




Metabolic Regulation of Tendon Inflammation and Healing Following Injury

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Abstract

Purpose of Review This review seeks to provide an overview of the role of inflammation and metabolism in tendon cell function, tendinopathy, and tendon healing. We have summarized the state of knowledge in both tendon and enthesis.

Recent Findings Recent advances in the field include a substantial improvement in our understanding of tendon cell biology, including the heterogeneity of the tenocyte environment during homeostasis, the diversity of the cellular milieu during in vivo tendon healing, and the effects of inflammation and altered metabolism on tendon cell function in vitro. In addition, the mechanisms by which altered systemic metabolism, such as diabetes, disrupts tendon homeostasis continue to be better understood.

Summary A central conclusion of this review is the critical need to better define fundamental cellular and signaling mechanisms of inflammation and metabolism during tendon homeostasis, tendinopathy, and tendon healing in order to identify therapies to enhance or maintain tendon function.

Keywords Tendon · Enthesis · Inflammation · Metabolism · Tendinopathy · Tendon healing

Introduction

The basal metabolic rate of tendon is relatively low, likely due to the high degree of quiescence observed in tendon cells during tissue homeostasis. However, multiple factors can lead to the loss of cell quiescence, alterations in tendon cell functions, and altered tendon function, with inflammation being a key regulator of these processes. This review will summarize the central inflammatory and metabolic mechanisms that regulate (i) tendon cell function; (ii) disruptions in tendon homeostasis (e.g., tendinopathy); and (iii) healing of acute tendon injuries and how inflammation can interact with and modulate metabolism in these contexts (Fig. 1). In addition, within the respective sections on tendinopathy and healing

of acute tendon injuries, a distinction is made between the tendon mid-substance and the tendon-bone junction (enthesis) as appropriate, given that there are important region-specific differences in the tendon response to inflammation and altered metabolic function.

Metabolism, Inflammation, and In Vitro Tendon Cell Biology

It has long been established that tendon cells are capable of both glycolysis and mitochondrial respiration for energy production [1, 2]. Recently, it has been shown that loss of *scleraxis* expression in tendon cells leads to upregulated oxidative phosphorylation and mitochondrion organization, suggesting that putative tendon markers affect tendon cell metabolism [3]. In contrast to tendon cells isolated during homeostasis, tendon cells isolated from injured human flexor tendons exhibited increased glycolytic pathway flux and the capacity for differentiation down both tenogenic and chondrogenic pathways [4•]. When treated with a glycolysis inhibitor, chondrogenic differentiation was inhibited and tenogenic differentiation was stimulated, suggesting that glycolysis pushed tendon cells along a non-tenogenic fate [4•].

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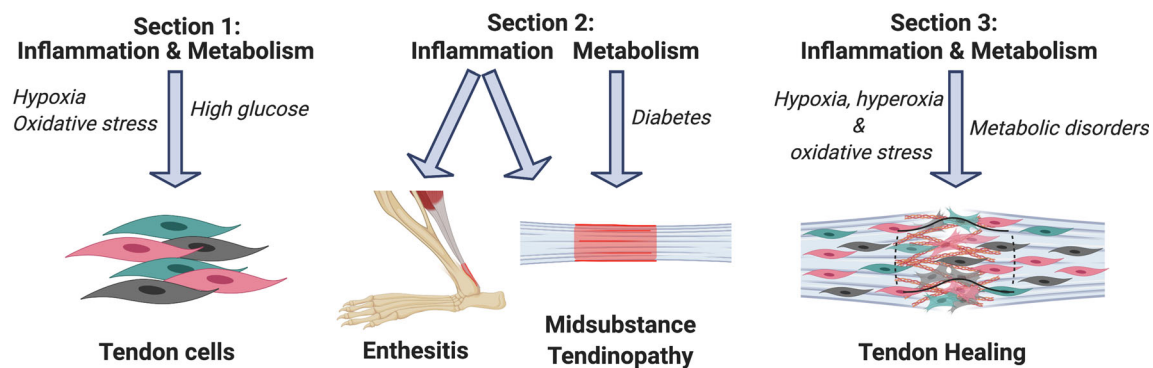


