ECM.

1.2 Functional anatomy of CLs

Knee stability is a complex interaction of passive (bony and musculotendinous structures, menisci and ligaments) and active (muscular forces and joint compression) forces (Slocum and Slocum, 1993, Korvick et al., 1994). Other knee structures act as complementary constraints to the CLs in various planes although differing in their primary functions (Markolf et al., 1981). The ACL acts to restrain anterior translation of the tibia as well as to limit tibial rotation

during flexion (Arnoczky and Marshall, 1977, Harari, 1993). The anteriomedial part of the ACL is taut throughout flexion and extension and the posteriolateral part taut in extension only. In extension, the collateral ligaments are the primary restraints as the CLs unravel in external rotation, and so no restraint is provided in this direction (Vasseur et al., 1985). Changes in joint flexion alters the tension in the separate bands as some fibres are stressed and others relaxed, allowing the CLs to withstand multi-axial stresses of normal function and range of motion (Kennedy et al., 1976, Takai et al., 1993). Muscular (active) support of the knee is a function of the quadriceps and patellar tendon anteriorly and the popliteal and hamstring muscles posteriorly (Markolf et al., 1981, Aron, 1988, Slocum and Slocum, 1993, Korvick et al., 1994). The importance of the PCL in knee stability is less clear, and prevention of posterior tibial translation appears to be the primary function (Arnoczky and Marshall, 1977).

CL ECM STRUCTURE

Overview

Ligaments are approximately two thirds water and one third solid (Boorman et al., 2006). The solid components are principally collagen (85% type I, the rest III, VI, V, XI and XIV), accounting for 75% of the dry weight (Frank, 2004). Collagen fibrils lie along the long axis of a ligament and display crimp along their length (Amiel et al., 1984, Yahia and Drouin, 1989, Amiel et al., 1995a). The remaining ligament consists of proteoglycans (PGs) (<1%), elastin, glycoproteins such as laminin, actin and the integrins, and other components of the dry weight which have not yet been identified (Frank, 2004).

1.3 Collagen

1.3.1 Collagen: Classification

Collagen is relatively inert, having a half life of 300-500 days, although this varies with tissue (Neuberger and Slack, 1953) and may be much longer (Thorpe et al., 2010). To date, there have been twenty seven collagens identified (**Table 1.1**) and they contribute to 30% of protein mass in the human body (Canty and Kadler, 2005). They have many important functions such as maintaining the structure of tissue, imparting tensile strength, cell adhesion, chemotaxis, cell migration and regulation of matrix biology in growth, healing and disease (Myllyharju and Kivirikko, 2004). Some collagens have a wide tissue distribution while others are very

restricted (Myllyharju and Kivirikko, 2004). Many collagens have non-collagenous domains which, when cleaved proteolytically, have important functions such as inhibition of endothelial cell migration, angiogenesis and tumour growth (Marneros and Olsen, 2001).

COLLAGEN TYPE	GENES	TISSUE AND SUPRAMOLECULAR ORGANISATION
I	COL1A1 COL1A2	Fibrils; Tendon, bone, skin, cornea, blood vessels.
II	COL2A1	Fibrils; Cartilage
III	COL3A1	Heterotypic fibrils with type I
IV	COL4A1 COL4A2 COL4A3 COL4A4 COL4A5 COL4A6	Network; Basement membrane
V	COL5A1 COL5A2 COL5A3	Heterotypic fibrils with type I
VI	COL6A1 COL6A2 COL6A3	Fine microfibrils, distinct from fibrillin microfibrils; Ubiquitous
VII	COL7A1	Fibrils; Dermal-epidermal junction
VIII	COL8A1 COL8A2	3D hexagonal lattice; Descemet's membrane
IX	COL9A1 COL9A2	Associated with type II fibrils
X	COL10A1	2D hexagonal lattice; Growth plate
XI	COL11A1 COL11A2	Heterotypic fibrils with type II
XII	COL12A1	Associated with type I fibrils
XIII	COL13A1	Transmembrane and cell adhesion
XIV	COL14A1	Associated with type I fibrils
XV	COL15A1	Specialised basement membranes, cleavage produces angiogenic fragment
XVI	COL16A1	Component of specialised fibrillin-rich microfibrils in skin, type II fibrils in cartilage
XVII	COL17A1	Transmembrane component of hemidesmosomes (cell-cell junctions) which attach epidermis to basement membrane
XVIII	COL18A1	Cleavage produces angiogenic fragment
XIX	COL19A1	Radially distributed aggregates
XX	COL20A1	May be associated with type I fibrils; Cornea, skin, cartilage, tendon
XXI	COL21A1	May be fibril-associated; Widespread
XXII	COL22A1	May be microfibril-associated; Tissue junctions
XXIII	COL23A1	Transmembrane in tumour cells
XXIV	COL24A1	Expressed in tissues containing type I
XXV	COL25A1	Transmembrane; neurons (cleaved form present in Amyloid plaques)
XXVI	COL26A1	Testis and Ovary
XXVII	COL27A1	Widespread, especially cartilage

Table 1.1: Collagen types, genes and supramolecular organisation and distribution (From Canty and Kadler 2005).

1.3.2 Collagen: Structure

Collagens are trimers of polypeptide chains, called α -chains. Each chain comprises a repeating Gly-A-B triplet, where A and B can be any residue, but usually proline and hydroxyproline respectively (Myllyharju and Kivirikko, 2004). This triplet motif results in a left-handed helix that can intertwine with two others to form a right-handed triple helix structure (van der Rest and Garrone, 1991).

Collagen fibrils are the principle source of tensile strength in mammalian tissue. They have a 67nm axial periodicity, are up to several millimetres long and have a wide range of widths from a few nanometers to 500nm (Canty and Kadler, 2005). They have a wide variety of tissue-specific arrangements, and in tendon and ligament they are organised in parallel arrays (Boot-Handford et al., 2003, Myllyharju and Kivirikko, 2004). In tendon, ligament, skin and bone, type I predominates and in cartilage type II is most abundant. Fibrils are usually heterotypic which is achieved through different α chains in a molecule: the six chains of type IV form at least three different molecules (Borza et al., 2001).

1.3.3 Collagen: Assembly

Formation of the collagen fibril begins with the synthesis of polypeptide chains on membrane bound ribosomes and secreted into the lumen of the endoplasmic reticulum (Myllyharju and Kivirikko, 2004). Here the main steps in biosynthesis take place:

- 1) Cleavage of the signal peptides
- 2) Hydroxylation of certain proline and lysine residues
- 3) Glycosylation of some of the hydroxylysine residues
- 4) Glycosylation of certain asparagine residues in the C, or C and N terminal propeptides
- 5) Association of three C propeptides, by specific recognition sequences
- 6) Formation of intra- and inter-molecular disulphide bonds

The C propeptide association forms the nucleus for triple helix formation, and following hydroxylation of around 100 proline residues, the triple helix is propagated toward the N-terminus (Canty and Kadler, 2005). The procollagen molecules are transported from the endoplasmic reticulum through the Golgi apparatus where they aggregate to form secretory vesicles. Following secretion, the N and C propeptides are cleaved and the molecules

spontaneously self-assemble into fibrils and form covalent crosslinks (Canty and Kadler, 2005).

1.3.4 Collagen: Crosslinks

Following fibril formation, crosslinking imparts mechanical strength and stability to the fibrils. The pattern of crosslinking varies between tissue and collagen types, and the pattern of strain and pathology of that tissue (Eyre et al., 1984, Paul and Bailey, 1996). There are three types: immature enzymatic, mature enzymatic and non-enzymatic (Kielty et al., 1993, Yamauchi et al., 1988, Kuypers et al., 1992).

1.4 Elastin, fibrillin and the elastic fibres

1.4.1 Overview

Although collagen provides tensile strength to ligament, other components may contribute to the overall mechanical function of the complex (Frank, 2004, Ujiie et al., 2008). Elastin fibres comprise a central cross-linked core of highly extensible elastin surrounded by a supporting sheath of microfibrils, with many other associated molecules (Kielty, 2006). Microfibrils (MFs) are polymers of fibrillins 1 and 2 and are considered to have a structural role in ligament and tendon. Bundles of MFs are known as oxytalan fibres. Collectively, oxytalan and elastin fibres are referred to as elastic fibres. Elastin has traditionally been considered a minor component of ligament tissue (Frank, 2004).

Elastic fibres have important mechanical, biochemical and cell-regulatory functions in tissue namely vascular and connective tissues such as intervertebral disc (Kielty, 2006). Reversible elasticity is a function of both elastin and oxytalan fibres (Eriksen et al., 2001) where elastic fibre distribution is considered to reflect function (Kielty et al., 2002a). MFs may have a key role in the extracellular regulation of transforming growth factor (TGF) β (Charbonneau et al., 2004, Feng and Derynck, 2005) as well as regulation of cell-matrix interactions (Ito et al., 1997, Wendel et al., 2000).

A wide distribution of elastic fibres in the human ACL has been described, with abundant elastin fibres and oxytalan fibres running with collagen bundles described using electron microscopy (Strocchi et al., 1992). In canine CLs, elastin fibres have only been reported at low levels (Paatsama, 1952, Vasseur et al., 1985). The presence of oxytalan fibres or microfibrils

has not been determined in the canine CL complex and the role of elastin fibres in the CL complex has not been determined in any species.

1.4.2 Molecular composition

1.4.2.1 Elastin core

Elastin is the most abundant component of elastic fibres (around 90% of the fibre) and has a very low turnover in healthy tissue (Petersen et al., 2002, Kielty, 2006). In man, coding for elastin is by a single copy gene on chromosome 7q11.2 and secreted as a 65-70kDa soluble tropoelastin precursor (Tamburro et al., 2005). Tissue specific functional properties may be achieved by splice variants of the original transcript (Kielty, 2006). Tropoelastin has a multi-domain structure with repeating hydrophobic and lysine-rich crosslinking domains each encoded by separate exons (**Figure 1.2**). The unique C-terminal domain contains two cysteine residues and may play a key role in elastin assembly. Crosslinked elastin is formed under the direction of lysyl oxidase (LOX) and other members of this enzyme family (Szauter et al., 2005). Proteoglycans, including biglycan, have been detected in the core, and heparan sulphate may also be present (Kielty, 2006).

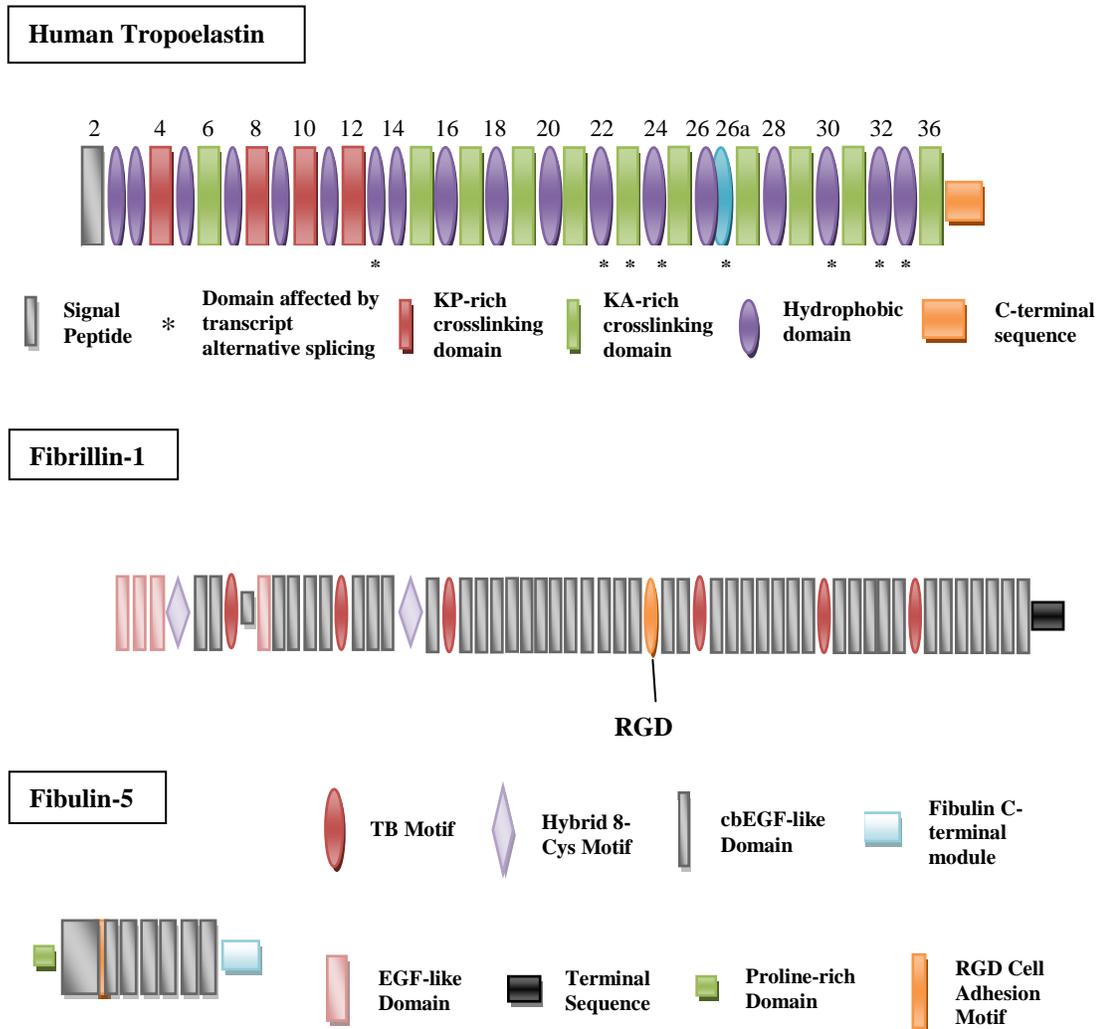


Figure 1.2: Domain structures of elastin, fibrillin-1 and fibulin-5. Abbreviations: calcium binding epidermal growth factor (cbEGF), Arg-Gly-Asp peptide sequence (RGD), eight-cysteine-containing motif (TB, also known as thrombospondin motif), epidermal growth factor (EGF). Adapted from Kielty 2006

1.4.2.2 Microfibrils

The structural framework of microfibrils (MFs) is comprised of fibrillins. These are large multi-domain glycoproteins of around 350kDa, and comprise 47 epidermal growth factor-like domains (EGF-like); 43 of which are calcium-bonding epidermal growth factor (cbEGF)-like domains (**Figure 1.2**). These domains are interspersed with 7 eight-cysteine-containing (TB) motifs and two hybrid motifs containing elements of both EGF-like and TB. There are 14-N-glycosylation sites, and the N terminus is a unique cysteine-containing motif (Kielty et al.,

2005). In man there are three fibrillin isoforms with fibrillins 1 and 2 having partially overlapping expression patterns. Fibrillin 1 is expressed throughout life and fibrillin 2 is expressed mainly in foetal tissues (Zhang et al., 1994, Charbonneau et al., 2003). Fibrillins 1 and 2 have been shown to co-localise within MFs and may overlap in function (Carta et al., 2006). There are many MF associated molecules, of which microfibril associated glycoprotein MAGP-1 is the most likely to have a structural component (Cain et al., 2006). (MAGP)-2 and latent TGF- β binding protein (LTBP)-1 also co-localise in certain tissues (Kielty, 2006).

1.4.2.3 Elastic fibre interface molecules

A number of molecules are found at the pericellular-elastic fibre or microfibril-elastin interface and are listed in **Table 1.2** (Reinhardt et al., 1996, Isogai et al., 2002). Collagen type VIII may also co-localise in vascular tissues forming hexagonal basement membrane associated networks.

Fibulin-5 is expressed by vascular smooth muscle cells and endothelial cells. It is a glycoprotein of around 55 kDa containing five cbEGF-like domains and an Arg-Gly-Asp (RGD) motif (**Figure 2**). It is involved in mediation of vascular cell adhesion through integrin receptors, smooth muscle cell proliferation and migration and in regulation of elastic fibre formation (Chu and Tsuda, 2004). Its expression is down-regulated in adult arteries, but markedly up-regulated in vascular cells following injury, angioplasty in neointimal cells, and in atherosclerotic cells. It is localised to the elastic lamina surfaces adjacent to endothelial cells and throughout the aortic media. Fibulin-4 is another factor essential for elastogenesis (McLaughlin et al., 2006)

Molecule	Elastic Fibre Location
Fibrillin-1	Microfibrils
Fibrillin-2	Microfibrils
Fibrillin-3	Unknown – microfibrils likely
MAGP-1	Microfibrils
MAGP-2	Some microfibrils
LTBP-1	Some microfibrils; also fibronectin
LTBP-2	Microfibrils, elastic fibres
LTBP-3	Fibrillar structures
LTBP-4	Fibrillar structures, fibrillin
Decorin	Microfibrils, microfibril-elastic fibre interface
Biglycan	Elastic fibre core
Versican	Some microfibrils
Heparin Sulphate	Microfibrils, elastic fibre core
Perlecan	Microfibrils
MFAP-1	Some microfibrils
MFAP-3	Some microfibrils
MFAP-4 (MAGP-36)	Some microfibrils
β lgH3	Elastic fibre-collagen interface
Tropoelastin	Elastic fibre core
LOX	Newly secreted tropoelastin, microfibril-elastin interface
LOXL	Microfibril-fibulin-5-elastin interface
Fibulin-1	Elastic fibre core
Fibulin-2	Elastin-microfibril interface
Fibulin-4	Unknown – likely in elastic fibre core
Fibulin-5	Elastic fibre-cell interface
Emilin-1	Elastin-microfibril interface
Emilin-2	Elastin-microfibril interface
Elastin binding protein	Newly secreted tropoelastin
Vitronectin	Some microfibrils in dermal tissues
Amyloid	Some microfibrils in dermal tissues
Collagen VIII	Vascular elastic tissues
Collagen XVI	Dermal microfibrils
Endostatin (C-terminus of collagen XVIII)	Vascular elastic fibres
Collagen VI	Some microfibrils

Table 1.2: Structural and associated molecules of microfibrils and elastic fibres. (Adapted from Kielty 2006). Abbreviations: β lgH3, also known as transforming growth factor- β -inducible gene-H3 and as keratoepithelin; LOX, lysyl oxidase; LOXL, lysyl oxidase-like; LTBP, latent transforming growth factor- β -binding protein; MAGP, microfibril associated glycoprotein; MFAP, microfibrillar associated protein.

**Pericellular Microfibril
Assembly**

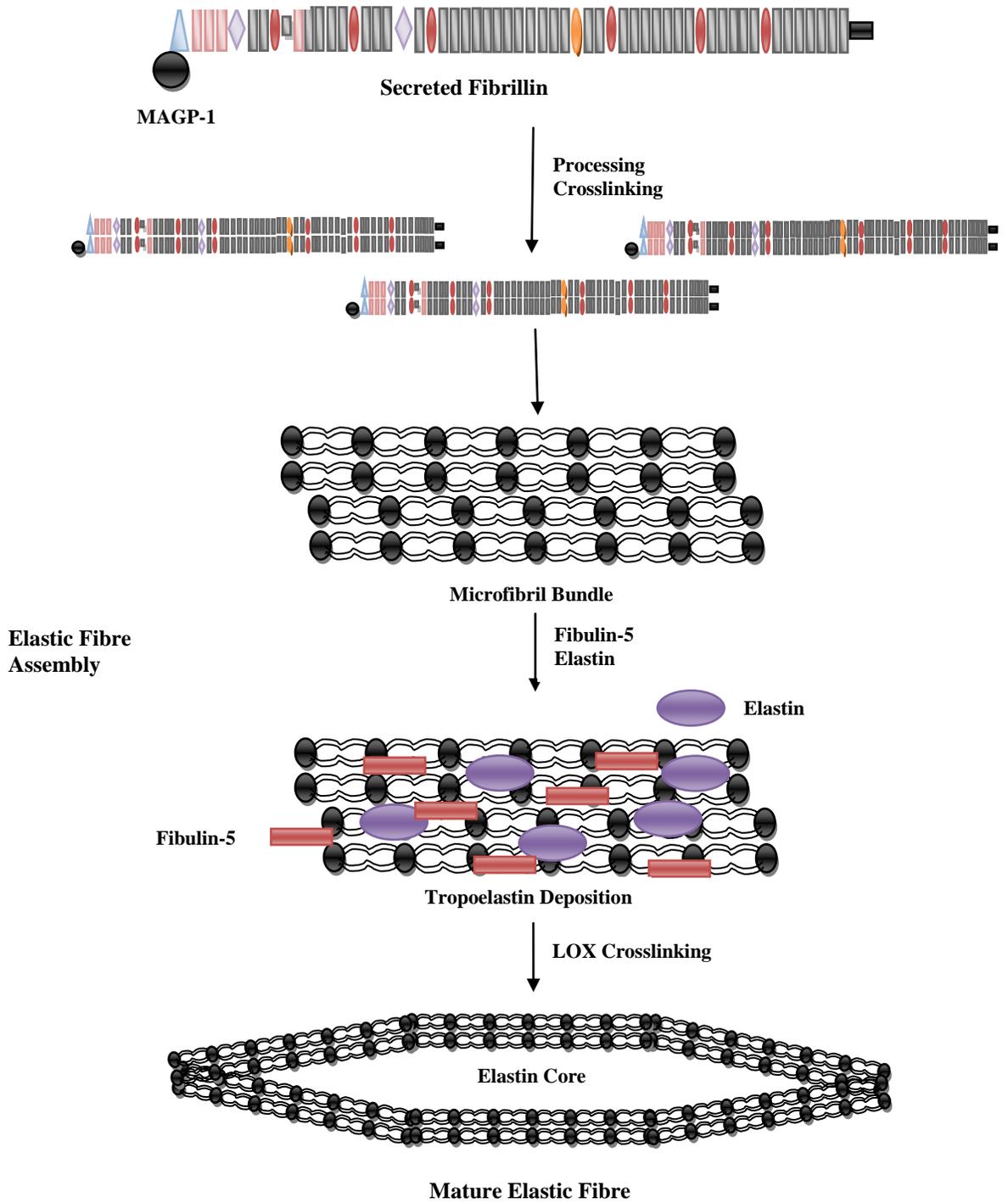


Figure 1.3: Schematic diagram of the assembly of microfibrils and elastic fibres

(Adapted from Kielty 2006)

1.4.3 Elastic fibre structure and assembly

1.4.3.1 Microfibril structure and assembly

Pericellularly secreted fibrillin-1 molecules assemble into beaded microfibrils linearly and laterally through specific N- and C-terminal interactions (**Figure 1.3**) (Cain et al., 2006, Kinsey et al., 2008, Ashworth et al., 1999a). The assembled microfibrils are then stabilised by transglutaminase crosslinks. Each microfibril probably has eight fibrillin-1 molecules in cross section (in **Figure 1.3** only two are shown for clarity).

Microfibril assembly involves interaction with many different molecules. MAGP-1 strongly binds at the N-terminal controlling N- and C-terminal interactions (Ramirez and Sakai, 2009). Heparin sulphate may have an *in vivo* role in regulating microfibril assembly by competing with MAGP-1 at the N-terminus and with tropoelastin at one of the central sites. Fibrillin-1 has been shown to interact with other molecules such as LTBP-1, fibulin-2, versican and with small chondroitin sulphate proteoglycans.

Microfibrils have been shown to be repeating globules on filamentous linear arrays using atomic force microscopy (Baldock et al., 2001, Sherratt et al., 2001), but more recent work has suggested in the physiological state they are cylindrical structures rather than beaded arrays and have inter-microfibril links (Davis et al., 2002).

1.4.3.2 Elastin fibre structure and assembly

Elastin fibres are deposited in early post-natal life in elastic connective tissues such as skin, lung, ligaments, articular cartilage, aorta and other elastic arteries (Kielty et al., 2005, Kielty et al., 2002b, Mithieux and Weiss, 2005). Tropoelastin is deposited into a fibrillin microfibril bundle template in the extracellular space (Czirok et al., 2006, Kozel et al., 2006). Elastin first appears inside microfibril bundles before coalescing to form the crosslinked elastin core of mature elastic fibres (Pasquali-Ronchetti and Baccarani-Contri, 1997, Kielty et al., 2002b). Elastin is then stabilised through formation of the elastin-specific crosslinks desmosine and isodesmosine to its insoluble form (Mithieux and Weiss, 2005). The widely accepted model is shown in **Figure 1.3**. Real time microscopy on an *in-vitro* cell culture has shown elastin globules, possibly associated with microfibrils may aggregate to form larger fibrillar structures (Kozel et al., 2006, Czirok et al., 2006). This process is coupled to cell motion, possibly through integrin bonding at the C-terminal and is known as coacervation (Ostuni et al., 2007).

Although mature elastic fibres have an outer mantle of microfibrils, some microfibrils appear to lie within the elastin core (Kielty, 2006).

Many other molecules have been associated with elastin fibres (**Table 1.2**). Lysyl oxidase (LOX) and a second isoform lysyl oxidase-like (LOXL) play significant roles in the integration process by mediating enzyme and tropoelastin binding (Thomassin et al., 2005). Cell binding may also play a role in fibre assembly (Broekelmann et al., 2005). There are two high-affinity binding sites for tropoelastin in fibrillin-1 but elastin also interacts with other molecules including MAGP-1 and biglycan (**Figure 1.3**). Fibulins 4 and 5 have critical but poorly understood roles in vascular elastic fibre formation in mice (Yanagisawa et al., 2002, Nakamura et al., 2002, McLaughlin et al., 2006). Fibulin-4 may be required to recruit LOX to facilitate tropoelastin crosslinking (Horiguchi et al., 2009). Fibulin-5 co-localises and binds with LOXL and may also have a role in elastin crosslinking (Yanagisawa et al., 2002).

1.4.4 Organisation of elastin fibres in tissue

The functional properties of elastin fibres strongly reflect the tissue-specific architecture (Kielty, 2006). In turn, this is dictated by the organisation of the microfibril template, the orientation of the cells that deposit them and the forces acting on the tissue. In the lung, elastic fibres are present throughout the respiratory tree in a fine highly branched network (Kielty et al., 2002b). In the aorta and elastic arteries, the elastic fibres form concentric fenestrated lamellar layers that intercalate with the smooth muscle. In the developing aorta, subendothelial microfibrils provide anchorage for endothelial cells and are aligned with the flow of blood (Davis, 1993a, Davis, 1993b). The skin contains an elastic fibre network varying from microfibril bundles at the dermal epidermal junction, elaunin fibres (small amounts of elastin) in the papillary dermis to thick horizontal elastic fibres in the reticular dermis (Kielty, 2006). Regions of tendon that undergo the greatest flex and strain deformation have the highest elastin content (Ritty et al., 2002). In ligament, elastin is abundant running parallel to collagen fibrils, but sparse in tendon. In cartilage, fibres surround chondrocyte lacunae and form a thin mesh running with interterritorial collagen fibrils (Kielty, 2006).

Thus the precise mechanical role of elastic fibres is thought to be a function of their chemistry, ultrastructure, arrangement and collective density relative to other ECM constituents

1.4.5 Elastic fibre functions

Elastic fibres have three recognised functions: 1) elastic recoil and resilience in dynamic connective tissue, 2) regulation of the activity of the TGF- β family of growth factors and 3) regulation of cell migration, survival and differentiation.

1.4.5.1 Elasticity

Elastin fibres comprised of both microfibrils and cross-linked elastin were an essential evolutionary advance to support vertebrate high-pressure circulatory systems and other elastic functions (Faury, 2001). Elastin fibres influence mechanical properties such as resilience and low-strain stiffness (Kielty et al., 2002c). Deformation of elastin acts as an energy store that is then used to drive recoil to the resting state (Gosline et al., 2002). Elastin is extremely insoluble due to extensive lysyl-derived crosslinks, and the crosslinked elastin core of the fibre provides the major contribution to tissue elasticity (Mithieux and Weiss, 2005).

Individual microfibrils and microfibril bundles also have elastic function (Sherratt et al., 2003). Individual microfibrils were shown to have a Young's modulus approximately two orders of magnitude stiffer than elastin and are highly resistant to axial tension (Glab and Wess, 2008). Elasticity in microfibril-containing tissue may arise primarily from reversible alterations in microfibril bundle reorganisations while individual microfibrils might act as reinforcing fibres (Sherratt et al., 2003). Untensioned microfibrils have a regular periodicity of approximately 56 nm, but isolated extended microfibrils up to 150nm have also been observed (Keene et al., 1991, Baldock et al., 2001). One study suggested a reversible extension of 56 to 100nm with irreversible extension above this (Baldock et al., 2001). Calcium and water are important in reversible elasticity (Haston et al., 2003, Wright et al., 1999).

1.4.5.2 TGF- β family activation

Fibrillin microfibrils may play a key role in the extracellular regulation of TGF- β activation and signaling (Charbonneau et al., 2004). The TGF- β family of growth factors are powerful regulators of cell survival, proliferation and differentiation, tissue morphogenesis and cellular responses to injury (Feng and Derynck, 2005). It remains unclear precisely how microfibrils regulate TGF- β . Current ideas include structural relationships with LTBPs, emilin-1 and possibly direct growth factor binding (BMPs).

1.4.5.3. Cell adhesion

Elastic fibres play an important role in cell-matrix interactions in elastic connective tissues. Interactions are mediated mainly through integrins, which are transmembrane receptors, that recognise RGD motifs within matrix molecules and link the vascular ECM directly to the cellular cytoskeletal framework (Mould and Humphries, 2004). Such integrin-mediated cell-elastic fibre interactions influence cell survival, phenotype, proliferation, migration and ECM expression and deposition. Major elastic fibre molecules involved are

a) **Elastin:** Cells bind through elastin binding protein which binds the tropoelastin hexapeptide VGVAPG. Interactions of VGVAPG peptides with this G-protein-coupled receptor stimulate actin polymerisation thereby influencing cell proliferation and migration (Karnik et al., 2003a, Karnik et al., 2003b). Certain elastin proteolytic fragments are highly chemotactic.

b) **Fibrillin-1:** A single RGD motif on fibrillin-1 mediates adhesion, cell behaviour and gene expression through integrin binding of human smooth muscle (Kielty, 2006). An RGD motif in fibrillin-2 is similarly active.

c) **Fibulin-5:** Fibulin-5 interacts directly with vascular cells in a RGD and cation-dependent manner which may contribute to its roles in elastic fibre deposition and modulation of smooth muscle phenotype (Chu and Tsuda, 2004).

1.4.6 Elastic fibre production and degradation

Fibrillins -1 and -2 are produced at low levels throughout life in human skin (Ashcroft et al., 1997). Elastin production peaks near birth and is nearly completely repressed by maturity (Perrin and Foster, 1997). Production is repressed in the adult cell mainly through post-transcriptional regulation. Expression of tropoelastin can be promoted (IL-1 β , IL-10, IGF-1) or repressed (TNF- α , bFGF) at the transcriptional level, as well as being promoted through stabilisation at the post-transcriptional level (TGF- β 1) (Duca et al., 2004). Strong evidence now exists that EFs are not replaced in adult life, and must function for the lifetime of the organism (Sherratt, 2009). As a result of this longevity, EFs are vulnerable to age-related damage (Ritz-Timme et al., 2003).

Crosslinked elastin is highly insoluble and resistant to degradation. However insoluble elastin is readily degraded by the serine proteinases (Novinec et al., 2007) and matrix metalloproteinase (MMP) classes especially MMPs -2, -7, -9 and -12 and neutrophil elastase

(Mecham et al., 1997, Ashworth et al., 1999b). Furthermore, EF morphology may be significantly altered through the action of MMP-14, leading to a loss of mechanical function (Sherratt, 2009). Elastin peptides produced enzymatically are able to influence the behaviour of a wide variety of cells including fibroblasts, macrophages and neutrophils (Duca et al., 2004). Elastin peptides are transduced through the elastin laminar receptor 1 (ELR1), a spliced variant of β -galactosidase (Hinek et al., 1993). Peptides containing VGVAPG and GXXPG sequences have been shown to bind to ELR1 (Brassart et al., 2001, Duca et al., 2004). They can directly influence chemotaxis, proliferation, protease release and even induce apoptosis (Duca et al., 2004, Privitera et al., 1998). Elastin degradation products and TGP- β 1 promoted myofibroblastic and osteogenic differentiation in dermal fibroblasts (Simionescu et al., 2007). Degraded elastin fragments have been suggested to contribute to the degenerative cascade through activation of matrix MMPs (Yu et al., 2007).

1.5 Proteoglycans (PG)

PGs make up around 1% of the dry weight of ligaments (Frank, 2004). PGs are proteins containing one or more glycosaminoglycan (GAG) chains. Most exist as aggregates and are non-covalently bound to a long chain of hyaluronate with link-protein (Heinegard and Hascall, 1974) They can also be small and non-aggregating.

1.5.1 Large aggregating PGs

The large aggregating PGs create a large osmotic swelling within cartilage, creating a water-swollen matrix critical to the properties of cartilage (Kiani et al., 2002). Aggrecan is the most abundant PG in cartilage and is required for chondrocyte survival and attachment in vitro (Lee et al., 2000). Versican is another important large proteoglycan with a wide range of actions. Versican forms an integral part of a pericellular matrix that organises the tendon cells in linear arrays between collagen fascicles (Ritty et al., 2003). It has a significant role in regulating cell phenotype (Kinsella et al., 2004) and may be involved in chondrogenic change in tissue (Zhang et al., 2001, Erdelyi et al., 2005).

In human Achilles tendon aggrecan mRNA was expressed in fibrocartilage and versican mRNA in the tendon midsubstance. This finding suggests the differing expression is a result of differing forces on the tendon: tension in the substance and compression in the insertion (Waggett et al., 1998). Versican expression in tendon has been shown to be site specific; predominant in tensile regions, while aggrecan is seen more in fibrocartilaginous regions or

where tendon wraps around bone (Waggett et al., 1998, Robbins and Vogel, 1994). The formation of fibrocartilage in ligaments and tendon has been characterised by the upregulation of aggrecan, versican, biglycan and type II collagen (Benjamin and Ralphs, 1998, Vogel and Meyers, 1999). Versican has been found to interact with fibrillin-1 and co-localisation of versican with microfibrils within the interlamellar space has been noted. Versican, along with other proteoglycans such as decorin and biglycan may play a critical role in the integration of microfibrils into the ECM (Kielty et al., 2002a).

1.5.2 Non-aggregating PGs

Small leucine-rich proteoglycans: This family of PGs are involved in the organisation of tissue through orientating and ordering collagen fibrils and elastic fibres during ontogeny, wound healing, tissue repair and tumour stroma formation (Iozzo, 1997). They are characterised by a core protein dominated by leucine-rich repeats, although they are genetically distinct (Heinegard, 2009, Neame et al., 2000). Decorin, biglycan and fibromodulin are all associated with elastic fibre assembly (Kielty, 2006).

Heparan sulphate proteoglycans: These may be involved in mediating interactions between cells and matrix molecules (Hook et al., 1986). One member, perlecan, is found in the basement membrane and may play a role in charge selectivity in glomerular filtration and is associated with microfibrils (Yanagishita, 1993).

