



Review Article

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Updates in Etiological Mechanism and Medical Treatment of Hepatic Fibrosis

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Abstract

Liver fibrosis is a pathological process triggered by chronic liver injury, characterized by abnormal accumulation of Extracellular Matrix (ECM), which can ultimately lead to cirrhosis and associated complications. During liver injury, Hepatic Stellate Cells (HSCs) become activated and trans-differentiate into myofibroblasts, resulting in an imbalance between ECM deposition and clearance. Without timely intervention, liver fibrosis may progress to cirrhosis, hepatocellular carcinoma, liver failure, or other life-threatening liver diseases. This review summarizes the primary etiological factors contributing to liver fibrosis and discusses the underlying pathogenic mechanisms, with a particular focus on HSC activation, inflammatory responses, and ECM remodelling. Additionally, the review provides an overview of diagnostic approaches for liver fibrosis, including histopathological assessments and non-invasive methods, such as serum biomarkers and advancements in imaging techniques. In terms of treatment and prevention, we highlight recent progress in antifibrotic strategies, including the potential of small molecule drugs and traditional Chinese medicine, and discuss the role of multi-omics research in advancing personalized and precision medicine. The review concludes by addressing key challenges and future directions in liver fibrosis research, particularly regarding the potential applications of personalized and precision therapies, thereby providing critical insights for future research and clinical application.

Keywords: Liver fibrosis, Pathogeny of Hepatic Fibrosis, Pathogenesis of Hepatic Fibrosis, Extracellular Matrix, Antifibrotic Therapy, Omics

Introduction

Hepatic Fibrosis is a pathological component of various chronic liver diseases. It is a repair response to chronic injury and inflammatory wounds, an imbalance between extracellular matrix deposition and clearance, leading to excessive ECM deposition and aggregation [1]. ECM is a non-cellular three-dimensional macromolecular network composed of collagens,

elastin, fibronectin, laminins and several other glycoproteins, which regulate diverse cellular functions, such as survival, growth, migration, and differentiation, and are vital for maintaining normal homeostasis [2]. Early Hepatic Fibrosis has a repairing effect on acute liver damage, but late Hepatic Fibrosis can lead to portal hypertension, liver failure, cirrhosis, and even liver cancer [3]. In the pathogenesis of hepatic fibrosis, the activation of HSCs is



a significant step in the process of liver fibrosis. When persistent liver damage occurs, HSCs are activated and continue to increase, transforming into myofibroblasts. These cells have been identified as the primary source of collagen production that damages the liver, leading to further aggravation of hepatic fibrosis [4]. Liver fibrosis, characterized by the abnormal accumulation of fibrillar collagen and scar matrix, is a result of dysfunctional wound repair in the presence of chronic liver damage [5]. The global incidence of liver fibrosis is 4.5-9% [6]. Cirrhosis, the end stage of progressive liver fibrosis, is the leading cause of liver-related deaths worldwide. In 2017, deaths due to cirrhosis accounted for 2.4% of total deaths globally and it was the 11th most common cause of death. Although fibrosis is a dynamic disease, it can be reversed if the underlying liver injury is removed or effectively treated [7]. However, advanced cirrhosis is generally irreversible, and there is currently no effective treatment other than hepatic transplantation. Hence, identifying effective and safe treatments for liver fibrosis is of utmost importance. The common causes of hepatic fibrosis are excessive drinking, viral hepatitis, and non-alcoholic fatty liver disease [8], autoimmune and hereditary diseases [9]. Hepatic fibrosis is partly a highly dynamic process [10]. The continuous process of hepatic fibrosis is the late stage of chronic liver disease, which can lead to ascites, jaundice, massive bleeding due to rupture of oesophageal and gastric varices, hepatic encephalopathy, liver cancer, and liver failure [11]. Some studies have shown that the clinical trials of antiviral therapy to block the progress of hepatic fibrosis provide key information for the treatment of hepatic fibrosis, so it is particularly important to actively treat patients early.

Pathogeny of Hepatic Fibrosis

Chronic Viral Hepatitis

Viral hepatitis is an acute inflammation of the liver caused by two sources of infection. It may be a virus named Virus A, which is the cause of infectious or epidemic hepatitis, or Virus B, which is the cause of serum hepatitis or homologous sero jaundice. The number of people infected with the hepatitis virus in the world annually is 10 times more than that of AIDS, and the number of deaths due to contracted hepatitis exceeds 1 million every year [12]. Hepatitis A and B viruses can cause acute liver disease, often accompanied by jaundice and systemic symptoms. Hepatitis B and C are at risk of developing cirrhosis and liver cancer. The Hepatitis E virus is easy to cause epidemic diseases in developing countries. Still, it is rare in the United States, and there is no evidence that it can cause acute or chronic liver injury. In HBV and HCV-related liver fibrosis diseases, the general pattern of fibrosis evolution is called post necrosis or bridging fibrosis, which is characterized by the increased deposition of ECM components in the form of portal vein central (vein) fibrosis diaphragm driving or portal vein central bridging necrosis [13]. Primary factors considered in the relationship between HBV or HCV and fibrosis: (1) The injury of liver parenchyma and the death of liver cells should be attributed to the immune response of trying to eliminate HBV and HCV virus. However, the overall response of virus-specific CD4 and CD8 T lymphocytes becomes inefficient over

time, and they cannot eliminate HBV or HCV in the liver, leading to chronic infection. (2) The circulation of low-level cell damage and persistent inflammatory reaction, the secondary recruitment of non-antigen specific monocytes (NK cells), which is crucial to the liver parenchyma damage, inflammatory reaction, and fibrosis progress [14]. (3) Direct action by specific HCV proteins such as core and NS3/NS5 proteins. Although HCV does not infect HSCs, these proteins can stimulate activated human HSCs in a ROS and redox-dependent manner, leading to upregulating pro-fibrosis and pro-inflammatory responses [15]. (4) In addition, HCV protein can induce increased ROS production in hepatocytes, which may lead to oxidative stress-mediated hepatocyte injury and activation/continuation of HSC phenotypic response [16]. Hepatic fibrosis is the initial central stage of viral hepatitis, developing into cirrhosis and liver cancer. Therefore, the focus of diagnosis and treatment of viral hepatitis is to eliminate the virus and prevent it from developing into cirrhosis or liver cancer.

