Intestinal Fibrosis and Liver Fibrosis: Consequences of Chronic Inflammation or Independent Pathophysiology?

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Introduction

Fibrosis is characterized by an exaggerated accumulation of extracellular matrix (ECM) proteins and an expansion of mesenchymal cells [1]. Fibrotic alterations can be recognized in several different organs of the human body such as the skin, lung, kidney, liver, and the gut. With regard to the latter, clinically apparent fibrosis is most frequently associated with Crohn’s disease (CD), a chronic inflammatory intestinal disorder with unknown etiology [2, 3]. More specifically, within the first 10 years after diagnosis, up to 50% of CD patients will develop a penetrating or strictureing course of disease [2, 4]. Patients suffering from strictureing CD may present with a persistent luminal narrowing that can lead to obstructive symptoms and an impaired quality of life. While inflammatory strictures may respond to anti-inflammatory medical treatment, fibrostenotic strictures do not resolve upon immunosuppressive therapy. Due to the paucity of anti-fibrotic drugs for intestinal fibrosis [5], CD-associated...
Fibrotic strictures are a major reason why approximately 75% of CD patients have to undergo surgery at least once during their lifetime [6].

In the context of chronic liver diseases, hepatitis B and C viruses (HBV and HCV) are among the most frequent causes for the development of liver fibrosis [7–9]. The progression from fibrosis to liver cirrhosis is of particular importance for affected patients, since the risk for hepatocellular carcinoma is significantly increased in the cirrhotic liver [8, 9]. Comparable to intestinal fibrosis in CD, where several mechanisms were identified to drive fibrogenesis, including cytokines, chemokines, or mesenchymal cells, the same players have been found during the manifestation of liver fibrosis.

In this review, we summarize the current knowledge on principal mechanisms shared by intestinal fibrosis and liver fibrosis. We furthermore discuss inflammation as the cause of fibrogenesis in both entities and depict unique features of intestinal and hepatic fibrosis.

**Cellular Basis of Fibrosis**

The main effector cell mediating intestinal fibrosis is considered the intestinal mesenchymal cell that is responsible for the excessive synthesis of ECM proteins. It exists in three distinct forms: the fibroblast [vimentin positive, α-smooth muscle actin (α-SMA) negative, desmin negative], the myofibroblast (vimentin positive, α-SMA positive, desmin negative), and the smooth muscle cell (vimentin positive, α-SMA positive, desmin positive). Mesenchymal cells can actively differentiate and de-differentiate between these cellular phenotypes.

The liver is unique in the sense that hepatic stellate cells (HSCs), fibroblast or myofibroblast precursor cells located within the space of Disse along the hepatic sinusoid, play the central role in hepatic fibrogenesis. HSCs have a characteristic feature in that they possess fat droplets containing vitamin A [10]. During the course of chronic liver injury and inflammation, HSCs activated by profibrotic mediators, such as platelet-derived growth factor or transforming growth factor-β (TGF-β), transform into myofibroblasts and deposit ECM in the liver parenchyma, resulting in liver fibrosis.

Such vitamin A-containing stellate cells were originally considered liver specific. However, the same types of cells are now recognized in the pancreas, kidney, and lung, contributing to organ fibrosis. From the viewpoint of the common origin of collagen-producing cells in the liver and intestine, abundant vitamin A-storing cells have been found in the lamina propria of the gastrointestinal mucosa in the lamprey, an observation that could be relevant to humans [11]. These vitamin A-storing cells may differentiate into the visceral type of fibroblasts that are distinguishable from dermal fibroblasts of somatic mesodermal origin (fig. 1). On the other hand, a recent gene expression analysis of fibroblasts present in various organs has indicated that human fibroblasts present in the gastrointestinal...
nal tract have a significantly different gene expression profile from the profiles in other organs including the liver [12]. These findings suggest that the local environment could have a role in shaping fibroblast function.

A key feature of fibrosis of the liver and intestine – aside activation of mesenchymal cells – is the expansion of the mesenchymal cell pool. This is highly relevant, since prevention of fibroblast accumulation could offer future therapeutic potential. A number of studies have reported that fibrocytes, which are circulating CD14+/CD45+ collagen-producing cells derived from bone marrow, act as a precursor of myofibroblasts and contribute to the progression of clinical and experimental liver fibrosis [13, 14]. However, subsequent studies using type I collagen reporter mice have shown that the direct contribution of fibrocytes and other blood-born collagen-producing cells to hepatic fibrogenesis is rather limited [15] or even negligible [16]. In the intestine, descriptive data are suggestive of a contribution of fibrocytes to fibrogenesis [17], but a functional evaluation is lacking.