Fig. 1 Overview of the review. Section 1 summarizes the effects of inflammation and metabolic alterations on in vitro tendon cell function, with a particular emphasis on the role of inflammation-induced hypoxia and oxidative stress, as well as the impact of high glucose on tendon cell function. Section 2 summarizes the impact of inflammation on both the tendon enthesis and mid-substance tendinopathy, as well as how

alterations in systemic metabolic function including diabetes can promote tendinopathy. Section 3 focuses on the effects of inflammation, including hypoxia, hyperoxia, and oxidative stress on the healing process, and the effects of systemic metabolic disorders on the tendon healing process. This figure was created using [Biorender.com](https://www.biorender.com)

Furthermore, treatment of tendon cells with the inflammatory cytokine IL-1 β drives lactate production, which results in decreased collagen and decreased expression of key tendon markers [5]. This demonstrates that inflammatory signaling can directly modulate tendon cell metabolism.

Hypoxia

Numerous studies have explored the effects of oxygen deprivation (hypoxia, typically 1–5% O₂) on tendon cells in vitro. Tendon cells cultured in hypoxic conditions have limited mitochondrial energy production accompanied by elevated glycolytic flux and lactate production, demonstrating that hypoxia drastically changes metabolic processes in tendon cells [6]. For example, hypoxia drives tendon cell proliferation and prevents multilineage differentiation potential while up-regulating stem cell markers such as NANOG and OCT-4, compared to normoxic culture conditions (~20% O₂) [7–11]. Hypoxia also drives increased expression of tendon markers, such as tenomodulin [7]. Importantly, tendon cells cultured in hypoxic conditions and subsequently cultured under normoxic conditions regain their multilineage differentiation potential, demonstrating that the effects of hypoxia on tendon cells are reversible [7]. Hypoxia also limits the formation of primary cilia on tendon cells, potentially limiting their mechanosensory properties [12]. Moreover, highly hypoxic conditions (~0.1% O₂) and anoxia lead to tendon cell death via apoptosis [13, 14].

Previous studies have demonstrated that aged tendon cells have impaired proliferative capacity, decreased expression of tendon markers, increased adipogenic differentiation potential, increased mineralized fibrocartilage phenotype, and increased senescence relative to young tendon cells [15–18]. Interestingly, culturing aged tendon cells with the conditioned media from young, hypoxic tendon cells increases proliferative capacity and migration rate of the aged cells, increases

expression of tendon markers, and decreases the number senescent cells, suggesting that paracrine signaling via secreted factors from hypoxic tendon cells reverses the aged tendon cell phenotype [17]. Furthermore, hypoxic tendon cell co-cultures drive other cell types, such as adipose-derived mesenchymal stem cells, along a tendon cell fate. Recent studies show that hypoxia triggers tendon cells to produce exosomes containing upregulated extracellular matrix-related genes, further supporting the idea of paracrine, cell–cell–mediated communication during hypoxia [19, 20].

Oxidative Stress

In addition to hypoxia, several other factors such as diet and environmental stimuli can drive oxidative stress. Many studies have examined the effects of oxidative stress in the form of reactive oxygen species (ROS) on tendon cells in vitro. Treatment of tendon cells with ROS results in decreases in cellular proliferation, migration, viability, and stemness [21, 22]. Activation of mitochondrial aldehyde dehydrogenase 2 (ALDH2), a known reliever of oxidative stress, in tendon cells prevents H₂O₂-induced cell death and prevents depolarization of mitochondrial membrane potential [23]. In addition, multiple vitamins have antioxidant properties. Treatment of tendon cells with vitamin C also decreases NO synthesis by tendon cells [24]. Treatment of tendon cells with low-dose vitamin C increases cell proliferation, viability, and migration [22, 24]. Tendon cell proliferation is further enhanced by co-treatment of vitamin C and thyroid hormone T₃ [24]. Similarly, treatment with vitamin D increases cell proliferation and reduces production of reactive oxygen species [25, 26]. However, vitamin D also reduces gene expression of type I collagen [25]. Retinoic acid, a metabolite of vitamin A, induces nuclear localization of scleraxis and aids in the maintenance of tendon stem cell properties [27].