Alcoholic Hepatitis

The normal metabolism of the liver of a person who takes a large amount of alcohol for a long time will be disrupted, which may lead to fatty liver (also known as steatosis), which is a disease that contains large vesicular triglyceride drops in liver cells. Although alcoholic fatty liver can be solved through abstinence, people who continue to drink are prone to liver fibrosis and cirrhosis. Although alcoholic fatty liver can be solved through abstinence, people who continue to drink are prone to liver fibrosis and cirrhosis. Compared with the general population, patients with alcoholic liver disease confirmed by biopsy have a nearly fivefold increased risk of death [17]. In addition, during the 2019 coronavirus disease pandemic, the association between heavy drinking and ALD (autoimmune liver disease) increased, and the consequences of alcohol are expected to continue [18]. It is worth noting that in patients with alcoholic liver disease, macrophages are enriched in the portal vein and can promote ethanol-induced inflammation *in vivo*, especially in severe alcoholic hepatitis, in which the increase of intestinal permeability and portal vein endotoxin level contributes to the accumulation and activation of macrophages in Ly-6Chi mice and the release of TNF, ROS, etc [19]. In addition, the metabolic pathway of alcohol produces reactive oxygen species, which are effective inducers of lipid peroxidation, leading to necrosis or apoptosis of hepatocytes. Notably, three liver enzymes are involved in the ethanol metabolism to acetaldehyde, which will produce a large amount of NADH during ethanol metabolism. NADH will inhibit the conversion of lactic acid into pyruvate in the process of gluconeogenesis, leading to lactic acid cumulative acidosis and hypoglycemia caused by the inhibition of gluconeogenesis. Acetaldehyde can participate in the modification of many proteins and enzymes in liver tissue. Long time and large amounts of ethanol intake will lead to the accumulation of acetaldehyde in liver tissue, which in the next step will cause functional problems of liver cells and cause damage and death of liver cells. Long-term damage and death of hepatocytes will lead to liver fibrosis and cirrhosis. It can be seen that it is indispensable

to control alcohol intake, especially for patients with primary liver diseases such as obesity and viral hepatitis.

Non-alcoholic Fatty Liver Disease

Nonalcoholic Fatty Liver Disease (NAFLD) is liver cell damage, inflammation, and fibrosis caused by liver lipid accumulation, which leads to more severe liver diseases. The global prevalence of NAFLD is 25.24%, with the highest prevalence in the Middle East and South America and the lowest in Africa [20]. Nonalcoholic Fatty Liver Disease (NAFLD) is one of the leading causes of liver cirrhosis in the world [21]. NAFLD is expected to increase exponentially in the next few years, significantly increasing the health system and economic burden [22]. In recent years, there is increasing evidence that NAFLD is a multisystem metabolic disease associated with hyperinsulinemia and genetic susceptibility, with hepatocyte triglyceride accumulation as a major pathologic change. In addition to liver related complications, NAFLD also increases the risk type 2 diabetes mellitus (T2DM), cardiovascular disease and chronic kidney disease. Conversely, obesity, T2DM, lifestyle changes, and drug-resistant genetic alterations also increase NAFLD morbidity and mortality.

Cholestatic Hepatitis

Cholestasis is the obstruction of bile secretion or excretion caused by various reasons, which leads to the inability of bile to flow normally into the duodenum and then back into the blood circulation. Bile transport dysfunction caused by injury of hepatocytes and bile duct cells induces proliferation of bile duct and peritubular myofibroblasts, eventually leading to liver fibrosis. Cholestatic liver disease, such as Primary Biliary Cholangitis (PBC) or Primary Sclerosing Cholangitis (PSC), is a chronic progressive disease, often leading to cirrhosis and subsequent complications. Congenital cholestasis syndrome, whose potential phospholipid invertase (ATP8B1) or bile salt outlet pump (BSEP; ABCB11) dysfunction may also rapidly progress to cirrhosis and often requires liver transplantation [23]. Although the pathogenesis of cholestasis varies, the systemic and hepatic accumulation of hydrophobic bile salts are standard pathogenic features [24]. For example, in PBC, it has been found that late-stage systemic bile salt levels increase by up to 20 times. This widely accepted hypothesis was proposed in the 1970s, and to this day, the accumulation of hydrophobic bile salts is still considered a driving force for fibrosis in cholestatic liver disease [25, 26].

Autoimmune Hepatitis

Autoimmune liver disease includes Autoimmune Hepatitis (AIH), characterized by necrotizing inflammation, primary biliary cirrhosis (PBC), and primary sclerosing cholangitis (PSC), all of which are characterized by progressive cholestasis [27]. Autoimmune hepatitis is an acute or chronic liver parenchymal disease characterized by a loss of tolerance [28] to liver cell-specific autoantigens, which is often caused by environmental and genetic

factors [29]. If left untreated, autoimmune hepatitis can usually lead to liver fibrosis, even cirrhosis and liver cancer. Previous studies on patients with autoimmune hepatitis have shown that approximately 30% of patients are diagnosed with cirrhosis. Immunomodulatory therapy can significantly improve the long-term prognosis of patients with autoimmune hepatitis [30]. Monoclonal antibodies targeting key cytokine pathways (monoclonal antibodies targeting CD3[31], CD20[32], and tumour necrosis factor-alpha) can now be used for molecular and cellular interventions. Recombinant molecules, such as cytotoxic T lymphocyte antigen-4 fused with immunoglobulin, may inhibit lymphocyte activation and cellular interventions, such as adoptive transfer of regulatory T cells and sensitization of disease-specific glycolipids to natural killer T cells [33], which may stimulate or inhibit counter-regulatory immune mechanisms. These interventions are now expected to improve treatment outcomes by targeting key areas of immune-mediated pathogenic responses precisely, potentially achieving faster, more complete, and lasting outcomes.

Schistosoma Infected Liver Disease

Liver fibrosis caused by schistosomiasis is essentially a severe parasitic infection caused by the deposition of schistosome eggs in the liver. Due to the host body's long-term immune response to schistosome egg antigen, hepatic stellate cells are abnormally activated and transformed into MFB, resulting in the deposition of ECM and liver fibrosis after a large number of value-added [34]. Epidemiological studies have shown that schistosomiasis is still the second most common tropical disease. Two hundred forty-nine million people in 73 countries worldwide suffer from different types of schistosomiasis, and nearly 800 million people are threatened by this infection [35]. Although various etiological factors may lead to chronic liver injury leading to liver fibrosis and even cirrhosis, such as drug abuse, alcoholism, and viral infection, schistosomiasis infection is one of the most common causes of liver fibrosis in some countries [36]. Some reports show that effective schistosome treatment could not prevent the development of egg granuloma inflammation and liver fibrosis [37, 38], possibly due to continuous pathological processes such as chronic inflammation. So far, the exact mechanism that mediates the constant activation of inflammation around egg granuloma in the liver during schistosomiasis infection remains unclear.

Pathogenesis of Hepatic Fibrosis

Cells and Hepatic Fibrosis

The liver is composed of parenchymal cells and non-parenchymal cells. Parenchymal cells refer to Hepatocytes (HC), while nonparenchymal cells include Sinusoidal Endothelial Cells (SEC), Kupffer Cells (KC), Hepatic Stellate Cells (HSC), and pit cells. Studies have shown that these cells are involved in developing liver fibrosis in different ways.