1.6 Glycosaminoglycans (GAGs)

GAGs are generally comprised of repeating disaccharide units containing an uronic acid (glucuronic acid or iduronic acid) and an N-acetylated sugar (N-acetyl glucosamine or N-acetyl galactosamine) (Funderburgh, 2000, Trowbridge and Gallo, 2002). At least one of the repeating units has a carboxylate or sulphate group, except in hyaluronan, which is not sulphated. They are synthesised intracellularly and sulphated prior to excretion.

Hyaluronan: Hyaluronan serves not only as a backbone for the large aggregating PGs, but aids in tissue hydration and reducing friction in moving surfaces. The smaller glycoproteins bind and link collagens to the cell surface and may also have an important role regulation of growth factors.

Chondroitin Sulphate (CS): This GAG is involved in articular and bone metabolism by controlling cartilage matrix integrity and bone mineralisation. It is comprised of disaccharide

repeats of glucuronic acid and N-acetyl glucosamine. The sulphate is present in every carbon 4 or 6 of the galactosamine of every disaccharide. Each side chain is 40-50 disaccharides, although there is some tissue variation. The disaccharide chain is O-linked to protein core via a serine-xylose-galactose-galactose structure (Bali et al., 2001).

Dermatan Sulphate (DS): This is a variant of chondroitin sulphate in which some of the glucuronic acid residues have become epimerised to iduronate. DS containing proteoglycans (DS-PG) are distributed in the ECM of tendon, cartilage, other connective tissues, skin and sclera (Rosenberg et al., 1986).

Keratin Sulphate (KS): KS comprises disaccharide repeats of galactose and N-acetyl glucosamine. Glucosamine is sulphated, but additional groups may be present. The length of the side chains varies. KS differs from other GAGs in that it does not contain an uronic acid.

1.7 Other ECM components

Water forms two thirds of a ligament and is essential for cell and matrix function, lubrication and viscoelastic properties (Amiel et al., 1995a).

EXTRACELLULAR MATRIX PHYSIOLOGY

The ECM undergoes continuous synthesis and degradation of its components. The balance achieved can reflect the mechanical and physiological environment as well as any tissue pathology. ECM degradation can be physiological in the case of uterine and mammary gland involution or adaptation of connective tissues to mechanical environment (Alexander and Werb, 1991). Degradation occurs in many pathological processes, and is a complex enzymatic process (Ries and Petrides, 1995). ECM remodeling is a fine balance between activation and inhibition of proteolytic and synthetic enzymes (Mauviel, 1993).

Proteolytic enzymes are the primary cause of ligament, cartilage and subchondral bone degeneration in joint disease (Cawston, 1998). Based on the amino acid or chemical group at the catalytic domain, the proteinases can be divided into four main groups, the metalloproteinases and the cysteine-, aspartate and serine-dependent proteases (Barrett, 1994). Collagen is very resistant to proteolytic attack. As crosslinks and close association with proteoglycans make it a poor target. Lysis is thought to begin extracellularly at a neutral pH through the action of the serine and metalloproteinases. This renders it susceptible to

proteolytic attack from other enzymes or phagocytosis and intracellular destruction (Klein and Bischoff, 2010). In this introduction we will concentrate on the MPs and cysteine proteases.

1.8 Metalloproteinases (MP)

MPs are part of a superfamily of zinc dependent proteases known as metzincins named after the conserved Met residue and zinc ion at the active site (Klein and Bischoff, 2010). Although there are many subfamilies, the matrix metalloproteinases, membrane-type MMPs and adamalysins are the most relevant in connective tissue.

1.8.1 Matrix metalloproteinases (MMPs)

MMPs are grouped by common properties as they share common sequences of amino acids, are secreted by inactive zymogens and are activated either proteolytically or by mercurial agents (Cawston, 1998). They are categorised into five subgroups:

- a) Collagenases (1, 8 and 13),
- b) Gelatinases (2 and 9),
- c) Stromelysins (3, 10 and 11),
- d) Membrane type-matrix metalloproteinases (MT-MMPs) (14-17, 24 and 25),
- e) A heterogenous group including matrilysin (7) and macrophage metalloelastase (12).

The important groups in ligament ECM turnover are the collagenases, gelatinases, stromelysins and MT-MMPs (Harper et al., 1989, Foos et al., 2001). Regulation of MMPs is a complex and poorly understood interaction of inflammatory cells, cytokines, growth factors and ECM molecules. Regulation of MMPs can occur at several stages including transcription, activation of proMMPs (Murphy and Docherty, 1992) and inhibition of active MMPs (Nagase, 1997).

Gelatinases (MMP-2 and -9) can cleave collagens type IV, V, VII and X, denatured collagens (gelatins) and elastin, but not fibronectin or laminin (Tyagi, 1997). MMP-2 is produced by all connective tissue cells making it the most widespread of all MMPs (Murphy and Crabbe, 1995). It is thought to be involved in ECM turnover through remodeling or removal of denatured collagen (Creemers et al., 1998, Kerkvliet et al., 1999, Yoshizaki et al., 2002, Morgunova et al., 2002). MMP-9 is of a similar structure to MMP-2 and is expressed in inflammation, angiogenesis and tissue repair (Tarlton et al., 1997).

1.9 Cysteine proteinases

These are subdivided into around 20 families, the main ones being calpain and papain. In inflammation, macrophages secrete cathepsins B, K, L and S, which are also identified as major damagers to tissue in chronic inflammation (Kielty et al., 2002b). Studies have implicated K, L and S in extensive degradation of elastic fibres that accompanies cardiovascular disease in man. Cathepsins have been shown to degrade elastin, and have preferences for particular tissue types (Novinec et al., 2007). Cathepsin S has a critical role in antigen presentation (Muir et al., 2005b). Cathepsin K has been associated with ACL rupture, and its expression appears to precede rupture (Barrett et al., 2005, Muir et al., 2005b, Muir et al., 2006). Caspases are another important family of cysteine proteases that play important roles in apoptosis, development and inflammation (Martinon et al., 2009).

1.10 Serine Proteases

These are the largest family of endopeptidases and are released from inflammatory cells (Milner et al 2001). The most important are those of the plasminogen/ plasminogen activator (PA) system (Murphy et al., 1993). Other serine proteases include cathepsin G, primarily intracellular in neutrophils and macrophages, and neutrophil elastase from neutrophils and active extracellularly, with broad spectrum action on collagen type III, elastin, PG, vitronectin and laminin (Doring, 1994).

1.11 Other collagenolytic agents

During chronic inflammation, reactive oxygen species generated by tartrate-resistant acid phosphatase (TRAP) as well as MMPs are important components of leucocyte-mediated collagenolysis. TRAP can also fragment the triple helix of collagen through generation of reactive oxygen species by the Fenton reaction (Muir et al., 2005b). Cysteine proteases, such as K, have a regulatory role in TRAP degradative activity by converting TRAP5a to the more active TRAP5b. Cathepsin K and TRAP cells have been co-localised within the ACL and are associated with ACL disease (Muir et al., 2002).

1.12 Role of proteinases

1.12.1 Function and mode of action

Traditionally, MMPs have been thought of as enzymes for degradation of structural components of the ECM. However, proteolysis can have other functions (Page-McCaw et al., 2007):

- Creation of space for cell migration
- Production of substrate-specific cleavage fragments with independent biological activity
- Regulation of tissue architecture through effects on ECM and intercellular junctions
- Direct and indirect activation, deactivation and modification of signaling molecules

1.12.2 Involvement of proteinases in degenerative conditions

There is considerable evidence for the role of proteases in joint diseases. Proteases have an important role in canine, equine and human OA (Wernicke et al., 1996, Clegg et al., 1997, Fernandes et al., 1998, Riley et al., 2002). Proteases are also important in diseases affecting ligaments and tendons. Elevated levels of proteases have been found in knee synovial fluid in ACL damage, meniscal damage, post-traumatic OA and pyrophosphate arthritis (Lohmander et al., 1993). The elevation occurs within days and can remain elevated for up to 18 years (Lohmander et al., 1994). However an elevation in proteases has also been noted in older animals without ACL damage (Muir et al., 2005a). MMP-2 has been implicated in the turnover of the periodontal ligament (Creemers et al., 1998). In the rabbit ACL an increase in collagenase activity has been correlated with a loss of collagen, a loss of cellularity and matrix disorganisation (Amiel et al., 1989). An increase in MMP-2 and -9 was seen in the ruptured ACL (Muir et al., 2005a) but not in an earlier study (Spindler et al., 1996b). An increase in proMMP-2 in the intact ACLs of a dog breed at high risk of ACL rupture was seen when compared to a dog breed with a very low risk to ligament rupture (Comerford et al., 2005). An association between elevation in proMMP-2 and AP laxity has been suggested (Comerford et al., 2005, Quasnichka et al., 2005). Mechanical stress can induce increased production of MMPs 2 and 9 and acid cathepsins in affected tissue in Dupuytren's disease which may contribute to tissue weakening (Tarlton et al., 1998).

CURRENT THOUGHTS ON THE AETIOPATHOGENESIS OF CANINE CRUCIATE LIGAMENT RUPTURE

1.13 CL idiosyncrasies

The CLs are unique for many reasons. Their trans-articular location is shared only by one other ligament in the body, the ligament of the femoral head also known as the teres ligament. However, although both these ligaments experience multiaxial stresses, only the CL complex is subject to intrinsic compression where the ACL wraps around the PCL (Arnoczky and Marshall, 1977). The transarticular location means the soft tissue support of the CL complex is minimal, comprising a thin layer of epiligamentous tissue (Clark and Sidles, 1990). As blood supply to the CL complex arises from the epiligament, the blood supply is also subject to compromise through epiligament damage or compression (de Rooster et al., 2006). Differences exist in the properties of the ACL and MCL on the knee in many aspects such as response to loading and injury (Amiel et al., 1984, Amiel et al., 1995b, Lo et al., 2002b).

1.14 Histologic and ultrastructural changes in CL disease

1.14.1 Histologic overview

An early study of partially and completely ruptured ACLs showed an absence of bundle structure and hyalinisation at the point of ligament failure (Paatsama, 1952). A later study demonstrated similar changes in dogs over 8 years of age without rupture of the ACL with calcium deposits, hyalinisation and loss of normal fibre architecture described (Zahm, 1965). Subsequent studies have described deterioration in collagen architecture and a fibrocartilagenous appearance of the matrix, **Figure 1.4** (Vasseur et al., 1985, Narama et al., 1996, Comerford et al., 2006b). Cell changes are characterised by loss of cell density, cells becoming rounded and undergoing chondroid transformation, and formation of chains of chondroid cells (Vasseur et al., 1985, Narama et al., 1996, Comerford et al., 2006b). Enlargement of CL fibroblast nuclei and perinuclear halo formation has been described in dogs kept under laboratory conditions (Narama et al., 1996), but lack of exercise may have influenced these results as immobilisation can alter the biochemical composition of connective tissue including increasing mucopolysaccharide content (Akeson et al., 1973) Changes are most readily observed at the mid substance where the ACL wraps around the PCL. Furthermore these changes were associated with a loss of mechanical properties (Vasseur et al., 1985). Changes in the PCL are also observed and although similar, are less severe

(Vasseur et al., 1985). Decreased cellularity has been described in the core of ruptured ACL, with an increase in spheroid cells and a chondroid transformation, and is thought to be driven by apoptosis (Hayashi et al., 2003b, Krayter et al., 2008, Gyger et al., 2007). A grading system for these changes has been developed (0=no changes, 1=mild, 2=moderate and 3=severe degenerative changes) which positively correlated with increasing age (over 5 years) and increasing weight (over 15kg), but not with gender (Vasseur et al., 1985).

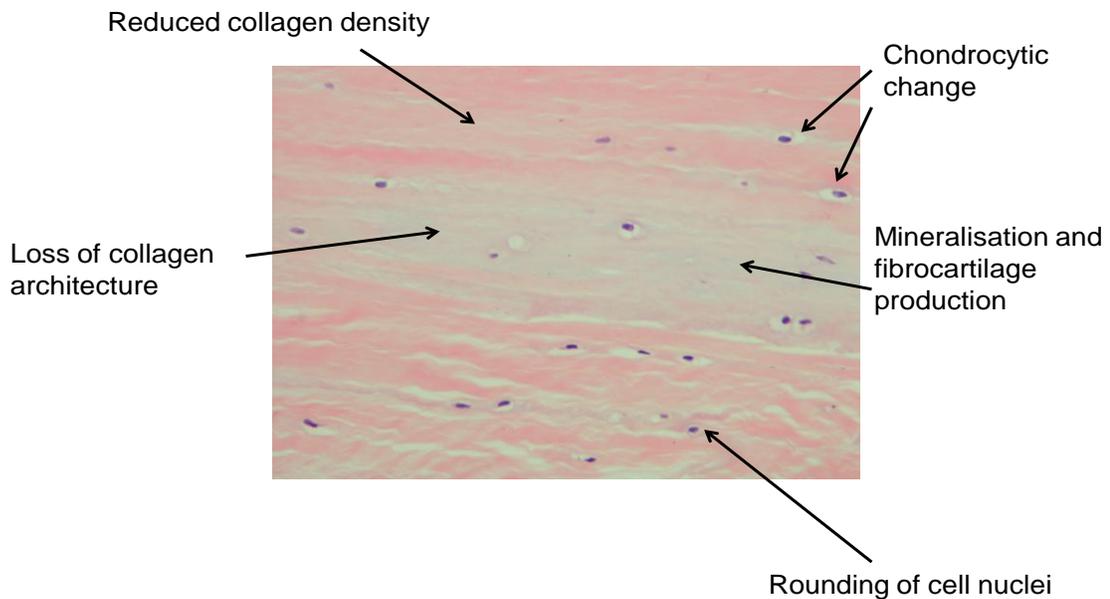


Figure 1.4: Typical changes associated with degeneration of the ACL. LR, x40, H&E

The fibrocartilaginous change described within the midsubstance of the ACL may represent a degenerative or adaptive change (Zahm, 1965, Vasseur et al., 1985, Comerford et al., 2006b). The fibrocartilage is thought to form in response to either the ligament compression as it wraps around the PCL or from the anteriomedial and posteriolateral bands tightening on each other (Comerford et al., 2006b). The mechanism for these changes is not understood, but may be a protective metabolic response to hypoxia or metabolic stress suggested for OA cartilage (Connor et al., 2001, Schipani et al., 2001, Grimshaw and Mason, 2001), an adaptation to compressive load (Benjamin and Ralphs, 1998) or a degenerative change preceding ACL

rupture (Vasseur et al., 1985). Fibrocartilage forms in breeds of dog at risk of ACL rupture (e.g. LR), in those with a low incidence (e.g. GH) and in older dogs in which the ACL remains intact (Comerford et al., 2006b, Vasseur et al., 1985, Zahm, 1965). However it may be produced in response to repair to stress/exercise in the low risk breeds and therefore may be a positive finding in these dogs (Comerford et al., 2006b).

1.14.2 Cell population changes

Inflammation of the synovium and of the ACL epiligament develops in the early phase of the naturally occurring disease and may precede knee instability and rupture of the ACL (Muir et al., 2005b). Osteophyte production is mediated by the synovial macrophage and osteophytes are often present at the time of surgery, suggesting synovitis precedes ACL rupture (Muir et al., 2005a). Degenerative changes are often found in the contralateral joint at the time of surgery. Inflammation of the epiligament in patients with rupture of the ACL has been demonstrated, with a significantly increased number of cathepsin K⁺ and TRAP⁺ macrophage-like cells (Muir et al., 2005b). Synovitis characterised by lymphocytic-plasmocytic infiltration has been described in a majority of dogs with ACL rupture, with a mononuclear synovitis present in the remainder (Galloway and Lester, 1995).

Higher numbers of TRAP⁺ macrophages were seen in ruptured canine ACLs than in intact canine ACLs or ruptured human ACLs suggesting the pathogenesis of ACL rupture may differ to human ACL rupture (Barrett et al 2005). In the human ACL, three distinct fibroblast morphologies, fusiform, ovoid and spheroid, have been described based on nuclear aspect ratios (Murray and Spector, 1999). Regional variation in their distribution was noted giving rise to three distinct proximal, middle and distal zones (Duthon et al., 2006). Ovoid, chondroid cells were noted proximally and distally with abundant spindle-shaped fibroblasts in the middle part of the PCL. This pattern differed from that described in the dog where rounded cells were found more commonly in the middle part of the ACL (Vasseur et al., 1985). Furthermore, fibrocartilage formation was noted in the proximal human ACL where it is in contact with the intercondylar notch, but not in the middle part of the ACL as seen in the dog (Duthon et al., 2006, Paatsama, 1952).

1.14.3 Collagen changes

Comparing ruptured ACLs of Labrador retrievers (LRs, a breed of dog with increased risk of ACL rupture) to ACLs from intact LRs, significant differences were seen in immature

crosslinks, with the ruptured ACL dogs having a higher concentration. The ratio of immature to mature was 4:96 in intact and 57:43 in ruptured ACL dogs (Comerford et al., 2004). No significant differences in total collagen content were seen between intact ACLs and ruptured ACLs in the LR (Comerford et al., 2004). LRs had a higher proportion of smaller fibres whereas GH had a broader distribution, and the fibre diameter and pattern of fibre distribution was significantly different indicating greater turnover of fibrils in the LR, although exercise may have influenced this result (Comerford et al., 2006b).

In a guinea pig model of spontaneous OA, the Dunkin-Hartley strain develops CL laxity prior to the onset of OA. It was proposed that this laxity was an initiating event in OA and may result from increased turnover of collagen (Quasnichka et al., 2006). An increase in type II collagen as well as altered PG deposition was noted in the midsubstance of the PCL prior to onset of OA in the Dunkin-Hartley strain (Young et al., 2002). Collagen type has yet to be studied in the canine CL complex but it is assumed the fibrocartilaginous changes seen will have an increased collagen type II relative to normal ligament.

1.14.4 GAG and PG changes

In ruptured CLs from LRs, sulphated GAG and total GAG content was significantly higher when compared to normal ACLs from the same breed (Comerford et al., 2004). However, comparing the at-risk LR to the GH prior to ACL rupture, there was no significant difference in total or sulphated GAGs (Comerford et al., 2005).

Little is known of PG changes in disease of the canine CL complex. Healing ligament has been shown to have higher total and aggregating PGs (Plaas et al., 2000, Bishop and Bray, 1993). The formation of fibrocartilage in ligaments and tendon has been characterised by the up-regulation of aggrecan, versican and biglycan (Benjamin and Ralphs, 1998, Vogel and Meyers, 1999). Decreased lubricin in synovial fluid has been associated with ACL injury in humans (Elsaid et al., 2008). 7-fold greater aggrecan content in the ACL compared to the PCL was noted in dogs, but was not related to histological changes (Valiyaveettil et al., 2005).

1.14.5 Gene expression profile

When comparing breeds at risk of ACL rupture with those considered not at risk, no significant differences were seen in mRNA expression profile in dogs with intact ACLs (Clements et al., 2008). In ruptured ACL there was an increase in protease (such as cathepsins

B and D and MMPs 2 and 9) and ECM (such as Col1a2 and Col3a1) gene expression relative to intact canine CL and these changes were similar to ruptured human ACLs (Clements et al., 2008). In the dog, mRNA profiling has revealed a ruptured ACL-specific mRNA profile in synovial fluid compared to knees with OA (Muir et al., 2007).

1.15 Concepts of aetiology

1.15.1 Weight, breed and age

Cruciate disease is likely to be multi-factorial with no single factor accounting for all aspects of its progression (Cook, 2010). Furthermore, pathogenesis may differ for different subgroups of cruciate patients (Bennett et al., 1988).

ACL rupture has an increased incidence in obese dogs (Barnes, 1977, Vasseur et al., 1985, Bennett et al., 1988, Whitehair et al., 1993). Although occurring in all sizes of dogs, larger breed animals are more commonly affected and at a younger age (Whitehair et al., 1993, Duval et al., 1999). An increased incidence of ACL rupture has been shown in certain breeds such as the Newfoundland, Rottweiler and LR (Whitehair et al., 1993, Duval et al., 1999). Other breeds, such as the Greyhound (GH), the Basset Hound and the Old English Sheepdog, are considered to have a low incidence of the disease. GHs and LRs or Rottweilers have formed the basis for comparison studies (Wingfield et al., 2000a, Wingfield et al., 2000b, Comerford et al., 2005, Comerford et al., 2006b), despite marked differences in conformation and exercise.

An increased incidence of ACL rupture has been noted in neutered animals, particularly females (Doverspike et al., 1993, Whitehair et al., 1993, Harasen, 1995, Slauterbeck et al., 2004). This may be related to an increased incidence of obesity noted in neutered females (Edney and Smith, 1986). Alternatively, endocrine changes may affect ACL metabolism (Takeda et al., 1975, Dannucci et al., 1987).

Histological changes associated with aging have been noted in ACLs from healthy dogs (Paatsama, 1952, Zahm, 1965, Vasseur et al., 1985, Bennett et al., 1988, Narama et al., 1996). These changes have been correlated with deterioration in mechanical properties (Vasseur et al., 1985).

1.15.2 Compromise of blood supply

Inadequate blood supply may contribute to the development of ACL rupture, particularly where the CLs twist on each other, the most common site for non-traumatic rupture (Paatsama, 1952, Tirgari and Vaughan, 1975b, Arnoczky et al., 1979, Vasseur et al., 1985). The failure of the ACL to undergo repair may be due in part to poor vascularity (O'Donoghue et al., 1971). ACL blood flow may be influenced by joint fluid (Kobayashi et al., 2006) so changes in joint fluid composition could conceivably affect blood flow. The observed loss of cellularity seen in the ACL core may result through the development of hypoxia from poor blood flow and the chondroid phenotype may be an adaptation to hypoxia (Hayashi et al., 2003b).

1.15.3 Cellular alterations

ACL fibroblast phenotype change is known to precede rupture in dogs (Vasseur et al., 1985, Narama et al., 1996). Pairs of spheroid fibroblasts observed were similar in appearance to clones of articular chondrocytes seen in late stage OA. The mechanism was unclear, but it may be similar to protective metabolic response to hypoxia or metabolic stress suggested for OA cartilage (Connor et al., 2001, Grimshaw and Mason, 2001, Schipani et al., 2001). Decreased cellularity in has been described in the core but not epiligament of ruptured ACLs, with an increase in spheroid cells, a chondroid transformation (Hayashi et al., 2003a). Cellular hypoxia in the epiligament following ACL rupture has been demonstrated through expression of HIF-1 α in synoviocytes (Hayashi et al., 2003b). However, there was no HIF-1 α staining in the substance of the ACL, and it was suggested that HIF-1 α may be an acute phase change missing from the chronic disease. As cells play key roles in maintenance of the ECM loss of cell number and phenotype may lead to a failure to maintain ACL matrix resulting in rupture.

1.15.4 Mechanobiological aetiologies

Mechanoresponsiveness is a fundamental feature of all living tissues, including tendons and ligaments (Wang et al., 2006). ACL fibroblasts appear to respond to their mechanical environment (Hannafin et al., 2006) and mechanical force may be required for cell arrangement (Henshaw et al., 2006). These responses are mediated by the integrin family of cell surface receptors which link the extracellular matrix to the interior of the cell through the cytoskeleton (Janmey, 1998). Tendinopathy has been traditionally considered to occur through a repetitive mechanical loading below the injury threshold of the tendon leading to an inadequate and degenerate cell-matrix response. This results in transient weakness in the

tendon and the continued loading exceeds the tendon's healing capacity leading to an overuse injury (Jones et al., 2006). In this model, cell overstimulation is thought to initiate the degenerative cascade leading to tendinopathy (Wang et al., 2003). However, *in-vitro* studies have shown that mechanobiological under stimulation of tendon cells can produce a pattern of catabolic gene expression that results in ECM degeneration and subsequent loss of tendon material properties (Egerbacher et al., 2008, Lavagnino et al., 2006). At extremes of physiologic loading, fibril damage can occur which alters normal cell-matrix interaction, which in turn can lead to under stimulation of tendon cells leading to catabolic changes (Lavagnino et al., 2006). This has been proposed as the mechanism for tendinopathy (Arnoczky et al., 2007). A similar mechanism may be possible in the ACL where cell-matrix coupling is compromised leading to under stimulation of fibroblasts and a degenerative cascade.

1.15.5 Ligament response to disease or injury

Conventional ligament healing occurs in three phases of healing: inflammation, proliferation and matrix remodelling (Shrive et al., 1995). However following healing, differences in structure and function persist such as altered proteoglycan (Plaas et al., 2000), altered collagen types (Amiel et al., 1987), failure of collagen crosslinks to mature (Frank et al., 1995), persistence of small diameter collagen fibrils (Frank et al., 1997), altered cell connections (Lo et al., 2002b), increased vascularity (Bray et al., 1996) and incomplete resolution of matrix 'flaws' (Frank, 2004).

Poor healing in the ACL has been noted by a number of authors, suggested as a reason for high failure rate of primary repair of ACL rupture (Arnoczky et al., 1979, Frank et al., 1985). Studies comparing human ACL and medial collateral ligament (MCL) following rupture have shown similar cell proliferation, cell migration and vascularisation in both ACL and MCL (Woo et al., 2000, Murray et al., 2000, Frank et al., 1983). However a provisional scaffold (fibrin-platelet plug) connecting the ruptured ends in the ACL was missing. Thus, although ACL cells may be biologically capable of healing the ACL, they are unable to fill the wound site. Subsequently a model using central ligament defects was created in the dog that was stable enough to allow potentially provisional scaffold formation and facilitate comparison of ACL and MCL healing responses (Spindler et al., 2006). The ACL, in comparison with the MCL, showed lack of provisional scaffold formation as well as reduced levels of key ECM proteins and cytokines within the wound (Murray et al., 2007). The failure of the provisional

scaffold in the ACL may result from increased urokinase plasminogen activator (uPA) within the joint following trauma, a response thought to prevent arthrofibrosis (Rosc et al., 2002). This premature loss of the provisional scaffold has been proposed as a mechanism for failure of the ACL to heal (Murray, 2009).

Other mechanisms proposed for poor ACL healing include deficiencies in stimulation or intrinsic deficiencies of cell migration and proliferation (Nagineni et al., 1992, Geiger et al., 1994, Amiel et al., 1995b, Schreck et al., 1995, Spindler et al., 1996c).

ELASTIC FIBRE -ASSOCIATED DISEASE

Failure of elastic fibres has been implicated in many disease processes (Kielty, 2006). Disease can arise from failure to form or function properly or through degeneration and be heritable or degenerative.

1.16 Heritable disorders:

1.16.1 Marfan syndrome (MFS) and related fibrillinopathies

MFS, an autosomal dominant hereditary disorder of connective tissue manifests as major cardiovascular, ocular and skeletal defects notably laxity of ligaments. It is caused by mutations in the gene for fibrillin-1, and causes premature death usually by acute aortic dissection (Robinson et al., 2006). Mutations in fibrillin-1 may cause MFS as a direct consequence of altered or reduced secretion or assembly of mutant molecules, and increased susceptibility of defective fibrillin-1-containing microfibrils to proteolytic damage. MFS has also been associated with excess TGF- β signaling. This results in developmental defects such as mitral valvulogenesis, as well as increased cell proliferation and altered ECM deposition and turnover (Neptune et al., 2003). Mutations in TGF- β receptors have been associated with aortic and cardiovascular defects with overlapping phenotypicity to MFS. Mutations in fibrillin-2 cause congenital contractual arachnodactyly, characterised by flexion contractures of tendon, arachnodactyly, kyphoscoliosis, abnormal pinnae and muscular hypoplasia (Robinson et al., 2006).

1.16.2 Supravalvular aortic stenosis (SVAS) and Williams-Beuren syndrome (WBS)

SVAS is inheritable in an autosomal dominant manner or as part of the complex developmental disorder WBS. The symptoms of SVAS are similar in both cases and involve decreased deposition of elastin associated with increased vascular cell proliferation (Li et al.,

1998). SVAS is caused by mutations in the elastin gene, and WBS by multiple microdeletions in up to 28 genes.

1.16.3 Cutis laxa

Cutis laxa is a heterogeneous group of disorders characterised by excess, sagging and inelastic skin (Milewicz et al., 2000). Although acquired cutis laxa is caused by dermal inflammation and associated elastic fibre degeneration, its pathogenesis involves an underlying genetic susceptibility. The interaction of specific elastin and fibulin-5 gene alleles (elastin, heterozygous for alleles A55V and G773D and the fibulin-5 allele G202R) render elastic fibres susceptible to inflammatory destruction (Hu et al., 2006). It is an unusual disease illustrating a tissue specific phenotype through failure of a molecule widely distributed.

1.16.4 Other conditions

Other heritable conditions associated with elastic fibre disease include age-related macular degeneration (mutations in fibulin-5), (Lotery et al., 2006), pseudoxanthoma elasticum (mutation in ABCC6), (Miksch et al., 2005) and arterial tortuosity syndrome (mutation in SLC2A10) (Coucke et al., 2006).

1.17 Degenerative disorders

All the major elastic fibre molecules may be degraded by proteases of the serine and MMP classes (Ashworth et al., 1999b). Consequently loss of elastic fibre architecture and function is a pathological feature of a number of degenerative and inflammatory diseases of man, including pulmonary emphysema and COPD, vascular aneurysm and photo- and chrono-aged skin. In damaged tissues the proportion of microfibrils to elastin declines and then the elastin core is degraded (Lewis et al., 2004).

1.17.1 Vascular proliferative disease

Vascular proliferative diseases are a diverse group of disorders that lead to arterial narrowing and arise in response to injury by multiple factors (Ross, 1995; Lusis, 2000). Elastin has a role in regulation of vascular smooth muscle cells (Karnik et al., 2003a). Loss of elastin fibres is thought to contribute to the pathogenesis of atherosclerosis (Robert et al., 1998). Inhibitors of elastin degradation may prevent pathological changes associated with atherosclerosis

(Basalyga et al., 2004). Elastin degradation products may be involved in the pathogenesis of atherosclerosis and have been reported to stimulate conversion of dermal fibroblasts to myofibroblasts, as did TGF β 1 (Simionescu et al., 2007).

1.17.2 Chronic obstructive pulmonary disease (COPD)

COPD is characterised by airflow obstruction arising from chronic bronchitis and emphysema (Devereux, 2006). Elastin degradation is a key step in the pathogenesis of COPD (Shifren and Mecham, 2006). Both neutrophils and macrophages secrete proteases that degrade elastin and other matrix proteins. Elastin fragments liberated during elastic fibre degradation recruit more inflammatory cells leading to a cycle of increasing destruction (Hautamaki et al., 1997). Oxidative stress has also been implicated in the disease, but it is thought to act by modification of protease activity rather than direct destruction of elastin (Shifren and Mecham, 2006).

1.17.3 Intervertebral disc disease (IVDD)

The elastin content of the annulus fibrosus (AF) of the human intervertebral disc has been extensively studied as the loss of elastin resilience and elasticity is a major contributing factor in IVDD. The change of disc height in compression or extension is thought to depend on the slip of adjacent lamellae (Szirmai, 1970) and there is a strong suggestion that elastin has a role in the recoil of lamellae following deformation (Yu et al., 2007). Degeneration of the annulus fibrosus was associated with a five-fold increase in elastin over that seen in healthy AF (Cloyd and Elliott, 2007) but no increase in EFs (Olczyk, 1994, Smith and Fazzalari, 2006). Furthermore, a decrease in elastin crosslinks with age (which has been shown to correlate with degeneration (Cloyd and Elliott, 2007) suggested the increase in elastin was non-fibrous in nature (Osakabe et al., 2001). In scoliotic discs, elastic fibres were sparse, and the collagen-elastic fibre networks were disorganised with loss of lamellar structure. It was suggested that normally elastin has a biochemical role in the human IVD, and its loss could be involved in the progression of the spinal deformity (Yu et al., 2005).

1.17.4 Other articular structures

In the human interspinous ligament, age-related changes were seen in the elastic fibre composition, with older patients showing an accumulation of mature elastic fibres which were short and fragmented and appeared to be arranged in clusters (Barros et al., 2002). The greater number of fragmented and degenerated fibres promoted a loss of normal compliance and there

was also a loss of oxytalan fibres. Patients with uni- and multidirectional shoulder instability had increased elastic content compared to normal shoulders, coupled with more stable and reducible collagen crosslinks and a greater mean collagen fibril diameter (Rodeo et al., 1998). Elastin plays an important role in the temporomandibular joint (TMJ); altered distribution of elastin was seen within the TMJ of operated rabbits, with increases and decreases in differing locations (Sato et al., 1998). A loss of function and altered distribution of elastic fibres followed disc displacement in the rabbit TMJ (Gu et al., 2002). Although an increase in elastin in some parts of the disc suggests that elastic fibre macroassembly is possible in the mature animal, the nature of the fibres was not categorised, and they may have been non-functional.

1.17.5 Periodontal ligament

It has been suggested that oxytalan fibres confer resilience on the periodontal ligament (Ujiie et al., 2008). The oxytalan fibres that connect the blood vessels to the teeth may cushion the pressure on the vessel walls, reducing the dislocation of vessels when the teeth occlude. Thus, the oxytalan meshwork may contribute to maintaining the integrity of the vascular system in the periodontal ligament (Chantawiboonchai et al., 1998).

1.17.6 Canine CLs

The involvement of elastin in pathology of the canine ACL has not been assessed. Elastin has been found in large amounts in the human ACL where it has been credited with permitting extreme distance changes and homogenous distribution of multiaxial stresses involved in ACL movement (Neurath and Stofft, 1992, Strocchi et al., 1992). To date, elastin has not been quantified biochemically in the cruciate ligament complex of any species. Estimates have ranged from 0-6% elastin (Suzuki et al., 2008, Paatsama, 1952, Strocchi et al., 1992).

1.18 Hypothesis and aims

The hypothesis of this thesis is that elastin has a mechanical and a biological role in the canine cruciate ligament complex. It is further hypothesised that the distribution and function of elastic fibres will vary between three breeds of dog with differing risk of ACL rupture: the greyhound with a low risk, the beagle with a low-to-moderate risk and the Labrador retriever with a high risk of ligament rupture.

1.18.1 Mechanical role for elastin in the canine CL complex

The mechanical function of elastin fibres is thought to reflect distribution in tissue (Kielty et al., 2002b). In order to examine the mechanical role of elastin in canine CL tissue it is proposed to study the distribution of elastic fibres using histology and immunofluorescence. Nomarski Differential Interference Contrast (NDIC) microscopy will be used to study CL microanatomy in unfixed, hydrated CL tissue and relate the observed anatomy to the distribution of elastic fibres.