High Glucose

Supplementation of culture medium with high levels of glucose modulates many aspects of tenocyte function. More specifically, tenocytes in high glucose demonstrate altered inflammatory signaling via IL-6 and COX2, increased ROS production, decreased proliferation, decreased migration, decreased mitochondrial membrane potential, and increased apoptosis [28–34]. Moreover, high glucose decreases the expression of tendon genes, promotes simultaneous decreases in type I collagen and increases in type III collagen expression, increases the expression and activity of multiple MMPs and TIMPS, and stimulates adipogenic transdifferentiation [28–34]. In addition, the effects of high glucose are exacerbated in the presence of advanced glycation end products (AGEs), as the combination leads to reduced proliferation, reduced ATP production, decreased electron transport efficiency, and alterations in collagen and MMP gene expression [34••]. Consistent with the effects of high glucose supplementation, tendon cells isolated from diabetic rat patellar tendons exhibit decreased proliferative ability, decreased expression of tendon markers, and increased osteogenic and chondrogenic differentiation ability [35], mimicking many effects of high glucose culture conditions and suggesting that diabetic cells retain some “memory” and, therefore, functional alterations of the in vivo environment. Interestingly, mechanotransduction can suppress several aspects of the effects of high glucose on tendon cells. Mechanical stretch prevents adipogenic transdifferentiation, increases tendon cell migration, and enhances fibroblastic-like morphology of cells under high glucose conditions, suggesting that mechanotransduction can ameliorate some of the negative outcomes associated with hyperglycemia [30•].

Other Metabolic Mediators of In Vitro Tendon Cell Function

Cholesterol can also affect tendon cell function. Recent studies have shown that high cholesterol suppresses tendon cell proliferation, cell migration, and *scleraxis* gene expression, while increasing ROS production [36, 37]. The effects of female sex hormones on tendon cells have also been investigated. Estrogen increases cell proliferative ability and decreases adipogenic differentiation potential of tendon cells [38]. Additionally, treatment with estradiol-17 β increases tendon cell proliferation but decreases *Scleraxis* gene expression [25]. Finally, there is some evidence that treatment with platelet-rich plasma (PRP) induces inflammation-related changes in tendon cell function. PRP treatment of tendon cells upregulates TNF- α -induced NF- κ B signaling pathway, downregulates the expression of extracellular matrix genes, and induces the expression of autophagy-related and ROS-related genes [39].

Inflammation and Metabolism in Tendinopathy

Inflammation and Mid-substance Tendinopathy

Loss of tendon homeostasis, resulting in tendinopathy, is characterized by inflammation and degeneration of the native tendon tissue. Tendinopathy represents a major clinical burden, decreases patient quality of life, and increases the risk of tendon rupture [40]. Our current understanding of the pathophysiology of tendinopathy is a combination of supraphysiological overloading and inflammation [41–43]. Rather than a single overloading event, cyclic overloading is required for the rapid expression of inflammatory mediators that are observed in tendinopathies [44]. These inflammatory mediators include alarmins (such as HIF-1 α , IL-33, S100a8/9, and HMGB1), which have been implicated in the earliest phases of tendinopathy development [43, 45, 46]. Moreover, there is strong evidence that many of the pathological changes observed in tendinopathy are driven largely by extrinsic factors [47•]. More specifically, immune cell involvement is thought to play a central role in tendinopathy development, although there is limited direct evidence for this contribution. Indeed, a recent review of inflammatory mediators of tendinopathy [48] defines a high degree of heterogeneity in terms of the extent of the inflammatory cell environment that has been characterized in tendinopathy studies. However, samples from both Achilles and supraspinatus tendinopathy patients demonstrate an influx of pro-inflammatory monocyte-derived macrophages, with increased macrophage content occurring with increased disease severity [41, 49], suggesting that severity of tendinopathy may be characterized by examination of the inflammation signature.