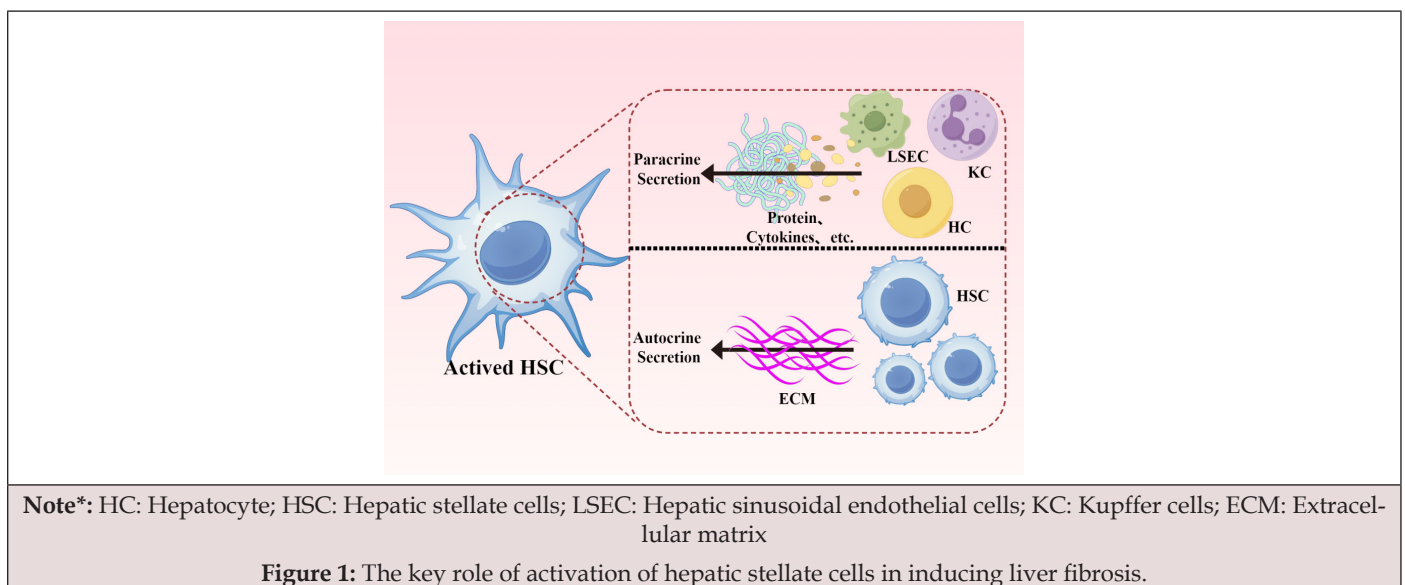
Hepatocyte

There are a lot of microvilli on the surface of hepatocytes. During the formation of liver fibrosis, hepatocytes synthesize collagen type I, III, IV, V, but the amount of collagen synthesis is less than that of HSC. The leading role of HC is to activate HSC. At present, it is believed that the activation mechanism of HC is that the normal hepatocyte membrane has a contact inhibition effect on the proliferation of HSC. In contrast, virus infection, steatosis, or alcohol-induced hepatocyte apoptosis and damage of hepatocyte membrane will lead to the loss of contact inhibition effect on HSC and activate HSC. Apoptotic hepatocytes have been shown to promote the secretion of pro-inflammatory and pro-fibrosis cytokines by macrophages and directly promote HSC activation [39,41]. Recent studies have also demonstrated that inflammatory stressed hepatocyte cytokine (such as IL-33) can promote fibrosis [42].

Hepatic Stellate Cells

HSCs are located in the interspace of the dice around the sinuses and the intercellular fossa of hepatocytes. The cytoplasm is rich in vitamin A-like lipid droplets. HSCs are the central source cells of the Extracellular Matrix (ECM) during liver fibrosis, and their activation is the vital and central event of liver fibrosis [40]. In the fibrotic liver, resting HSC transdifferentiate into proliferative, migratory, and contractile myofibroblasts, which exhibit fibrosis transcription and secretion characteristics (called "cell activation") and secrete ECM molecules, which accumulate in hepatocytes and form scar tissue. Storage of (retinoid in cytoplasmic droplets is a unique feature of fixed HSC, which is gradually lost during transdifferentiation.

However, its causal relationship with HSC activation remains uncertain [39,44]. Endothelin-1 (ET-1) is a potent vasoconstrictor secreted by activated HSC cells that promotes cell proliferation, fibrosis and contraction and may be associated with cirrhotic portal hypertension [43]. Activation includes two main stages: the initiation stage and the sustained stage. The initiation stage refers to the changes in early gene expression and the phenotypic changes in cells produced by cytokines and other stimulating factors. The initial changes in HSC may result from paracrine stimulation from nearby cells, including SEC, HC, KC, etc. During the sustained phase, these stimulating factors maintain the activation phenotype of HSC, resulting in the formation of liver fibrosis. Both autocrine and paracrine pathways regulate the activation of HSC at this stage. As the primary executor of fibrosis, HSC can receive a wide range of signals from injured liver cells and disturbed liver microenvironment. HSC interacts with liver cells, macrophages, lymphocytes, and endothelial cells to promote fibre formation. In addition, activated HSC interacts with natural killer cells, leading to HSC death and termination of fibrosis response. HSC derived from mice or humans exhibit high reactivity to pro-inflammatory cytokines and LPS, activating pro-inflammatory signalling pathways such as nuclear factor-kB and AP-1 and producing chemokines and cytokines (Figure 1). However, compared to the crucial role of macrophages and other white blood cell populations, HSC may have a relatively small contribution to overall liver inflammation. HSC may play a role mainly as receptors for inflammatory signals rather than promoting the overall inflammatory state of the liver [45].