Epithelial-mesenchymal transition (EMT) or endothelial-mesenchymal transition is a process in which fully mature epithelial or endothelial cells acquire a spindle-shape morphology, lose their typical markers, and acquire a mesenchymal phenotype and function [18]. Contribution of EMT to the progression of liver fibrosis has been reported in parenchymal hepatocytes and biliary epithelial cells [19, 20]. This finding, however, is controversial: a series of cell fate mapping experiments questioned EMT as an underlying mechanism of hepatic fibrogenesis [21–23]. One report is available suggesting colonic EMT contributing to intestinal fibrogenesis [24], and inflammation-induced endothelial-mesenchymal transition has been shown to occur in vitro, in situ, and in vivo in intestinal inflammation [25], however, to a limited extent.

**Innate Immunity**

*Pathogen-Associated Molecular Patterns*

The majority of our microbes are found in the gut and influence health and disease. Alterations in the immune system, the microbiota, or the barrier function of the intestine leading to bacterial translocation into the intestinal mucosa or portal circulation can change host-microbiome interaction and lead to inflammation. Essentially, all intestinal as well as hepatic immune and nonimmune cells sense microbe-derived pathogen-associated molecular patterns (PAMPs) through pattern recognition receptors [26].

Evidence about the influence of microbial composition or sensing on intestinal fibrosis can be derived from human genetic studies. Gene variants affecting innate immunity, located in or near genes involved in bacterial recognition and processing, such as NOD2 [27], are associated with fibrostenosing inflammatory bowel disease (IBD). Comparable genes have been linked to complicated IBD, and the risk of these complications increases proportionally with the number of variants carried by the patient [28]. Circulating antibodies against microbial peptides are qualitatively and quantitatively associated with and predictive of a more complicated disease phenotype [29]. In addition, essentially all animal models of intestinal fibrosis appear to be dependent on the presence of a microflora to initiate or perpetuate gut inflammation and fibrosis [25]. Although direct evidence about the innate immunity gene variants in liver fibrosis is lacking, several studies have shown that NOD2 is associated with the increased mortality in nonalcoholic liver transplant patients [30] or the increased risk of spontaneous bacterial peritonitis in patients with liver cirrhosis [31, 32].

On a molecular level, fibroblasts and stellate cells in the intestine and liver express multiple pattern recognition receptors such as Toll-like receptors (TLRs) and Nod-like receptors [33–36] that can activate them towards a profibrogenic phenotype. Most information about innate immune signaling and fibrosis is available in the liver. Translocation of bacteria or their products to the liver, lipopolysaccharide in the portal vein, or activation of TLR4 promote liver fibrogenesis [34, 37, 38]. Lipopolysaccharide also downregulates the TGF-β1 decoy receptor BAMBI and hence indirectly contributes to fibrogenesis increasing HSC susceptibility to TGF-β1 [34]. A different PAMP, the TLR5 ligand flagellin (present in all flagellated bacteria), induces a proinflammatory and profibrotic phenotype in primary human intestinal fibroblasts [36], as well as promotes fibroblast proliferation [39]. Colonic EMT occurs after ligation of the TLR4 variant D299G [40]. This not only suggests a different location of action of PAMPs on fibrogenesis in the liver and intestine but also indicates that different microbial peptides may be active in the liver and intestine.

**Monocytes/Macrophages**

Hepatic macrophages play divergent roles in both the progression and regression of liver fibrosis, depending on the stage of fibrosis development [41]. Macrophage plasticity and polarization appear to be critical in this process. In early liver fibrosis, macrophages can act in a profibrogenic fashion through the production of profibrotic me-
diators, such as TGF-β, leading to HSC activation. In contrast, macrophages have been implicated in liver fibrosis resolution as well, possibly through the production of matrix metalloproteinases (MMPs), induction of apoptosis in HSCs, elimination of apoptotic cells, or secretion of IL-10 [41–47]. Hepatic recruitment of macrophages also plays a pivotal role in the fibrosis progression of experimental steatohepatitis [48, 49].

In contrast to the abundant data from the liver, information is lacking on an effect of macrophages and macrophage subsets on the initiation or resolution of intestinal fibrosis.

Cytokines

TGF-β and its intracellular mediators, Smad proteins, are the most potent signal to promote collagen production in the liver and intestine. Several factors increasing the expression of TGF-β or its activation, including MMPs, plasmin, plasminogen activators, αvβ6 integrin, and thrombospondins, have been reported [50].

TGF-β exerts a wide variety of functions in liver and intestinal fibrosis. It not only promotes collagen production, but also activates HSCs and myofibroblasts and suppresses MMP expression (fig. 2). During the past two decades, a large number of studies have been focusing on the molecular mechanisms responsible for the TGF-β-stimulated type I collagen gene transcription and its pathological roles during the fibrotic process [50, 51]. Our group has succeeded in suppressing liver fibrosis by promoting the nuclear translocation of YB-1 that is a downstream effector of interferon-γ [52]. Among many factors that interact with the TGF-β/Smad signal, bone morphogenetic protein (BMP) 7 [53] and hepatocyte growth factor [54] have been shown to suppress liver fibrosis by counteracting TGF-β action on the type I collagen gene.