In terms of modulating inflammation to treat tendinopathy, anti-inflammatory approaches have had mixed success but have also highlighted the effects of treatment timing on treatment efficacy. While some in vitro studies have suggested that aspirin can promote tenogenesis via GDF7/Smad1/5 signaling [50], in vivo studies have not demonstrated the same promise. For example, Heinemeier et al. found that 1-week treatment with ibuprofen elicited no change in adult human chronic tendinopathic tissues [51], while Bittermann et al. showed that ibuprofen treatment during the inflammatory phase actually blunted healing in a model of murine Achilles tendinopathy [52]. A potential alternative anti-inflammatory approach involves the use of lipid mediators of aspirin-induced eicosanoid metabolism, such as 15-epi LXA₄. These have proven effective in other chronic inflammatory diseases such as pulmonary inflammation and eczema and, therefore, may also hold promise in resolving tendon inflammation without the damaging side effects [41, 53].

Inflammation and Enthesitis

In addition to impacting the integrity and homeostasis of the tendon mid-substance, inflammation also impacts the integrity of the tendon-bone interface (enthesitis). Enthesitis, or inflammation of the enthesis, is predominantly associated with arthritis [54] but can also be initiated by repeated bouts of high mechanical stress, which induces an inflammatory response [55], consistent with mid-substance tendinopathy development. For example, while mice overexpressing TNF are highly susceptible to enthesitis and arthritis, hindlimb suspension inhibited arthritis development and blunted the TNF-mediated inflammatory response, compared to controls [56]. Importantly, enthesitis is a hallmark of spondyloarthritis (SpA), particularly juvenile cases [57••], which often go undiagnosed due to lack of diagnostic criteria. As such, increasing our fundamental understanding of the cellular and molecular mechanisms that drive and initiate increased enthesitis in the context of SpA could have a substantial impact of both disease diagnosis and treatment.

Metabolism and Tendinopathy

In addition to inflammation and mechanical overloading, a variety of metabolic diseases can also initiate tendinopathy development. While metabolic syndrome encompasses many comorbidities, including high blood pressure, high blood glucose, and obesity, very few studies have examined the collective effect of metabolic syndrome on tendon pathology but have more commonly looked at the impact of individual comorbidities. However, there is a strong association between metabolic syndrome and the development of trigger finger [58]. In terms of individual conditions, type II diabetes greatly increases the risk of tendon pathology [59–62] (summarized in Fig. 2), likely due to a combination of chronic, low-grade inflammation and a

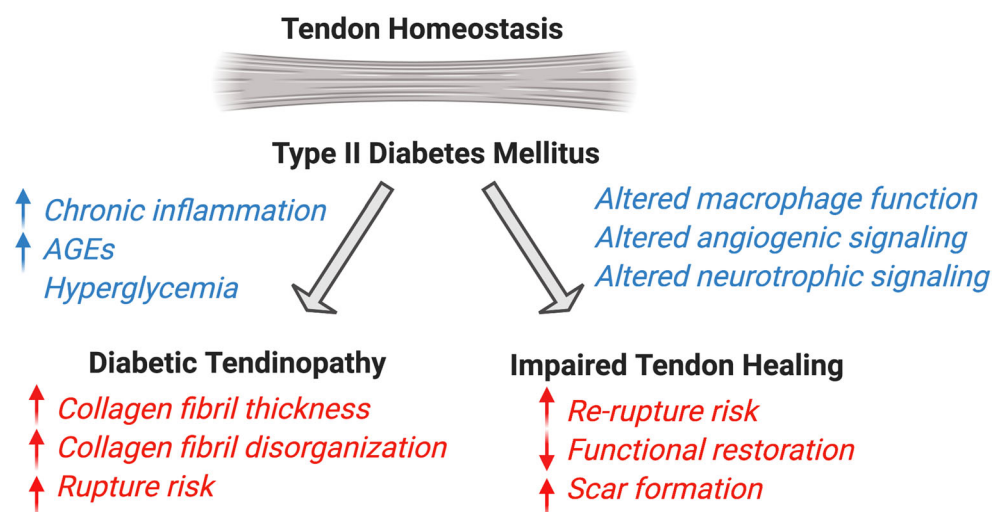
high glucose environment. Consistent with this, there is some evidence that alterations in glucose metabolism may represent a metabolic marker of tendinopathy [6••]. Moreover, diabetic tendons are thicker and demonstrate fibril disorganization at homeostasis, changes that are thought to be due to a combination of hyperglycemia and accumulation of AGEs [30, 34]. AGEs promote increased cross-linking with age or diabetes and inhibit normal sliding of tendon fascicles, leaving them prone to pathology [63]. Other perturbations in metabolism have also been shown to influence development of pathological conditions, such as elevated cholesterol levels [37, 64, 65] though these effects are less well-characterized.