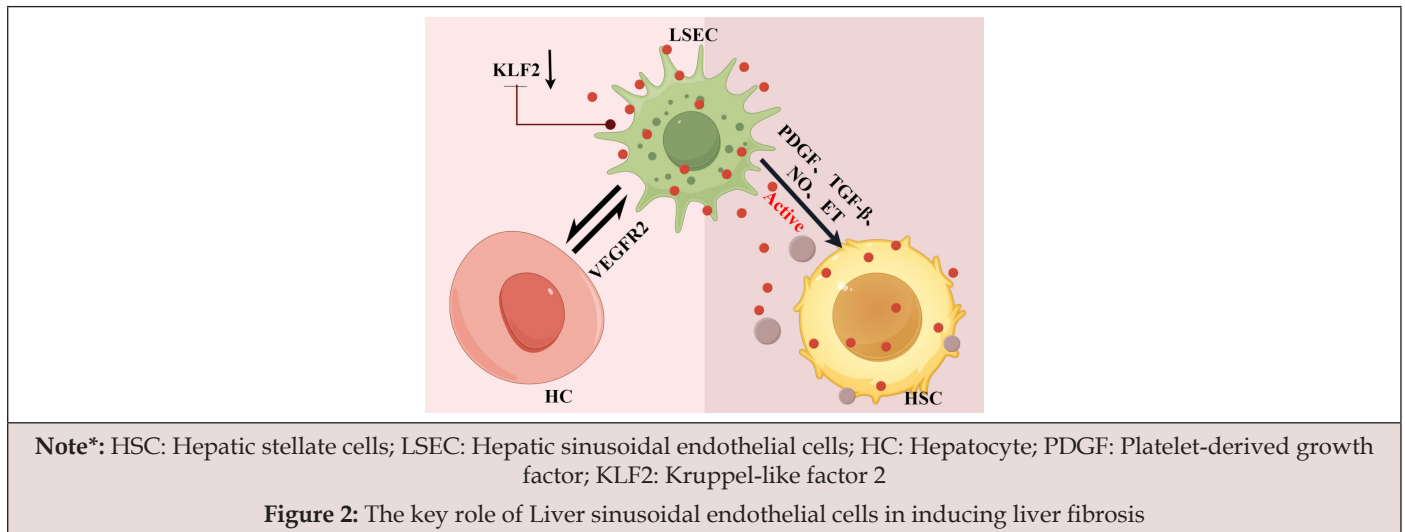


Liver Sinusoidal Endothelial Cell

Hepatic Sinusoidal Endothelial Cells (LSEC) are damaged, swollen, and even necrotic, narrowing the hepatic sinuses and leading to the contraction of LSEC and HSC. Expressing adhesion

molecules makes inflammatory cells gather in the hepatic sinuses and makes LSEC phenotypic transfer to vascular endothelial cells. Liver fibrosis can be caused by the crosstalk between LSEC and hepatocytes [46]. Hepatocytes and LSEC interact through

the VEGFR2 pathway. CD147 is simultaneously expressed in hepatocytes and LSEC. It is a transmembrane glycoprotein associated with liver fibrosis [47] and plays a regulatory role in the information exchange between hepatocytes and LSEC (Figure 2).



Interestingly, the anti-CD147 antibody inhibits angiogenesis through the VEGF-a/vegfr2 pathway, thereby improving the progress of liver fibrosis. However, due to the inability to protect adjacent cells, inhibition of hepatocyte apoptosis after injury is not beneficial to preventing liver fibrosis [48]. In addition, the interaction between hepatocytes and LSEC promotes the capillary effect of LSEC, reduces portal vein angiogenesis, and promotes the progress of liver fibrosis. The above effect is achieved by combining leukocyte-derived chemokine 2 (LECT2) expressed by hepatocytes and Tie1 produced by LSEC [49]. Liver fibrosis can also be caused by crosstalk between LSEC and HSC. In continuous liver injury, LSEC is capitalized and has the phenotypes of promoting vasoconstriction, inflammation, angiogenesis, and fibrosis [50]. Capillary LSEC does not promote HSC quiescence but activates HSC by secreting Platelet-Derived Growth Factor (PDGF), TGF- β , and reducing Kruppel-Like Factor 2 (KLF2), a transcription factor that is a protective molecule of liver endothelial vessels. In addition, it has been proved that LSEC-derived fibronectin can also affect the phenotype of HSC and promote its activation [51]. Subsequently, HSC began to increase, contract, and deposit many collagen fibres and extracellular matrix molecules in the liver parenchyma, leading to organ sclerosis and interfering with all cell functions [52].

Studies have found that Nitric Oxide (no) and Endothelin (ET) secreted by LSEC can change HSC and induce liver fibrosis. LSEC can mediate acute and chronic liver inflammation and activate cytokines that promote ECM matrix synthesis, such as TGF- β . At the same time, it causes lipid metabolism disorder in vivo, promotes HSC to transform into fibroblasts, and synthesizes and secretes a variety of extracellular matrix. The phenotype of hepatic sinusoidal endothelial cells changes during liver fibrosis. The two main characteristics of hepatic sinusoidal capillarization are the loss of fenestration of hepatic sinusoidal endothelial cells and the formation of a continuous basement membrane under the endothelium [53]. These changes will interfere with the transport

of substances from the hepatic sinuses to hepatocytes and the blood circulation of the liver and become a precursor of liver fibrosis. The loss of fenestration of LSEC is characterized by the reduction of the number and diameter of LSEC, which can occur in the early stage of liver fibrosis. The loss of fenestration of LSEC is related to the cells themselves and the changes in the extracellular matrix. Standard ECM plays an essential role in maintaining the morphology of LSEC. The deposition of ECM during liver fibrosis may lead to the change of LSEC fenestra. Early studies have shown that the number of fenestra decreases with interstitial collagen deposition during liver fibrosis and is parallel to the deposition of ECM in the Disse cavity. It emphasizes the importance of the phenotypic transformation of hepatic sinusoidal endothelial cells, provides a reference index for the clinical detection of hepatic sinusoidal capillarization, and provides a new idea for preventing and diagnosing liver fibrosis.