TGF-β also drives intestinal stricture formation in IBD [55]. Levels of TGF-β1 are increased in the intestinal mucosa of ulcerative colitis and CD patients [56]. TGF-β1 is also elevated in models of chronic experimental colitis and accompanies increased collagen production [57]. In agreement with these observations, our recent study has shown that a small compound interfering with the TGF-β signal suppresses experimental intestinal fibrosis as it did for murine liver fibrosis [52]. Surprisingly, TGF-β1 has not been shown to have a definitive role in promoting the numerical expansion of intestinal fibroblasts in vitro, despite doing so in other tissues and organs [59]. In a mouse model of colonic TGF-β1 overexpression achieved via adenoviral transfer, animals developed a rapid accumulation of fibroblasts and thickening of the bowel wall [60]. These effects could be a direct effect of TGF-β1 itself or, alternatively, be an indirect effect of this growth factor by inducing the synthesis of connective tissue growth factor, platelet-derived growth factor, and insulin-like growth factor I. The presence of an indirect

![Fig. 2. TGF-β/Smad signal and its antagonists in organ fibrosis. TGF-β and its intracellular mediators, the Smad proteins, are the most potent accelerators of collagen production in the liver and intestine. TGF-β not only promotes collagen production but also activates HSCs and myofibroblasts and suppresses MMP expression, leading to the progression of liver and intestinal fibrosis. Among the Smad/BMP superfamily members, Smad 3 plays a pivotal role in stimulating collagen gene transcription, while Smad 7 inhibits the TGF-β signal by suppressing the phosphorylation and the subsequent nuclear translocation of Smad 3. BMP7 and hepatocyte growth factor (HGF) have been reported to suppress organ fibrosis by counteracting TGF-β action on the type I collagen gene.]
effect is supported by gene expression studies that have revealed an upregulation of TGF-β1 and connective tissue growth factor transcript levels in strictured CD tissue [61].

While the scientific rationale for TGF-β1 in fibrosis is strong, it needs to be considered that it is not only crucially involved in fibrogenesis but also functions as a key regulator of cellular differentiation, proliferation, transformation, tumor suppression, as well as immunoregulation, and its actions may be context dependent. TGF-β1-deficient mice develop severe multiorgan inflammation [62, 63], and targeted deletion of Smad 2 and Smad 4 is associated with early death in mice [64, 65], indicating the need for caution in using anti-TGF strategies as an anti-fibrotic therapy. Interestingly, a recent clinical study has reported that a Smad 7 antisense oligonucleotide (enhancing the TGF-β signal) is effective as anti-inflammatory treatment of human CD without promoting stricture formation in the short term [66].

A change in the CD4+ T-cell profile towards Th2 predominance, as indicated through elevations in IL-4, IL-5, and IL-13, has been reported during the course of fibrosis in various organs [67]. IL-13 mediates, through binding to its IL-13Ra, an increased production of TGF-β [68]. In the liver, Th2-associated alternative Kupffer cell activation promotes liver fibrosis [69]. Blockade of IL-13 significantly suppressed experimental liver fibrosis by repressing TGF-β production [70].

In the intestine, an increased amount of IL-13 transcripts was detected in fibrotic intestinal samples from CD patients compared to noninflamed areas of the same patients [71]. Inhibition of IL-13 signaling by the administration of a small interfering RNA targeting the IL-13-α2 receptor attenuated inflammation-associated murine intestinal fibrosis [72]. To the contrary, a study showed no difference in IL-13 production in the mucosal explants and lamina propria mononuclear cells between patients with strictureing CD and control subjects [73].

In experimental studies, IL-17A was involved in the pathogenesis of liver fibrosis by activating both Kupffer cells and HSCs [74, 75]. Likewise, IL-17 expression levels were elevated in experimental intestinal fibrotic tissues [72, 76]. In CD patients, IL-17A but not IL-17E was overexpressed in tissue samples from intestinal strictures as compared to samples from nonstrictured areas in the same patients and compared to healthy colonic samples from control subjects [77]. While targeting of IL-12/IL-23 by anti-IL-23 p19 monoclonal antibody [78] ameliorated acute and chronic murine colitis, administration of the human anti-IL-17A antibody secukinumab failed to induce remission in CD patients [79]. Further studies are necessary prior to any clinical application of an anti-IL-17-based strategy for the treatment of intestinal fibrosis.