1.18.2 Biological role for elastin fibres in the canine CL complex

In order to examine the biological role of elastin fibres in the canine CL complex, the following studies are proposed:

1. Elastic fibres are thought to be important in cell-matrix coupling. It is proposed to study cell morphology in the canine CL complex in the greyhound and Labrador retriever using histology, immunofluorescence and confocal laser scanning microscopy. The results will be related to elastic fibre distribution, as described above.
2. It is proposed to quantify elastin biochemically and compare the results with subjective histological analysis of CL degeneration and elastic fibre distribution in the greyhound.
3. Failure of elastin fibres has been associated with disease in ligament. It is proposed to use histology and immunofluorescence to study the distribution of elastic fibres in the CL complex in canine CL tissue with evidence of degenerative change and make comparisons between breeds with differing risk of ACL rupture.
4. To assess whether fragments of degraded elastin exert a biological effect on canine anterior cruciate ligament cells in an *in vitro* culture system.

CHAPTER 2: Variations in cell morphology in the cruciate ligament complex

Abstract

Outline: To describe the morphology and regional variations of cells in the cruciate ligaments (CLs) of two breeds of dog with a differing predisposition to non-contact anterior cruciate ligament (ACL) injury.

Method: Macroscopically normal CLs were harvested from dog breeds at a high (Labrador retriever) and low (greyhound) risk to ACL rupture. Antibodies against the cytoskeletal components vimentin and alpha tubulin were used to identify cell morphology. Nuclei were stained with DAPI. Following counterstaining with a fluorescent marker, images were collected using conventional and confocal microscopy.

Results: Both anterior and posterior CLs contained cells of heterogenous morphologies. Cells were arranged between collagen bundles, and frequently had cytoplasmic processes. Some of these processes were of great length, while other cells had shorter, thicker and more branched processes, and some cells had no processes at all. Processes were frequently shown to contact other cells, extending longitudinally and transversely through the CLs. Cells with longer processes had fusiform nuclei, and cells without processes had rounded nuclei. The cells with rounded nuclei were more frequent in the midsubstance of both CLs. Cells with long processes were apparently more commonly noted in the CLs of the breed at low risk of ACL rupture.

Conclusions: Local variation in cell types may reflect the complex biomechanics of the CLs. Contact between cells may facilitate direct communication. Differences in cell morphology between breeds with differing risk of ACL rupture may reflect fundamental differences in CL physiology possibly through altered cell-to-cell communication.

2.1 Introduction

Ligaments consist of sparse fibroblasts surrounded by dense collagenous matrix. The extracellular matrix (ECM) determines the mechanical properties of ligament and the cells maintain the matrix, controlling responses to altered mechanical load and injury (Frank, 2004). The mechanical environment of the cell has been shown to influence cell morphology in other normal connective tissues (Giori et al., 1993, Matyas et al., 1994, Ralphs et al., 1998). Cells in tensile load in the absence of significant compressive load have cytoplasmic processes which may be long and extend parallel perpendicularly or transversely through the collagen fibres (Bruehlmann et al., 2002, Lo et al., 2002a). The detection of gap junctions in association with these cell connections suggests the potential to coordinate cellular and metabolic responses throughout the tissue through cell-to-cell communication (Bruehlmann et al., 2002, McNeilly et al., 1996, Lo et al., 2002b). This elaborate three-dimensional structure has been termed the cellular matrix, and has been described in tendon (McNeilly et al., 1996, Ralphs et al., 1998), meniscus (Hellio Le Graverand et al., 2001a), intervertebral disc (Bruehlmann et al., 2002) and medial collateral ligament (MCL) of the knee (Lo et al., 2002b). The cellular matrix is dynamic, and changes have been noted in healing (Lo et al., 2002b), injury and degenerative joint disease (Hellio Le Graverand et al., 2001d).

Cruciate ligaments (CLs) are the primary stabilisers of the knee joint. The two components are defined as anterior (A) and posterior (P) CLs (Arnoczky and Marshall, 1977). Groups of collagen fibres are referred to as bundles, divided by interbundle regions and bundles are grouped as fascicles, divided by interfascicular regions (Clark and Sidles, 1990). Within the knee joint, the ACL and PCL are in intimate contact, with the ACL twisted around the PCL. They are considered to function as a unit hence the term CL complex (Arnoczky and Marshall, 1977, Harari, 1993). The existence of a cellular matrix has been demonstrated in the ovine ACL (Lo et al., 2002b). It is not known whether this cellular matrix exists in the CLs of other species such as the dog nor if there is any variation between population subtypes, between the ACL and PCL or regionally within each ligament.

Cellular changes in the canine ACL include apoptosis and metaplasia of surviving fibroblasts to chondrocytes (Vasseur et al., 1985, Krayer et al., 2008). These changes have been described as degenerative and are associated with deterioration in collagen architecture, with a fibrocartilaginous appearance of the matrix and loss of mechanical properties. They are most readily observed at the mid substance where the ACL twists around the PCL (Vasseur et al.,

1985, Narama et al., 1996, Comerford et al., 2006b). Changes in the PCL are also observed, and although similar, are less severe (Vasseur et al., 1985). The mechanism for these changes is not understood, but may be a protective metabolic response to hypoxia or metabolic stress suggested for OA cartilage (Connor et al., 2001, Schipani et al., 2001, Grimshaw and Mason, 2001), an adaptation to compressive load where the CLs twist (Benjamin and Ralphs, 1998) or a degenerative change preceding ACL failure (Vasseur et al., 1985). Although such changes have been described in both the Greyhound (GH) and Labrador Retriever (LR), ACL failure is extremely rare in the GH but common in the LR (Whitehair et al., 1993).

Cruciate ligaments are subject to complex multiaxial stresses (Takai et al., 1993, Kennedy et al., 1976), yet little is known of the cells responsible for maintaining the ligaments. Previous canine CL studies have used limited histochemical staining to assess the cell population (Vasseur et al., 1985, Narama et al., 1996, Paatsama, 1952). In this study, we use histology, immunofluorescence and confocal laser scanning microscopy to methodically detail cell morphology in the canine CL complex. By reporting variations in anatomy of cells in the CL complex in two breeds of dog, at altered risk of ACL failure, we hope to provide a basis from which to understand the physiology of these complex structures in health and disease.

2.2 Materials and methods

2.2.1 Sample collection and preparation

ACL and PCLs from both knee joints were harvested by sharp dissection from six skeletally mature greyhounds (GH) and four skeletally mature Labrador retrievers (LR), with no gross evidence of knee joint pathology. The animals were euthanized for reasons other than musculoskeletal disease and informed consent, in accordance with University guidelines, was obtained in each case for tissue removal. Ligaments were divided into proximal, middle and distal sections before sagittal bisection to allow longitudinal and transverse sections to be cut. Sections were immediately embedded on cork discs in Tissue-Tek OCT (Sakura Finetek; Torrance, CA, USA) and snap-frozen in isopentane, then stored at -80°C until required for analysis.

2.2.2 Histology and immunofluorescence

Sections of 15µm (H&E) or 30µm (immunofluorescence) were cut on a cryostat (Bright OTF5000) and transferred to poly-L-lysine slides (Poly-L, VWR, Batavia, IL). Tissue was

fixed overnight in 100% methanol at -20°C then slides were rinsed in distilled water and allowed to dry prior to staining. The immunostaining protocol was optimised using a range of section widths, fixatives, autofluorescence blocker, and type, duration, concentration, pH and diluent of primary and secondary antibodies to optimise imaging of the cells. Negative controls were performed where either the primary or secondary antibody was omitted with no labeling observed. Positive controls were performed on cultured ligament fibroblasts. The sections were rehydrated in phosphate-buffered saline, pH 7.4 (PBS, P5493, Sigma-Aldrich, St. Louis, MO) for 5 minutes, drained then incubated with 5% normal goat serum (PCN5000, Invitrogen, CA, USA) for 1 hour at room temperature. The serum was removed and the sections incubated with a solution of the primary antibodies (1:400 vimentin, mouse IgG and 1:5000 α -tubulin, mouse IgG, Abcam, UK) overnight at 4°C in a humidity chamber. Following incubation, the sections were rinsed three times in PBS, and incubated for one hour at room temperature with 1:500 anti-mouse IgG conjugated Alexafluor 488 (A12379, Invitrogen, CA, USA). Again the sections were washed three times in PBS before mounting in DAPI-containing medium (H-1500, Vectashield, Vector Laboratories, UK). The specificity of these antibodies in the dog was confirmed by Western blotting (results not shown). The use of antibodies raised against vimentin and α -tubulin to stain cells has been validated previously (Lo et al., 2002b).

2.2.3 Imaging

Two dimensional images were collected on a dedicated microscope (Nikon Eclipse 80i) using x10 or x40 objective lenses. A confocal laser scanning microscope (Zeiss LSM 510 META NLO) was used to obtain three dimensional images. Either x63 oil immersion or x10 water immersion lenses were used, and laser power and gain, and z interval adjusted for each section. Two dimensional images are presented as projections of three dimensional stacks.

2.3 Results

2.3.1 Animals

Animal data is summarised in **Table 2.1**.

Breed	Total Number	Age Range (months)	Median Age (months)	Male	Female
Greyhound	6	32-60	35	4	2
Labrador	4	22-126	35	2	2

Table 2.1: Summary data for animals in study

GHs used in the study had all been in training within the last 6 months. However precise details of exercise regimes were not available for either greyhounds or LRs.

2.3.2 General observations

The canine CLs contained a heterogenous population of cells. Nuclei varied from round to strap-like and cells were frequently observed to have processes branching from the nuclei. In order to avoid confusion over phenotype, classification of cells in this study were based on morphology. While the cells are likely to be predominantly fibroblasts in the ECM and interfascicular regions, and synoviocytes and fibroblasts in the epiligament, the exact phenotype was not determined, and may include a range of other cell types. To avoid assumptions about the phenotype, the cells will be referred to by location only: epiligament, interfascicular region or substance.

2.3.3 Epiligament

The cells of the epiligament formed a dense meshwork of shorter processes which were frequently branched, and had rounded nuclei (**Figure 2.1**).

No overall orientation in cell processes was apparent. An increase in the cellularity of the epiligament was observed in the region of the midsubstance of the ACL. This was confirmed in conventional histology, and was most marked in the ACL at the region of contact with the PCL (**Figure 2.2**). No marked change was seen regionally on the surface of the PCL.

Where the epiligament was of normal thickness (2-4 cells), the underlying ligament had abundant cells, usually with cell processes (**Figure 2.1**). In areas of thickened epiligament (>6 cells), the underlying ligament was characterised by decreased cell density and rounded cells lacking processes, considered typical of degenerative change (**Figure 2.2**). Cell morphology in the epiligament of LR CLs was similar to that described in the GH.

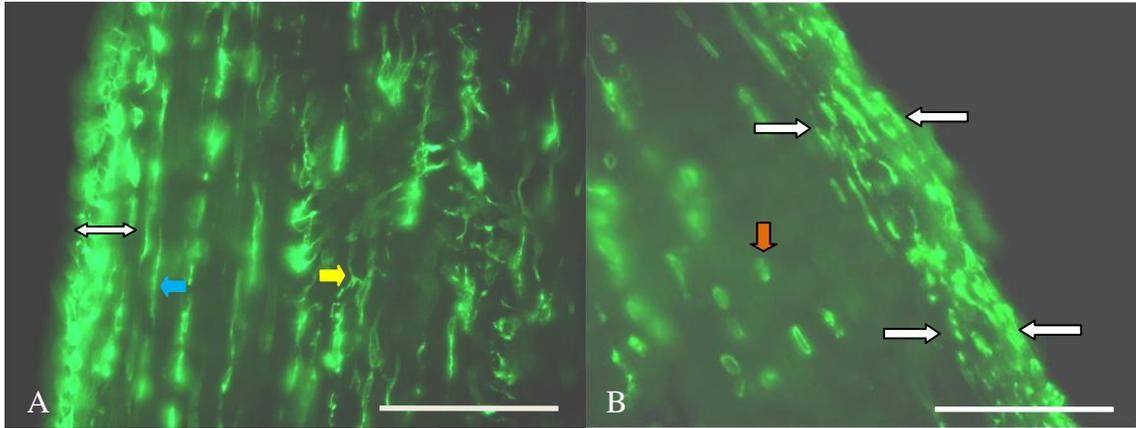


Figure 2.1: Variation in cell morphology in the epiligament: Greyhound ACL 30 μ m longitudinal sections at x40 magnification stained with vimentin and α -tubulin to show cytoskeleton. (A) The epiligament (delineated by white arrow) is around 2-4 cells thick. Underlying ligament has cells, with a mix of parallel (blue arrow) and branched (yellow arrow) processes. (B) Epiligament is 6-8 cells thick (delineated by white arrows). Cellularity of underlying ligament is sparse and generally without branching (orange arrow). Magnification bars =100 μ m

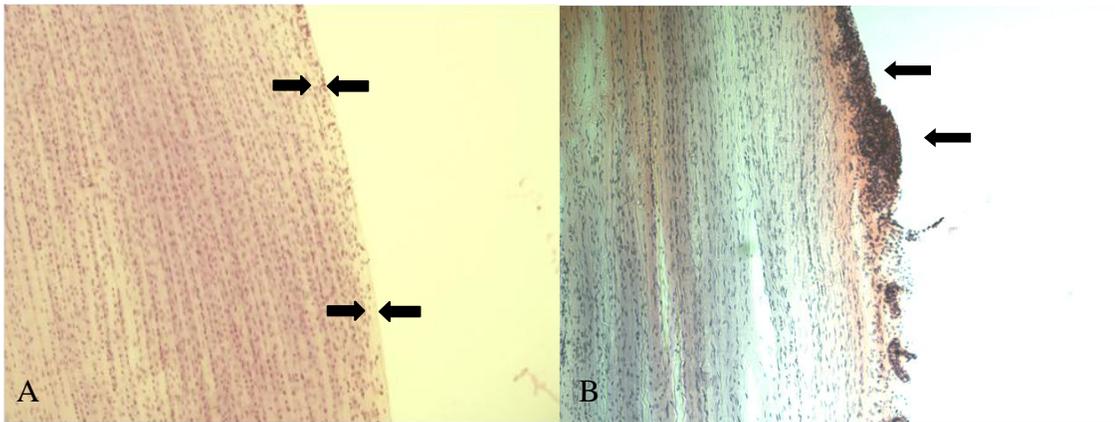


Figure 2.2: Variations in epiligament cellularity. (A) Proximal greyhound ACL with regular, thin epiligament (between arrows) of only a few cells in thickness. The epiligament overlies healthy ligament with only minimal signs of degenerate change. (B) Mid ACL of greyhound showing irregular cellular thickening of the epiligament (arrows). Such changes were common where the epiligament overlay ligament substance that showed changes associated with degeneration. Magnification bar in A and B= 100 μ m

2.3.4 Interfascicular regions

The dense meshwork of cells seen on the epiligament was mirrored in the interfascicular regions of the ligament. These cells of the interfascicular region had thick processes of widely varying length, which often branched markedly, forming complex meshworks of irregular, stellate cells (**Figure 2.3**). In some locations, the predominant cell orientation was parallel to the long axis of the ligament, but in others there appeared to be no overall orientation and again nuclei were rounded. There appeared no marked variation between the LR and GH.

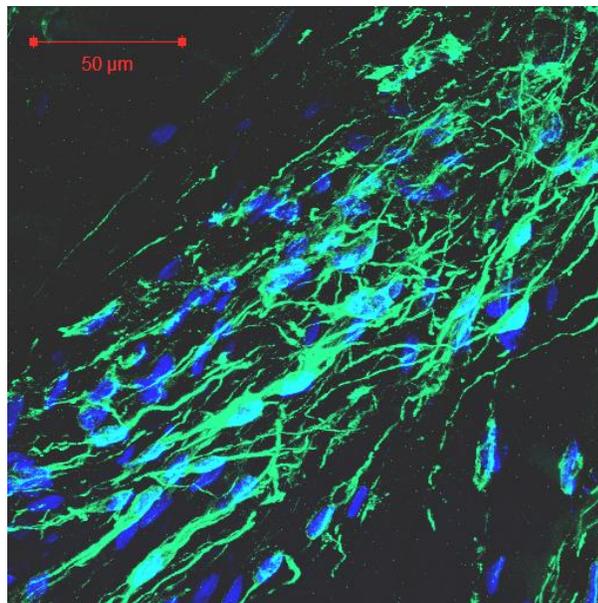


Figure 2.3: Confocal microscopy images of cells of the interfascicular region showing dense cellularity with cells showing multiple branching processes with variation in width, length and orientation. Greyhound ACL, Zproj=30µm, x63 magnification, staining: vimentin and α -tubulin to show cytoskeleton (green) and with DAPI to show nuclei (blue)

2.3.5 Substance: variations in cell morphology

Cells of the fascicular regions were of three broad types (**Figure 2.4**). One group of cells within the ligament had long, thin cytoplasmic processes extending mainly parallel to collagen bundles, denoted as type A cells (**Figure 2.4A**). Individual processes could be very long, frequently 100µm and sometimes extending in excess of 200µm. The processes were rarely

straight, with some conformation to collagen crimp, but with undulation in all planes, and frequently extended to adjacent cells or processes from nearby cells. The nuclei were narrow, frequently very long (up to 70 μm), ranging from slightly oblique to the collagen bundles to parallel with them. Transverse and perpendicular processes were noted, and always originated from the nucleus. These cells were seen in all regions of both CLs.

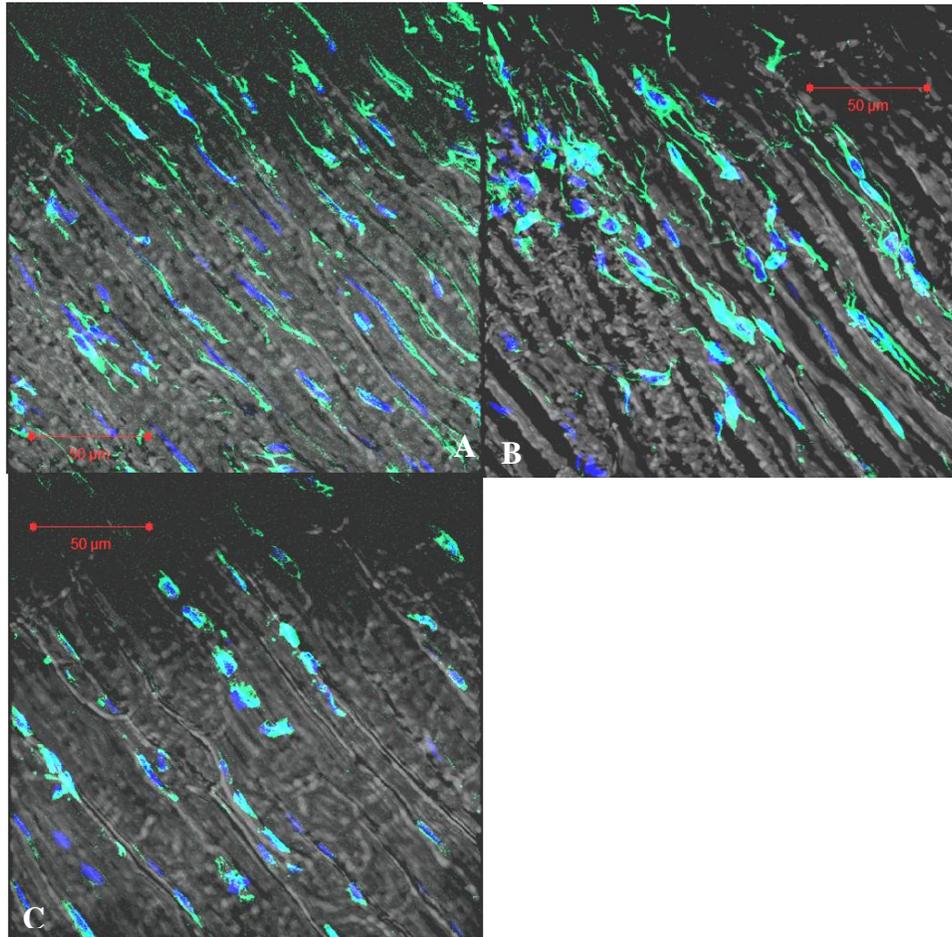


Figure 2.4: Variations in cell morphology of the fascicular region of the CL complex: Confocal microscopy images, greyhound ACL, Zproj 30 μm . Staining: vimentin and α -tubulin to show cytoskeleton (green) and with DAPI to show nuclei (blue) (A) Type A cells showing thin longitudinal processes of moderate length with minimal branching. (B) Type B cells showing shorter thicker processes with branching (C) Type C cell morphology showing rounded nuclei and an absence of processes.

A second type of cell, type B, had shorter, thicker, frequently branching processes, with rounder nuclei (**Figure 2.4B**). This second type was the most common morphology and was

frequently found in association with type A cells. The branching processes extended in all directions, and like type A cells, contact with other cell processes was common. It must be noted that the transition from type A to B was indistinct and some cells were of mixed type A and B morphology.

The third cell type, type C, had rounded nuclei, and no cytoplasmic processes in any direction (**Figure 2.4C**). The overall density of these cells did not appear to differ from types A and B. Although these cells were most frequently found in isolation, columns of closely apposed type C cells were also noted, with frequent cell-to-cell contact.

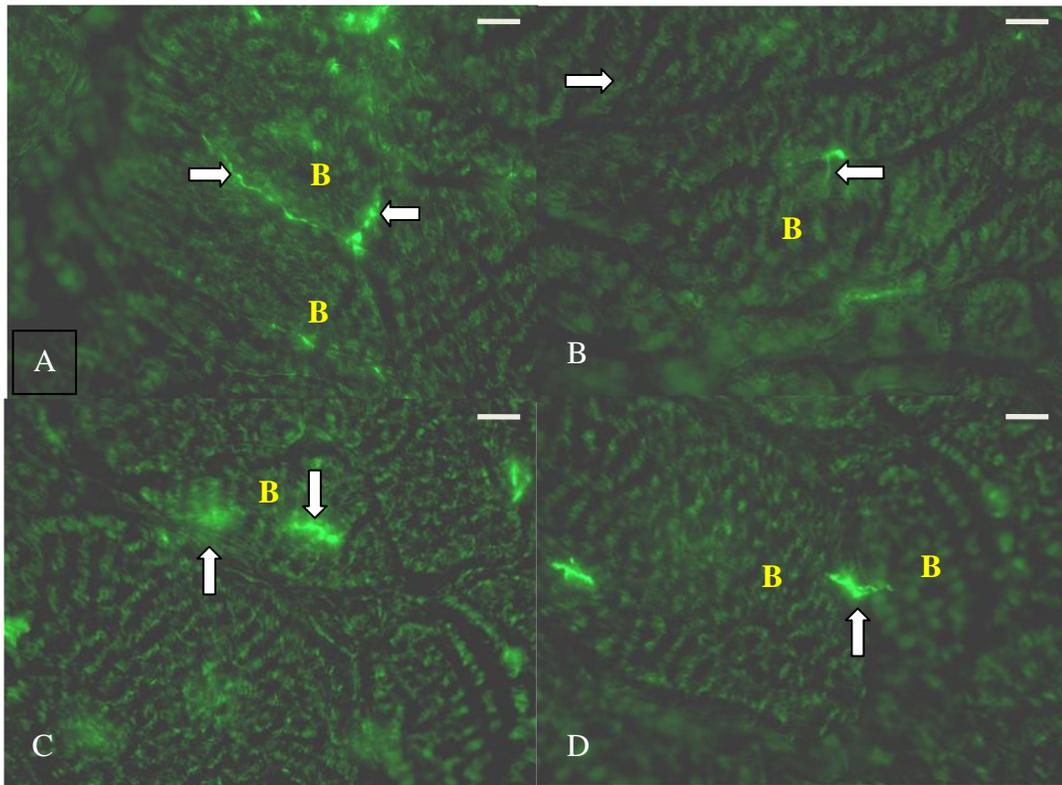


Figure 2.5: Relationships of transverse processes to collagen bundles. Greyhound PCL, 30 μ m sections, x40. Collagen bundles are marked with a yellow B (A) Transverse processes (arrows) within interbundle region encircling a collagen bundle. (B) Transverse processes (arrow) extending from an interbundle region penetrating a collagen bundle. (C) Transverse processes (arrows) within a collagen bundle. (D) Transverse processes (arrow) passing between two collagen bundles. Magnification bars =10 μ m

Cells of the three morphologies described in the GH were also noted in the LR. Although morphology was similar, the proportion of each type appeared to differ. Subjective assessment of the overall CL cellularity between breeds appeared similar. Again subjectively assessed, a smaller proportion of type A cells was noted in the LR, giving way to a greater proportion of type B cells. The overall proportion of type C cells appeared similar between breeds.

On transverse sections of GH CL, cell processes were seen extending from interfascicular regions to interbundle regions (**Figure 2.5**). Furthermore, cells in the interbundle regions frequently encircled collagen bundles and also processes were seen to penetrate the interior of the collagen bundle. Although processes on transverse sections were observed in both ligaments in all regions, there appeared to be fewer in the midsubstance of both CLs than in proximal and distal CLs. In the LR CL, there appeared to be greater numbers of processes on transverse sections, in all regions of both CLs (**Figure 2.6**), but with similar distribution to that described in the GH.

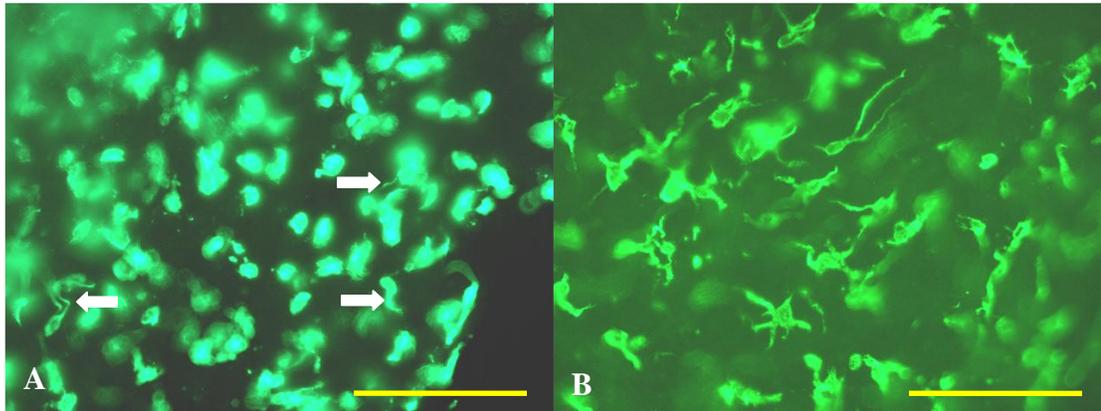


Figure 2.6: Breed variation in processes seen in transverse sections. Staining: vimentin and α -tubulin to show cytoskeleton (green). (A) Greyhound PCL x40. Only small numbers of transverse processes are noted (white arrows). (B) LR PCL x40. Many transverse processes are noted giving cells a 'stellate' appearance. Magnification bars = 100 μ m.

2.3.6 Substance: regional variations in the GH

Assessed subjectively, in the GH, the majority of each ligament appeared to comprise either:

- a population of cell type A (**Figure 2.7A**),
- a mixed population of cell types A and B (**Figure 2.7B**),
- type C alone (**Figure 2.7C**).

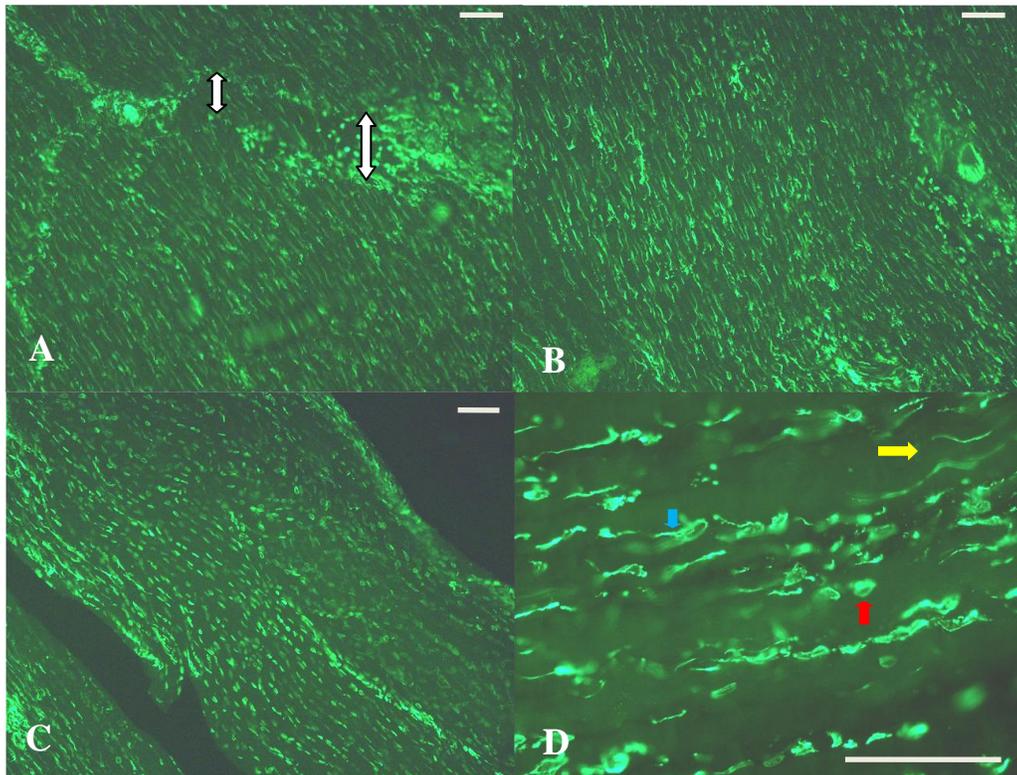


Figure 2.7: Regional and local variations in cell morphology in the greyhound. 30µm sections. Staining: vimentin and α -tubulin to show cytoskeleton (green). (A) ACL x10. Large areas of type A cells with interposing interfascicular region delineated by white arrows. (B) PCL x10 Mixed populations of types A and B cells. (C) PCL x10 Homogenous population of type C cells. (D) PCL x40 Adjacent collagen bundles show populations of types A (yellow arrow), B (blue arrow) and C (red arrow). Type A cells manifest as long thin processes (yellow arrow). Magnification bars = 100µm

However, a juxtaposition of bundles containing differing cell morphologies was often seen too, subjectively most commonly in the proximal and distal ligament. Cells of type A and B were commonly seen in apposition to cells of type C, varying between alternate collagen bundles (**Figure 2.7D**). In the midsubstance of the CLs, there appeared to be an increase in the relative proportion of type C cells compared to proximal and distal CLs. Where collagen bundles appeared regular and healthy, cells of all three morphologies appeared to predominate. Where there was loss of collagen density, identified by reduced collagen bundle density, there appeared to be only morphologies B and C, although in these samples it was rare to find large areas of reduced collagen density.

2.3.7 Substance: regional variations in the LR

While the LR had cells of similar morphologies described in the GH, regional variation appeared to differ. Assessed subjectively, the LR lacked regions of predominant cell type A seen in the GH. Instead most of the ligament appeared to comprise:

- mixed populations of types A and B (most common proximally and distally of both CLs) (**Figure 2.8A**)
- regions of predominantly cell type B (most common in the middle region of both CLs) (**Figure 2.8B**)
- predominantly cell type C (most common in the middle region of both CLs) or of all three morphologies (found throughout both CLs) (**Figure 2.8C**).

Large regions of predominantly cell type B appeared to corresponded with regions of reduced collagen density where there appeared to be a loss of collagen architecture. Furthermore, marked juxtaposition was commonly seen between regions where one cells of type C morphology would appear in one bundle adjacent to another of cell type B, changing from one collagen bundle to the next (**Figure 2.9 A,B**). Despite the contrast in cell morphology, the bundles were of similar orientation and appearance. Although seen in both ACL and PCL, these juxtapositions were more common and extended over a larger area in the LR ACL (**Figure 2.9 C,D**). Juxtaposition of all three cell morphologies on a local level, as described in the GH, appeared more common in the LR.

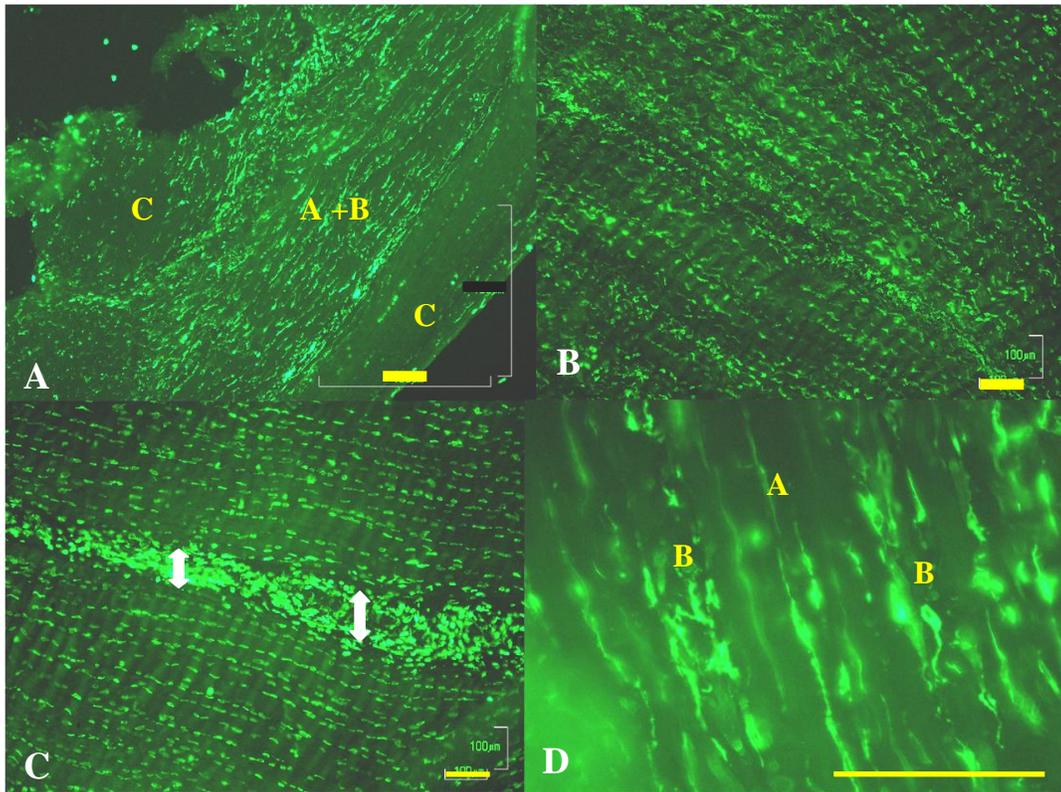


Figure 2.8: Typical regional variation of cell morphology in the LR. Staining: vimentin and α -tubulin to show cytoskeleton (green). (A) ACL, x10. Three regions are represented across the width of the ligament. There are two outer regions of type C and a central region of mixed type A+B. (B) PCL, x10. A large region of predominantly type B. (C) ACL, x10 Typically large regions of type C with highly cellular interbundle region (delineated with white arrows). (D) ACL, x40 Typical variation in interbundle cell morphology in the LR: A bundle of type A morphology is flanked by bundles of type B morphology. Magnification bars (yellow) =100 μ m.