Kupffer Cell

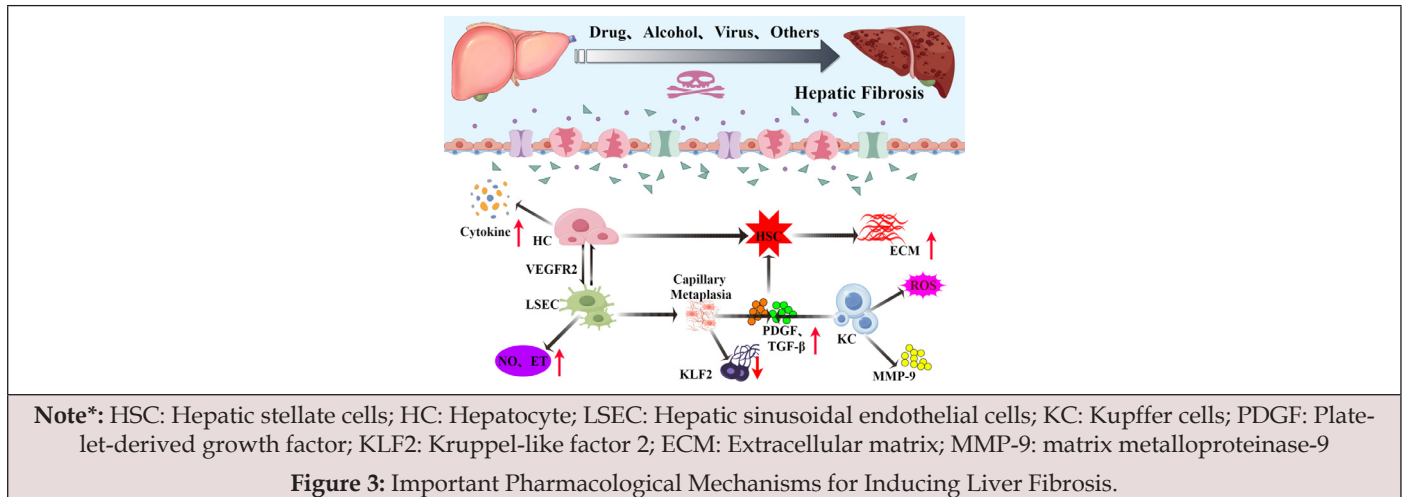
Kupffer Cells (KC) are macrophages located in the hepatic sinuses. Their processes can extend into the Disse space under the sinusoidal endothelial cells, sec pores, and direct contact with HC and HSC. With the injury of liver parenchymal cells, the original Kupffer cells in the liver and the monocytes in the blood began to infiltrate and increase into the liver tissue. A KC-conditioned medium can accelerate the activation of HSC in the early stage of culture, increase its proliferation, and promote its secretion of ECM. KC can stimulate the synthesis of extracellular matrix, cell proliferation, and activation of HSC by secreting cytokines, especially TGF- β . TGF- α secreted by KC can induce cell proliferation, while TGF- β secreted by KC stimulates HSC to secrete many ECM. In addition, KC can also affect HSC by secreting matrix metalloproteinase-9 (MMP-9, gelatinase B). MMP-9 can activate TGF- β , which stimulates HSC to secrete collagen. KC is also a significant source of reactive oxygen species intermediates (ROS) in the liver: ROS can promote HSC activation and collagen secretion, whether in or outside the

cell. Nitric oxide produced by KC can reduce the proliferation of HSC and inhibit the activation of ROS on HS [54].

Lacunar Cell

Lacunar cells are large granular lymphocytes with natural killer activity in the liver. Their morphological characteristics are

cell polarity and aniline blue granules in the cytoplasm. The role of lacuna cells in developing liver fibrosis is unclear. Lacuna cells are significantly increased in the rat model of viral hepatitis. It is speculated that lacuna cells indirectly affect the formation of liver fibrosis by interacting with other non-parenchymal cells (Figure 3).



Extracellular Matrix and Hepatic Fibrosis

The Extracellular Matrix (ECM) includes collagen, non-collagen glycoprotein, proteoglycan, and elastin, constituting the matrix of stromal cells and the basement membrane of epithelial cells and blood vessels. During liver fibrosis, the changes in ECM include quantitative changes (excessive collagen formation and deposition in the liver) and qualitative changes (local reconstruction or redistribution of ECM), resulting in the imbalance of ECM synthesis and degradation, excessive accumulation and deposition in the liver, especially the reduction of degradation in the later stage, which is the primary mechanism of liver fibrosis formation. It is believed that the cells that produce matrix in the liver are HC, HSC, SEC, KC, and pit cells, and HSC is the primary source of ECM. In the process of liver injury, HSC is activated and transformed into myofibroblast-like cells, accompanied by increased ECM synthesis and expression of α -SMA. Other ECM components synthesized by HSC include laminin, etc.

Collagen: Collagen protein accounts for 5% -20% of the total protein in the normal liver, but it can increase to about 50% in the case of liver fibrosis. There are several main types of collagen in the case of liver fibrosis, namely, interstitial collagen (type I, III, V collagen), stromal membrane collagen (type IV collagen), and short chain or microfilament collagen (type VI collagen), of which type I and III collagen are the main types. In the normal human liver, type I and III collagen account for about 80% of the total collagen in the liver. In contrast, in the case of liver fibrosis, type I and III collagen can account for more than 95% of the total collagen in the liver. Liver fibrosis was mainly type III in the early stage and type I in the late stage. In the early stage of liver fibrosis, interstitial

collagen was deposited in the disease space, which destroyed the functional basement membrane of type IV collagen. The disease space was capitalized, resulting in blood supply and nutrition supply disorders. In the normal rat liver, the collagen around the hepatic sinuses is composed of newly synthesized type IV collagen.

In contrast, the portal area mainly comprises mature types I and III, also essential components of fibrosis. In liver fibrosis, type I and III collagen is a necessary source of liver fibrous septum. The newly formed type I and III collagen form fibrous bundles and distribute along these cells in a dendritic manner to create a new fibrous septum. In addition, the collapse of reticular fibrous scaffold caused by cell degeneration and necrosis of mature type I and III collagen aggravates liver fibrosis. The role of type IV in liver fibrosis is to participate in the capillarization of hepatic sinusoid. Only discontinuous type IV collagen can be seen on the wall of normal hepatic sinusoid. During liver fibrosis, the type IV collagen on the wall of the hepatic sinusoid becomes continuous, and laminin begins to appear, which destroys the functional basement membrane of type IV collagen and forms capillarization. The capillarization of the hepatic sinusoid is one of the foundations for developing liver fibrosis to cirrhosis [55].

Glycoprotein: Non-collagen glycoproteins include Fibronectin (FN), Laminin (LN), and Endokine (EN). FN is a multifunctional macromolecular non-collagenous glycoprotein first found in the body through plasma and cell types. FN can be detected in all the extracellular spaces of the liver. In the early stage of liver fibrosis, plasma FN is suspected to increase, which occurs before the deposition of other extracellular matrix components. Still, this increase is not related to the severity of the disease and other

biochemical indicators. It is believed that FN first appears on the fibrous septum and hepatic sinusoid wall in the early stage of liver fibrosis and then is replaced by other interstitial components. The role of FN in liver fibrosis is mainly to play the role of matrix skeleton, conditioning, and cell movement matrix. FN through fibroblasts indicates that adhesion molecules (receptors) will transfer polypeptides into cells, which plays a pacemaker-like role in liver fibrosis. LN is the main glycoprotein in the process of liver fibrosis. Ln mainly exists in the basement membrane and is a unique basement membrane component. LN in the liver comes from HSC, HC, etc. LN and type IV collagen are distributed in the vascular wall, bile duct wall, and other parts. In the early stage of experimental liver fibrosis, the rough endoplasmic reticulum of HSC was dilated, and there was apparent Ln staining, suggesting that the cells synthesized a large amount of LN. In the late stage of liver cirrhosis, the intense positive staining on the sinusoidal wall was distributed with type IV collagen, indicating that both were involved in liver sinusoidal capillarization. Electron microscopy also confirmed a specific binding point between LN and type IV collagen. Thus, LN is involved in the formation of liver fibrosis. The role of LN in liver fibrosis is mainly to connect the macromolecular components of the matrix and jointly participate in the formation of the basement membrane and the capillarization of the hepatic sinusoid. Ln can crosslink with type IV collagen, non-collagen glycoprotein, heparin sulfate, and other matrix components and can crosslink itself, which plays a vital role in forming basement membrane.