Metabolic and Inflammatory Mediators of Acute Tendon Healing

Metabolic Disorders

Metabolic disorders, such as diabetes mellitus, dramatically affect the tendon healing process (Fig. 2). For example, rotator cuff repairs in diabetic patients are up to two times more likely to re-rupture than nondiabetic patients and show evidence of diminished healing [66, 67], including slower or decreased improvements in pain and function [68]. Moreover, murine and rat models of type II diabetes demonstrate decreased biomechanical properties following tendon injury, compared to nondiabetic animals [69–71]. Healing type II diabetic tendons also exhibit increased scar tissue formation [71], aberrations in macrophage polarization [71], and detrimental effects to angiogenic and neurotrophic signaling pathways [72]. Interestingly, type I diabetes also disrupts tendon healing. Streptozotocin-induced type I diabetes suppresses immune cell infiltration (including macrophages), decreases cell proliferation, and decreases biomechanical properties during

Fig. 2 Summary of the impact of type II diabetes on tendon homeostasis and healing of acute tendinopathy. This figure was created using [Biorender.com](https://www.biorender.com)



healing, relative to nondiabetic controls [73–75]. To better understand the effects of high blood glucose on Achilles tendon healing, nondiabetic rats were supplemented with a high glucose diet, which did not elevated blood glucose levels even after 4 weeks of treatment. Glucose supplementation resulted in increased tendon thickness and stiffness, as well as an altered gait pattern [76]. In addition, cell proliferation was increased, as was expression of chondrogenic markers (*Sox9*, *Col2a1*, *Acan*, *Comp*), and cartilage-like areas were detected within the repair tissue [76], suggesting that high glucose alters in vivo tenogenic function and may promoter aberrant chondrogenic differentiation in the healing tendon.

Interaction of Metabolism and Inflammation

Acute tendon injuries contain elevated levels of metabolites, including glutamate, lactate, and pyruvate, compared to uninjured tendon [77–79]. Recently, ¹³C-glucose labeling was utilized to examine glycolytic and TCA cycle activity during murine Achilles tendon healing following transection. While glycolysis was elevated at both 1 and 4 weeks post-injury, the flux through the TCA cycle was significantly increased at 1 week post-repair [78••]. This indicates that multiple metabolic processes are highly active during the early, inflammatory phases of healing, while glycolysis persists into the later remodeling phase. A clinical study in humans demonstrated that glutamate, lactate, and pyruvate are elevated at 2 weeks post-Achilles tendon injury [79]. Weight-bearing mobilization of the injured tendon further increased glutamate, which was significantly correlated with elevated procollagen type I levels [79]. Interestingly, metabolic activity, as measured by glucose uptake remains elevated in healing human tendon through at least 12 months post-injury, with metabolic activity being highest shortly after injury and slowly decreasing over time [80••]. Patient-reported tendon functional scores were negatively correlated with high glucose uptake, suggesting that sustained metabolic activity may negatively impact healing [80••]. In contrast, an observational study examining human Achilles tendon rupture duration of operating time (DOT) found that patients who experienced longer DOT had higher levels of glutamate, which was significantly associated with improved functional outcomes [81].