2.3.8 Cell density

Although there appeared to be no overall regional variation in cell density, marked local variation existed, assessed subjectively, with small areas completely devoid of nuclear or cytoskeletal staining (**Figure 2.10**). The collagen architecture in these regions appeared normal, and these changes were noted in both CLs, and in both breeds of dog.

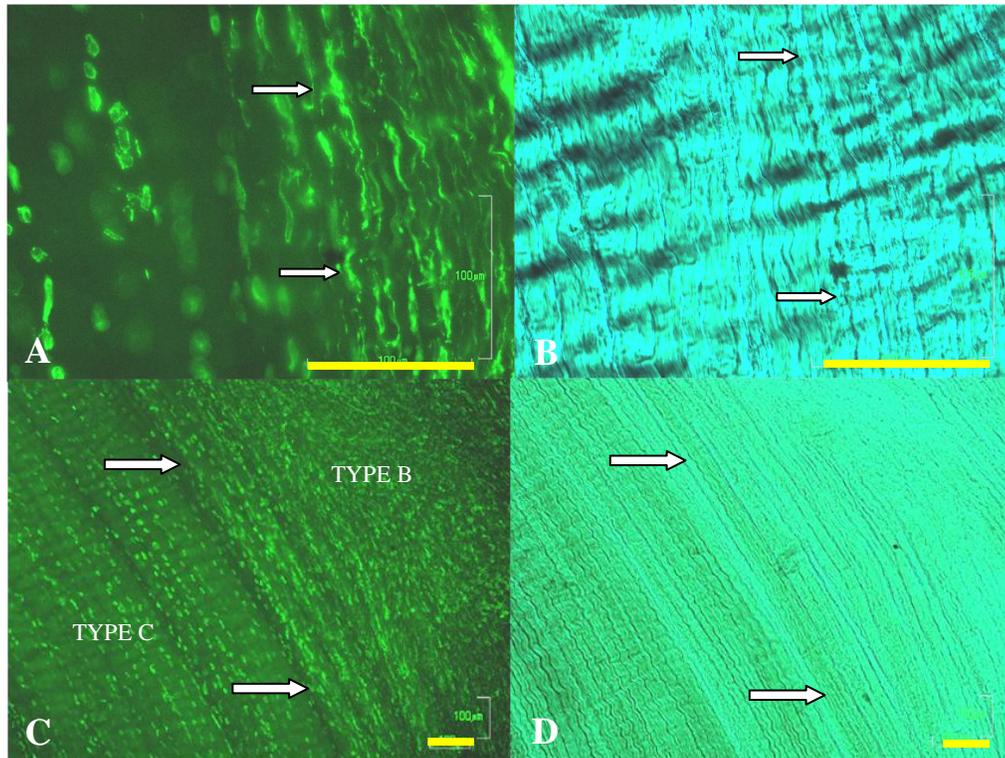


Figure 2.9: Striking juxtaposition of regions of differing cell morphologies in the LR ACL. 30 μ m sections, x40 (A) Staining: vimentin and α -tubulin to show cytoskeleton (green). Left side of image shows cells of type C morphology, right side shows predominantly type A morphology and the distinction is marked with arrows. (B) Polarized image of collagen in figure (A) showing bundles of similar orientation and density despite containing cells of differing morphologies. Arrows mark juxtaposition of cell morphologies. (C) Staining: vimentin and α -tubulin to show cytoskeleton (green). Juxtaposition of cell types B and C extending over a large area. (D) Polarized image of collagen architecture shown in (C). Magnification bars =100 μ m.

2.4 Discussion

In this study we have described what may be a three dimensional network of cells in the CL complex of the dog, consistent with the cellular matrix described in other load-bearing connective tissues and in the ACL of the sheep. However, we have suggested heterogeneity in cell morphology throughout the ligament and suggested previously undescribed regional and interbreed differences. Understanding variations in cells within the CL complex is likely to be important in improving our understanding of the role of cell communication in ligament

disease and may be important in the development of bioengineered constructs for the treatment of cruciate ligament failure.

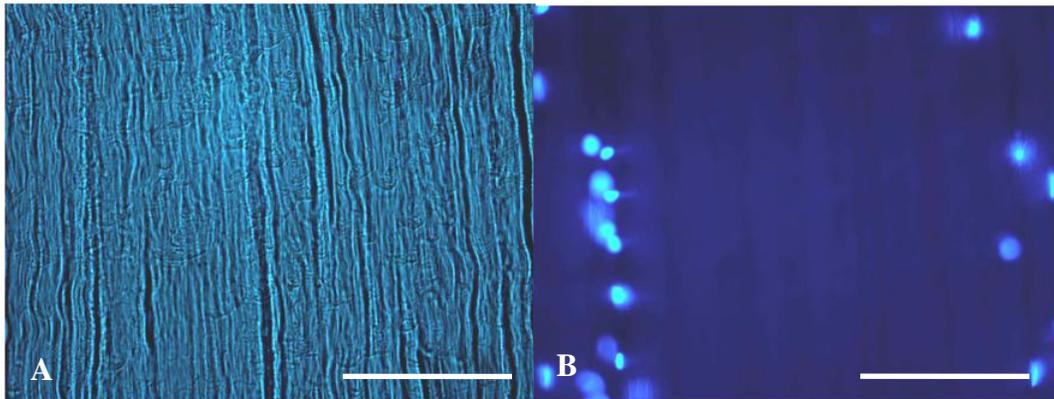


Figure 2.10: Areas devoid of nuclear staining. Greyhound PCL, 30 μ m sections, x40. A and B are light and fluorescent images respectively of the same section of tissue (A) Regular, organised collagen bundles that (B) show no staining for cell nuclei (blue, stained with DAPI). Magnification bars = 100 μ m

2.4.1 General differences in cell morphology

Cell morphology has been considered an adaptation to ligament mechanics or physiology (Ralphs et al., 1998). Cellular processes are considered a feature of connective tissue under tensile load (Lo et al., 2002a) with tension important in maintaining their presence (Baschong et al., 1997, Hellio Le Graverand et al., 2001a). Fusiform cells with long cytoplasmic processes (similar to the type A found in our study) were described in meniscal regions under tension and cells with a lack of processes and rounded nuclei (similar to the type C found in our study) were noted in regions under compression. In the human intervertebral disc, correlation with increasing compression and decreasing cell process length has been described (Bruehlmann et al., 2002). Type C cells appeared to be seen most commonly in the midsubstance of the ACL which is a site where the ligament is thought to undergo compression (Comerford et al., 2006b), therefore these morphological changes may reflect an adaptation to compression (Milz et al., 2005). It remains possible that variation in the composition of the ECM may lead to the described morphological variation. However cells of all phenotypes were seen in CL tissue with normal collagen architecture but only cells of types B and C were described where there had been disruption. This would suggest that the cell phenotype precedes tissue morphology change.

Long cytoplasmic processes may facilitate cell to cell communication through contact with other cells (Lo et al., 2002). Additionally these processes provide a mechanism whereby cells apparently distant from blood vessels may acquire nutrients (Bruehlmann et al., 2002). A seemingly gradual transition in cell morphology has been noted in the intervertebral disc and meniscus, where cells gradually change shape between regions, losing their processes and becoming rounder (Bruehlmann et al., 2002, Hellio Le Graverand et al., 2001a). Thus cells of type B may be a transitional morphology from types A to C.

Local variation in cell morphology was seen both between groups of parallel collagen bundles and also along the length of groups of collagen. The canine ACL is considered to be subject to multiaxial stresses (Kennedy et al., 1976, Takai et al., 1993) and local variation in cell morphology may reflect variation in stresses and strains. Other extrinsic factors that may also influence forces acting on the ligament include the dimensions of the intercondylar notch of the femur (Comerford et al., 2006a), joint pathology (Hellio Le Graverand et al., 2001b, Hellio Le Graverand et al., 2001c) and muscular support and limb usage (Colborne et al., 2005).

Areas of acellularity in the canine ACL have been described by a number of authors (Vasseur et al., 1985, Comerford et al., 2006b), and have been considered early degenerative changes. Recent work has ascribed this loss of cellularity to apoptosis (Gyger et al., 2007, Krayer et al., 2008), but it is not known whether the decreased cellularity leads to degenerative change in the matrix. Here we demonstrate normal collagen architecture in these hypocellular regions and suggest loss of cells may be an initiating event in this change.

2.4.2 Breed differences

Large areas of cell type A were seen in the GH but not the LR. This difference may be in part explained through the lack of large areas of collagen loss in the GH. In the sheep ACL poor healing has been associated with failure of long cytoplasmic processes to span wounds (Lo et al., 2002b). In the canine ACL, the formation of rounded cells without processes may mean the ligament is unable to repair damage. More long cell processes in the GH CL may aid healing and partly explain the lower risk of ACL rupture in this breed. The formation of columns of type C cells may also reflect an attempt to re-establish cellular communication.

Juxtaposition of differing cell morphologies from one longitudinal area to another appeared marked in the LR but not the GH and may reflect a marked change in mechanics from one bundle to the next. This possible breed-specific opposition of cell morphologies may reflect

physiological banding. Local juxtaposition of all cell types was also appeared more common in the LR and may reflect differing CL factors from the GH such as functional differences (Wingfield et al., 2000a, Wingfield et al., 2000b), exercise (Patterson-Kane et al., 1998, Stanley et al., 2008) or intrinsic susceptibility to CL disease (Bennett et al., 1988). Further work is required to investigate these findings in other breeds of dog.

This study did have certain limitations. The lack of quantifiable data made some of our observations subjective, rendering comparisons within and between breeds similarly subjective. The process of examining each entire sample in the required detail to quantify cell types was not feasible within the life time of the fluorophore and further work must address this technical issue to validate our observations. Furthermore, lack of clear exercise history may influence the ability to compare these results. Although our study population was heterogeneous with differences in age, nutrition, and exercise, there appeared to be little variation in results between dogs of the same breed. Future quantification of our observations will enable us to assess the validity of low sample numbers. Due to the twisting nature of the CLs, precise longitudinal sectioning was not possible, and frequently long processes could not be followed completely. Cell processes were consistently demonstrated through a variety of fixatives and antibodies, and were also seen to a lesser degree on the H&E sections. Given the technique used is similar to previous cell morphology studies (Lo et al., 2002b, Bruehlmann et al., 2002), we do not consider the cell morphologies we have demonstrated artefactual. Although failure of antibody penetration can result in artefacts, staining, measured on confocal microscopy, was consistent and complete. Future work will involve relating cell morphology to intraligament forces and the identification of gap junctions.

In summary, we have suggested regional variation in the cell morphology of the canine CL complex as well as juxtaposition and differences in type of cell morphology in dog breeds at a differing risk to non-contact ACL injury. The possibility of a three dimensional network of cells has ramifications for cell nutrition, mechanical sensing and coordinated response to injury in the CL complex. Clearly more work is required to understand the importance of these morphological variations and to develop a better understanding of their role in the pathology and degeneration of the CL complex.

CHAPTER 3: The organisation of elastin and fibrillins 1 and 2 in the cruciate ligament complex

Abstract

Outline: While elastin fibres and oxytalan fibres (bundles of microfibrils) have important mechanical, biochemical and cell regulatory functions, neither their distribution nor function in the canine ligaments have been investigated.

Method: Twelve pairs of cruciate ligaments (CLs) were obtained from 10 adult greyhound dogs with no evidence of knee osteoarthritis. Elastic fibres were identified using Verhoeff's and Miller's staining. Fibrillin 1 and 2 were immunolocalised and imaged using confocal laser scanning microscopy. Hydrated, unfixed tissue was analysed using Nomarski differential interference microscopy (NDIC) allowing structural and mechanical analysis.

Results: Microfibrils and elastin fibres were widespread in both CLs, predominantly within ligament fascicles, parallel to collagen bundles. Although elastin fibres were sparse, microfibrils were abundant. We described abundant fibres composed of both fibrillin 1 and 2 which had a similar pattern of distribution to oxytalan fibres. NDIC demonstrated complex interfascicular and interbundle anatomy in the CL complex.

Conclusions: The distribution of elastin fibres is suggestive of a mechanical role in bundle reorganisation following ligament deformation. The presence and location of fibrillin 2 in oxytalan fibres in ligament differs from the solely fibrillin 1-containing oxytalan fibres previously described in tendon and may demonstrate a fundamental difference between ligament and tendon.

3.1 Introduction

Cruciate ligaments (CLs) are dense bands of collagenous tissue that are the primary stabilisers of the knee (femorotibial) joint. The two components are anterior and posterior cruciate ligaments, with the anterior cruciate ligament (ACL) twisted around the posterior cruciate ligament (PCL) forming the CL complex (Arnoczky and Marshall, 1977). Each CL is comprised of multiple fascicles containing bundles of collagen fibres (Kennedy et al., 1974, Yahia and Drouin, 1989, Amis and Dawkins, 1991). Collagen fibres are not recruited isometrically during knee joint motion and each change in knee joint position recruits fibres differently (Amis and Dawkins, 1991, Butler et al., 1992). Although collagen provides tensile strength to the ligament complex, other structural components likely contribute to the overall mechanical function of the complex (Frank, 2004). Microfibrils (MFs), polymers of fibrillins 1 and 2, are considered to have a structural role in ligament and tendon. Bundles of MFs are known as oxytalan fibres. Elastin fibres comprise a central cross-linked core of highly extensible elastin surrounded by a supporting sheath of MFs, with many other associated molecules (Kielty, 2006). Collectively, oxytalan and elastin fibres are referred to as elastic fibres. Elastin has traditionally been considered a minor component of ligament tissue (Frank, 2004). A wide distribution of elastic fibres in the human ACL has been described (Strocchi et al., 1992). In canine CLs, elastin fibres have only been reported at low levels (Paatsama, 1952, Vasseur et al., 1985).

Elastic fibres have important mechanical, biochemical and cell-regulatory functions in tissue. Reversible elasticity is a function of both elastin and oxytalan fibres and is dependent on water and calcium (Eriksen et al., 2001). MFs are stiffer than elastic fibres (Sherratt et al., 2003) and are highly resistant to axial tension (Glab and Wess, 2008). Distribution of elastic fibres in tissue is considered to reflect function (Kielty et al., 2002a). Regions of canine superficial digital flexor tendon (SDFT) that undergo the greatest strain deformation have the highest regional elastin content (Ritty et al., 2002). MFs also have key roles in extracellular regulation of transforming growth factor (TGF) β (Charbonneau et al., 2004) and cell adhesion (Ito et al., 1997, Wendel et al., 2000).

In the canine SDFT, fibrillin 1 was predominantly found in fibre-form with elastin and fibrillin 2 predominantly found pericellularly. Fibrillin 2 is commonly found in MFs in foetal tissues but has been considered to have limited distribution in adult tissue (Cain et al., 2006). A recent study has suggested microfibrils in post natal tissue may comprise of a fibrillin 2 core and a

fibrillin 1 outer sheath (Charbonneau et al., 2010b). Failure of elastic fibres has been implicated in a number of serious diseases (Kielty, 2006).

In this study, we use histology and immunofluorescence to methodically detail the distribution of elastic fibres and fibrillins 1 and 2 in the canine CL complex. We also use micromechanical manipulation and enzymatic digestion to explore CL microanatomy. By understanding the distribution and function of these molecules in the CL complex, we intend to gain a greater understanding of CL physiology providing valuable insight into the aetiopathogenesis of non-contact ACL injury and information for future ligament engineering projects in mammalian species.

3.2 Materials and methods

3.2.1 Sample collection and preparation

Twelve pairs of ACLs and PCLs were harvested from ten skeletally mature Greyhounds with no macroscopic evidence of any knee joint pathology. The animals were euthanized for reasons not related to this study with informed consent obtained according to standard University ethical review. Six pairs of CLs from five dogs were sectioned into proximal, middle and distal sections, embedded on cork discs in Tissue-Tek OCT (Sakura Finetek; Torrance, CA, USA) and immediately snap-frozen in isopentane using liquid nitrogen then stored at -80°C until immunofluorescence staining and NDIC analysis. Six pairs of CLs from five dogs were fixed in 4% paraformaldehyde (P6148, Sigma-Aldrich, UK) for 24 hours then embedded in paraffin before sectioning for histological examination.

3.2.2 Histology

Sequential sections of 4µm from paraffin-embedded samples were stained with haematoxylin and eosin (H&E), Verhoeff's iodine-iron haematoxylin (EVH) (elastin fibres only) and Miller's stain (M) (both elastin and oxytalan fibres) (Barros et al., 2002). This staining methodology was used to allow assessment of tissue architecture, and through comparison of EVH and M-stained sections, differentiation of elastin and oxytalan fibres. Images were recorded on a dedicated microscope (Nikon Eclipse 80i). H&E sections were assessed by two observers blinded to sample information (KDS and EJC) for signs of CL degeneration according to criteria detailed previously (Vasseur et al., 1985). Briefly, this involved awarding a score from 0-3 according to the following criteria. Grade 0: healthy ligament, Grade 1: mild

degenerate changes with focal loss of collagen architecture, Grade 2: moderate degeneration with regional disruption to collagen architecture and Grade 3: severe degeneration affecting large sections of ligament.

3.2.3 Antibodies

Two antibodies were used to immunostain fibrillins 1 and 2 and elastin. Antibodies against fibrillin 1 were rabbit polyclonal antibodies raised against the proline-rich domain (Trask et al., 1999) and toward the carboxyterminal domain of human fibrillin 1 (Ritty et al., 1999) respectively. Antibodies against fibrillin 2 were both rabbit polyclonal antibodies raised against the glycine-rich region of human fibrillin 2 and against amino acids Met-1 to Thr-1114 of human fibrillin 2 respectively (Trask et al., 1999). All four fibrillin antibodies have been affinity purified and do not cross react (Ritty et al., 2002) and were kindly donated by Dr. Timothy Ritty. Immunolabeling of elastin was achieved using two commercial antibodies (ab9519, monoclonal mouse IgG, Abcam, UK, 1:100, and E4013, monoclonal mouse IgG, Sigma-Aldrich, UK).

3.2.4 Immunofluorescence

Longitudinal and transverse sections of 30µm were cut from six pairs of CLs from five dogs on a cryostat (Bright OTF 5000) and transferred to poly-L-lysine slides (Polysine, VWR, UK). Tissue was fixed overnight in 100% methanol at -20°C then slides were rinsed in distilled water and allowed to dry prior to staining. Sections were rehydrated in phosphate-buffered saline, pH 7.4 for 5 minutes, then incubated with hyaluronidase (4800 IU/ml in PBS, H3884, Sigma-Aldrich, UK) with a protease inhibitor cocktail (P2714, Sigma-Aldrich, UK) for 24 hours at room temperature and then rinsed for 5 minutes, 3 times in PBS. A second incubation with collagenase (30 IU/ml in PBS, C2674, Sigma-Aldrich, UK) for 30 minutes was followed by a further PBS rinsing. Sections were covered with 5% normal goat serum (PCN5000, Invitrogen, CA, USA) for 1 hour at room temperature then incubated with a solution of two antibodies raised against elastin (ab9519 at 1:100, Abcam, UK and E4013 at 1:5000, Sigma-Aldrich, UK) overnight at 4°C in a humidity chamber. A second incubation with antibodies against fibrillin 1 or 2 (1:50) overnight at 4°C was followed by rinsing in PBS and incubation for one hour at room temperature with anti-mouse IgG conjugated Alexafluor 488 (1:500, A11001, Invitrogen, CA, USA) and anti-rabbit IgG conjugated Alexafluor 568 (A11011, Invitrogen, CA, USA). Sections were rinsed in PBS before mounting in DAPI-containing

medium (H-1500, Vectashield, Vector Laboratories, UK). Negative controls were achieved by omitting the primary or secondary antibody and one or both enzymatic digestions (data not presented).. Positive controls for elastin were performed on elastin-rich vascular tissue where the distribution of elastin could be compared to histochemically stained sections (data not presented). All four fibrillin antibodies have been validated previously in canine connective tissue (Ritty et al., 2002).

3.2.5 Nomarski differential interference contrast optical microscopy (NDIC)

Unfixed cryosections of 30µm from six pairs of CLs from five dogs were transferred to slides and kept moist using lactated Ringer's solution (Aquapharm No.11, Animalcare, UK). Fabric tabs were attached to the sections using superglue to allow application of transverse and longitudinal strain. Strain was measured relative to the width of the tissue sample in the direction of strain application and strain was progressively applied to 300%. Lactated Ringer's was applied to the section before applying a coverslip. Images were recorded on a dedicated microscope (Nikon Eclipse 80i). This facilitated manipulation of unfixed tissue within a physiological solution where the elasticity of elastic fibres was preserved as described previously (Pezowicz et al. 2005; Pezowicz et al 2006).

3.2.6 Imaging

Two dimensional images were collected on a dedicated microscope (Nikon Eclipse 80i) using x10 or x40 objective lenses. A confocal laser scanning microscope (CLSM) (Zeiss LSM 510 META NLO) was used to obtain three dimensional images. Either x63 oil immersion or x10 water immersion lenses were used, and laser power and gain, and z interval adjusted for each section. Two dimensional images are presented as projections of three dimensional stacks

3.3 Results

3.3.1 Animals

Summary data are provided in **Table 3.1**. All dogs had been in training for racing prior to being subjected to euthanasia.

	Dogs	Pairs of CLs	Age Range (months)	Median Age (months)
Male	6	7	34-68	35
Female	4	5	32-36	34
TOTAL	10	12	32-68	35

Table 3.1: Summary data for animals

3.3.2 H&E sections

All ACLs and PCLs from six knees were assessed as having a low level of degenerative change (Grade 1 by both assessors according to Vasseur scale). Staining of collagen throughout the ligaments demonstrated no loss of density or disruption to the collagen architecture.

3.3.3 Elastin fibres (EVH)

Elastin fibres appeared to be distributed consistently throughout both ACLs and PCLs with little apparent variation within CLs, between ACLs and PCLs or between CLs of differing joints or dogs. There appeared to be increased numbers of elastin fibres in the epiligament. This increased staining appeared uniform in degree and pattern in the surface layers of both CLs. Within the substance of both CLs, elastin fibres appeared to vary in width and orientation and most were orientated parallel to collagen bundles (**Figure 3.1A**). Elastin fibres seemed to be found more commonly on the surface of rather than within collagen bundles. There appeared to be increased elastin fibre staining in the interbundle regions (**Figure 3.1B**). Interbundle elastin fibres seemed orientated either perpendicular or oblique to collagen bundles and were either straight or tortuous (**Figure 3.1C**). Elastin fibres appeared abundant in interfascicular regions and were loosely organised. Elastin fibres could not be imaged without dual enzymatic digestion.

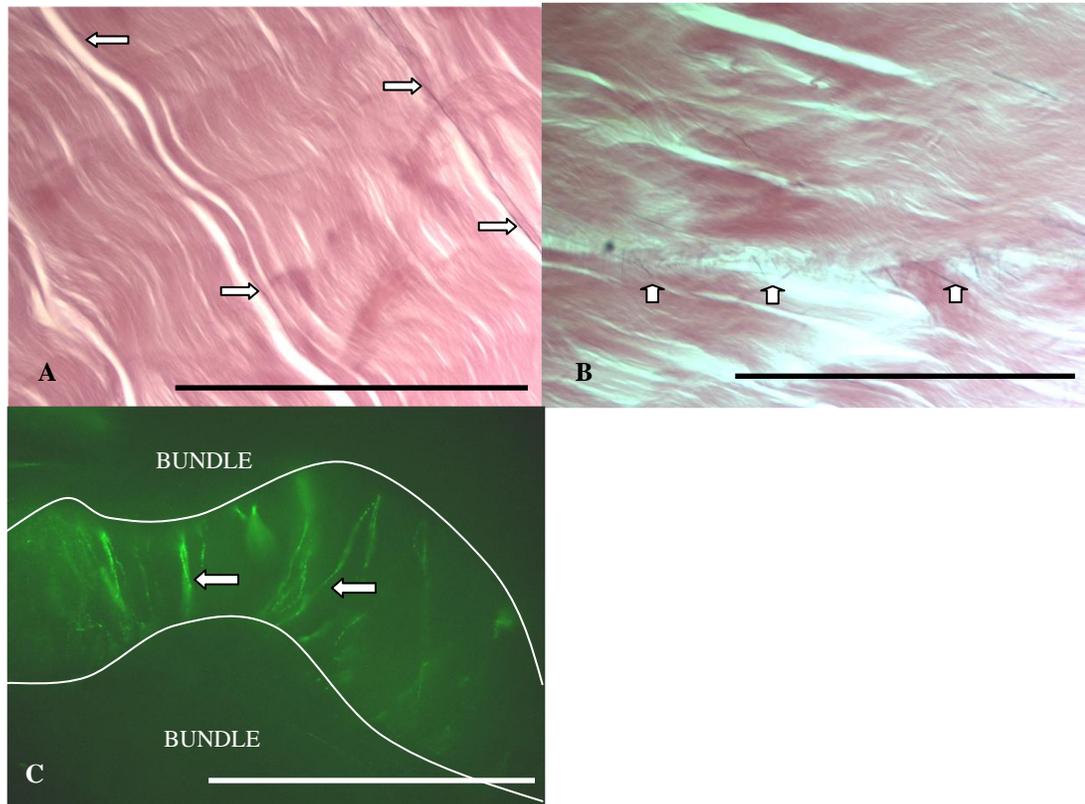


Figure 3.1 Distribution of elastin fibres in the CL complex (A) ACL x100 Verhoeff's stain. Sparse elastin fibres (arrows) orientated parallel to collagen bundles. Magnification bar = 40 μ m. (B) PCL x100 Verhoeff's stain. Longitudinal section showing short elastin fibres (arrows) obliquely between two collagen bundles. Magnification bar = 40 μ m. (C) ACL x100 stained with antibodies to elastin. Perpendicular fibres of elastin (green; arrows) spanning an interbundle region (edge of collagen bundles marked by white lines). Notice the similarity to the interbundle fibres stained histochemically in (B). Magnification bar = 40 μ m.

3.3.4 Oxytalan fibres (Miller's stain)

Oxytalan fibres appeared more numerous and finer than elastin fibres throughout both CLs. Assessed subjectively, no differences were noted in oxytalan staining between proximal, middle and distal CLs, between ACLs and PCLs or between knees or dogs. Oxytalan fibres were found in the epiligament, at apparently greater frequency than that observed in the substance. Fibres seemed broadly orientated along the axis of the underlying collagen bundles, but significant numbers of fibres appeared oblique and perpendicular. Within the substance of the CL, large numbers of fibres appeared to run parallel to collagen bundles, and often

appeared only just visible at the very limit of resolution of the microscope at 100 times oil-immersion magnification (**Figure 3.2A**). The majority of fibres in the interfascicular region appeared to be arranged in a fine, tortuous meshwork with no overall orientation, although some ran perpendicularly between bundles. Oxytalan fibres were commonly found both on the outside and within collagen bundles, where again there appeared to be marked variation in diameter (**Figure 3.2B**).

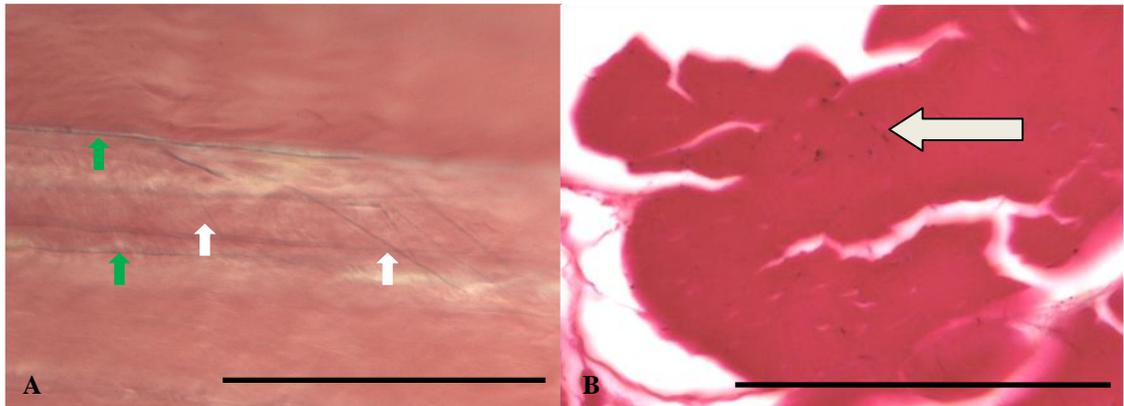


Figure 3.2: Distribution of microfibrils in the canine CL complex. (A) PCL, Miller's stain: Detail of image of x100 magnification. Larger elastin fibres (green arrows) are seen parallel to collagen bundles. Finer microfibrils (white arrows) run obliquely and vary in size. The field contains many fine microfibrils, many of which are only just visible at this magnification. Magnification bar = 40µm. (B) Transverse section, ACL, Miller's stain. Detail of image of x100 magnification. Microfibrils (arrowed black dots) abundant within the collagen bundles. Magnification bar = 40µm.

3.3.5 Fibrillins

Fibrillin 1 appeared to be found consistently throughout both ACL and PCLs. In sections that had not undergone enzymatic digestion, fibrillin 1 appeared to form fibres throughout both CLs (**Figure 3.3A**). These fibres appeared often markedly branched. Dense fibre staining was seen throughout interfascicular regions where fine fibres formed a dense, irregular meshwork, similar to that previously described for histochemical microfibril staining. Where nuclei were round, fibrillin 1 would be found only pericellularly (**Figure 3.3B**). Co-localisation with elastin was commonly seen pericellularly, but fibrillin 1 was rarely seen when the elastin was part of a fibre (**Figure 3.4**). In the epiligament fibrillin 1 appeared more abundant than was seen in the ligament substance. No interligament or regional variation was apparent. The

staining intensity of fibrillin 1 appeared reduced slightly by collagenase but not by hyaluronidase enzymatic digestion.

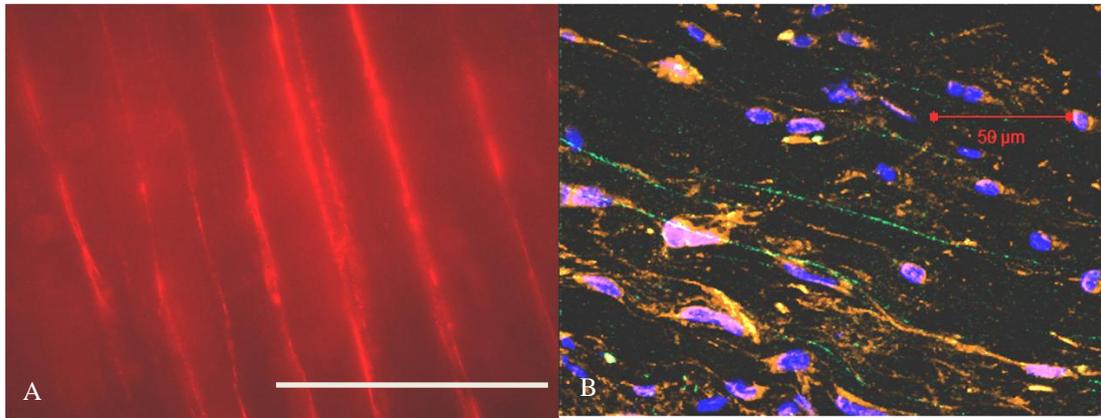


Figure 3.3: Distribution of fibrillin-1 in the canine CL complex (A) Longitudinal ACL, x100. Fibrillin-1 (red) forms fibres. Magnification bar = 40µm. (B) ACL x63: Fibrillin 1 (red) is found pericellularly where nuclei are rounded (nuclei in blue). Staining of fibrillin 1 is also seen extending parallel to collagen in a fibre-like structure (arrows) from an elongated nucleus. Although some co-localisation (yellow) is seen with elastin (green), elastin fibres were generally found to contain little fibrillin 1

Fibrillin 2 appeared to be organised in numerous fibres, broadly orientated parallel with collagen bundles, with occasional oblique fibres (**Figure 3.5A**). These fibres were observed throughout ACL and PCLs and could also be highly branched (**Figure 3.5B**). The pattern of distribution appeared similar to that of fibrillin 1. Where elastin fibres were seen, fibrillin 2 was commonly co-localised (**Figure 3.6**). However, the majority of fibrillin 2 fibres did not appear to stain for elastin. Weak pericellular staining was occasionally noted throughout both CLs. No interligament or regional variation was apparent. Fibrillin 2 fibres were present on sections without collagenase and hyaluronidase treatment and staining appeared unaffected by enzymatic digestion.

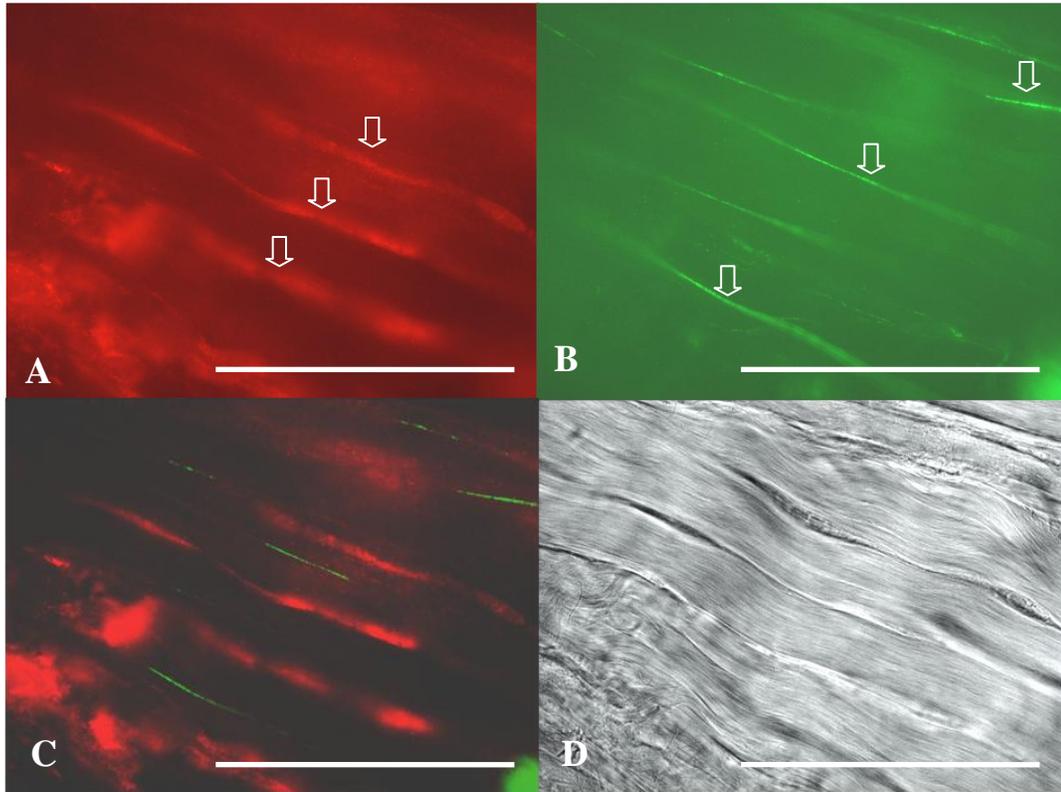


Figure 3.4: Elastin and fibrillin 1 rarely co-stain when in fibre form. Greyhound ACL, x100 (A) Fibrillin 1 (red) forming fibres (arrows). (B) Elastin (green) forming fibres (arrows). (C) Combined image of A and B showing lack of colocalisation of fibrillin 1 and elastin in fibre form. (D) Light image showing detail of collagen bundles. Both fibrillin 1 and elastin fibres run parallel with collagen bundles. Magnification bars=40µm

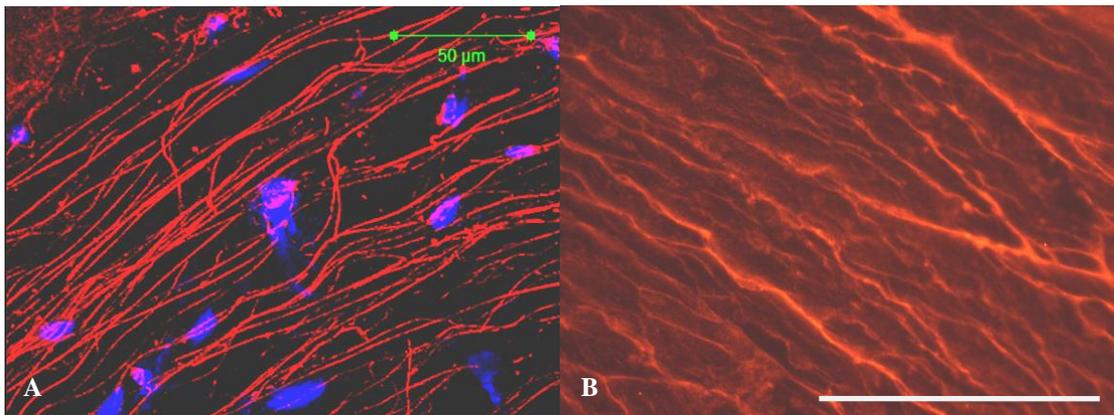


Figure 3.5 Distribution of fibrillin-2 in the canine CL complex. (A) ACL x63 CLSM image from fascicular region of ACL showing fibrillin-2 (red) in long and dense fibres broadly aligned with collagen bundles, with some branching. Nuclei are stained with DAPI (blue) (B) Longitudinal PCL, x40. Fibrillin-2 (red) shows a highly branched pattern. Magnification bar = 100µm.