Matrix Metalloproteinases and Their Inhibitors: ECM degradation is mainly mediated by Matrix Metalloproteinases (MMPs). Studies have shown that in the early stage of liver fibrosis, proenzyme activity changes little, and the rapid degradation system mainly degrades the increased collagen synthesis at this stage in cells. In the process of fibrosis development, the activity of MMP gradually increased, and in the late stage of liver fibrosis, the activity gradually decreased. Further studies showed that the activity of MMPs could be inhibited by Tissue Inhibitors of Metalloproteinases (TIMPs), which were specific inhibitors of MMPs. TIMPs were the corresponding side regulators of MMPs in the metabolic regulation of ECM and were synthesized and secreted in the same cells as MMPs. TIMPs combine with active MMPs to form a 1:1 chemical complex, which specifically inhibits the activity of MMPs and promotes cell proliferation. The main effect is to inhibit the activation of MMP zymogen and its degradation of EC [56].

Cytokines and Hepatic Fibrosis

Platelet-Derived Growth Factor

During the onset and progression of liver fibrosis, PDGF and its receptor levels were notably greater than normal liver tissue, with the activity of PDGF intensifying as liver fibers worsened. The expression level in HSC was significantly positively correlated with the degree of liver fibrosis. PDGF is the mitogen with the most potent effect on HSC among the known polypeptide growth factors.

Hepatocyte damage caused by various factors, such as viruses, chemicals, and mechanical injury, can cause cells, including Kupffer cells, to synthesize and release PDGF and activate the corresponding signal molecules in the cells through specific receptors on the HSC membrane. Under some transcription factors, it transduces signals into the nucleus. It promotes the transcription and expression of the corresponding target genes to activate HSC, increases change in the cytoskeleton distribution in HSC, and stimulates its transformation into myofibroblasts. HSCs acquire chemotaxis in the process of transforming to MFB, migrating, and gathering in the inflammatory damaged area; Activated HSCs can secrete PDGF and overexpress the corresponding receptors, which can synthesize and secrete a large number of extracellular matrix and promote the occurrence and development of liver fibrosis. PDGF can also up-regulate tissue metalloproteinase inhibitors and inhibit collagenase activity, reducing the degradation of secreted ECM.

Transforming Biochemical Factor Beta (TGF-β)

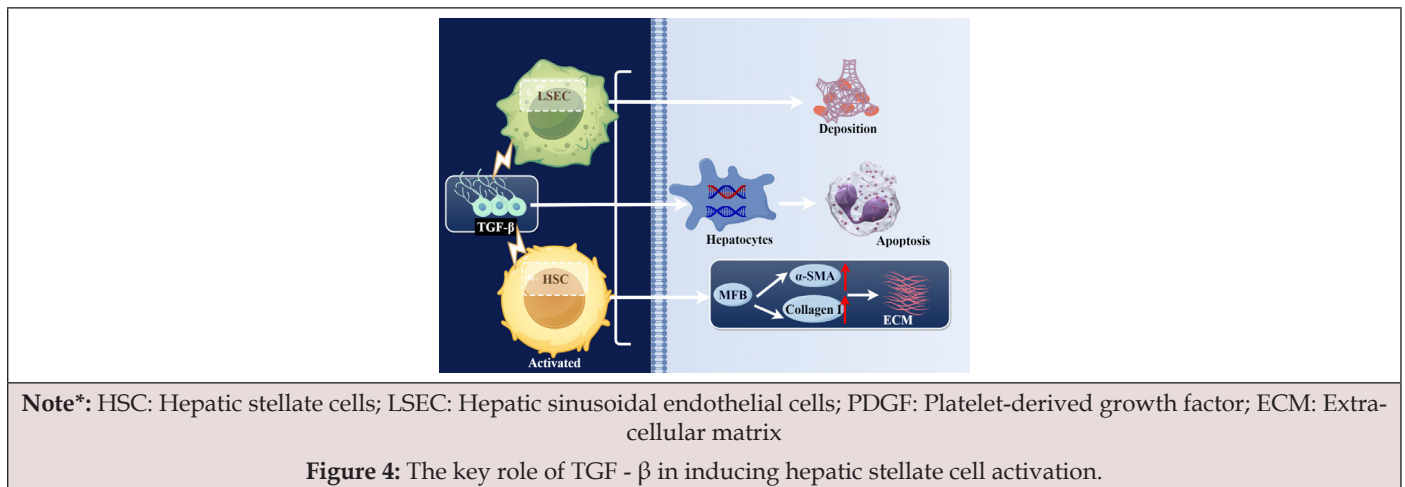
The primary producers of TGF-β in human tissue include platelets, monocyte macrophages, fibroblasts, smooth muscle cells, endothelial cells, and lymphocytes. In the liver, it is synthesized primarily by hepatic stellate cells/myofibroblast-like cells, Kupffer cells, and sinus endothelial cells. TGF-β is synthesized and secreted in an inactive form in all types of cells, and it must be activated and bound to receptors before it can show biological activity. TGF-β subfamily is composed of at least six structurally related molecules (TGF-β 1, β 2, β 3, β 4, β 5, β 6). TGF-β 1 plays a crucial role in the initiation and sustainable development of liver fibrosis. The following are: (1) The activated HSC can proliferate, swim, enhance contraction, and synthesize many growth factors, cytokines, and extracellular matrix. The activated HSC also synthesizes and secrete TGF-β 1. This positive feedback regulation of autocrine is essential to developing liver fibrosis. (2) Induction of matrix gene expression. (3) Inhibit matrix degradation so that collagen synthesis is more than degradation, leading to excessive deposition of collagen fibres, hepatic sinusoidal capillarization, and collagen, thus promoting the development of liver fibrosis. The collagenous environment of the stroma is not only related to the occurrence of liver fibrosis but also promotes the transformation of cells. (4) It can induce hepatocyte apoptosis by inhibiting the synthesis of hepatocyte DNA and indirectly inhibit hepatocyte division and proliferation by inhibiting the secretion of Hepatocyte Growth Factor (HGF) by stromal cells and inhibiting the promotion of GF on hepatocyte mitosis [57] (Figure 4).

