Review: Emerging concepts in the pathogenesis of tendinopathy

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A B S T R A C T

Tendinopathy is a common clinical problem and has a significant disease burden attached, not only in terms of health care costs, but also for patients directly in terms of time off work and impact upon quality of life. Controversy surrounds the pathogenesis of tendinopathy, however the recent systematic analysis of the evidence has demonstrated that many of the claims of an absence of inflammation in tendinopathy were more based around belief than robust scientific data. This review is a summary of the emerging research in this topical area, with a particular focus on the role of neuronal regulation and inflammation in tendinopathy.

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Introduction

Tendinopathy is a common clinical problem, the three most common sites affected are the Achilles, patellar and rotator cuff tendons.1 The rotator cuff tendons are the most commonly affected with an annual incidence of over 1% that increases with age2,4 and consequently there is a rising rate of surgery for rotator cuff tears.5 Others include the tendons around the elbow (golfer’s and tennis elbow) and the tendons around the wrist. There is a significant disease burden attached to painful tendinopathy, not only in terms of health care costs, but also for patients directly in terms of time off work and impact upon quality of life. The primary purpose of this review is not to give an overall summary relating to all tendinopathies, it is to summarise specific emerging areas relating to tendinopathy pathogenesis research in which the authors have a particular expertise while giving a brief overall context this recent research.
Aetiology and pathogenesis

The pathogenesis of tendinopathy is certainly multifactorial and complex. Increased age is a key risk factor for the development of tendinopathy, although the commonly affected tendons all experience high levels of mechanical stress and over-use is a frequently implicated risk factor. The mechanism of overuse has been well demonstrated in animal models, while both metabolic and vascular risk factors are associated with the development of tendinopathy. Inactivity and the unloading also have an effect on tendon collagen homeostasis.

Recent systematic reviews have clearly demonstrated that patients with high cholesterol and diabetes are at significantly higher risk of developing tendinopathy, while recent review has demonstrated that an association exists between the metabolic-hormonal imbalances and tendon degeneration. Hypercholesterolaemia has also been demonstrated to have a significant impact upon tendon repair in vivo, while the clear link between hypercholesterolaemia and inflammation has been long known. This emerging link between metabolic dysregulation and chronic inflammation in tendinopathy has also been supported by a recent study using Achilles tendon biopsies from a group of patients.

The historical context relating to the rotator cuff provides an interesting insight into the frequent debates and changing viewpoints as regards tendinopathy pathogenesis. Codman initially proposed in 1934 that degeneration within the tendon was the ‘intrinsic’ primary causal factor. The ‘extrinsic’ theory relating to tendon damage secondary to attrition by surrounding structures was popularised by Neer and the term ‘impingement’ was coined. Broadly the modern consensus recognises both the role of intrinsic and extrinsic factors, but sees the intrinsic factors as being more dominant overall with the use of the term ‘impingement’ appearing increasingly baseless. It is probable that different patients have different disease phenotypes with different intrinsic and extrinsic factors playing variable roles. Certainly not all tendinopathies are identical, as represented by differences in both the tendon’s local anatomy and epidemiological profile.

Histopathology and clinical features

Tendinopathy has characteristic histopathological, clinical and radiological findings. The histopathological changes include collagen disorganisation, the increased deposition of mucoid ground substance, increased overall cellularity, as well as the appearance of round and plump ‘chondroid’ type cells. These features of apparent attempted healing diminish as degree of tendon degeneration increase. The overall picture is one of pathological chondroplasia in which tissue which normally exhibits a tensitional morphology is replaced by tissue of a fibrocartilage-like phenotype. Historically several different words have been used to described tendon related pathology including ‘tendinosis’ (implying degenerative aetiology), ‘tendinitis’ (implying inflammatory aetiology) and the more recently favoured and less aetologically specific ‘tendinopathy’. This diversity of language reflects a historical disagreement within the scientific community as to the exact role of inflammation in the aetiology of ‘tendinopathy’. Recent evidence has shown that tendon overload is linked to alterations in cell shape, as well as increased markers of inflammation and matrix degradation. The way in which the cells interact with the extracellular matrix is an area of much interest; inflammation and damage-induced matrix remodelling seem to be concentrated in, or in the vicinity of, the highly cellular interfascicular matrix. It may be therefore be postulated that interactions between the tendon, the interfascicular matrix and adjacent fat pads are instrumental in the development of tendinopathy, with the latter being a key potential source of key cytokines and inflammatory cells. This may help explain the presence of persistent inflammation in tendinopathy, a phenomenon which has been shown to have important effects on tendon cells in vitro.

Clinical symptoms including pain are frequently poorly matched to the histopathological and radiological findings, meaning that a high proportion of patients with a tendon that is both histopathologically and radiologically degenerate experience have no related pain or symptoms. The reasons for this mismatch between pathology and perceived pain are poorly understood, however recent research has identified the peripheral and central pain processing pathways as good candidates for an explanation. It appears that the presence of pain in tendinopathy not only relates to mechanical changes in the tendon but also changes to the ways in which the local cells and the peripheral nerves react to this change, thus contributing to the nociceptive pathways to higher centres being activated. Overall the vast majority of tendon ruptures (97%) occur in patients with histopathologically abnormal tendons.