3.3.6 NDIC

Perpendicular stretching of hydrated unfixed ligament under NDIC revealed complex interconnections between collagen bundles and fascicles when compared to unstrained tissue. These changes appeared consistent between CLs and between dogs. When the whole diameter of the ligament was subjected to lateral stress, all of the strain was observed in the interfascicular regions (**Figure 3.7A, B**). The collagen bundles remained tightly opposed while the interfascicular region showed loose but organised fibres.

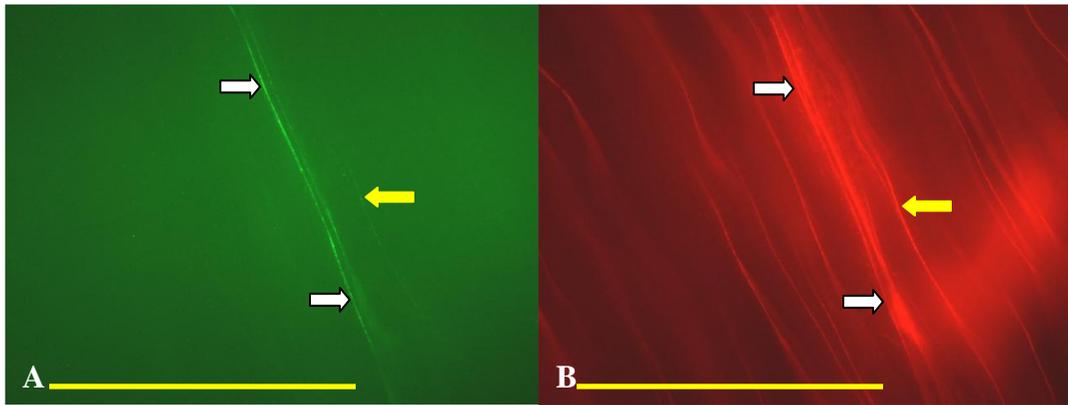


Figure 3.6: Co-staining of elastin and fibrillin 2. Greyhound ACL (A) Staining for elastin, x100. Two brightly staining parallel elastin fibres are noted (white arrows) with one solitary faint staining fibre (yellow arrow). (B) Two fibrillin 2 fibres corresponding to the elastin fibres are noted (white arrows). There is a fibrillin 2 fibre corresponding to the faint elastin fibre (yellow arrow). The majority of fibrillin 2 fibres do not co-stain for elastin. Magnification bars =40 μ m.

When individual fascicles were manipulated, perpendicular strain allowed observation of interbundle anatomy. With minimal strain, residual fibres were seen running transversely (**Figure 3.8A**), obliquely in one direction (**Figure 3.8B**) or obliquely in both directions. Following the removal of the lateral stress the bundles would recoil spontaneously. Further separation of collagen bundles separated the oblique fibres into S-shaped subdivisions which were again reversible following the removal of lateral stress. Fibres that spanned interbundle regions transversely could be found with or without oblique fibres and could be anchored by a single point on a collagen fibre or a more diffuse and complex attachment deeper into the bundle.

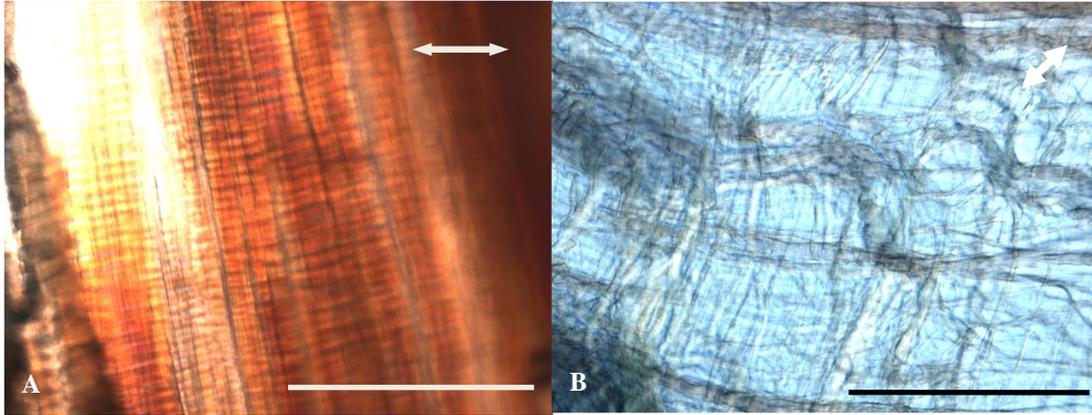


Figure 3.7: Variation in ligament strain during constant stress in the ACL in fully hydrated, unfixed tissue. Images (A) and (B) taken simultaneously from adjacent regions under identical stress. Double headed arrows indicate direction of applied stress. (A) Tightly adherent collagen bundles with no lateral separation with stress applied perpendicularly to collagen bundles. (B) Interfascicular region demonstrating loose but organised tissue following application of stress perpendicular to ligament fascicles. Interfascicular fibres run obliquely to fascicles in both directions. Magnification bars A and B=100 μ m

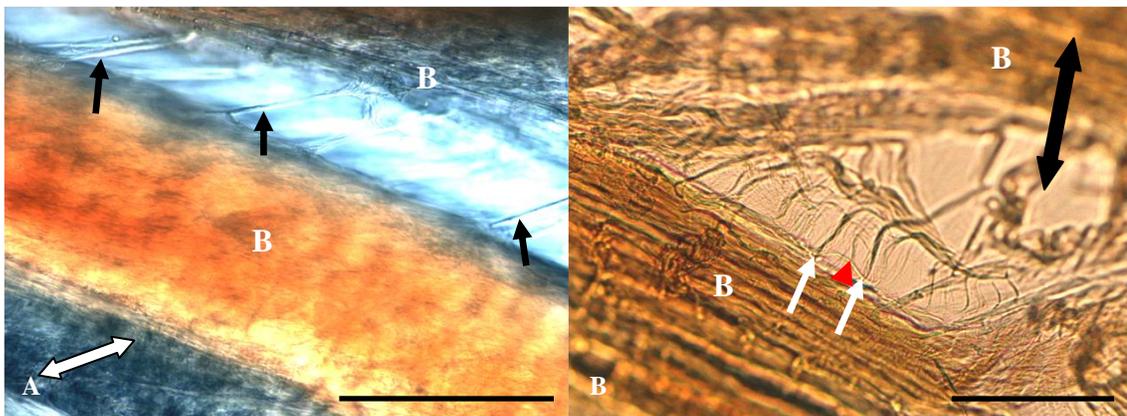


Figure 3.8: Variation in interbundle strain during application of perpendicular stress in the CLs in fully hydrated, unfixed tissue. (A) PCL x40 Direction of applied strain is shown by white arrow. Collagen bundles are marked as B. Straight, thick transverse interbundle fibres (black arrows) (B) ACL, detail of x40. Direction of applied strain is shown by black arrow. Collagen bundles are marked as B. Unilateral oblique interbundle fibres (white arrows). Subdivision of these oblique fibres is noted (red arrow). Magnification bars A=100 μ m and B=40 μ m

3.4 Discussion

In this study we have suggested a widespread presence of elastic fibres, elastin and fibrillins 1 and 2 in the canine CL complex. Analysis of subdivisions of the CL complex has suggested abundant oxytalan fibres and that these may have a similar pattern of distribution to the fibres comprising of fibrillins 1 and 2.

Elastin fibres were found throughout both ACL and PCLs and appeared abundant in the interbundle and interfascicular regions. Sparse fibres within bundles seem unlikely to contribute to the mechanical strength of the ligament but may provide additional stiffness at low strain and stress (Smith and Fazzalari, 2006). In the periodontal ligament, elastin fibres were suggested to provide mechanical support to the vascular network, restoring vessel shape following deformation (Sawada et al., 2006). During ligament micro-movement, blood vessels are thought to be occluded (O'Donoghue et al., 1971, Kobayashi et al., 2006) and such elastic mechanical support may also be important in the CL. While elastin fibres were only rarely found within bundles, oxytalan fibres were abundant. Certainly the high density of oxytalan fibres within collagen bundles would suggest some role in mechanical ligament function such as absorption of low strain stiffness, complementing the tensile strength of collagen fibres, or restoration of longitudinal conformation following longitudinal strain (Oxlund and Andreassen, 1980, Oxlund et al., 1988, Oakes and Bialkower, 1977, Lee et al., 2001, Karlinsky et al., 1976).

Where elastin formed part of an elastin fibre, fibrillin 2 but not fibrillin 1 appeared to commonly co-localise. The authors believe the pattern of fibrillin 2 staining was similar to both fibrillin 1 and oxytalan fibre distribution throughout the CLs. The density and distribution of oxytalan fibres appeared similar in histochemically stained and immunofluorescence sections (compare density in **Figure 3.2B** and **3.5B**). Thus fibrillin 2 appears to be a significant component of microfibrils in canine CL. As the MF is a key component of the elastin fibre, this explains the co-localisation of elastin and fibrillin 2. However immunostaining of fibrillin 1 revealed fibres likely to be microfibrils (Kielty, 2006, Yu et al., 2007) yet here was very little fibrillin 1 staining associated with elastin fibres. Collagenase digestion has been shown to remove fibrillin 1 from the MF potentially exposing the fibrillin 2 core (Charbonneau et al., 2010b) and in this study resulted in a reduction in the intensity of fibrillin 1 staining. As collagenase digestion was required to immunostain elastin, this may have disrupted the fibrillin 1 staining pattern.

In the canine CL complex, the distribution of fibrillin 1 and 2 differs from that described in other tissues. Fibrillin 2 has been considered to have limited distribution in adult tissue (Cain et al., 2006) but MFs may comprise of an inner core of fibrillin 2 surrounded by fibrillin 1 (Charbonneau et al., 2010b). In canine SDFT, only fibrillin 1 was observed to form fibres (Ritty et al., 2002). However the unmasking of the fibrillin 2 epitope may be a marker for MF degeneration (Charbonneau et al., 2010a). Canine CLs commonly have histologic changes considered to be degenerative (Vasseur et al., 1985, Comerford et al., 2006b) thus the widespread fibre pattern may reflect degeneration of the CL ECM unmasking fibrillin 2.

When imaging the variation in interfascicular and interbundle anatomy in unfixed, hydrated tissue the fibres of the interfascicular region appear loosely organised and may allow fascicles to move freely in relation to each other. Furthermore, the highly deformable nature of the interfascicular fibres could offer some stress protection for other structures such as blood vessels and nerves that run within these regions. Our observations suggested a similar hierarchical subdivision of interbundle fibres to that described in the human annulus fibrosus (Pezowicz et al., 2005). We have suggested interbundle and interfascicular fibres contain elastin and oxytalan fibres histochemically and through immunofluorescence. These elements show strikingly similar anatomical distribution to the interbundle and interfascicular fibres we demonstrated on unfixed, hydrated CL using NDIC (compare **Figure 3.1C** and **3.8A**). We have also suggested the interbundle and interfascicular fibres have elastic properties. Therefore it is not unreasonable to assume that these fibres may contain elastin and oxytalan fibres. A passive recoil system formed from elastin and oxytalan fibres offers a mechanism whereby ligament anatomy can be restored following deformation. Such a mechanism has been proposed in the heart valves (Vesely, 1998) and annulus fibrosus (Cloyd and Elliott, 2007, Yu et al., 2007).

This study was limited by the lack of quantifiable data. Thus our comparisons within and between ligaments and dogs were largely subjective. Lack of detailed information on husbandry and exercise history may affect comparison of these results. Although our study population was heterogeneous with differences in age, there appeared to be little variation in results between dogs and we believe our conclusions to be valid. Future quantification of our observations will enable us to assess the validity of low sample numbers.

In summary we have suggested that elastic fibres may have potentially important mechanical roles in this ligament complex. We described abundant oxytalan fibres composed of both

fibrillin 1 and 2 and have suggested this differs from the solely fibrillin 1-containing oxytalan fibres previously described in tendon (Ritty et al., 2002). The presence and location of fibrillin 2 in MFs in ligament may demonstrate a fundamental difference between ligament and tendon.

CHAPTER 4: Elastin in the cruciate ligament complex: a correlative histological and biochemical study

Abstract

Outline: To determine whether variation in elastin content might relate to histological and biochemical properties of the canine cruciate ligament (CL) complex. Such variation may provide insight into cruciate ligament physiology.

Method: Eleven pairs of CLs were harvested from seven skeletally mature Greyhounds and paired samples taken for histology and biochemical analysis. Sequential histology sections were assessed for extracellular matrix degeneration (haematoxylin and eosin), elastin fibre staining (van Gieson) and microfibril staining (Miller's stain). Novel scoring systems were developed to quantify extracellular matrix degeneration and microfibril staining. Samples for biochemical analysis were freeze-dried and dry weight, elastin, collagen and sulphated glycosaminoglycan content determined.

Results: Increasing degeneration of the extracellular matrix (ECM) was associated with increased microfibril staining pericellularly and within the ECM. The average value for elastin content in the anterior CL was 9.86 ± 3.97 % dry weight and $10.79 \pm 4.37\%$ in the posterior CL and this did not alter with advancing degeneration.

Conclusions: Production of microfibrils may reflect a healing response in degenerate CL ECM. Elastin forms a far greater proportion of canine CLs than previously suggested and may have non-mechanical roles.

4.1 Introduction

Cruciate ligaments (CLs) are the primary stabilisers of the knee (femorotibial) joint. The CL complex is comprised of the anterior cruciate ligament (ACL) twisting around the posterior cruciate ligament (PCL) (Arnoczky and Marshall, 1977). Anterior cruciate ligament (ACL) rupture is the most common orthopaedic condition to affect dogs and has a high incidence in people (Bennett et al., 1988, Renstrom et al., 2008). Canine ACL disease is recognised as degeneration of the CL extracellular matrix (ECM) which eventually leads to ligament rupture through non-contact injury. The ACL will not heal following rupture and attempts at primary repair have been unsatisfactory (Feagin and Curl, 1976). Little is known as to why the ACL does not heal following injury but it is thought that the mechanical environment of the knee in combination with exposure to synovial fluid and lack of soft tissue support all contribute (O'Donoghue et al., 1971, Arnoczky et al., 1979, Frank et al., 1985, Cameron et al., 1994).

Alterations to ACL matrix and cellular metabolism are implicated in the pathogenesis of non-contact ACL failure (Vasseur et al., 1985). Fibroblast phenotype changes and a loss of cells have been shown to occur prior to ACL failure (Comerford et al., 2006b, Vasseur et al., 1985). Through characterisation of cell and matrix changes, degeneration of CLs can be categorised on a scale from 0 (no degeneration) to 3 (severe degeneration) providing a numerical assessment of the degree of CL degeneration (Vasseur et al., 1985). Increased matrix turnover has been observed in the CLs of breeds at a high risk of ACL failure (Labrador retriever) when compared to breeds such as the greyhound at a low risk of failure (Comerford et al., 2004).

Elastin has traditionally been considered a minor component of the ECM of ligament tissue (Frank, 2004). Elastin fibres (EFs) are comprised of an amorphous elastin core within a fibrillin-containing microfibril (MF) scaffold and impart reversibly extensibility and resilience to soft tissue (Kielty, 2006). Although the role of EFs in the CL complex is not well understood, we have recently shown a wide distribution throughout the CLs and proposed a role in collagen reorganisation following ligament deformation (Chapter 3). To date, elastin has not been quantified biochemically in the CL complex of any species. Estimates have ranged from 0-6% elastin (Suzuki et al., 2008, Paatsama, 1952, Strocchi et al., 1992) and quantitative data on elastin content in ligament is also extremely limited. Elastin was shown to comprise 13.8% of the dorsal human posterior dorsal ligament (Nakagawa et al., 1994) and up to 9.3% of the human annulus fibrosus (Cloyd and Elliott, 2007). An elastin content of 22%

was reported in canine ligamenta flava using crosslink analysis, with the content correlating with EFs observed histologically (Ponseti, 1995).

Bundles of MFs are known as oxytalan fibres (OFs) and collectively, MFs, OFs and EFs are known as elastic fibres. Failure of elastic fibres has been implicated in a number of serious diseases (Kielty, 2006). The importance of MFs in ligament and tendon biology is highlighted where mutations in fibrillin 1 or 2 disrupt MF assembly, resulting in Marfan syndrome (MFS) or congenital contractile arachnodactyly (CCA) respectively (Kielty, 2006). Changes in elastic fibres have been associated with degeneration in other ligamentous tissue, such as the annulus fibrosus. In the human, degenerate annulus fibrosis had nearly a five-fold increase in elastin over that seen in healthy annulus fibrosus (Cloyd and Elliott, 2007). However, other observers, using histological methods, did not see increases in EFs in annulus fibrosus degeneration (Olczyk, 1994, Smith and Fazzalari, 2006). An age-related accumulation of mature but fragmented elastic fibres was observed in the human interspinous ligament, with a loss of OFs (Barros et al., 2002).

We have described previously the distribution of elastic fibres in the normal canine CL complex (Chapter 3). As elastic fibres may have important mechanical and biological functions, we aim to quantify elastin biochemically and compare the results with detailed histological analysis of CL degeneration and elastic fibre distribution. Elastin has never been quantified biochemically in the CL complex in any species, and its role in CL ligament ECM is unknown.

4.2 Materials and methods

4.2.1 Tissue preparation

Eleven pairs of CLs were harvested from seven skeletally mature greyhounds with no macroscopic evidence of knee pathology. The animals were euthanatized for reasons not related to this study and informed consent, according to University guidelines, was obtained in each case prior to tissue removal. Each CL was sectioned into proximal, middle and distal sections. Of each of these subdivisions, further division into thirds through longitudinal sectioning allowed one third to be formalin fixed for histology and the remaining two thirds were stored at -80°C until required for biochemical analysis.

4.2.2 Histology: Staining

Sequential sections of 4µm from paraffin-embedded samples were stained with 1) haematoxylin and eosin (H&E), 2) Verhoeff's iodine-iron haematoxylin (EVH) to show only EFs and 3) Miller's stain, M (also referred to as Weigert's stain), to show both elastin and OFs. This series allowed assessment of tissue architecture, and through comparison of Verhoeff's- and Miller's-stained sections, differentiation of elastin fibres and microfibrils. Images were recorded on a dedicated microscope (Nikon Eclipse 80i). All sections were read by two observers blinded to section location or animal (KDS and EJC) on two separate occasions at least one week apart.

4.2.3 Histology: Scoring methods

H&E: H&E sections were assessed for signs of CL degeneration. All GH samples were graded 0-3 according to criteria previously described (Vasseur et al., 1985). The broad grade 1 category was subdivided by the authors in this study with a more detailed scoring system (**Table 4.1**). A score from 0-4 was awarded based on the extent of the changes for each of eight factors giving a range of possible scores from 0-24. These results are referred to as modified Vasseur Score (mVS).

Miller's Stain: A scoring system was developed to quantify changes in microfibril staining (**Figure 4.1**). Increased staining in interfascicular and interbundle regions, ligament substance (inrabundle), as well as the extent and degree of pericellular staining, could be awarded up to two points giving a score range of 0-10. These results are referred to as Miller's Score (MS).

Rounding of cell nuclei (without halo formation)
Loss of cell density from regions of normal collagen architecture
Loss of cell density from regions of reduced collagen density
Loss of cell density from regions of abnormal collagen architecture
Formation of cell chains
Fibrocyte chondroid metaplasia (rounded nucleus within halo)
Fibrocyte chondroid metaplasia associated with mineralisation
Fibrocyte chondroid metaplasia associated with mineralisation and abnormal adjacent collagen architecture

Table 4.1: Criteria for modified Vasseur scoring. A score from 0-4 was awarded based on the extent of the changes for each of the eight factors: absent=0, isolated=1, affecting up to 25% of tissue=2, affecting 25-50% of tissue=3, affecting >50% tissue=4.

	Low (Normal)		Moderate		Numerous
Interfascicular oxytalan fibre staining	<input type="checkbox"/>				
Interbundle oxytalan fibre staining	<input type="checkbox"/>				
Substance oxytalan fibre staining	<input type="checkbox"/>				
	Absent	Isolated	Low	Moderate	Frequent
<u>Pericellular deposition</u>	<input type="checkbox"/>				
	Minimal (orN/A)		Moderate	Marked	
<u>Degree of pericellular deposition</u>	<input type="checkbox"/>				

Figure 4.1: Miller’s stain scoring system sheet. Each of the 5 categories scores up to 2 points, with 0, 1, or 2 respectively for categories 1-3 and 5, and 0-2 in 0.5 increments for category 4, giving a range of 0-10.

4.2.4 Biochemical analyses

Paired ligament samples from proximal, middle and distal regions were used for elastin, collagen and sulphated glycosaminoglycan (sGAG) assays. Samples were weighed, dried overnight at 60°C and reweighed again to obtain % dry weight.

Elastin: Elastin was measured using the Fastin™ dye-binding assay (Biocolor Ltd, N. Ireland). The average dried weight of samples used for elastin assay was 14.2mg (4.4-24.9mg). Insoluble cross-linked elastin was converted to a soluble form (α - and κ -elastin polypeptides) by heating the ligament to 95°C in 0.25M oxalic acid (35295, Sigma-Aldrich, UK) for 1 hour. They were centrifuged at 3000g for 10 minutes and the supernatant extracted. Preliminary analysis showed this needed to be repeated 6 times to extract all elastin. Pooled extracts were incubated with the dye 5,10,15,20-tetraphenyl-21,23-porphine tetrasulphonate for 90 minutes to bind α - and κ -elastin, lathyrogenic elastins, and soluble tropoelastin. Following centrifugation (10000g) for 10 minutes, the residue was resuspended in a dissociation agent (guanidine HCl and propan-1-ol) and absorbance read in a microwell plate reader (Multiskan EX, Thermo) at 440nm. Samples were analysed in quadruplicate for each sample analysed.

Collagen: Ligament samples were digested for 24 hours with papain (300 μ g/ml, P4762, Sigma-Aldrich, UK) in PBS with 5mM Cysteine HCl and 5mM EDTA at 60°C. Collagen was measured using a colourimetric assay to determine hydroxyproline (OHPro) content (Jamall et al., 1981) assuming 13.7% of total collagen as OHPro. Briefly, samples were hydrolysed in 6N HCl for 24 hours at 110°C before freeze-drying. Following reconstitution in water, quadruplicate aliquots were thoroughly mixed with a solution containing sodium acetate trihydrate, tri sodium citrate dehydrate, citric acid and, propan-2-ol and chloramine T. The colour reagent, containing dimethylamino benzaldehyde, perchloric acid and propan-2-ol was added, then heated at 70°C for 20 minutes. Standards from stock hydroxyproline were used to calculate the standard curve from 0-10 μ g/ml.

sGAG: Ligament samples were digested for 24 hours with papain (300 μ g/ml, P4762, Sigma-Aldrich, UK) in PBS with 5mM Cysteine HCl and 5mM EDTA at 60°C. Total sGAG was assessed using the 1,9-dimethylmethylene blue (DMMB) dye binding assay (Farndale et al., 1986). Quadruplicate aliquots of papain-digested ligament samples were immediately analysed at 540 nm following the addition of DMMB. The assay was calibrated by use of standards up

to 40 µg/ml shark chondroitin sulphate (Sigma-Aldrich, UK), and sGAG concentration obtained by comparison with the standard curve.

4.2.5 Data and statistical analysis

Elastin, collagen and sGAG were normalised to total ligament dry weight. Additionally, elastin was normalised to collagen. A two factor ANOVA was used where the two factors were CL (2 levels: ACL and PCL) and location (3 levels: proximal, middle and distal) to assess differences in biochemical and histological data. Where significance at 5% was met, a Bonferroni *post hoc* test was applied and t-tests were used for direct comparisons. Pearson’s correlations were used to assess relationships between factors, with significance set at 5%. Results are presented as mean values ± SD. Kendall’s coefficient of concordance was calculated for intra- and inter-observer concordance of both modified Vasseur and Miller’s scoring. Kendall’s coefficient of concordance ranges from 0 to 1 with values closer to 1 suggesting a high degree of concordance. Data were analysed using Minitab Statistical Software (Minitab, UK).

4.3 Results

Animals: Summary data are shown in **Table 4.2**. All dogs had been in training within the last 6 months.

	Dogs	Pairs of CLs	Age Range (months)	Median Age (months)
Male	4	6	32-60	39
Female	3	5		
Total	7	11		

Table 4.2: Summary data for animals

4.3.1 Histology: H&E

All CL samples stained with H&E were graded as grade 1 according to the published system (Vasseur et al., 1985). All were also graded on the modified scoring system (mVS), with an average score of 11.9 ± 3.3 (range 5.5-21.5, median 12). There were no significant differences in mVS within or between ligaments. Kendall’s coefficient of concordance for inter-observer concordance was 0.85 for mVS. Intra-observer concordance for KDS was 0.94 and for EJC was 0.95 (**Table 4.3**).

	Intra-observer variation		Inter-observer variation
	KDS	EJC	KDS and EJC
Observers			
Modified Vasseur Score	0.94	0.95	0.85
Miller's Score	0.91	0.94	0.85

Table 4.3: Kendall's coefficients of concordance for histology scoring methods.

4.3.2 Histology: Elastic fibre staining

General: In the majority of tissue EFs showed no differences in either size or distribution, but where there was loss of collagen architecture a reduction in EFs was noted. If mineralisation was present, EFs were only rarely found. In contrast, the samples with higher grade 1 degeneration showed a marked increase in OF quantity, with changes in distribution. Increased OF staining was differentially noted within three subdivisions of the ligament substance in all CLs with high grade 1 changes.

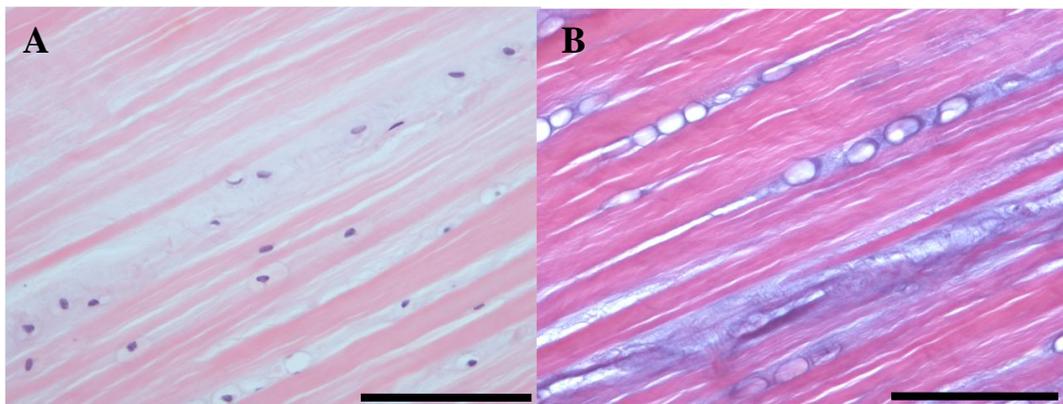


Figure 4.2: Histologic changes in CLs with degeneration. (A) H&E, ACL, x40. Loss of cell density, collagen density, collagen architecture along with chondrocytic change are all clear. (B) Miller's stain, ACL, x40. Image from same section as A and shows marked pericellular staining of most chondrocytic cells, with moderate interbundle and substance staining. Magnification bars = 100µm

Interfascicular: Staining of OFs as previously described (Chapter 3) was loose and mesh-like. Cells that had undergone chondrocytic change commonly, although not ubiquitously, had increased pericellular OF staining, which was also occasionally seen in fibrocytes with rounded nuclei (**Figure 4.2**). CLs with more advanced degeneration had increased interfascicular OF density through increased numbers of small OFs in a mesh pattern.

Interbundle: Staining of OFs showed marked differences from low-grade 1 sections (**Figure 4.3A,B**). Where collagen bundle integrity remained largely intact, the increased OF staining took the form of small fibres running largely parallel or obliquely to the collagen. The increase in staining could be marked, often seeming to separate collagen bundles. If there was loss of collagen architecture, then these areas of loss commonly showed increased OF staining with much less regular organisation, more similar to the mesh seen in the interfascicular regions (**Figure 4.4 C,D**).

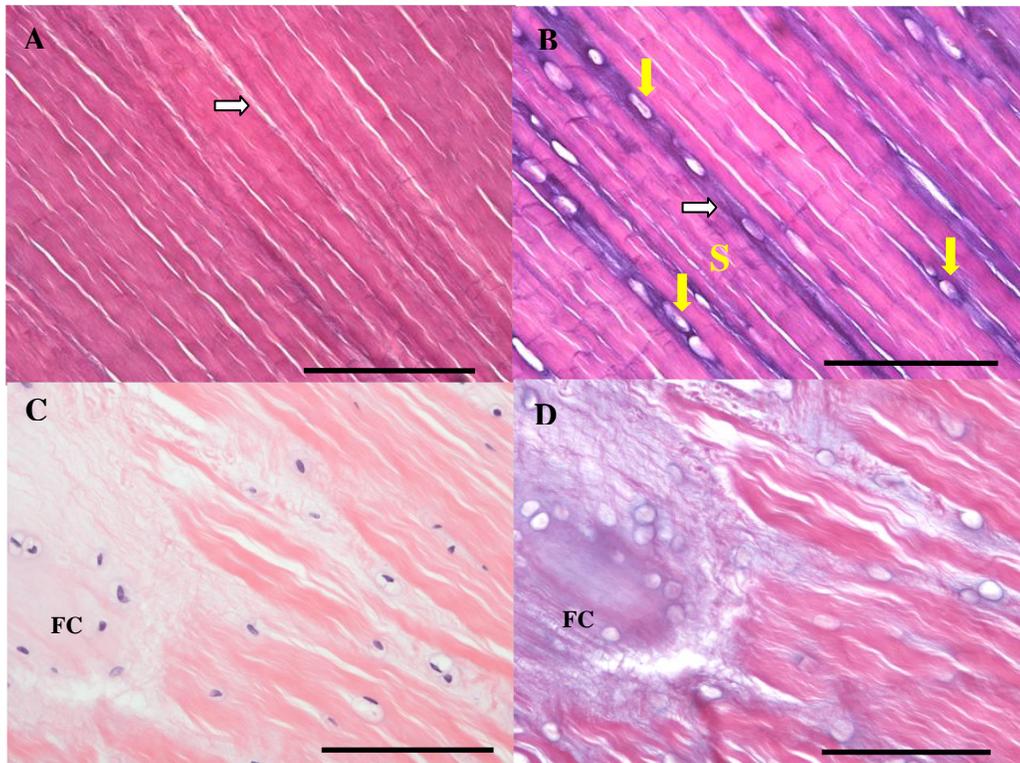


Figure 4.3: Variation in interbundle staining. (A) Miller's stain, ACL, x40. Minimal interbundle OF staining (example arrowed) is seen in this section with low grade 1 (Vasseur) changes. (B) Miller's stain, ACL, x40. Marked interbundle staining is seen (white arrow) along with widespread and marked pericellular staining (yellow arrows) and a moderate degree of CL substance staining (in the region marked S). (C) H&E, PCL, x40. Area of complete loss of collagen architecture with mineralisation is seen (marked FC). This would be considered fibrocartilaginous change. (D) Miller's stain, PCL, x40. Image from same section as C. Area of fibrocartilaginous change shows dense and fine meshwork of MFs, manifesting as increased purple staining. Magnification bars = 100µm

Bundle substance: Increased OF staining was also sometimes present in regions with a loss of collagen density. These OFs were usually aligned with the collagen, and were very fine.

Score and inter/intra-observer data: Statistically significantly higher MS were seen in the ACL (5.3 ± 1.9) than the PCL (4.0 ± 1.4), $p=0.02$ (**Figure 4.4**). Kendall's coefficient of concordance for inter-observer concordance was 0.85 for MS. Intra-observer concordance for KDS was 0.91, and for EJC was 0.94. The scoring system for the Miller's staining showed a strong correlation with the mVS ($r=0.690$, $p<0.001$), confirming the histological observation (**Figure 4.5**).