Connective Tissue Growth Factor

The study confirmed that CTGF can induce proliferation and migration of HSC and upregulate the expression of Type I collagen and α-SMA in rats. Further studies showed that it promotes the proliferation of rat HSCs by activating the ERK1/2 signalling pathway. It has been found that the expression of CTGF mRNA increased in patients with liver disease, animal models of liver

cirrhosis, and cultured rat hepatic stellate cells, and the degree was positively correlated with the degree of liver fibrosis. TGF- β 1 can directly induce the expression of CTGF mRNA in hepatic fibrosis

model, and vascular endothelial growth factor (VEGF), PDGF\lipid peroxide, and acetaldehyde can also up-regulate the expression of CTGF mRNA in HS [58].



Insulin-like Growth Factor

Insulin-like Growth Factor (IGF) is a mitogen-like peptide in cells. There are three peptide hormones in the IGFs family, namely Insulin (INS), IGF- I , and IGF- II , which bind to IGFs binding proteins and regulate the role of IGFs through IGFBP. IGFs can promote cell proliferation, differentiation, and maturation, inhibit cell apoptosis, mediate most of the effects of growth hormone, promote growth and anabolism, reduce blood glucose, and regulate immunity. During the recovery period of liver fibrosis, activated HSCs are reduced mainly through apoptosis rather than phenotypic transformation. Cell growth depends on various growth factors. IGF and its receptors are the central regulators of inhibiting cell apoptosis. Experiments have corroborated that IGF-I can significantly inhibit the apoptosis of cycloheximide-induced HSC. In vitro experiments showed that PDGF could substantially enhance the secretion and release of IGF-1 and IGF-1 binding protein in rat hepatic stellate cells and strengthen the activity of IGF-1 by increasing the binding with IGF-1. IGF-1 and PDGF jointly stimulated the DNA synthesis in rat hepatic stellate cells and promoted their division and proliferation, suggesting that the interaction between IGF-1 and other cytokines, such as PDGF, contributes to the formation of liver fibrosis.

Tumor Necrosis Factor Alpha

Tumour necrosis factor- α (TNF- α) is mainly produced by monocyte macrophages, HSC, Kupffer cells, and so on, which have pro-inflammatory activities and cytotoxic effects. In the process of liver fibrosis, TNF- α plays an essential role in the proliferation and activation of HSC, the synthesis of ECM, and the release of matrix metalloproteinases and tissue inhibitors. Studies have found that TNF- α can enhance the proliferation of HSC, promote the transformation of HSC to MFB, and increase the effect of TGF- β on the synthesis of ECM. However, some scholars directly intervened

in primary cultured HSCs with TNF- α and found that TNF- α inhibited the proliferation and apoptosis of HSC and induced the "activation" of HSC. The effect of TNF- α on the synthesis of the HSC matrix occurs at the gene transcription stage. The results showed that TNF- α could induce apoptosis of HSC, which was related to its activation state. Its role in promoting HSC apoptosis may be linked to inhibiting the expression of Bcl-2, promoting the expression of Bax, and down-regulating the value of Bcl-2/Bax. In a word, the effect of TNF- α on HSC is multifaceted, and the mechanism of action is complex, which not only promotes liver fibrosis but also inhibits liver fibrosis. TNF- α and other cytokines such as TGF- β , PDGF, and IL-1 form a regulatory network vital in initiating and regulating liver fibrosis.

Angiotensin II

Angiotensin II (Ang II) is the primary bioactive substance secreted by the Renin-Angiotensin System (RAS). Ang II can exert its biological effects only by binding to specific receptors on tissue cell membranes, mainly angiotensin II type 1 receptor (AT1R). Active hepatic stellate cells have local RAS. The effects on activated hepatic stellate cells include: (1) Promote the activation and proliferation of hepatic stellate cells. Studies have reported that exogenous Ang II can induce a dose-dependent increase of intracellular Ca²⁺ concentration in cultured human hepatic stellate cells mainly through L-type calcium channels, and the increase of intracellular Ca²⁺ concentration is an essential condition for the activation and proliferation of hepatic stellate cells. (2) Promote the contraction of hepatic stellate cells. Studies have shown that Ang II can induce the contraction of hepatic stellate cells in a dose-dependent manner, which is closely related to the change of intracellular Ca²⁺ concentration. The contraction intensity was approximately equal to that of ET-1 and could be inhibited by losartan, suggesting that AT1R mediated this effect. (3) Promote

the synthesis of ECM by hepatic stellate cells. Studies have found that Ang II can promote the synthesis of collagen and the secretion of hyaluronic acid and laminin in hepatic stellate cells in a dose-dependent manner.

It is speculated that Ang II may activate a series of signal transduction pathways by binding to AT1R on the membrane of hepatic stellate cells and increasing the expression of TGF- β 1 and PDGF, thereby stimulating the synthesis of collagen in hepatic stellate cells. (4) Inhibit the degradation of ECM by hepatic stellate cells. It was found that Ang II could promote the production of TIMP-1 in activated rat hepatic stellate cells in a dose and time-dependent manner. Ang II binds to AT1R on the membrane of hepatic stellate cells and promotes TIMP-1 production by activating the protein kinase C (PKC) signalling pathway, which AT1R antagonists and PKC inhibitors can inhibit. (5) It is involved in the network regulation of cytokines by hepatic stellate cells. In cultured activated hepatic stellate cells, II can induce the expression of TGF- β 1 mRNA through AT1R, which can be completely inhibited by candesartan, an AT1R antagonist. (6) Inhibition of hepatic stellate cells. Studies showed that Ang-II upregulated Bcl-2 expression but not Bax expression, increased Bcl-2/Bax ratio and inhibited apoptosis of hepatic stellate cells.

Interleukin

Interleukin-1 (IL-1) comes from KC and sec. Its particular biological activity is to promote the proliferation of fibroblasts and HSCs, increase collagen synthesis, increase ECM production, and inhibit MMP synthesis. It is involved in the process of liver fibrosis in many links. Interleukin-6 (IL-6) is also mainly secreted by KC. Activated HSC cells and myofibroblasts are also the primary sources of IL-6, which can stimulate the proliferation of HSC, induce the production of a variety of acute phase proteins, and promote the deposition of ECM by promoting matrix degeneration or interacting with its adhesion receptors. Recent studies have shown that interleukin-10 (IL-10) may be an anti-hepatic fibrosis factor. Active HSC can negatively regulate liver fibrosis by autocrine IL-10, inhibiting type I collagen transcription and stimulating collagenase production.

Platelet Activating Factor

Platelet-Activating Factor (PAF) belongs to phospholipids, which can regulate inflammation, vasoconstriction, and other physiological functions. PAF mediates the interaction between cells and causes the change in cell phenotype. It is speculated that PAF may also be involved in the activation of HSC. Studies have found that KC and HSC can secrete PAF in the early stage of acute liver injury, and Ca²⁺ carriers, thrombin, and LPS can stimulate the secretion of PAF. Studies have shown that PAF is the primary inflammatory mediator secreted by HSC after liver injury by carbon tetrachloride or free radicals.