There has been relatively little exploration into how modulation of metabolic pathways affects tendon healing. However, inhibition of lactate synthesis (a metabolite of the glycolysis product pyruvate) was positively correlated with decreased width and cross-sectional area of the healing tendon, increased biomechanical properties, improved collagen fiber alignment, and reduced mineralization of the injury site, suggestive of improved healing [78••]. Interestingly, the effects of acute injury and healing on

mitochondrial function are unclear. For example, enhanced mitochondrial activity is observed during rat rotator cuff healing [82], while diminished mitochondrial activity was observed following rotator cuff tenotomy in a sheep model [83]. Moreover, photobiomodulation, a nonionizing laser therapy that stimulates mitochondrial energy production, resulted in mild improvements in mechanical properties of Achilles tendons after injury [84]. Collectively, these studies demonstrate the need for additional work to understand the specific contributions of metabolic processes throughout the various stages of tendon healing and to determine how modification of mitochondrial function may impact tendon healing.

Hypoxia, Hyperoxia, and Oxidative Stress

While alterations in tendon cell function as a result of oxygen concentration are well-characterized in vitro, the in vivo effects are less clear. However, the importance of oxygen concentration and consumption is demonstrated by clinical studies in the rotator cuff, which demonstrate decreased cellular activity and oxygen consumption in large tears and are associated with worse outcomes, relative to smaller tears [85]. As such, a few studies have examined the effects of hyperbaric oxygen treatment on tendon healing, although no consensus has yet emerged. In a rat patellar tendon injury model, hyperbaric oxygen supplementation did not alter the healing process in one study [86], while another reported increased collagen gene expression at 1–2 weeks post-injury [87]. Recent work demonstrated increased fibrotic tissue during Achilles tendon healing following supplementation with hyperbaric oxygen in a rat model [88].

Given the effects of oxidative stress on tendon cell function in vitro, substantial investigation has been conducted into the efficacy of vitamin C due to its antioxidant functions and role in collagen formation [89]. Vitamin C supplementation increases angiogenesis and type I collagen deposition in a rat Achilles tendon injury model [90]. Moreover, administration of a supplement containing mucopolysaccharides, vitamin C, and collagen during Achilles tendon healing did not alter biomechanical properties or collagen production but did increase cell proliferation and TGF- β 1 production in endotenon fibroblasts [91]. Local administration of vitamin C following Achilles tendon repair improved mechanical properties [92] and functional outcomes [93] of healing tendons, while a combination of vitamin C and thyroid hormone T₃ also enhanced healing [94].

While oxygen concentration and consumption, as well as oxidative stress, impact tendon cell function and in vivo healing, more work is needed to determine if and how these pathways can be utilized to enhance tendon healing.

Conclusions and Key Knowledge Gaps

Given the frequency of tendon injuries, including those induced by tendinopathy, and the substantial subsequent complications that can occur, understanding the underlying mechanisms that regulate healing is critical to the identification of therapeutic targets. Inflammation is central to successful healing; however, aberrant, excessive, or insufficient inflammation all has profound effects on the tendon healing process. As such, future work is needed to better define (i) the central inflammatory signaling cascades; (ii) how inflammatory and immune cells interact with resident tendon cells and extrinsic cells to mediate the healing process, and (iii) how inflammation modulates cell metabolism and therefore cell function during healing. In terms of tendon homeostasis, while the effects of altered systemic metabolism (e.g., type II diabetes) on the tendon are clear (Fig. 2), there is a gap in knowledge in terms of our fundamental understanding of how metabolism regulates tendon cell function to maintain homeostasis and how different aspects of metabolism may be targeted to restore normal tenogenic function and therefore tendon homeostasis in the context of altered systemic metabolism (e.g., metabolic syndrome or diabetes). Collectively, while much remains unknown about the molecular and cellular level impact of altered inflammation and metabolism on tendon homeostasis and healing, the recent and rapid maturation of the tendon field suggests it is only a matter of time until these processes are more clearly defined and can therefore be leveraged to prevent or reverse tendon pathology and improve healing.

Compliance with Ethical Standards

Conflict of Interest Jessica Ackerman, Katherine Best, Samantha Muscat, and Alayna Loiselle declare that they have no conflicts of interest.

Human and Animal Rights and Informed Consent This article does not contain any primary data with human or animal subjects performed by any of the authors.

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