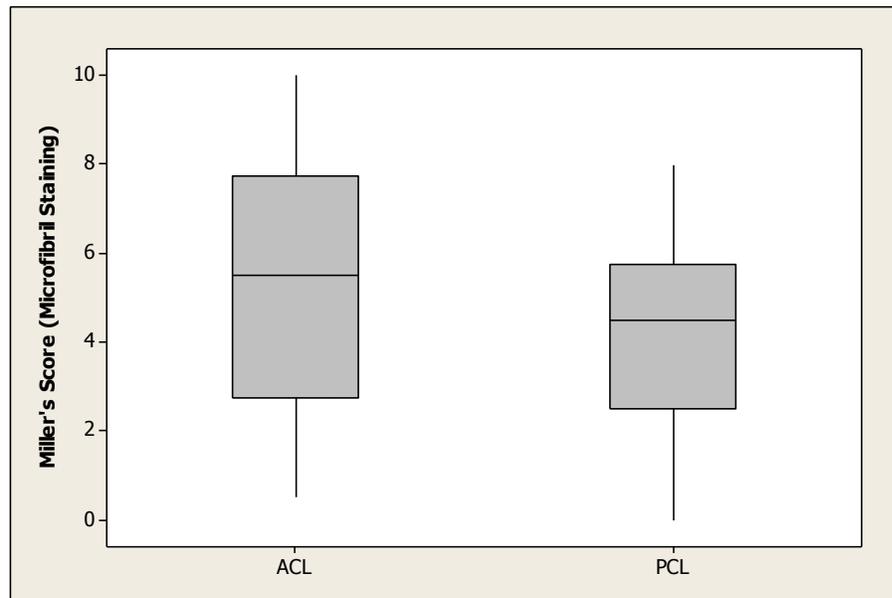


Figure 4.4: Boxplot summary of ACL and PCL results for Miller's score. The ACL has a significantly greater MS than the PCL ($p=0.02$)

4.3.3 Biochemical analyses

Water Content: Water content in the ACL averaged $59.3\% \pm 3.1$ in the ACL and $61.9\% \pm 2.2$ in the PCL (**Figure 4.6**). The ACL had a significantly lower water content than the PCL ($p=0.002$). Water content demonstrated statistically significant difference according to location within ligament ($p=0.035$). The mid PCL had significantly higher water content than proximal and distal PCL and mid ACL. No correlations with other factors (elastin, collagen, sGAG, mVS and MS) were seen.

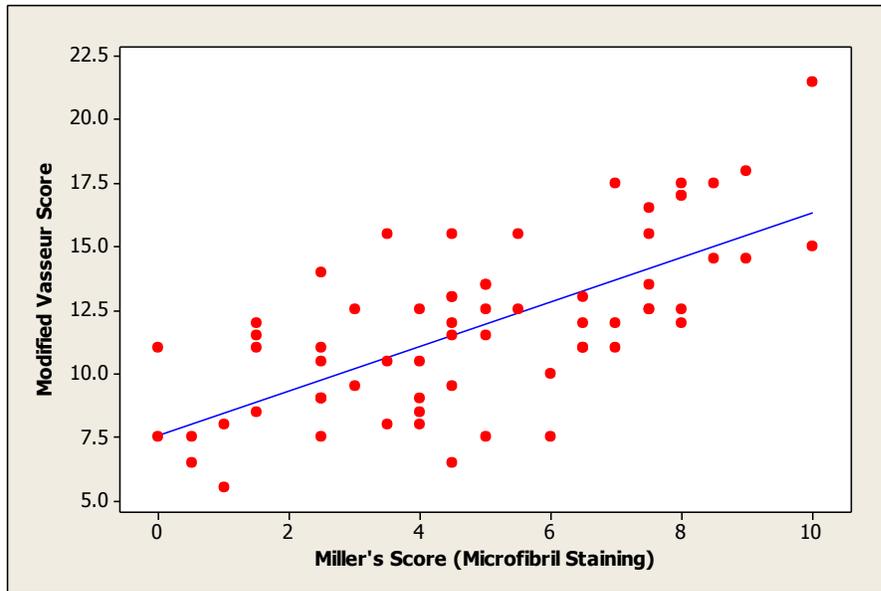


Figure 4.5: Significant correlation between modified Vasseur score and Miller's score ($r=0.690$, $p<0.001$). As degeneration of the ligament advances increased MF staining is noted.

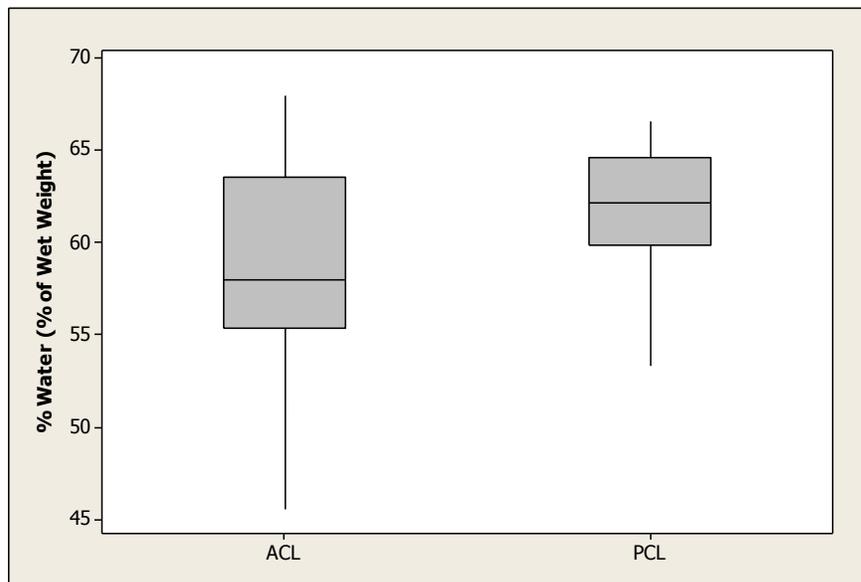


Figure 4.6: Boxplot summaries of ACL and PCL results for water content (%). The PCL has a significantly greater water content than the ACL ($p=0.002$).

Elastin: Elastin averaged 9.86 ± 3.97 % (total ligament dry weight) in the ACL and 10.79 ± 4.37 % in the PCL, but this difference was not statistically significant ($p=0.28$). A marked variation in elastin content in both ligaments was noted; ranging from 5.9% to 19.4%. Despite this large range, pairs of ACLs and PCLs had very similar intra- and inter-ligament elastin content (**Figure 4.7**); there were no significant statistical differences between pairs of ACLs and PCLs or between left and right knees. There was not a statistically significant variation in elastin content according to location (proximal, middle or distal), age, sGAG or % dry matter. No correlations were noted between elastin and mVS (Pearson correlation ($r= -0.016$, $p= 0.897$) and MS (Pearson correlation = -0.105 , $p= 0.403$) in the whole or ligament subsections. Normalising elastin to collagen rather than dry weight did not result in any significant variations or relationships with location or other parameters.

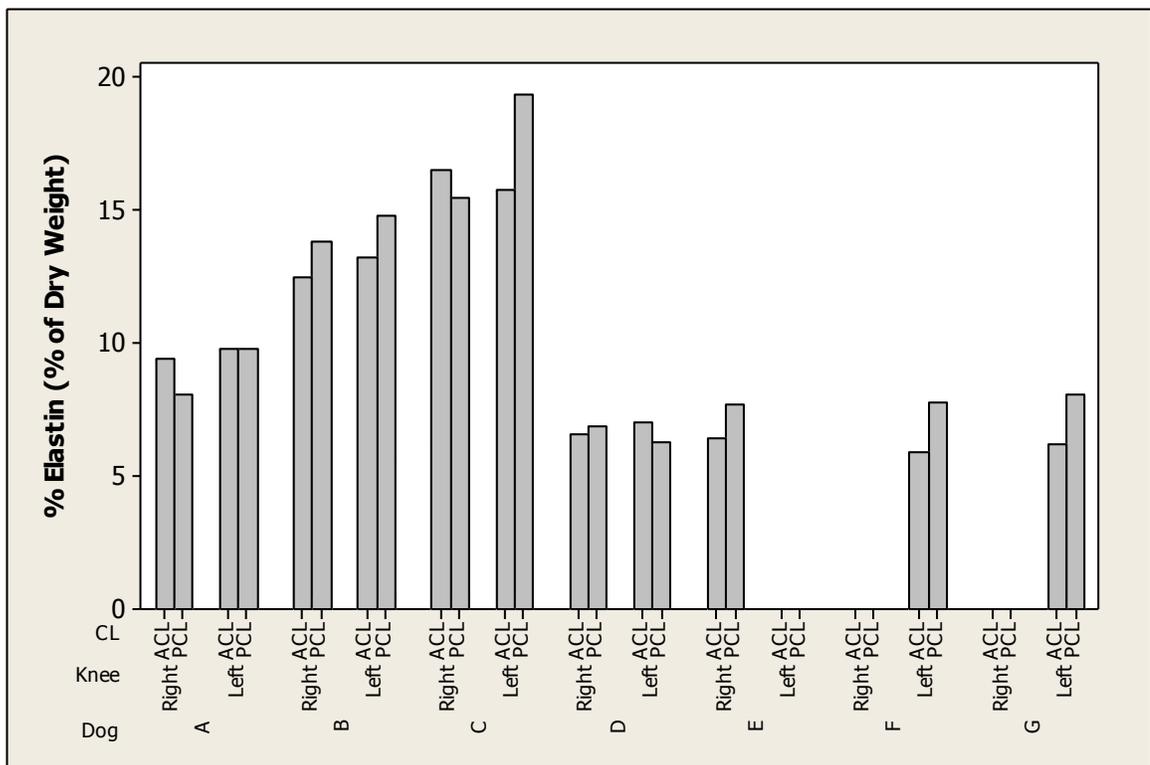


Figure 4.7: Elastin content (%) of canine CLs. In dogs A-D, both left and right knees were studied. Pairs of ACLs and PCLs share similar elastin content, and this is conserved between left and right knees. However large variation between dogs can be seen.

Collagen: Collagen as a percentage of dry weight averaged $75.34\% \pm 5.19$ (range 62.2%-86.7%). A significant positive correlation with age was noted ($r=0.813$, $p<0.001$), but no other significant variations or relationships with location or other parameters (elastin, water content, sGAG, mVS and MS) were observed.

sGAG: sGAG content as a percentage of dry weight (% sGAG) ranged from 0.012 to 0.148%, (average $0.062\% \pm 0.031$). Significantly greater % sGAG was noted in the ACL ($0.081\% \pm 0.019$) than the PCL ($0.043\% \pm 0.010$, $p<0.001$), a difference also significant for location within the ligaments ($p=0.016$). Within the ACL, the %sGAG distally was significantly higher than both mid ($p=0.012$) and proximal ligament ($p= 0.025$), while in the PCL, the middle ligament had significantly higher %sGAG than that found both proximally ($p<0.001$) and distally ($p<0.001$). Significant correlation with mVS ($r=0.389$, $p=0.002$) and MS ($r=0.607$, $p<0.000$) were also observed (**Figure 4.8**).

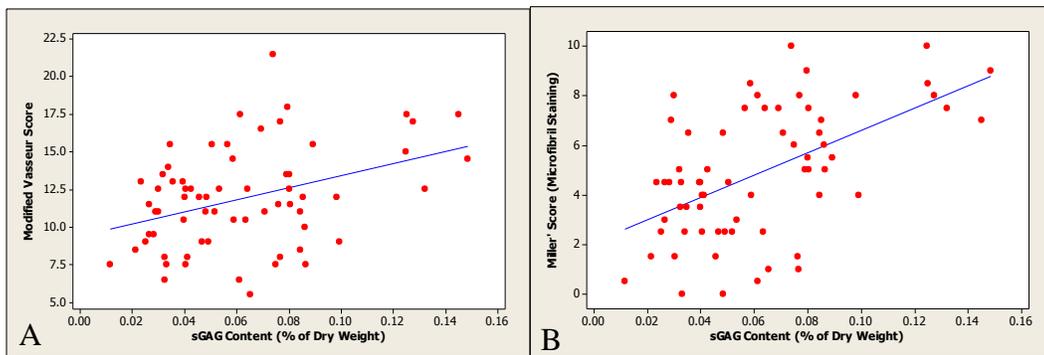


Figure 4.8: Relationships between sGAG content and histological data. (A) Significant positive correlation between degeneration (modified Vasseur score) and sGAG content ($r=0.607$, $p<0.001$). (B) sGAG content correlates positively with Miller's score (oxytalan fibre) staining ($r=0.389$, $p=0.002$).

When the results of the elastin, collagen and sGAG assays were combined, the total percent of dry weight for each ligament averaged $84.7\% \pm 4.68$ (range 72.4%-94.7%).

4.4 Discussion

In this study we have shown that elastin fibres appear to be lost with CL degeneration, but as this degeneration advances there is a marked production of oxytalan fibres. We have developed two novel scoring systems to assess ligament degeneration and OF staining that have shown excellent repeatability. Using our histological scoring systems, we have shown the

increase in oxytalan fibres to be proportional to the degree of degeneration. Furthermore, we have shown the elastin content of canine CLs to be far greater than previous estimates and to have no relationship with progressive degeneration of the CL, or with the other ligament components sGAG, collagen and water.

It is not fully understood why the canine ACL undergoes degeneration and fibrocartilage formation prior to rupture. Multiaxial stresses experienced by CLs likely include compression where they twist or contact other structures within the knee, such as the intercondylar notch of the femur (Arnoczky and Marshall, 1977, Harari, 1993, Comerford et al., 2006a). Fibrous connective tissue under compressive load may form fibrocartilage (Milz et al., 2005). Alternatively the changes may be a protective metabolic response (Connor et al., 2001, Schipani et al., 2001, Grimshaw and Mason, 2001) or reflect tissue injury (Vasseur et al., 1985). Degeneration has been identified throughout the CL complex and the mid-ACL has been shown to undergo the region of greatest degree of degeneration, proposed to lead to rupture of the ligament in some breeds (Paatsama, 1952, Vasseur et al., 1985). In choosing the greyhound as our study breed, a breed with an extremely low incidence of ACL rupture, we hoped to obtain tissue that could be considered 'undiseased'. However degeneration may be present in all breeds of dog and as a correlation with age and degeneration has been described, obtaining non-degenerate tissue may not be possible (Comerford et al., 2006b, Vasseur et al., 1985). Changes consistent with mild degeneration were found in every section examined in our study. As no significant differences in the pattern and distribution of change was observed between the ACLs and PCLs examined in this study, our results suggest degeneration in canine CLs are generalised.

As far as we are aware, increased production of MFs or OFs with advancing degeneration has not been described in any ligament. The lack of correlation between elastin content and MS in the dog would appear to suggest the OF production is not associated with either elastin production or subsequent development to EFs, suggesting that it is entirely fibrillin driven. OFs may have a number of roles in the CL complex including provision or maintenance of elasticity, stabilisation of blood vessels, anchoring tissue or guidance of cell migration (Fullmer et al., 1974, Chantawiboonchai et al., 1998, Everts et al., 1998, Tashiro et al., 2002). Studies on the developing rat periodontal ligament showed concurrent expansion of the OF network and development of the vascular system, and it was proposed OFs have a role in the maintenance of integrity of the vascular system (Tashiro et al., 2002). A loss of OFs has been described with age and degeneration in the human AF (Barros et al., 2002) and fibrillin is lost

from ageing human ligamenta flava (Osakabe et al., 2001). However, assembly of OFs is commonly seen in healing responses in artery (Sinha et al., 2001), myocardium (Vracko et al., 1990), muscle (Fricke et al., 2008) and skin (Tsuji and Sawabe, 1987). As the tissue in our study was from a breed in which the observed degenerative changes almost never progress to ACL rupture, these studies would suggest that the increased quantity of OFs observed in greyhound CLs reflects a healing response.

Previous studies have estimated elastin content in the CL complex from 0-6% using histochemical staining of EFs, crosslink analysis (demosine and isodesmosine) or electron microscopy (Paatsama, 1952, Vasseur et al., 1985, Strocchi et al., 1992). These methods may have underestimated elastin content by only analysing intact EFs or have used crosslink analysis on formalin-fixed tissue, known to significantly affect the concentration of these crosslinks (Abe et al., 2003). In this study we show the elastin content to be significantly greater than previously reported, ranging from 5.9 to 19.4% of ligament dry weight. The lack of EFs in areas of mineralisation would suggest that as CL degeneration progresses, there would be an overall loss of EFs. As EFs are not produced in adult tissue, these changes reflect an irreversible loss, and may have significant implications for ligament physiology (Sherratt, 2009). There appeared to be no correlation between mVS and elastin content suggesting that loss of EFs may not affect total elastin content. Despite EFs being sparse in the CL, elastin formed approximately 10% of dry weight of CL tissue suggesting that not all elastin in CL tissue is contained within EFs. Levels of elastin are relatively homogenous within and between pairs of CLs, and left and right knees, but vary between dogs. Such idiosyncrasy may arise geno- or phenotypically.

Elastin has been suggested to have a mechanical role in the CL complex. It has been proposed that EFs absorb recurrent maximal stresses and OFs distribute complex multiaxial stresses through the collagen architecture (Strocchi et al., 1992). We have previously suggested a role for EFs in bundle reorganisation following CL deformation (Chapter 3), a mechanism also proposed in the aortic valve (Vesely, 1998) and AF (Cloyd and Elliott, 2007, Yu et al., 2007). However, variation in elastin content did not reflect increased numbers of EFs and bore no relationship to OF production. Degeneration of the annulus fibrosus was associated with a five-fold increase in elastin over that seen in healthy annulus fibrosus (Cloyd and Elliott, 2007) but no increase in EFs (Olczyk, 1994, Smith and Fazzalari, 2006). Furthermore, a decrease in elastin crosslinks with age (which has been shown to correlate with degeneration (Cloyd and Elliott, 2007) suggested the increase in elastin was non-fibrous in nature (Osakabe

et al., 2001). It would therefore seem plausible that the variation in CL elastin observed between dogs may have non-mechanical functions. Although our study lacked samples of tissue free of degenerate changes and of high grade degenerative tissue, the absolute lack of any relationship with mVS and elastin within grade 1 ligament degeneration suggests elastin does not play a significant role in its progression once initiated. By expressing elastin as a percentage of dry weight and as a ratio to collagen confirmed changes in other ligament constituents were not masking any possible relationship.

We found significant differences in biochemical results observed within and between CLs such as higher sGAG in the ACL than the PCL and in the mid section of each ligament when compared to the distal section. Increasing sGAG could be associated with fibrocartilage production in the CLs and a correlation with degeneration (mVS) was noted (**Figure 4.7**). Increasing sGAG with degeneration has been well documented in other connective tissue such as tendon (Chard et al., 1994, Fu et al., 2007). Although the distal ACL had higher sGAG than other regions of the ACL, mineralisation at the ligament insertion site may have lead to this higher than expected result (Wang et al., 2006).

Our study has several limitations. In our study, the elastin assay used was unable to distinguish between insoluble and degraded elastin peptides. It may be that the elastin measured comprises significant elastin peptides and monomers trapped within the matrix. Further work is required to determine whether the elastin measured is the tropoelastin precursor, insoluble elastin or degraded peptides. Although the Fastin™ assay has not been used in the dog previously, we did not feel it necessary to validate it with traditional methods of elastin extraction such as the hot alkali method (Mecham, 2008). The Fastin™ assay has been used in a number of tissues and species, where results of quantification have correlated with other biochemical methods of quantification (Romanowicz and Sobolewski, 2000, Cloyd and Elliott, 2007). Our samples lacked more advanced degeneration that becomes more apparent in the mid CLs (Vasseur et al., 1985, Narama et al., 1996). Evaluation of CL elastin and OFs through high grade degenerate change in a range of domestic breeds will be required to confirm the conclusions of this study.

In this study we have demonstrated elastin forms a far greater proportion of the canine CLs than has previously reported. We have shown mild degenerative ligament changes to be associated with increasing prevalence of OFs which has never been described in any other

species or ligament. We believe this may reflect an attempt to repair the CL and is part of the CL healing response preventing ligament rupture in a breed of dog at low risk of CL rupture.

CHAPTER 5: The effect of elastin degradation peptides on canine anterior cruciate ligament cell cultures

Abstract:

Outline: To determine whether elastin degradation peptides (EDPs) containing the VGVAPG motif exert a biological effect on the cells of the canine anterior cruciate ligament (ACL)

Method: ACL cells were harvested from four skeletally mature greyhounds. Cells were cultured to confluence in 6 well plates then treated with TNF- α , TGF- β 1, EDPs and a combination of TGF- β 1 and EDPs for 6 and 24 hours. mRNA was extracted from the cells and using real time quantitative RT-PCR the effect of each treatment on a panel of 16 genes was assessed.

Results: Increased collagen 2a1 gene expression was seen following 6 (p=0.025) and 24 hours (p=0.05) of exposure to TGF- β 1. Increased transcription of fibrillin 2 following 24 hours of exposure to EDPs alone (p=0.032) and in combination with TGF- β 1 (p=0.018) was seen. There was a synergistic effect of EDPs and TGF- β 1 resulting in increased transcription of elastin laminin receptor 1 (p=0.014).

Conclusions: TGF- β 1 may be involved in collagen type II gene transcription which may result in ACL fibrocartilage production. Increased transcription of elastin laminin receptor 1 from treatment with EDPs in combination with TGF- β 1 may increase cell receptivity to EDPs. EDPs may play a key role in ACL response to injury through production of fibrillin 2.

5.1 Introduction

The anterior cruciate ligament (ACL) is the primary stabiliser of the knee and serves to limit anterior translation and rotation of the tibia. ACL rupture is a major cause of knee morbidity in man and the dog (Bennett et al., 1988, Frank and Jackson, 1997). In the canine ACL, alterations to ligament morphology have been described characterised by loss of collagen architecture and fibrocartilage production. The mechanism for these cellular changes is not understood, but may be an adaptive change (Connor et al., 2001, Schipani et al., 2001, Grimshaw and Mason, 2001, Comerford et al., 2006b, Benjamin and Ralphs, 1998) or a degenerative change preceding ACL failure (Vasseur et al., 1985).

The ACL will not heal following rupture and attempts at primary repair have been unsatisfactory to date (Feagin and Curl, 1976). Little is known as to why the ACL does not heal following injury but it is thought that the hostile mechanical environment of the knee in combination with exposure to synovial fluid and lack of soft tissue support all contribute (O'Donoghue et al., 1971, Arnoczky et al., 1979, Frank et al., 1985, Cameron et al., 1994). Healing following injury is dependent on achieving the appropriate balance of degradation and synthesis (Schultz and Wysocki, 2009). Ligament healing is primarily associated with the production of more collagen type III in relation to type I (Shrive et al., 1995). Growth factors such as TGF- β 1 stimulate proteoglycan and collagen synthesis, down regulating matrix degrading enzymes (Taipale et al., 1998) and cytokines such as TNF- α trigger production and activation of matrix degrading enzymes (Brenner et al., 1989). Healing depends on the interaction between ACL fibroblasts, the extracellular matrix (ECM) and growth factors (Schultz and Wysocki, 2009, Iozzo, 1998). Knowledge of how ACL fibroblasts respond to these factors is fundamental to understanding the limited regenerative ability of the ACL.

Cruciate ligament (CLs) are comprised of approximately two thirds water and one third solid (Boorman et al., 2006, Comerford et al., 2005). The solid component is principally collagen type I, accounting for 75% of the dry weight, the rest being proteoglycans, microfibrils, elastin and other proteins (Frank, 2004). Microfibrils are polymers of fibrillins 1 and 2 and are considered to have a structural role in ligament (Kielty et al., 2002a). Elastin fibres comprise of a central cross-linked core of highly extensible elastin surrounded by a supporting sheath of microfibrils (Kielty, 2006). Crosslinked elastin is insoluble and highly resistant to degradation. However during inflammation, insoluble elastin is degraded by cathepsins B and K (Novinec et al., 2007, Boudier et al., 1991) and matrix metalloproteinase (MMP) classes especially

MMPs -2, -9 and -12 (Mecham et al., 1997, Ashworth et al., 1999b). The resultant elastin peptides are able to influence the behaviour of a wide variety of cells including fibroblasts, macrophages and neutrophils (Duca et al., 2004, Privitera et al., 1998). Elastin peptides are transduced through the elastin laminin receptor 1 (ELR1), a spliced variant of β -galactosidase (Hinek et al., 1993). Peptides containing VGVAPG and GXXPG peptide sequences have been shown to bind to ELR1 (Brassart et al., 2001, Duca et al., 2004).

Biologically active fragments generated from ECM through protease action have been described in other tissues including lung and skin (Gaggar et al., 2008). Modes of action include chemoattraction of cells displaying ELR1 on their surface, triggering differentiation and activation of protein production (Gaggar et al., 2008, Mochizuki et al., 2002, Simionescu et al., 2005). The potential importance of EDPs in disease pathophysiology has been highlighted in a study that showed chronic lung disease was prevented in knockout mice with the inability to produce EDPs (Hautamaki et al., 1997). Given that the proportion of elastin in the canine ACL ranges from 6-20% (Chapter 4), the potential consequences of unregulated protease activation could be greatly underestimated (Djekic et al., 2009).

The production of enzymes that can hydrolyse elastin, such as MMP 2 and cathepsin K, has been demonstrated in both healthy and CLs with fibrocartilaginous change (Muir et al., 2002, Comerford et al., 2005, Barrett et al., 2005). However, nothing is known of the potential effects of the resultant degraded elastin fragments on the physiology of ACL cells. Enzymes associated with the hydrolysis of elastin have also been shown to degrade elastin-associated glycoproteins releasing TGF- β 1 (Sinha et al., 2002). Mechanical trauma, such as may occur where the CLs twist around each other, may also lead to damage and subsequent catabolism of elastin fibres (Arnoczky and Marshall, 1977, Vasseur et al., 1985, Lavagnino et al., 2006). In this study we aim to assess whether elastin peptides exert a biological effect on canine anterior cruciate ligament (ACL) cells. Furthermore we will compare the effect of elastin peptides to TGF- β 1 and TNF- α and look at the effect of elastin peptides in combination with TGF- β 1 mimicking what may occur following elastin hydrolysis of the ACL.

5.2 Materials and methods

5.2.1 Elastin peptides

Elastin degradation products (EDPs) were obtained commercially (CB573 Elastin Products Company, MO, USA). EDPs are a mixture of highly soluble fragments with a range of

molecular weight of 1000 to 60,000 Da, prepared by hydrolysing then purifying bovine nuchal ligament. EDPs have been shown to contain the VGVAPG repeating sequence.

5.2.2 Donors, extraction and preparation of cells

ACLs from paired knee joints were harvested by sharp dissection from four skeletally mature Greyhounds (GH) with no macroscopic evidence of knee joint pathology. The animals were euthanized for reasons other than musculoskeletal disease and informed consent, in accordance with University ethical guidelines, was obtained in each case for tissue removal. Each ligament was placed directly into Hank's balanced salt solution (HBSS, Gibco, MD, USA) and washed twice with 10 ml HBSS prior to processing.

ACLs were sharply sectioned into small pieces of around 1mm^3 and placed into a filtered solution of 0.1 % collagenase (C2674, Sigma-Aldrich, MO, USA) in Dulbecco's Modified Eagle's Medium (DMEM, Gibco, MD, USA) also containing 5% foetal bovine serum (FBS, Sigma-Aldrich, MO, USA). The solution was incubated at 37°C overnight. Following centrifugation at 1000g for 4 minutes, 10mls DMEM with 10% FBS was added. This step was repeated and cell density assessed using a haemocytometer.

5.2.3 Preparation of 6-well plates

Cells were seeded in 6 well plates at a density of $5000/\text{cm}^2$ with 2mls of DMEM in each well. Cells were incubated in a humidified incubator at 37°C with 5% CO_2 until confluent, changing the medium every 3 days. Medium comprising DMEM, 10% FBS and 100U/ml penicillin and 100 U/ml streptomycin (Penstrep, Gibco, MD, USA) was prepared containing a) 10 ng/ml recombinant human TGF- β 1 (100-21, PeproTech, NJ, USA), b) 100 $\mu\text{g}/\text{ml}$ elastin degradation products (EDP, CB573 Elastin Products Company, MO, USA), c) 10 ng/ml TGF- β 1 and 100 $\mu\text{g}/\text{ml}$ EDP and d) 10 ng/ml TNF- α (T0157, Sigma-Aldrich, MO, USA). Once the cells were confluent the medium was changed in each well to allow cells to be exposed to the four combinations of cytokines and EDPs and a medium only control. Each cell harvest was prepared in triplicate for each time point.

5.2.4 Harvesting of cells

Cells were harvested at 6 and 24 hours following exposure to the cytokines and/or EDPs. The medium was aspirated and 1ml of phenol and guanidine isothiocyanate solution (Trizol

Reagent™, Invitrogen, CA, USA) added to lyse the cells. Following brief agitation of the plate the trizol and cell lysate was removed from each well and frozen at -80 °C until analysis.

5.2.5 mRNA extraction and real time RT-PCR

To each cell lysate 200µl of chloroform (Sigma-Aldrich, MO, USA) was added, shaken then left at room temperature for 10 minutes. Following centrifugation (13000g, 15 minutes, 4°C) the top aqueous phase was added to 500µl of propan-2-ol, mixed and precipitated for 10 minutes at room temperature.

Following repeated centrifugation (13000g, 15 minutes, 4°C) the supernatant was removed leaving the mRNA pellet. 1ml of 70% ethanol (Sigma-Aldrich, MO, USA) in RNA free (diethylpyrocarbonate-treated, DEPC) water was added and the tubes centrifuged again (13000g, 10 minutes, 4°C). The ethanol was removed and the pellets left to air dry before re-suspension in 12.4µl of DEPC water. The mRNA concentration was assessed prior to freezing at -80°C until required. Levels of RNA were measured using the ND-1000 spectrophotometer (Nanodrop Technologies). The absorbance of 1µl of RNA at 260nm and 280nm was determined with a 260:280 ratio between 1.9-2.1 was indicative of 'pure' RNA.

cDNA was obtained by reverse transcription of the mRNA pellet. Reverse transcription was achieved by adding 1µl random hexamers (Promega, WI, USA) to the mRNA solution and heating for 5 minutes at 70°C (Px2 Thermal Cycler, Thermo Scientific, MA, USA). Following this 5µl 5x RT buffer (Promega, WI, USA), 1.25µl each of 10mM nucleotides (Promega, WI, USA), 24 units RNAase inhibitor (Promega, WI, USA) and 200 units M-MLV RT enzyme (Promega, WI, USA) were added making a total volume of 25µl. Following incubation at 37°C for 60 minutes, the mixture was heated to 90°C for 5 minutes then immediately cooled and stored at -80°C until required.

5.2.6 Primer design

Transcript sequences were obtained from the National Centre for Biotechnology Information (Bethesda, MD, USA). Canine sequences were aligned to human, bovine and rat sequences using online software (www.ebi.ac.uk/Tools/clustalw/, www.ensembl.org/) to predict canine exon boundaries. Primers were designed using Primer Express 3.0 (Applied Biosystems, CA, USA) and selected to span predicted exon boundaries where possible. BLAST searches were performed for all sequences to confirm gene specificity. Target and reference gene primers

were synthesised by Eurogentec (Seraing, Belgium). All primers were validated using a standard curve of five serial dilutions so that all primer efficiencies were between 95-105%.

Gene	Name	Forward Primer	Reverse Primer	Reference
<i>GAPDH</i>	Glyceraldehyde phosphate dehydrogenase	CTGGGGCTCACTGAAAGG	CAAACATGGGGGCATCAG	(Clements et al., 2006)
<i>B2M</i>	Beta-2 macroglobulin	CCTTGCTCCTCATCCTCCT	TGGGTGTCGTGAGTACACTTG	(Ayers et al., 2007)
<i>Col1a2</i>	Collagen Type 1 A2	CTATCAATGGTGGTACCCAGTTT	TGTTTTGAGAGGCATGGTTG	(Clements et al., 2006)
<i>Col2a1</i>	Collagen Type 2 A1	CTGGTGAACCTGGACGAGAG	ACCACGATCACCTTGACTC	(Clements et al., 2006)
<i>Col3a1</i>	Collagen Type 3 A1	GGATGGTGGCTTCCAGTTT	CCAGCTGGACATCGAGGA	(Clements et al., 2006)
<i>Fbn2</i>	Fibrillin 1	TGCCCTGGATGGAAAACCT	GGAATGCCGGCAAATGG	
<i>Fbn1</i>	Fibrillin 2	CTTTTGCAAGTGTCTCCTGGTT	TGCTCTGATGGGACACATCTCA	
<i>Eln</i>	Elastin	TGTGGCCGGAAGAGAAAGTG	CGTTGATGAGGTCGTGAGTCAG	
<i>ASMA</i>	α Smooth Muscle Actin	CGCGGGATCCTGACCTT	GTCGTCCCAGTTGGTGATGAT	
<i>SOX 9</i>	Sex Determining Region Box 9	CAAGAAAGACCACCCGGAT	GGAGGAGGAGTGC GCGGAGT	
<i>MMP2</i>	Matrix Metalloproteinase 2	ACCTGCAAGGCAGTGGTC	TCCAAATTCACGCTTTCA	(Clements et al., 2008)
<i>MMP9</i>	Matrix Metalloproteinase 9	CCGGAGGTTACCCAAGTCA	ATCCACGTCTGCGTCTT	(Clements et al., 2008)
<i>CTSB</i>	CathepsinB	CGGCCTTACCCTGTACT	GTGACGTGCTGGTACTCC	(Clements et al., 2006)
<i>CTSK</i>	CathepsinK	AACCTGGTGGACTGTGTCTCTGA	CTGCACATACTGGAAGGCATTG	(Muir et al., 2005b)
<i>ELR 1</i>	Elastin Laminin Receptor 1	ACGTACGTGCCCTGGAACCTT	GCTCCCCAGAAAAGTGTACTG	
<i>Vcan</i>	Versican	GAAGACACACAAGACACGGTATCG	GGTTGCCGCTCTGTAGTGAAA	

Table 5.1 List of primer sequences used for reference and target genes.

5.2.7 Absolute quantification of mRNA expression

Quantitative PCR was conducted on a 7900HT Fast Real Time PCR System (Applied Biosystems, CA, USA). cDNA was first diluted 6-fold in DEPC water. 384-well plates were prepared with each well containing 4 μ l cDNA, 1 μ l 3.3 μ M forward and reverse primer mix and 6 μ l of SYBR Power Green (Applied Biosystems, CA, USA) giving a total well volume of 10 μ l. Each sample was normalised to the expression of GAPDH as a housekeeping gene and compared to control samples (medium alone) using the $2^{-\Delta\Delta CT}$ method (Livak and Schmittgen, 2001).

5.2.8 Statistical analysis

Mixed effects regression was used to assess the significance of the effect of treatment on each gene when compared with a control with significance set at $\alpha=0.05$. A Bonferroni *post hoc* test was used to correct for multiple comparisons. Where significance was achieved, direct comparisons were made using a Mann-Whitney U-test. S-PLUS (TIBCO Software Inc., MA) was used for mixed effects regression analysis and Minitab Statistical Software (Minitab, UK) for Mann-Whitney U-tests. Graphical presentation of the data was produced using Excel (Microsoft Office 2007).

5.3 Results

5.3.1 Animals

All of the dogs were male and two were 4 years old and two 5 years old. The average weight was 36kg (range 30-38kg) and all dogs were in training prior to euthanasia.

5.3.2 Reference genes

B2M was detected at very low levels or not at all in our cultures (results not shown). Therefore only *GAPDH* was used as the reference gene in all $2^{-\Delta\Delta CT}$ calculations.