Neuronal pathways and glutamate

The neuronal response to tendon injury involves nerve in growth during the initial inflammatory phase; the subsequent proliferative and remodelling phases are regulated by sensory nerves, as well as the glutaminergic and autonomic systems. Glutamate is an important amino acid involved in many key physiological processes including cell metabolism, pain sensitization and collagen synthesis. Glutamate receptors can be broadly broken down into two major types: inotropic, which are glutamate-gated ion channels (iGlu), and metabotropic, which are G-protein coupled receptors that modulate signal transduction cascades (mGlu). The inotropic receptors include Kainate (KA) receptors, α-amino-3-hydroxy-5-methyl-4-isoxazole propionic acid (AMPA) receptors and N-Methyl-D-Aspartate (NMDA) receptors. Glutamate has been shown to induce pain and hyperalgesia when injected around human tendon tissue. An upregulation of the glutaminergic signalling has been linked to inflammatory change in a rat supraspinatus model.

The first study to recognize the presence of glutamate in tendinopathy used a microdialysis technique in chronic painful Achilles tendinopathy. Glutaminergic changes have since become increasingly described in painful tendinopathy, including an increase in extracellular glutamate
concentration and the up-regulation of N-Methyl-D-Aspartate (NMDA) receptors. Recently the upregulation of the glutaminergic system has been confirmed to be present in rotator cuff tendinopathy for the first time. This histological and immunohistochemical study demonstrated that an increase in glutamate staining was present in the painful tendinopathic rotator cuff tendons alongside the classical histological changes which included increased collagen disorganization and cellularity. Glutamate staining was distinctly expressed in resident cells within the tendon. The release of glutamate from tendon cells was first hypothesized by Scott et al., who detected vesicular glutamate transporter expression in cells localized in tendon tissue in lower limb tendinopathies. This was later detected by Schizas et al. Because of the mechanical and structural differences between tendon locations, it cannot be assumed that cell behaviour is uniform in both upper and lower limb tendinopathies. This study has also shown significant staining of certain ionotropic and metabotropic receptors on tendon cells residing in damaged rotator cuff tissue, for example NMDAR1 and mGluR1. This study did not find any correlation with the severity of patient pain symptoms. This is not unsurprising as previous studies have failed to detect this link.

Sensory neuropeptide expression has also been shown to be associated with both failed healing and pain in an animal model of tendon injury, they have also been causally linked with tendinopathy-like changes in animal models. Neuropathic conditions have pro-proliferative, angiogenic and stem cell-stimulating properties in vitro, however little work had been carried out to investigate the effects of glutamate on tendon derived cells. Dean et al. exposed tendon derived cells from both healthy controls and patients with tendinopathy to different concentrations of glutamate and specific glutamate receptor inhibitors. Tendon derived cell viability was reduced after 72 h of exposure to relatively low concentrations of glutamate (0.05 mM and 1.875 mM) and this deleterious effect was attenuated by NMDAR antagonism. Higher concentrations of glutamate reduced cell viability at 24 h in tendon tear derived cells and not in control cells. A reduction in collagen (COL1A1 and COL3A1) and increase in aggregcan gene expression were seen after both 24 and 72 h of glutamate exposure. Overall the in vitro effects of glutamate in terms of reducing cell viability, decreasing collagen gene expression and increasing aggregcan gene expression suggested that the raised levels of glutamate contributes to the pathogenesis of tendinopathy.

**Clinical meaning of neuronal changes**

Tendon samples from patients with persistent pain demonstrated increased levels of metabotropic glutamate receptor 2 (mGluR2), Kainate receptor 1 (KA1), Protein Gene Product 9.5 (PGP9.5), CD206 (macrophage marker) and CD45 (pan-leucocyte marker) versus pain-free controls. Notably the painful and pain-free patient groups were matched in terms for basic demographic and tendon structure, while there were no differences between groups in terms of the basic histology.

NMDAR1 co-localised with CD206 positive cells, whereas PGP9.5 and glutamate were predominantly expressed by resident tendon cells. Within the gene expression data related to cells derived from rotator cuff years there were strong correlations between CD206 expression and glutamate receptor expression. This work demonstrated an association between glutaminergic and pro-inflammatory changes in tendon cells and pain. To our knowledge this is the first histological study that has used structurally and age matched tendon from pain-free tendinopathic patients as a control. The co-localisation of NMDAR1 and CD206 suggests that certain glutamate receptors are predominantly expressed on ‘inflammatory’ type cells within tendon.