Matrix Turnover

The imbalance between the production and degradation of ECM is observed commonly in various fibrotic diseases and represents a promising target for antifibrotic approaches [5]. Degradation of the ECM components is regulated by the enzymatic activity of MMPs [80], which is affected by the expression of tissue inhibitors of metalloproteinase (TIMPs). One mechanism, by which TGF-β1 leads to fibrosis, is the creation of an imbalance of TIMP-1/MMP expression in favor of TIMP-1 [81]. TIMP-1 also exerts an antiapoptotic effect on HSCs [82]. Transgenic overexpression of TIMP-1 in the murine liver attenuates the spontaneous recovery from carbon tetrachloride-induced liver fibrosis by decreasing the active form of MMP-2 [83]. Along the same line, the use of MMP-9 mutant proteins as TIMP-1 scavengers reduced fibrosis by enhancing matrix resorption [84].

In intestinal strictures, TIMP-1 expression is increased in the mucosa overlying fibrotic areas compared to those in the nonfibrotic area, while MMP-3 and MMP-12 are downregulated in CD [55]. On the other hand, it has been reported that colonic tissue expression of MMP-1, MMP-2, MMP-3, and MMP-9 is significantly increased in samples from the inflamed mucosa as compared to noninflamed mucosal samples in IBD [85]. Serum MMP-9 levels are increased and correlate with disease activity in pediatric CD patients [86]. Since MMP-2 and MMP-9 also possess proinflammatory potential, their use as anti-fibrotic reagents in liver and intestinal fibrosis may be hampered.

Future Perspectives

The ultimate treatment goal for fibrotic complications would be to stop or even reverse the fibrotic process. Great attention has been devoted to the phenomenon of reversibility of fibrosis in the field of hepatology [87]. Liver fibrosis, especially cirrhosis, was originally considered progressive and irreversible. However, a number of clinical studies have shown that it can be reversed if the causative agents are adequately removed or the patients are effectively treated [88–90]. With the advent of novel antiviral compounds that effectively inhibit HBV prolifera-
tion [91], or even completely eradicate HCV [92], the impact of ‘trigger elimination’ on the reversibility of fibrosis/cirrhosis can be easily studied in human patients. Indeed, there are several studies demonstrating that effective virus suppression in HBV-positive patients resulted in cirrhosis regression as proved by repetitive liver biopsies. Interestingly, this effect was independent from the antiviral agent used [93–95]. The same holds true for HCV infection, where studies report cirrhosis reversibility rates of up to 70–90% upon treatment with modern direct-acting agents [90, 96]. Consistent with such clinical evidence for the reversibility of liver fibrosis, we have previously reported that bone marrow-derived cells migrating into fibrotic liver contribute to the regression of experimental liver fibrosis by expressing MMP-13 and MMP-9 [97]. The concept of reversibility of intestinal fibrosis is in concordance with various observations from other organs such as the skin, kidney, lung, or heart [98–103].

In contrast to virus-induced liver fibrosis, CD-associated fibrostenotic strictures are likely a result of ongoing intestinal inflammation, which cannot be attributed to a single pathogenic factor. When fibrostenotic strictures are already present, they are considered permanent and nonreversible. This notion, however, has to be challenged. Recently, a large combined analysis assessing the efficacy of strictureplasty in CD found an overall symptomatic recurrence rate of 39% for jejunoileal strictures and 36% for ileocolonic strictures. Strikingly, only 3 or 20%, respectively, of CD patients’ strictures were found at the previous site of strictureplasty [104]. This finding indicates that surgical intervention has the potential to stop progressing or even to reverse intestinal fibrosis [105]. Serial ultrasound examinations in patients after strictureplasty found a reduced thickness of the intestinal wall [106] also suggesting that intestinal fibrosis is not only a one-way street.

As long as no single pathogenic factor for CD has been identified, potential antifibrotic treatment approaches for structuring CD should focus on receptor-ligand interactions, pathways of fibrosis regression (e.g. TGF-β), or stimulate the resolution of fibrosis (e.g. TIMP antagonists). As discussed above, multiple distinct signaling cascades and mechanisms have been identified to contribute to fibrosis in the intestine and liver. Therefore, future therapeutic approaches should not only address a single but multiple targets in the form of a combined approach. The complexity of their interactions is high. It is obvious that quality, quantity, and timing of profibrotic mediators deserve thoughtfull attention. Targeting individual cell types over mediators could be a promising approach. Activating subpopulations of macrophages to increase MMP expression and the induction of myofibroblast apoptosis may be suitable and may serve as an antifibrotic therapeutic armamentarium [107]. Another approach could be the stimulation of MMP production, which has been successfully performed in animal studies [5, 108, 109].

We have summarized the current knowledge on mechanisms of fibrosis which are shared by intestinal and liver fibrosis as well as their unique features. A collaborative effort in the field of fibrosis, covering multiple organ systems, will have the highest chance of leading to the development of a successful antifibrotic intervention.

Disclosure Statement

F.R. is on the advisory board of AbbVie. The other authors have no conflicts of interest to declare.

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