5.3.3 Treatment with EDPs

Although treatment with **EDPs** for 6 hours stimulated marked increases in *Col2a1* and *CTSB* these were not statistically significant (**Figure 5.1**). Following 24 hours of treatment **EDP** exposure resulted in a statistically significant increase in *Fbn2* ($p=0.032$).

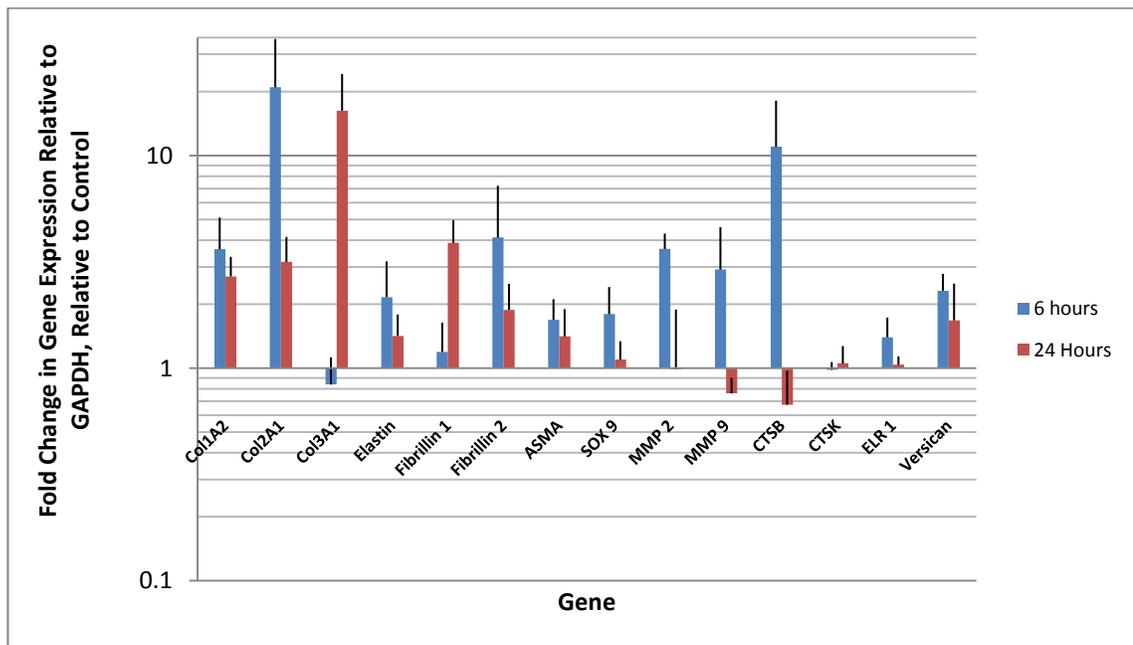


Figure 5.1: Fold change in mRNA transcription relative to GAPDH following 6 and 24 hours of treatment with EDPs alone.

5.3.4 Treatment with TGF- β 1

A statistically significant increase in *Col2a1* was seen following treatment with **TGF- β 1** for 6 hours ($p=0.025$). Application of **TGF- β 1** for 24 hours resulted in a statistically significant increase in *Col1a2* ($p=0.035$) and *Col2a1* ($p=0.005$) (**Figure 5.2**).

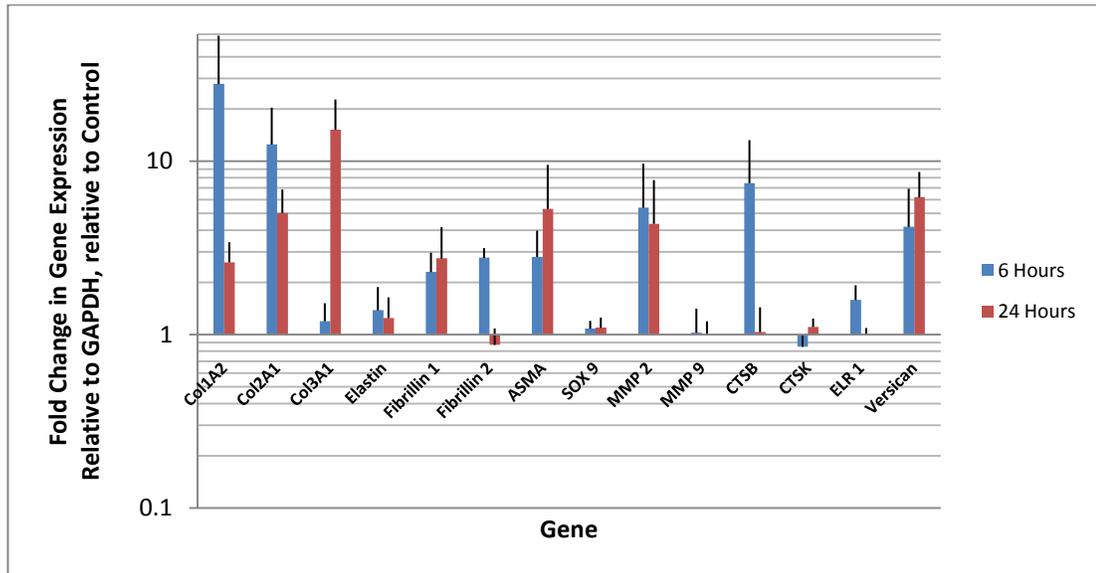


Figure 5.2: Fold change in mRNA transcription relative to GAPDH following 6 and 24 hours of treatment with TGF- β 1 alone.

5.3.5 Treatment with TNF- α

Application of **TNF- α** for 6 hours resulted in a statistically significant increase in *MMP2* ($p=0.004$) (**Figure 5.3**). After 24 hours a statistically significant reduction in mRNA expression of *CTSK* ($p=0.038$), *Eln* ($p=0.009$), *ELR1* ($p<0.001$), *MMP9* ($p=0.004$) and *SOX9 genes* ($p=0.009$), in response to **TNF- α** treatment, was seen (**Figure 5.4**).

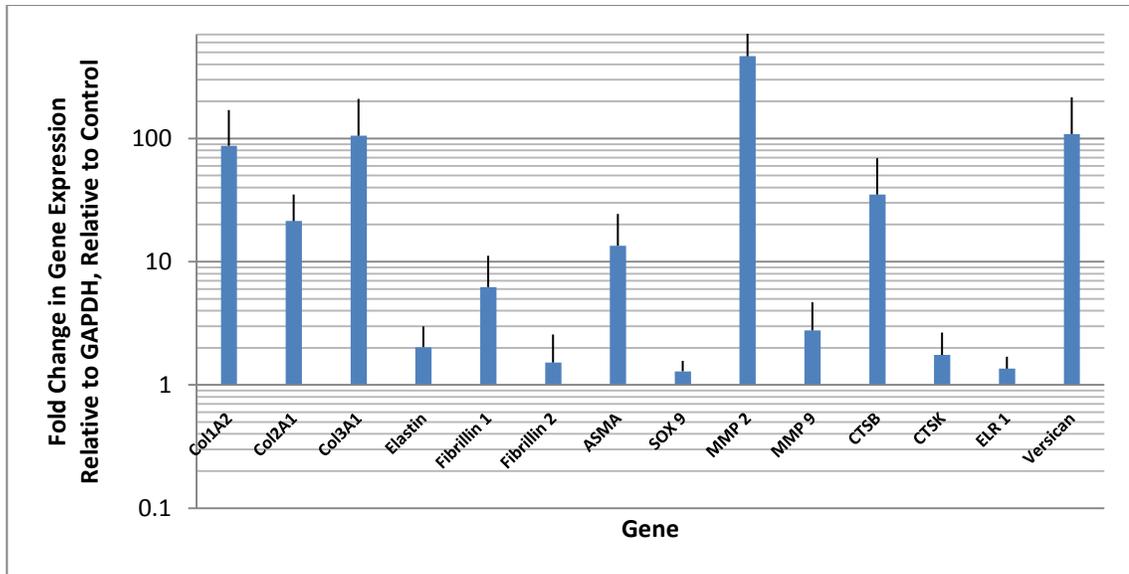


Figure 5.3: Fold change in mRNA transcription relative to GAPDH following 6 hours of treatment with TNF- α .

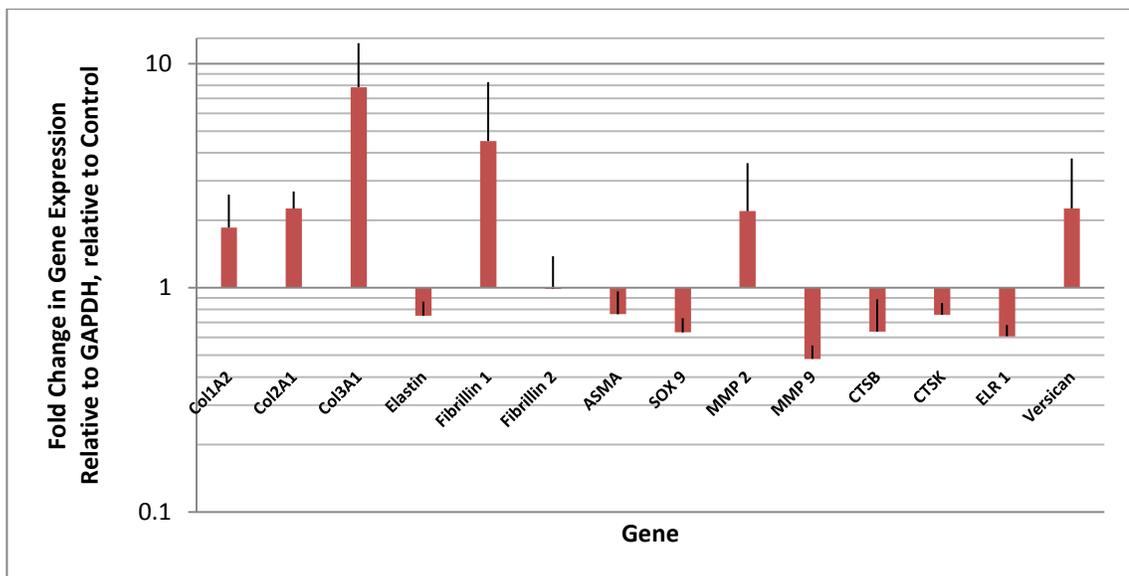


Figure 5.4: Fold change in mRNA transcription relative to GAPDH following 24 hours of treatment with TNF- α .

5.3.6 Treatment with EDPs and TGF- β 1

Following 6 hours of treatment with **EDPs** in combination with **TGF- β 1** a statistically significant increase in *ELR1* was seen ($p=0.014$) (**Figure 5.5**). After 24 hours of treatment

with both a statistically significant increase in *Fbn2* was seen ($p=0.018$) but did not differ significantly from the values seen for treatment with EDPs alone ($p=0.595$) (Figure 5.6).

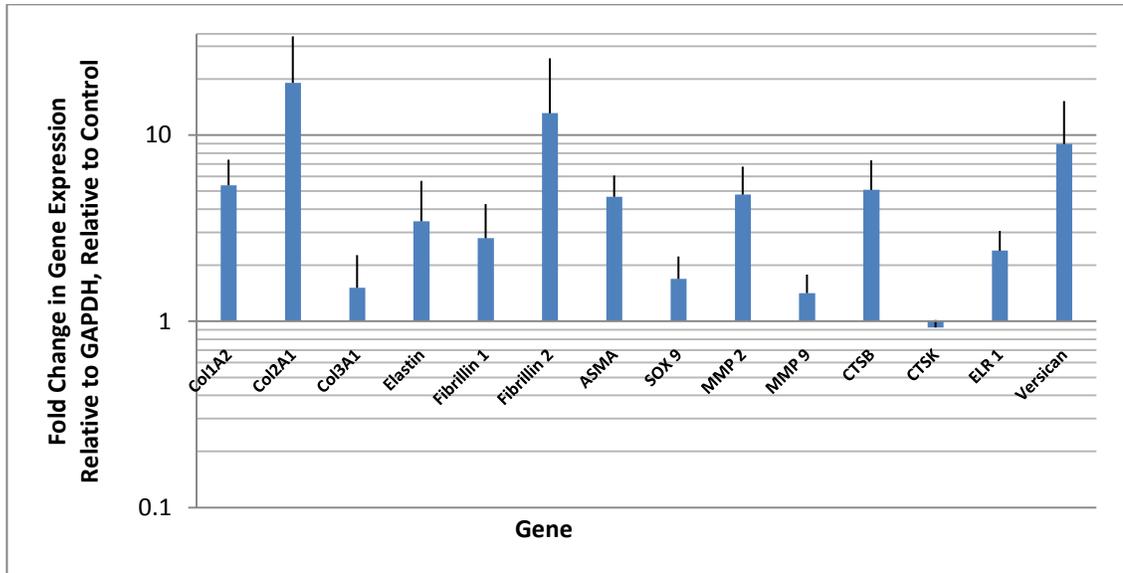


Figure 5.5: Fold change in mRNA transcription relative to GAPDH following 6 hours of treatment with TGF- β 1 and EDPs in combination.

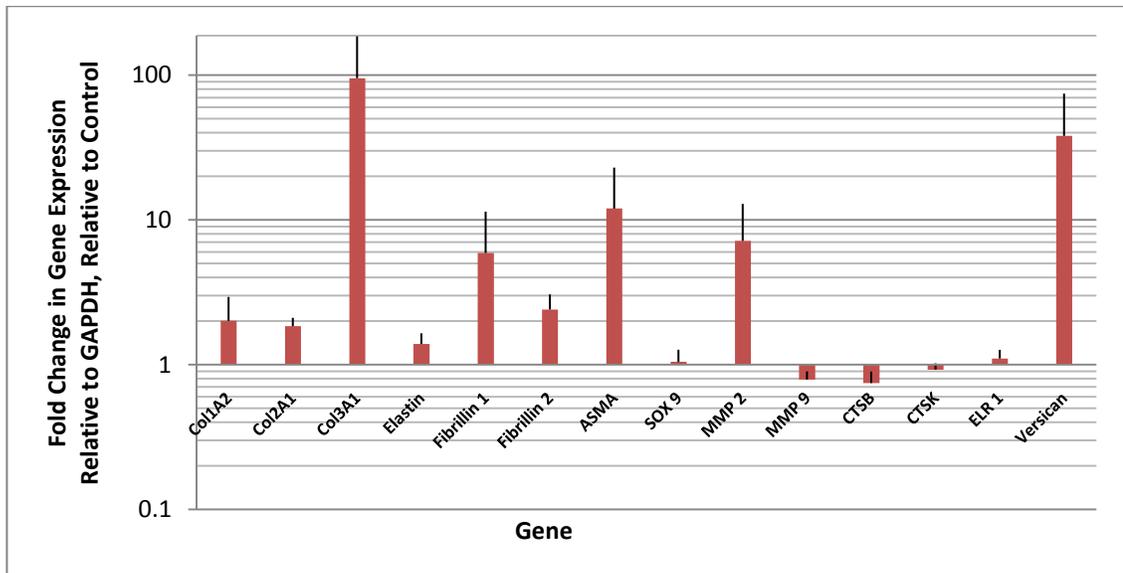


Figure 5.6: Fold change in mRNA transcription relative to GAPDH following 24 hours of treatment with TGF- β 1 and EDPs in combination.

5.4 Discussion

We have shown that fragments of elastin containing the VGVAPG motif exert a biological effect on ACL cells *in vitro* resulting in a significantly increased transcription of *Fbn2*. Furthermore there was a synergistic effect of EDPs and TGF- β 1 resulting in increased transcription of *ELRI*. TGF- β 1 appears to stimulate ACL cells to increase *Col2a1* mRNA transcription which could suggest a role in ACL fibrocartilage production.

EDPs are produced by enzymatic or mechanical action (Duca et al., 2004). We have shown the loss of elastin fibres from regions of fibrocartilaginous change in the canine ACL (Chapter 3). Enzymes with elastolytic ability such as MMPs and cathepsins have been previously identified in normal ACLs with fibrocartilaginous change (Comerford et al., 2006b, Muir et al., 2005b). Whether the loss is through enzymatic or mechanical action, it is possible EDPs containing the known active VGVAPG or GXXPG motifs may have been present in these regions.

TGF- β 1 induced significant increases in *Col2a1* mRNA transcription following 6 hours of treatment. Transcription of matrix proteins such as collagen types 1 and 3 by ligament cells in response to TGF- β 1 would be expected (Chang et al., 2002, Silverio-Ruiz et al., 2007). Fibrocartilage production in the canine ACL has been considered a degenerative change (Vasseur et al., 1985). However recently these changes have been described as adaptive in response to compression or hypoxia (Comerford et al., 2006b) and TGF- β 1 may play a role in this adaptation. Following 24 hours of exposure to TGF- β 1, there was a significant increase in *Colla2*, more typical of the expected ligament healing response (Shrive et al., 1995). As with 6 hour exposure there remained an increased expression of *Col2a1* which again may suggest a role in ligament fibrocartilage production.

No significant changes in mRNA transcription of any gene were seen following 6 hours of EDP treatment alone. Exposure to EDP over 24 hours resulted in a statistically significant increase in *Fbn2*. We have shown that microfibrils are comprised of fibrillin 2 in canine CL tissue (Chapter 3). Furthermore, we have also demonstrated a proportional increase in oxytalan fibres (bundles of microfibrils) in greyhound CL tissue as the degenerative/adaptive fibrocartilaginous change progresses (Chapter 4). As EDPs may be produced as a result of fibrocartilaginous change, oxytalan fibre production may be a direct consequence of cell response to EDP stimulation. Sensitisation early in inflammation (6 hours) to EDPs through upregulation on *ELRI* may lead to later production of oxytalan fibres. We considered the

production of oxytalan fibres part of the response to ligament injury in the greyhound and EDPs may play an important role in this process.

In combination EDPs and TGF- β 1 resulted in a statistically significant increase in *ELRI* following 6 hours of treatment. As TGF- β 1 alone had no effect on *ELRI* mRNA transcription, this change may demonstrate synergism between these two treatments. TGF- β 1 may be produced during inflammation and repair (Schultz and Wysocki, 2009). If replicated at the protein level, an increase in *ELRI* may result in increased ligament cell sensitivity to EDPs thus EDPs may have a role to play as part of the response to injury of the canine ACL. EDPs have been shown to directly influence chemotaxis, proliferation, protease release and even induce apoptosis (Duca et al., 2004). The increase in *Fbn2* following 24 hours of treatment may result from the effect of EDPs alone as the magnitude did not differ significantly from that seen with treatment with EDPs alone.

TNF- α significantly increased transcription of *MMP-2* following 6 hours of treatment. TNF- α is associated with degradation of the ECM and inhibition of repair thus is considered pro-inflammatory. Elevations in TNF- α have been found in the synovial fluid of dogs with ACL rupture (Fujita et al., 2006). However it has been demonstrated to improve wound healing and may be angiogenic (Mooney et al., 1990). MMP-2 is involved in the release of active TGF- β 1 from latent TGF binding proteins (Jian et al., 2003) thus TNF- α may actually lead to an increase in TGF- β 1. Following 24 hours of treatment, TNF- α resulted in a statistically significant decrease in mRNA transcription of *CTSK*, *Eln*, *ELRI*, *MMP-9* and *SOX9*. Downregulation of the proteases *CTSK* and *MMP-9* might not be expected with TNF- α , usually considered pro-inflammatory. The categorisation of TNF- α and TGF- β as pro- and anti-inflammatory agents must be regarded with caution as many of the molecules they promote appear to have positive and negative effects on articular tissue health (Clements et al 2003). Both have multimodal biological activities including growth stimulation and inhibition, modulation of ECM production and regulation of cell proliferation (Massague, 1990).

There is currently no information available on the *in vitro* concentrations of EDP. Our selection of 100 μ g/ml of EDP is based on results from MMP-2 expression on rat smooth muscle cells (Simionescu et al., 2005). While we have measured transcription of mRNA of the examined genes in this study, we have not assessed the protein production. Furthermore, many of the molecules we have studied are regulated *in vivo* through cell-matrix interactions and the limits of *in vitro* cultures without surrounding matrix must also be appreciated. The ACL is

considered a low oxygen environment (O'Donoghue et al., 1971). Our conditions for cell culture may in effect hyperoxygenate ligament cells and this may have affected the validity of the results. However oxygen tension has not yet been measured within the ACL and markers of hypoxia have not been demonstrated in the cells of the ACL core (Hayashi et al., 2003b). Future work will involve investigation as to whether significant increases in mRNA are replicated at the protein level and examine the role of these peptides *in vivo* (this work is currently ongoing in our laboratory).

In conclusion this preliminary study demonstrated potential biological activity of EDPs which may play a role in ligament healing through production of fibrillin 2. Furthermore we suggest TGF- β 1 may be involved in stimulating ACL cells to produce fibrocartilage. As a working model we suggest protease release in the injured ACL could release EDPs from elastin fibres. As part of the initial inflammatory response, EDPs in combination with TGF- β 1 increase cell ELR1, sensitising the ACL fibroblasts to EDPs. Subsequently EDPs increase fibrillin 2 expression leading to microfibril formation as part of the response to injury. This is the first time EDPs have been studied in the mammalian ACL. Their potential importance in ACL pathophysiology is huge and further work into the effect on ACL healing may help provide a solution to the failure of healing in this tissue.

CHAPTER SIX: Comparison of elastic fibre distribution in the anterior cruciate ligament in dogs at a differing risk of anterior cruciate ligament rupture

Abstract:

Outline: An increase in oxytalan fibre (bundles of elastin microfibrils) staining with advancing degeneration has been described in the cruciate ligaments (CLs) of the greyhound, a breed with a very low incidence of anterior cruciate ligament (ACL) rupture. This study aims to investigate whether this change is found in two other breeds of dog (beagle (low to moderate risk) and Labrador retrievers (LR, high risk) with a greater risk of ACL rupture.

Methods: Macroscopically normal ACLs were harvested from 6 beagles, and 6 LR. Sequential histological sections were assessed for extracellular matrix degeneration (haematoxylin and eosin) using Vasseur and modified Vasseur score. Oxytalan fibre staining (Miller's stain) was assessed using novel scoring systems. Each section was scored twice by two observers blinded to breed type.

Results: There was increased ACL degeneration in the beagle ($p=0.03$) and LR ($p=0.045$) in comparison to the greyhound. There was a statistically significant increase in oxytalan fibre staining with advancing ACL degeneration in both beagle ($p=0.001$) and LR ($p=0.001$). Statistically significantly less oxytalan fibre staining was seen in the beagle ($p=0.03$) and LR ($p=0.001$) ACLs than in the greyhound.

Conclusions ACLs of dog breeds at a differing susceptibility to cruciate ligament disease vary in the production of oxytalan fibres with advancing ligament degeneration. For both breeds the ability to produce oxytalan fibres may affect CL healing and reflect their differing risk to ACL rupture.

6.1 Introduction

ACL rupture is a major source of morbidity in the dog, leading to severe osteoarthritis of the knee joint (Bennett et al., 1988). To date, the pathogenesis of this debilitating condition is poorly understood. In previous studies, we have shown that high risk breeds such as the Labrador retriever (LR) demonstrate a higher incidence of ACL rupture associated with altered ligament extracellular matrix (ECM) metabolism (Comerford et al., 2005, Comerford et al., 2006b). Histological changes within the ACL such as loss of collagen architecture, loss of ligament cells and chondrogenic change are considered degenerative (Vasseur et al., 1985). However such changes are seen in both the LR and in the greyhound (GH), a breed with a low incidence of ACL rupture (Comerford et al., 2006b) and it has been suggested that these changes may reflect a healing response to microinjury in the GH (Chapter 4).

Microfibrils are polymers of fibrillins 1 and 2 and bundles of microfibrils are known as oxytalan fibres (OFs). Elastin fibres comprise an outer scaffold of microfibrils with a central cross-linked core of elastin with many other associated molecules (Kielty, 2006). Collectively, oxytalan and elastin fibres are referred to as elastic fibres. Elastin has traditionally been considered a minor component of ligament tissue (Frank, 2004) but it has been shown recently to form up to 20% of the canine CL complex (Chapter 4). A wide distribution of elastic fibres in the canine CL complex has been described, with abundant elastin fibres and OFs running with collagen bundle (Chapter 3). Elastic fibres have important mechanical, biochemical and cell-regulatory functions in tissue. Reversible elasticity is a function of both elastin and oxytalan fibres and is dependent on water and calcium (Eriksen et al., 2001). Microfibrils may have a key role in the extracellular regulation of transforming growth factor (TGF) β activation and signaling (Charbonneau et al., 2004) thereby playing an important role in tissue morphogenesis and cellular responses to injury (Feng and Derynck, 2005).

Compared to other ligaments, the canine ACL has poor intrinsic healing capacity (Frank et al., 1985, Arnoczky et al., 1979). The canine ACL, unlike the medial collateral ligament of the knee, does not form a provisional scaffold (fibrin-platelet plug) and also has reduced levels of key ECM proteins and cytokines within the wound (Murray et al., 2007, Spindler et al., 2006). This failure of the provisional scaffold has been proposed as a mechanism for poor ACL healing (Murray, 2009). Other mechanisms proposed for poor ACL healing include deficiencies in stimulation, intrinsic deficiencies of cell migration and proliferation and overproduction of degradative enzymes (Tang et al., 2009, Nagineni et al., 1992, Geiger et al.,

1994, Spindler et al., 1996a) and inadequate blood supply (Vasseur et al., 1985, Hayashi et al., 2003a, Arnoczky et al., 1979, Tirgari and Vaughan, 1975a, O'Donoghue et al., 1971). It has been suggested ACL healing in breeds with a low incidence of ligament rupture may be superior to those with higher incidence (Arnoczky et al., 1979, Frank et al., 1985). We have demonstrated a proportional increase in OFs with advancing ACL degenerative/adaptive change in the GH and this increased production may form part of the CL healing response (Chapter 4).

In this study we will examine ACL tissue from two breeds of dog with differing incidence of ACL rupture (the beagle and LR) and compare the results with those of the GH (a breed with very low incidence of ACL rupture) ACLs we examined in Chapter 4. The beagle has a moderately low incidence of ACL rupture (relative risk 2.34%), the Labrador retriever has a high incidence (relative risk 3.81%) and the GH a very low incidence (relative risk 0.51%) (Whitehair et al., 1993). By quantifying ACL degeneration (using Vasseur and modified Vasseur scoring) and OF staining (using Miller's scoring) we aim to assess whether previously described OF production with advancing ACL degeneration is consistent in dog breeds with a differing risk to ligament disease and rupture. This information may help to explain this differing incidence of ACL rupture in different dog breeds.

6.2 Material and methods

6.2.1 Animals

Beagles: Six ACLs were harvested by sharp dissection from six beagles with no gross evidence of knee pathology. One sample was taken at random from each ACL.

Labrador retrievers: 9 ACLs were harvested by sharp dissection from 6 skeletally mature LRs with no gross evidence of knee joint pathology. Two ACLs, obtained from the USA, were sectioned into proximal, middle and distal sections providing 6 samples. Of the remaining 7 ACLs, obtained from the UK, tissue was sampled randomly from the length of each ligament, providing 10 samples of which no more than two were from an individual ACL.

Greyhounds: 11 ACLs were collected and processed as detailed in Chapter 4

6.2.2 Sample collection

All ACLs were harvested from dogs which were euthanatized for reasons other than musculoskeletal disease and informed consent, in accordance with University ethical guidelines, was obtained in each case for tissue removal. All samples were obtained within 24 hours of death and then fixed in formalin and embedded in paraffin blocks.

6.2.3 Histology: Staining

Sequential sections of 4 μ m from paraffin-embedded samples were stained with 1) haematoxylin and eosin (H&E), 2) Verhoeff's iodine-iron haematoxylin (EVH) to show only elastin fibres and 3) Miller's stain, MS (also referred to as Weigert's), to show both elastin fibres and OFs. This series allowed assessment of tissue architecture, and through comparison of Verhoeff's- and Miller's-stained sections, differentiation of elastin fibres and OFs. Images were recorded on a dedicated microscope (Nikon Eclipse 80i). All sections were read by two observers blinded to section location or animal (KDS and EJC) on two separate occasions at least one week apart.

6.2.4 Histology: Scoring methods

H&E: H&E sections were assessed for signs of CL degeneration. All samples were graded 0-3 according to criteria previously described (Vasseur et al., 1985). The broad grade 1 category was subdivided with a more detailed scoring system previously developed by the authors (Chapter 4). Briefly, a score from 0-4 was awarded based on the extent of the changes for each of eight factors resulting in a range of possible scores from 0-24 being awarded. These results are referred to as modified Vasseur Score (mVS).

Verhoeff's: No scoring system was developed as initial observations suggested these fibres were sparse and did not change with degeneration.

Miller's Stain: A scoring system developed previously by the authors was used to quantify changes in OF staining (Chapter 4). Increased staining in interfascicular and interbundle regions, ligament substance (intra-bundle), as well as the extent and degree of pericellular staining, could be awarded up to two points giving a score range of 0-10. These results are referred to as Miller's Score (MS).

6.2.5 Statistical analysis

Scores for histological sections are presented as mean values \pm SD. The relationship between VS and MS in the LR was examined using Pearson's correlation and linear regression analysis was used to examine the relationship between mVS and MS. Significance was set at 5%. Direct comparisons between groups of unequal number were made using a Mann-Whitney test. Comparisons between Vasseur scores were made using a Kruskal-Wallis test using breed as the factor. Kendall's coefficient of concordance was calculated for intra- and inter-observer concordance of both modified Vasseur and Miller's scoring. Kendall's coefficient of concordance ranges from 0 (no agreement) to 1 (complete agreement). For the results of MS in the Labrador, Fleiss' Kappa was used to calculate agreement for the MS results, where scores above 0.7 are considered to represent a high degree of concordance. Data were analysed using Minitab Statistical Software (Minitab, UK).

6.3 Results

6.3.1 Animals

Age range for beagles was 10-23 (mean 14 ± 4 , median 11) mths and for the Labradors was 13-144mths (mean 60 ± 44 , median 60), summarised in **Table 6.1**. LRs ($p=0.007$) and GHs ($p=0.001$) were significantly older than the beagles but there was no statistically significant differences between age of LRs and GHs ($p=0.45$). Of the beagles, 4 were female and 2 male and of the LRs 3 were female and 3 male.

BREED	Greyhound	Beagle	Labrador
AGE (months)	42 ± 9.7	14 ± 4	60 ± 44
Degeneration (mVS)	12.1 ± 1.6	16.8 ± 4.6	16.3 ± 4.4 *
Oxytalan Fibre Staining (MS)	5.3 ± 1.9	3.1 ± 1.9	0.31 ± 0.85
mVS vs MS Regression	$r^2=0.44, p<0.001$	$r^2=0.98, p=0.001$	$r^2=0.0, p=0.98$

Table 6.1: Summary data for all breeds. *mVS scores for the LR are not directly comparable as many LR ACLs scored greater than 1 on the Vasseur scale thus could not be scored on the mVS scale.

6.3.2 Degeneration Scoring

Results for ACL degeneration scoring are summarised in **Table 6.1**.

A) Greyhounds: results detailed in Chapter 4.

B) Beagles: All ACL samples stained with H&E were graded as grade 1 according to the published system (Vasseur et al., 1985). All were also graded on the modified scoring system (mVS, Chapter 4), with an average score of 16.8 ± 4.6 (range 9.8-22.5, median 17.1). There was a statistically significantly higher mVS in the beagle than in the GH ($p=0.03$) but no significant difference compared to the LR ($p=0.54$)

C) Labradors: Degenerative change was seen in all LR ACLs. 11 of 16 ACL samples stained with H&E were graded as grade 1 according to the published system (Vasseur et al., 1985). Of the remaining ACLs, 3 were grade 2 and 2 were grade 3. LR ACL Vasseur score was significantly greater than the GH ACL Vasseur score ($p=0.045$). Those graded 1 were also graded on the modified scoring system (mVS), with an average score of 16.3 ± 4.4 (range 8-22.5, median 17).

6.3.3 Miller's Scoring (Oxytalan Fibre Staining)

A) Greyhounds: results are detailed in chapter 4. For the greyhound ACL, a statistically significant positive regression was seen between mVS and MS ($r^2=0.44$, $p<0.001$, **Table 6.1**).

B) Beagles: MS averaged 3.1 ± 1.9 (range 0.25-5.5, median =3.5) in the beagle. This was significantly greater than the LR ($p=0.001$) but significantly lower than the GH ($p=0.03$). Regression analysis of the scoring systems (MVS and MS) showed a significant positive regression ($r^2=0.98$, $p=0.001$) (**Figure 6.1**).

C) For all ACLs, MS averaged 0.31 ± 0.85 (range 0-25, median 0) in the LR. This is significantly lower than the MS for the beagle ($p=0.001$) and GH ($p=0.001$). For ACLs graded on the mVS scale, all MS scores were 0. No significant relationship was seen between mVS and MS. However a statistically significant positive regression was seen between Vasseur score and MS ($r^2=0.48$, $p=0.007$, **Figure 6.2**).

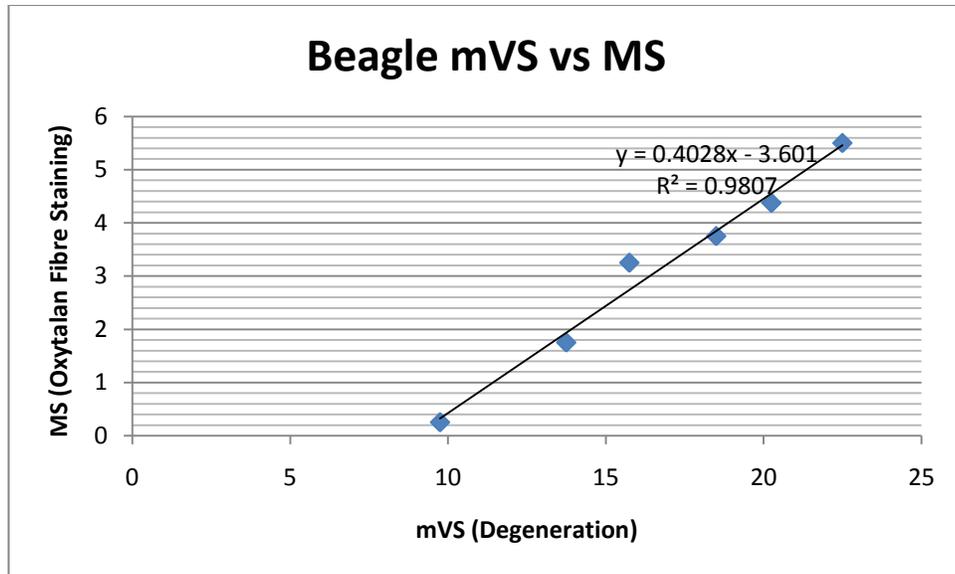


Figure 6.1: Relationship between ACL degeneration and oxytalan fibre staining in the beagle.