Recent *in vivo* work by Valkering et al. has investigated the role of glutamate in Achilles tendon healing following acute Achilles tendon rupture. Patients were randomised into two groups, and those in the functional weight bearing group had significantly higher levels of glutamate than the non-weight bearing group, while the higher glutamate levels correlated with both the level of the marker of procollagen type I (PINP) and improved functional outcome at six months. Substance P enhances cell proliferation and sensory nerve in growth in the rat, conversely joint immobilisation has been shown to reduce the expression of sensory neuropeptides in the rat and this is associated with significantly poorer tendon healing. In clinical terms glutamate appears important in normal tendon healing, while its upregulation in tendinopathy appears consistent with an environment of potentially failing healing and perhaps a failure of inflammation to resolve.

**The emergence of inflammation**

The dwindling of popularity of the term ‘tendinitis’ represented skepticism regarding the role of inflammation in tendon degeneration. Recent systematic analysis of the evidence has demonstrated that many of the claims of an absence of inflammation in tendinopathy were more based around belief than robust scientific data. Several studies have stated that ‘no inflammatory cells’ were present in samples from tendinopathic patients but had only looked for neutrophils, not other types of ‘inflammatory cell’. In fact the evidence to support the role of inflammation in tendinopathy pathogenesis has become increasingly overwhelming in recent years with a majority of studies demonstrating increased numbers of macrophages in diseased tendons.

Macrophages are known to play an essential role orchestrating inflammation and tissue repair. The signalling pathways underpinning activation of macrophages to become M1 or M2 subtypes have been revised to identify the signalling pathways underpinning macrophage activation interferon, NF-κB, (STAT6) and glucocorticoid receptor activation pathways. These macrophage activation pathways have recently been identified in samples of diseased human rotator cuff tendons. Tendon tissues from patients with early stage disease showed increased expression of genes and proteins induced by Interferons and NF-κB. Conversely, tendons from patients with advanced stage disease (large to massive tears) showed expression of genes and proteins induced by STAT6 and glucocorticoid receptor activation pathways. These
findings highlight the complexity of inflammatory processes in diseased tendons and how they might change with the stage of disease.

Increasing evidence has shown that inflammatory mechanisms and the innate immune system are activated within the tendon matrix microenvironment during tissue injury and dysregulated homeostasis. An essential component of the regulatory process that drives tendon remodelling includes cytokines that dictate cell type and tissue specificity of responses that ultimately balance a reparative versus degenerative process. Endogenous expression of TNFα, IL-1β, IL-6, IL-10, VEGF and TGFβ has been demonstrated in tenocytes65–68 while diseased human tendon has shown expression of Such expression is functionally implicated in vivo: for example, the mechanical properties of healing tendons in IL-6−/− mice were inferior compared with littermate controls.69

Recent mechanistic dissection has highlighted a role for the cytokine IL-33, a member of the IL-1 cytokine family that plays a major role in innate and acquired immune responses, in matrix/inflammatory crosstalk in tendon damage.70 IL-33 message and protein expression was significantly increased in early human tendinopathy compared to both established tendinopathy and normal tendon while the addition of exogenous IL-33 to in vitro human tenocyte cultures resulted in a reduction of collagen mRNA/protein. Moreover, an in vivo patellar tendon injury model the addition of IL-33 significantly reduced the tendon strength (load to failure) of WT mice by ~30% at early time points, likely as a consequence of the concomitant collagen 3 matrix changes, which result in mechanically inferior tendon. Taken together these studies demonstrate a key functional role for IL-33 in early injury induced matrix dysregulation and subsequent cytokine feedback mechanisms that have an ultimate biomechanical and clinical effect.

Emerging studies highlight microRNAs (miR) as key regulators of leukocyte function and the cytokine network while orchestrating proliferation and differentiation of stromal lineages that determine extracellular matrix composition.71 We identified reduced expression of miR-29a, which directly targets numerous extracellular matrix genes and is implicated in the regulation of innate and adaptive immunity, in human biopsies and demonstrated that its reduction leads to development of tendinopathy. In human tenocytes, miR-29a was only capable of influencing the expression of collagen type 3 and not type 1. This work has been expanded to an equine tendinopathy model,72 considered equivalent to human disease, reinforcing that injury-induced loss of miR-29a is responsible for the over-expression of col3 seen in equine tendinopathy. Collectively these data suggest that the reintroduction of miR-29a to the injury-induced miR-29a deficiency in tendon could reverse the key collagen switch that remains a core pathological feature of tendinopathy. Thus while the functional contribution of cytokine biology and its downstream consequences in tendinopathy remains to be established there is now a convincing scientific rationale towards translational immunobiology to benefit tendinopathy patients.73 The short term benefits of anti-inflammatory drug treatments in tendinopathy have been demonstrated in the short term,74 however there remains a distinct lack of effective long term treatments of any form.75

Conclusions

Fundamentally a better understanding of the pathogenesis of tendinopathy and the underlying mechanisms is essential if we are to develop more effective long term treatment strategies for the management of tendinopathy.76,77 Our improved understanding which includes this work relating to the emerging roles of the glutaminergic and inflammatory systems means that more effective novel treatments may be just around the corner.

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