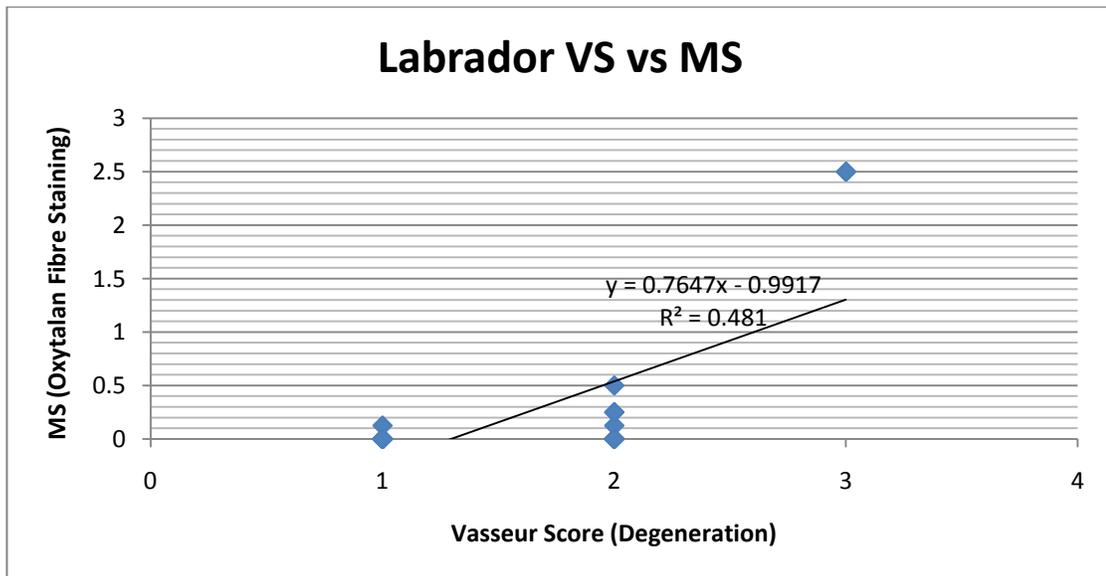


Figure 6.2: Relationship between ACL degeneration and oxytalan fibre staining in the Labrador retriever. Note the x axis is Vasseur score not mVS.

6.3.4 Inter/intra-observer data

Beagle: Kendall’s coefficient of concordance for inter-observer concordance was 0.91 for mVS. Intra-observer concordance for KDS was 1 and for EJC was 0.92 (**Table 6.2**). Kendall’s coefficient of concordance for inter-observer concordance was 0.96 for MS. Intra-observer concordance for KDS was 1, and for EJC was 0.95.

Labrador: Kendall’s coefficient of concordance for inter-observer concordance was 0.90 for mVS. Intra-observer concordance for KDS was 0.97 and for EJC was 0.91 (Table 6.1). For MS, Kendall’s coefficient of concordance for inter-observer variation was 0.76 showing good agreement (**Table 6.2**). Intra-observer concordance for both KDS and EJC was 1.

Variation		Intraobserver		Inter-observer
Breed	Scoring	KDS	EJC	
Beagle	mVS	1	0.92	0.91
	MS	1	0.95	0.96
Labrador	mVS	0.97	0.91	0.90
	MS	1	1	0.76

Table 6.1: Intra- and inter-observer agreement in mVS and MS scoring. Kendall’s coefficient of concordance was used for all data except Labrador MS where Fleiss’ Kappa was used.

6.3.5 Descriptive Histology

Elastin Fibre Staining (Verhoeff’s)

A) Greyhounds: results detailed in Chapter 4

B) Beagles: EFs were distributed consistently throughout the ACL with little variation within or between ACLs of differing joints or dogs. Within the ACL substance elastin fibres were mostly orientated parallel to collagen bundles. Interfascicular elastin fibres were orientated either perpendicular or oblique to collagen bundles. A reduction in EFs was noted where mineralisation was present.

C) Labradors: In CL tissue with minimal degeneration, elastin fibres were distributed in the same manner already described for the beagle. Again, as with the beagle, a reduction in

elastin fibres where mineralisation was seen. In areas of marked degeneration (Vasseur grades 2 and 3), seen only in the LR, EFs were often absent.

Oxytalan Fibre Staining (Miller's)

A) Greyhounds: results detailed in Chapter 4

B) Beagles: OFs were found throughout the ACL and were seen in both the epiligament and substance. Within the substance of the ACL, large numbers of fibres were observed running parallel to collagen bundles. The majority of fibres in the interfascicular region were arranged in a fine, tortuous meshwork with no overall orientation, although some ran perpendicularly between bundles. OFs were commonly found both on the outside and within collagen bundles. Beagle ACL samples with higher grade 1 (>mVS of 10) degeneration showed a marked increase in OF staining (**Figure 6.3A, B**). Increased OF staining was differentially noted within all three subdivisions of the ligament substance in all ACLs with high grade 1 changes.

Interfascicular: Staining of OFs was loose and mesh-like (**Figure 6.3C**). Increased pericellular OF staining was commonly seen around cells that had undergone chondrocytic change. ACLs with more advanced degeneration had increased interfascicular OF density through increased numbers of small OFs in the mesh pattern.

Interbundle: Increased OF staining in the form of small fibres running largely parallel and obliquely to the collagen was seen where collagen bundle integrity remained largely intact (**Figure 6.3B**). OF staining was much less regularly organised if there was loss of collagen architecture where increased OF staining was more similar to the mesh seen in the interfascicular regions

Bundle substance: Increased OF staining was also sometimes present in regions with a loss of collagen density (**Figure 6.3D**). These OFs were usually aligned with the collagen, and were very fine.

C) Labradors: In ACL with minimal degeneration (<mVS of 10), OFs were found throughout the ACL with a similar distribution. As degeneration became more severe (>mVS of 10) there was only occasional very slight increase in OF staining in the LR samples of very high grade (Vasseur grade 2 and 3) examined (**Figure 6.4**). In order to ensure the alteration in

OF and EF staining was not an artefact, staining as a positive control was confirmed as normal in areas unaffected by degenerate change such as blood vessels in synovium.

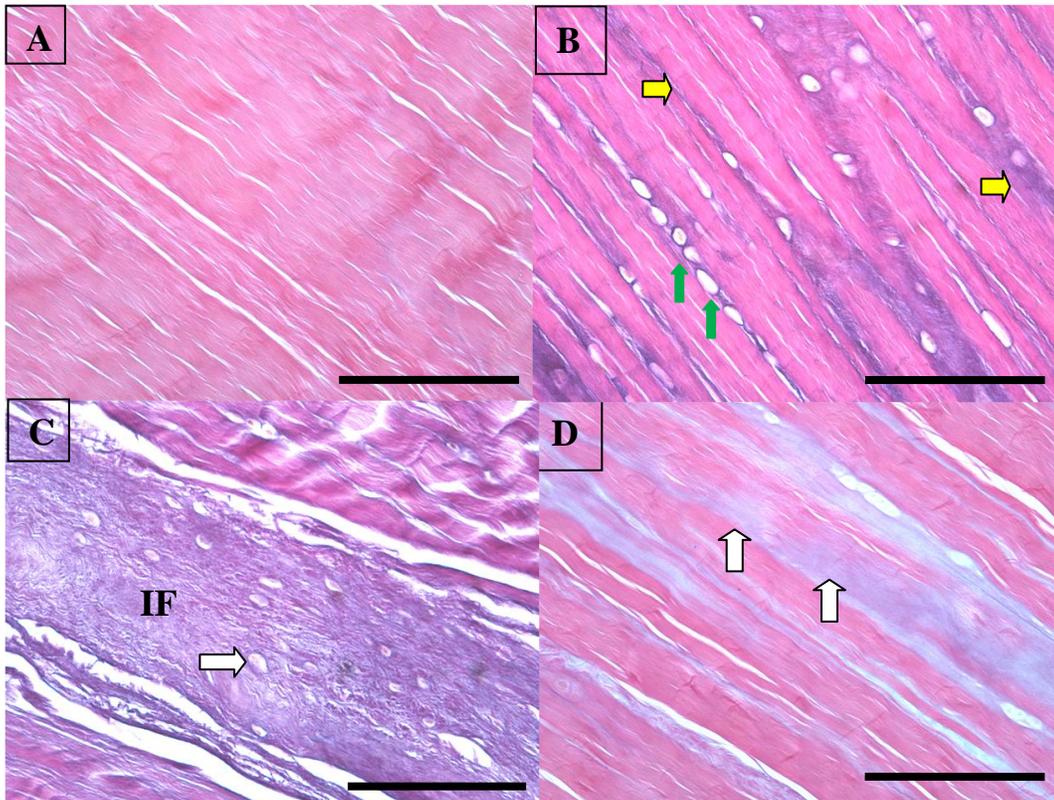


Figure 6.3: Variation in OF staining in beagle ACL. (A) Miller's stain, ACL, x40. Minimal interbundle OF staining is seen in this section with low grade 1 (Vasseur) changes. (B) Miller's stain, ACL, x40. Image from same section as A showing marked pericellular staining of chondrocytic cells (green arrows) and moderate interbundle staining (yellow arrows). (C) Miller's stain, ACL, x40. Moderately increased OF staining in the interfascicular region (IF). Chondrocytic cells (arrow) show pericellular deposition of OFs. (D) Miller's stain, ACL, x40. Increased OF staining with loss of collagen density (arrows). Magnification bars = 100µm

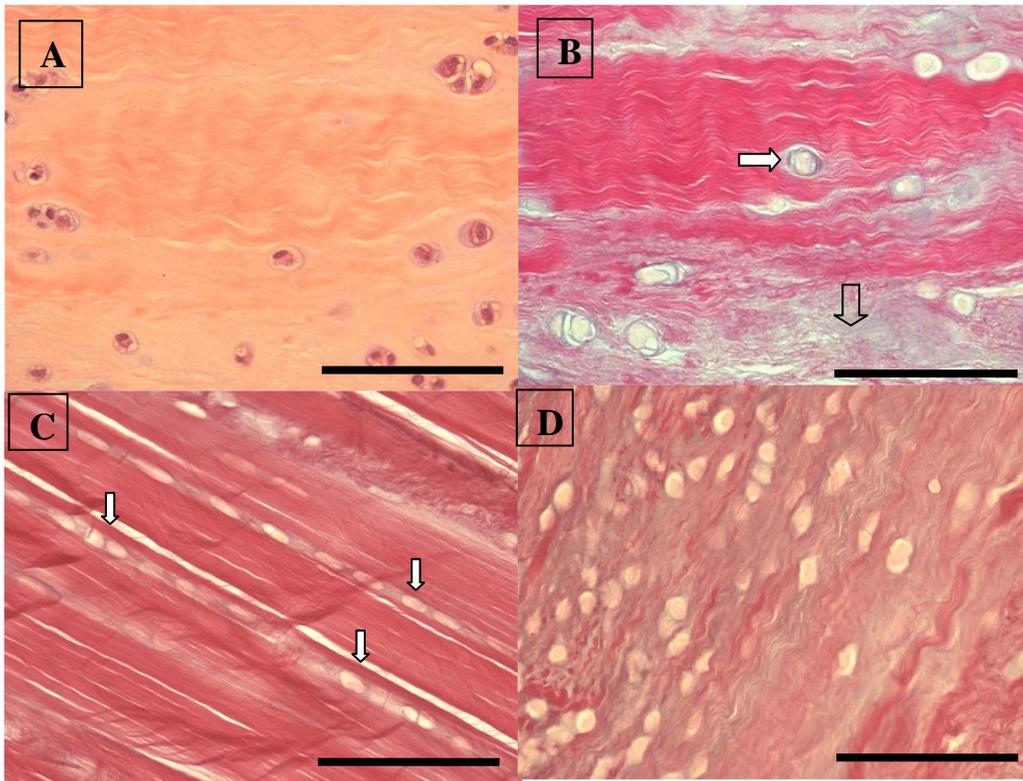


Figure 6.4: Minimal increase in OF staining in high grade degenerate LR CL. (A) H&E stain showing marked disruption to collagen architecture and chondrocytic change of fibrocytes. This was given a Vasseur score of Grade 3. (B) Corresponding section stained with Miller's. Some slight increase in pericellular staining (block arrow) but no increase is seen in interbundle regions (open arrow) or bundle substance. (C) High Vasseur grade 1 degeneration with numerous chondrocytic cells (arrows) but no increase in OF staining. (D) High grade degeneration (Vasseur grade 2). There is widespread chondrocytic change and marked loss of collagen architecture but only very slight pericellular OF staining. Magnification bars = 100µm

6.4 Discussion

In this study we have shown that in both beagles and LRs elastin fibre staining appears to be absent in regions of ACL degeneration. There was a proportional increase in OFs with advancing ACL degeneration in both beagle and LR. However the increase in OF staining in the LR ACL and beagle was significantly less than that of the GH.

Degenerative change, as assessed on H&E sections, was confined to the Vasseur grade 1 category in the beagle and GH but in the LR was found to be more advanced than grade 1 in 5 of 16 sections. Degenerative change has been correlated with age in the dog by some authors (Vasseur et al., 1985, Narama et al., 1996) but not others (Comerford et al., 2005). As there was no significant difference between their ages, this suggests that degeneration may be more advanced in the LR ACL than in the GH. As the beagles were significantly younger than the LRs, it may be the lower degree of degeneration is a function of age. However although the beagles were significantly younger than the GHs they had significantly more advanced degeneration. Advancing degeneration has been correlated with mechanical weakness (Vasseur et al., 1985). As the incidence of ACL rupture is higher in the beagle than the GH (Whitehair et al., 1993), this difference may reflect a tendency to a higher degree of degeneration and consequent weakness. Grade 2 and 3 changes have not been described in the GH, a breed with an extremely low incidence of ACL rupture (Comerford et al., 2006b). Indeed generalised grade 1 changes were described in both ACL and PCL in the GH leading to the suggestion that these low grade changes are not in fact degenerative but adaptive (Comerford et al., 2006b). Thus we suggest in the beagle we are seeing ‘adaptive’ change but in the LR the change may be truly degenerative and indicative of disease and may be a precursor to rupture.

As with the GH (Chapter 4), OF production in the beagle and LR showed a statistically significant relationship with degeneration. Despite having the lowest mVS of all breeds in the study the GH had the highest MS. This suggests that the increased OF staining seen with advancing degeneration was most marked in the GH. Furthermore the beagle MS was significantly greater than the LR despite the mVS being lower. Thus the increase in OF staining seen with advancing degeneration in the beagle is not as marked as in the GH but more marked than in the LR. These findings are suggestive of a fundamental difference between the LRs, beagles and GHs and may reflect reduced production or assembly, or increased destruction of fibrillin.

In all breeds of dog, the distribution of EFs and OFs in ACL tissue with low grade degenerative changes was similar to that previously described in the GH. As degeneration progresses and variation in OF staining is noted, it is likely that this reflects genuine change rather than intrinsic breed variation in ACL OF content though further work would need to be done in age and gender matched samples to confirm this.

In the ACLs of all breeds, OF staining increased proportionately with advancing degeneration. This was first described in the GH CL (Chapter 4) and has not been described in any other ligament tissue. This role of OFs in ACL tissue is unknown and may involve provision or maintenance of elasticity, stabilisation of blood vessels, anchoring tissue or guidance of cell migration (Fullmer et al., 1974, Chantawiboonchai et al., 1998, Everts et al., 1998, Tashiro et al., 2002). Previously we proposed the increased quantity of OFs observed in GH CLs may reflect a healing response as assembly of OFs is commonly seen in healing responses in artery (Sinha et al., 2001), myocardium (Vracko et al., 1990), muscle (Fricke et al., 2008) and skin (Tsuji and Sawabe, 1987).

Assembly of OFs is commonly seen in healing responses thus this relative lack of OFs in degenerating LR CL tissue may contribute to the eventual rupture of the ligament through a failure of healing. Although poor healing has been described in the canine ACL (Frank et al., 1985, Arnoczky et al., 1979, Murray et al., 2007, Spindler et al., 2006) this does not mean that intrinsic healing capacity is uniform for all breeds in both CLs. Increased relative risk of ACL rupture may indicate reduced intrinsic healing capacity.

Limitations in tissue availability precluded a more thorough regional analysis of LR ACLs. However as no LR sample showed any evidence of increased OF staining, we are satisfied our data is representative of this breed. The lack of OF staining in LRs was checked by reference to known sites of normal OFs, blood vessels and epiligament, where OFs could be seen. Additionally the LR CLs were from two sources, one in the UK and one in the USA, again with consistent results. Further work will include analysis of other breeds with differing risk of CL rupture as well as beagle ACLs with a greater range of ages in order to validate our findings as a possible mechanism in the aetiopathogenesis of this condition.

In summary we shown in the GH ACL a corresponding increase in OF staining with degeneration (chapter 4). In this study we have shown similar findings in beagle ACLs, a breed at low to moderate risk of ACL rupture. Additionally we have showed that in LR ACLs, a breed at high risk of ACL rupture, there is only a very slight increase in OF staining in advanced degenerative change. These differences demonstrate that the CLs of differing breeds of dog vary in their response to ACL degeneration. For each breed the ability to produce OFs may affect ACL healing and reflect their differing risk of ACL rupture. Whether these differences arise through genotypic or phenotypic mechanisms remains to be elucidated. Tissue integrity is a balance of damage and repair and the balance may be progressively tipped

in favour of damage where OF production is diminished. This may ultimately manifest as ACL rupture and failure to generate sufficient OFs in CL tissue may be a key step in the aetiopathogenesis of this disease.

CHAPTER 7: General Discussion

7.1 Introduction

The main objective of this thesis was to examine the distribution and role of elastin and elastic fibres in the canine CL complex. The aims were to look at mechanical and biological roles for elastin and elastic fibres in CL tissue.

7.2 Cell morphology

The objective of this study was to describe the morphology and regional variations of cells in the CLs of two breeds of dog with a differing predisposition to non-contact ACL injury. The canine CLs have traditionally been considered to comprise of dense collagenous matrix with sparse, isolated cells. This study described cell processes extending from the cells of the CL complex that had not been described before in the dog. Cell processes have been described in connective tissue under tensile load and contact between cells via processes may facilitate direct communication. This has important implications in CL physiology. Traditionally considered isolated, the cells of the CL may actually be able to network and respond to changes in tissue environment rapidly as part of the joint organ. Furthermore, disruption to this network may interfere with normal ligament physiology. Indeed we suggested the chain formations of rounded cells described in the canine CL complex may be an attempt to re-establish this communication.

The range of cell morphologies described was a novel finding in CL anatomy. Three differing morphologies were identified: type A with long processes, type B with shorter, branching processes and type C with rounded nuclei and no processes. Variations in cell process morphology have been described in the meniscus and annulus fibrosus in rabbits and cattle respectively. In these studies the variations were considered to reflect mechanical forces whereby cell process loss was thought to occur in tissue normally under tension that undergoes compression. The canine ACL is considered to be subject to multiaxial stresses, and other extrinsic factors, such as intercondylar notch of the femur, joint conformation and gait, may also influence forces acting on the ligament. Such mechanical variation may lead to the phenotypic variation described in the canine CL. However, cell morphology may be influenced by many other factors such as variation in the composition of the ECM or joint

pathology. Variation in cell morphology based on nuclear morphology had already been established in the canine cruciate ligament complex. This study showed that nuclear shape could be used to estimate the shape and extent of cell processes.

It was suggested that the GH had large areas of cells with long processes but the LR did not. Poor healing has been associated with failure of long cytoplasmic processes to span wounds in the ovine ACL. Long cell processes in the GH ACL may aid healing and partly explain the lower risk of ACL rupture in this breed. Juxtaposition of differing cell morphologies from one longitudinal area to another appeared marked in the LR but not the GH and may reflect differing CL factors from the GH such as mechanical forces or ECM properties.

This study suggested marked regional variation in the cell morphology of the canine CL complex as well as juxtaposition of and differences in cell type morphology in dog breeds at a differing risk to non-contact ACL injury. The possibility of a three dimensional network of cells has ramifications for cell nutrition, mechanical sensing and coordinated response to injury in the CL complex. Understanding variations in cells within the CL complex is likely to be important in improving our understanding of the role of cell communication in ligament disease and may be important in the development of bioengineered constructs for the treatment of ACL failure.

7.3 Elastic fibres and fibrillins

The objective of this study was to investigate the distribution of elastin fibres, microfibrils and fibrillin 1 and 2 in the canine cruciate ligament complex and study their contribution to mechanical function. Elastin fibres were found throughout both ACL and PCLs and appeared abundant in the interbundle and interfascicular regions. They may provide additional stiffness at low strain and stress or provide elastic mechanical support to blood vessels. The high density of oxytalan fibres within collagen bundles may have mechanical functions such as absorption of low strain stiffness, complementing the tensile strength of collagen fibres, or restoration of longitudinal conformation following longitudinal strain.

In the canine CL complex, the distribution of fibrillin 1 and 2 appeared to differ from that described in other tissues. Fibrillin 1 immunostaining revealed fibres that may be microfibrils. The pattern of fibrillin 2 staining appeared similar to both fibrillin 1 and oxytalan fibre distribution throughout the CLs. Therefore fibrillin 2 appears to be a significant component of

microfibrils in canine CL. However collagenase digestion may have exposed fibrillin 2 in the microfibril core.

Using NDIC to image the variation in interfascicular and interbundle anatomy in unfixed, hydrated tissue allowed identification of previously undescribed anatomy in the CL complex. The fibres of the interfascicular region were loose and may allow fascicles to move freely in relation to each other or offer stress protection for fragile structures such as blood vessels and nerves. A hierarchical subdivision of interbundle fibres, similar to that described in the human annulus fibrosus, may regulate interbundle movement. The interbundle and interfascicular fibres that were demonstrated using NDIC revealed similar anatomical distribution to interbundle and interfascicular fibres containing elastin and oxytalan fibres described earlier. These may therefore contain elastic fibres. A passive recoil system formed from elastin and oxytalan fibres offers a mechanism whereby ligament anatomy can be restored following deformation.

This study has shown that elastic fibres may have potentially important mechanical roles in this ligament complex. Abundant oxytalan fibres composed of both fibrillin 1 and 2 were suggested. This may differ from the solely fibrillin 1-containing oxytalan fibres previously described in tendon and may demonstrate a fundamental difference between ligament and tendon.

7.4 Elastic fibres and CL degeneration

The objectives of this study were two-fold. Firstly the aim was to investigate the distribution of elastin fibres and microfibrils in canine CL tissue from a breed at low risk of ACL rupture and relate these findings to degenerative ligament change. Secondly it was intended to determine whether variation in elastin content might relate to histological and biochemical properties of the canine CL complex.

Changes consistent with mild degeneration were found in every histological section examined suggesting degeneration in canine CLs was likely to be generalised according to a previously published histological scoring system (Vasseur et al., 1985). It is not known why the canine ACL undergoes degeneration and fibrocartilage formation. Fibrous connective tissue under compressive load may form fibrocartilage but these changes may be a protective metabolic response reflecting adaptive rather than degenerative change.

Increased oxytalan fibre (OF) staining with advancing degeneration has not been described previously in any ligament. OFs may have a number of roles in the CL complex including provision or maintenance of elasticity, stabilisation of blood vessels, anchoring tissue or guidance of cell migration. However, assembly of OFs is commonly seen in healing responses in other Tissue such as skin, muscle and blood vessels. As the tissue in our study was from a breed in which the observed degenerative changes never result in ACL rupture, these studies would suggest that the increased quantity of OFs observed in greyhound CLs may reflect a healing response to injury especially as these are an athletic, racing breed .

Previous studies have estimated elastin content in the CL complex from 0-6%. In this study, elastin content was shown to be significantly greater than previously reported, ranging from 5.9 to 19.4% of ligament dry weight. Despite EFs being sparse in the CL, elastin formed approximately 10% of dry weight of CL tissue suggesting that not all elastin in CL tissue is contained within EFs. Levels of elastin are relatively homogenous within and between pairs of CLs, and left and right knees, but vary between dogs. Taken together this suggests elastin may have non-mechanical functions.

In this study it has been suggested that elastin forms a far greater proportion of the canine CLs than has previously reported. Mild degenerative ligament changes were shown to be associated with increasing prevalence of OFs which has never been described in any mammalian ligament. This may reflect an attempt to repair the CL and is part of the CL healing response preventing ligament rupture in an exercising low risk dog breed to ACL rupture.

7.5 Breed variation in elastic fibres

In Chapter 4 an increase in oxytalan fibre staining with advancing degeneration in the CLs of the GH was described. This study aimed to investigate whether this change is found in two other breeds of dog at a greater risk of ACL rupture. The beagle is at low to moderate risk and the LR is at high risk to ACL rupture

Degenerative change appeared significantly more advanced in the LR than the beagle or GH. Degenerative change has been correlated with age in the dog by some authors. As there was no significant difference between the ages of this group of dogs, degeneration may be more advanced in the LR ACL than in the GH. Although the beagles were significantly younger than the GHs they had significantly more advanced degeneration. Advancing degeneration has been correlated with mechanical weakness by Vasseur et al., 1985. As the relative risk of ACL

rupture is higher in the beagle and LR than in the GH, this difference may reflect weakness as a consequence of a higher degree of degeneration.

In the ACLs of all breeds, OF staining increased proportionately with advancing degeneration. It is likely that this reflects genuine change rather than intrinsic breed variation in ACL OF content as despite having the lowest degeneration score of all breeds in the study the GH had the highest OF staining. The increase in OF staining seen with advancing degeneration was most marked in the GH and least marked in the LR. This suggests a fundamental difference between the LRs, beagles and GHs and may reflect reduced production or assembly, or increased destruction of fibrillin. Assembly of OFs can be commonly seen in healing responses in skin muscle and blood vessels. The relative lack of OFs in degenerating LR CL tissue may contribute to the eventual rupture of the ligament through a failure of healing. Although poor healing has been described in the canine ACL, this does not mean that intrinsic healing capacity is uniform for all breeds in both CLs. Increased relative risk of ACL rupture in certain breeds may indicate reduced intrinsic healing capacity.

For each breed the ability to produce OFs may affect ACL healing and reflect their differing risk of ACL rupture. Whether these differences arise through genotypic or phenotypic mechanisms remains to be elucidated. Tissue integrity is a balance of damage and repair and the balance may be progressively tipped in favour of damage where OF production is diminished. This may ultimately manifest as ACL rupture and failure to generate sufficient OFs in CL tissue may be a key step in the aetiopathogenesis of this disease.

7.6 Elastin degradation peptides

The objective of this study was to determine whether elastin degradation peptides (EDPs) containing the VGVAPG motif exert a biological effect on the cells of the ACL of the dog *in vitro*. EDPS are enzymatically hydrolysed fragments of elastin. Enzymes with elastolytic ability such as MMPs and cathepsins have been previously identified in normal ACLs with fibrocartilaginous change. In Chapter 3 the loss of elastin fibres from regions of fibrocartilaginous change in the canine ACL was described.

TGF- β 1 induced significant increases in *Col2a1* mRNA transcription following 6 hours of treatment. Although fibrocartilage production in the canine ACL has been considered a degenerative change, it has also been described as adaptive response to compression or hypoxia. TGF- β 1 may play a role in this adaptation. Following 24 hours of exposure to TGF-

$\beta 1$, there was a significant increase in *Col1a2*, more typical of the expected ligament healing response, but there remained an increased expression of *Col2a1*.

Exposure to EDP over 24 hours resulted in a significant increase in *Fbn2*. In Chapter 3 it was shown that microfibrils contain fibrillin 2 in canine CL tissue. Furthermore, in Chapter 4, a proportional increase in oxytalan fibres in greyhound CL tissue as the degenerative/adaptive fibrocartilaginous change progresses was demonstrated. As EDPs may be produced as a result of fibrocartilaginous change, oxytalan fibre production may be a direct consequence of cell response to EDP stimulation. As the production of oxytalan fibres may be part of the response to ligament injury in the greyhound, EDPs may play an important role in this process.

In combination EDPs and TGF- $\beta 1$ resulted in a significant increase in *ELR1* following 6 hours of treatment. As TGF- $\beta 1$ alone had no effect on *ELR1* mRNA transcription, this change may demonstrate synergism between these two treatments. If replicated at the protein level, an increase in *ELR1* may result in increased ligament cell sensitivity to EDPs. EDPs have been shown to directly influence chemotaxis, proliferation, protease release and even induce apoptosis. The increase in *Fbn2* following 24 hours of treatment may result from the effect of EDPs alone.

In conclusion this preliminary study demonstrated potential biological activity of EDPs which may play a role in ligament healing through production of fibrillin 2. Furthermore TGF- $\beta 1$ may be involved in stimulating ACL cells to increase expression of *col2a1* which could lead to the formation of fibrocartilage. As part of the initial inflammatory response, EDPs in combination with TGF- $\beta 1$ increase cell ELR1, sensitising the ACL fibroblasts to EDPs. Subsequently EDPs increase fibrillin 2 expression leading to oxytalan fibre formation as part of the response to injury. The potential importance of EDPs in ACL pathophysiology may be underestimated and further work into the effect on ACL healing may help provide a solution to the failure of healing in this tissue.

7.7 Conclusions

- 1) Local variation in CL cell types may reflect the complex biomechanics of the CLs. Contact between cells may facilitate direct communication. Differences in cell morphology between dog breeds with differing risk of ACL rupture may reflect fundamental differences in CL physiology possibly through altered cell-to-cell communication.
- 2) The distribution of elastin fibres is suggestive of a mechanical role in bundle reorganisation following ligament deformation. The presence and location of fibrillin 2 in oxytalan fibres in ligament differs from the solely fibrillin 1-containing oxytalan fibres previously described in tendon and may demonstrate a fundamental difference between ligament and tendon.
- 3) In the greyhound CL there is a significant proportional increase in OFs with advancing CL degeneration. Production of microfibrils may reflect a healing response in degenerate/injured CL extracellular matrix. Elastin forms a far greater proportion of canine CLs than previously suggested. The majority of the elastin is not in fibre form and may have a non-mechanical role.
- 4) CLs of dogs breeds at a differing susceptibility to cruciate ligament disease vary in the extent of oxytalan fibre staining with advancing ACL degeneration. For all breeds the differential rate of production or destruction of OFs may affect CL healing and reflect differing interbreed risk of ACL rupture.
- 5) Fragments of elastin containing the VGVAPG motif appear to affect ACL cells *in vitro* resulting in a significantly increased transcription of *Fbn2*. Furthermore there was a synergistic effect of EDPs and TGF- β 1 resulting in increased transcription of *ELR1*. TGF- β 1 appears to stimulate ACL cells to increase *Col2a1* mRNA transcription which could lead to an increase in ACL fibrocartilage production.

CHAPTER 8: Future Studies

8.1 Cell morphology

It was speculated in Chapter 2 that cell processes may facilitate communication between cells over potentially long distances. Immunostaining of gap junctions has been used in ligament and tendon to establish the ability of cells in contact to communicate. Using antibodies to a range of connexins it is planned to investigate the distribution of gap junction in the canine CL complex and relate this to cell morphology. Demonstration of cell processes in the canine CL complex would provide additional support for their role in communication. It was suggested that chain formations of rounded cells were an attempt to re-establish communication following disruption of cell processes. Staining of gap junctions in these chains may add substance to this idea.

Investigation of local CL mechanics may aid in understanding cell morphology. A future proposed study will involve investigating cell-matrix coupling in *in situ* fresh canine CLs. Canine knees will be dissected leaving only collateral and cruciate ligaments. The CLs will be labeled with STYO 83 (nuclear stain) and 5-DTAF (matrix stain). A jig will be adapted to hold the knee joint during full range of motion (flexion and extension). Using a confocal laser scanning microscope (CLSM), a line will be photo-bleached perpendicular to the collagen bundle axis and the knee joint flexed or extended. Three dimensional images of the cell nuclei relative to repositioning of the photo-bleached line will then be collected using the CLSM. This will allow assessment of the variation in collagen bundle movement and calculation of relative strain on the cells. As we have shown how cell morphology can be established from nuclear shape, an understanding of how strain affects cell morphology can be gained.

8.2 Elastic fibres and fibrillins

Further understanding of bundle interconnections, as illustrated using NDIC, may aid in our understanding of the functional isometricity of the canine CL. The interconnections that were described could limit bundle movement relative to each other and potentially give rise to differential bundle recruitment depending on the direction of knee motion. This information may be gained from the aforementioned dynamic CLSM study

8.3 Elastic fibres and CL degeneration

As we found CL tissue to comprise of up to 20% elastin by dry weight, further work is required to determine in what form this elastin exists in the CL. Elastin immunostaining in Chapter 3 was frequently associated with high background staining that, despite several hundred variations to the technique, could not be eliminated. Given the high proportion of elastin in the CL it now seems plausible that this was not background staining but genuine elastin staining, suggesting a diffuse distribution of elastin throughout the ligament substance. Mass spectrometry would allow identification of tropoelastin as well as variations in elastin degradation peptides.

Understanding the role of oxytalan fibres in degenerating CL will form a major part of future work. It is hypothesised that OFs have three roles in the degenerating CL

- 1) To provide a framework to allow cell or cell processes to span the defect facilitating healing,
- 2) To regulate growth factors within the wound,
- 3) To provide additional mechanical support to a degenerate ligament through lateral stress dissipation

Investigation of parts 1 and 2 will involve *in vitro* studies of canine CL cells using microfibril scaffolds. Investigation of part 3 will involve immunostaining fibrillin 2 and investigating tissue dynamics using a CLSM. An *in vivo* model of ACL healing has been established in the dog (Spindler et al., 2006). The use of this model may allow validation of the hypothesis that oxytalan fibres aid in healing of the ACL.

8.4 Breed variation in elastic fibres

It was shown in Chapter 6 that the extent of OF staining with advancing CL degeneration was inversely proportional to risk of ACL rupture in the three breeds. This pattern needs to be further investigated in a range of breeds at varying risk. Further studies will investigate turnover of OFs in the CL complex of dog breeds at a varying risk to ligament rupture. Such studies may involve *in vitro* cell culture models to assess whether ACL cells from different breeds differ in their production of fibrillins or of degradative enzymes. There may be intrinsic variation in the ability of breeds to produce OFs or breeds at greater risk of ACL rupture have a more hostile intra-articular or intra-ligament environment that precludes the formation of

OFs or accelerates their breakdown. Additionally the role of OFs in response to injury in the ACL must be investigated.

8.5 Elastin degradation peptides

This study provided fundamental information of the biological activity of EDPs. However as the study was limited by lack of power, increasing the number of dogs used may provide additional information on this biological activity.

Although significant changes in mRNA transcription were demonstrated, it is not known whether this translated to increases in protein. Work is currently underway in our laboratory to quantify MMP-9 production and activity using Western blotting and gel zymography. Protein concentrations of collagen types 1 and 2, fibrillin 2 and elastin laminin receptor 1 need to be measured to confirm the model proposed in this chapter.

The dog breed investigated in this study was the greyhound, a breed at low risk of ACL rupture. As we proposed EDPs may contribute to OF production, it would be interesting to compare the results from the greyhound to those from the Labrador retriever, a breed at increased risk.

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