Leptin

Studies have found that HSC isolated in vitro and in vivo can produce leptin during activation. In the animal model of chronic liver injury induced by fatty liver and carbon tetrachloride, it was found that leptin-deficient rats had no liver fibrosis and typical liver structure. Liver endothelial cells and Kupffer cells are the primary target cells of leptin-induced liver fibrosis. Leptin can increase the expression of type I procollagen mRNA and TGF- β type II receptor mRNA and protein, enhance the effect of TGF- β 1 on collagen synthesis, and enhance the proliferation of PDGF-dependent HSCs through the PI3K/Akt pathway.

Activin

Activin (ACT) is a member of the large family of TGF β peptide factors, but it cannot bind to the receptor of TGF- β . HSC in resting state does not express ACT, but activated HSC does. In liver fibrosis, HSC expresses a large amount of ACT. There are act-a receptors on the surface of HSC. Smad Protein is vital in act-a signal transduction from cell surface silk and threonine kinase receptors to nuclear target genes. ACT expression levels of HSC at different locations in the hepatic lobule. The closer the hepatic lobule is to the fibrous connective tissue, the higher the gap expression level, showing a gradient distribution. ACT can promote HSC to secrete type I collagen, fibronectin, ECM, and so on in a dose-dependent manner within a specific concentration range, and act-a can play this biological effect in coordination with TGF- β . HSC secreting a large amount of act can also cause apoptosis of surrounding hepatocytes.

Endothelin

Endothelin-1 (ET-1) has a contractile solid effect on HSC. Activated HSC expresses the ET-1 receptor, and the binding of ET-1 with the receptor can increase the content of intracellular Ca²⁺, causing the contraction of myosin and sarcoplasmic protein. It also increases the expression of α -SMA and regulates the proliferation of HSC. An increase in intracellular Ca²⁺ concentration and cell contraction induced by ET-1 was observed at any stage of HSC activation. PDGF, TGF- β antagonists, and oxygen stress can increase the release of ET-1 from HSCs. ET-1 acts directly on the hepatic sinusoids, resulting in a decrease in portal vein blood flow in the body.

Epidermal Growth Factor

HSCs proliferate, and epidermal growth factor-r (EGF-R) expression increases in hepatic fibrosis. These findings suggest that EGF has a positive regulatory effect on the gene expression of MMP-3 and TIMP-1 in HSC and may be a synergistic regulation. The role of EGF in the occurrence and development of liver fibrosis depends on the effect of EGF on MMPs and TIMPs, the interaction between EGF and other cytokines, and the role of EGF in the balance of cytokines.

Diagnosis of Hepatic Fibrosis

Histopathology Diagnosis

Basic requirements for histopathological examination of liver biopsy: case histological examination is an essential basis for precise diagnosis, measurement of inflammatory activity, degree of fibrosis, and determination of drug efficacy. The liver biopsy specimens should be made into serial sections and routinely stained with hematoxylin-eosin, reticular fibres, and Masson trichrome to accurately judge the degree of inflammation, structural changes, and fibrosis in the liver. Immunohistochemical staining or in situ, viral antigen, or nucleic acid examination were added.

Noninvasive Diagnosis

Serum Markers: They should be relevant parameters that help to predict or detect inflammation and fibrosis in the liver, liver fibrogenesis, or degradation. At present, it is considered that hyaluronic acid (HA), type III procollagen peptide or its metabolic fragments (including P III P, P III NP, P III CP), type IV collagen or its metabolic fragments (including P IV-NP, P IV-NC1, P IV) and laminin (LN) which react to Extracellular Matrix (ECM) components; The combined detection of matrix protease inhibitor-1 (TIMP-1), which reflects the changes of ECM related enzymes, and transforming growth factor β 1 (TGF β 1), which demonstrates the formation of fibrosis, is more significant. Two or more of the above six indicators are abnormal, which has the diagnostic significance of liver fibrosis. **Related Liver Function and Immune Function:** In addition to child Pugh classification, albumin, prothrombin time, aspartate aminotransferase, chlorate aminotransferase, γ -glutamyltranspeptidase, apolipoprotein A1, α 2-macroglobulin, γ -globulin, 1GG, and alpha-fetoprotein can be used for auxiliary observation.

Imaging Evaluation: The reasonable selection of ultrasound, CT, and MRI and their mutual verification are helpful for dynamic observation. Quantitative or semi-quantitative criteria were used to observe the changes in liver elasticity, liver volume, liver surface edge, liver abdominal thickness, liver parenchyma, intrahepatic blood vessels, bile ducts, spleen, and splenic vein, gallbladder, and other indicators, which can provide valuable reference data for the diagnosis of liver fibrosis and the evaluation of the activity of the lesions. Transient elastography is also a standard imaging evaluation method. Transient Elastography (TE) is a non-invasive method specially used to measure liver fibrosis. The device has a probe that can emit ultrasound and an elastic wave that passes through the liver parenchyma, which measures Liver Stiffness (LS). Ultrasound (US) is usually the first-line radiological examination for patients with suspected liver cirrhosis because it is easy to obtain, noninvasive, well tolerated, cheaper than CT or MRI peers, provides real-time image acquisition and display, and will not expose patients to adverse reactions of intravenous contrast agent or radiation. The software on the machine processes the cross-velocity data of elastic waves in the liver and provides the value in

kilopascals for each measurement. The median of 10 measurements is the examination result. Since 2005, TE has entered the clinical practice of Hepatology, especially in evaluating patients with chronic viral hepatitis [59]. In 2015, the operation was also used to select hepatitis C patients with severe fibrosis and cirrhosis for new antiviral treatment [60].

Treatment of Liver Fibrosis

Liver fibrosis is potentially reversible. Patients receiving treatment for HCV infection experience significant reversal of fibrosis after achieving complete HCV negativity, but this improvement is mainly seen in the early stages (fibrosis grades 1 and 2). However, once fibrosis progresses to more advanced stages (grades 3 and 4) with the formation of cross-linked type I collagen, often accompanied by cell damage and inflammation, this reversibility becomes limited [13]. Therefore, addressing liver fibrosis is both a necessary and urgent task.

Research on the treatment of liver fibrosis has been steadily progressing. Current therapeutic approaches are primarily categorized into Western medicine and traditional Chinese medicine. Western medicine predominantly involves small-molecule chemical drugs, focusing on etiological treatment and anti-fibrotic therapy. Conventional Chinese medicine mainly utilizes herbal medicines to inhibit liver fibrosis.

Small Molecule Chemical Therapy

Etiological Treatment

The etiological treatment effectively prevents liver damage and is a viable method for treating liver fibrosis. In patients with chronic hepatitis C, significant fibrosis regression is observed following treatment with pegylated interferon α -2a or α -2b combined with ribavirin, particularly in mild to moderate cases, with women showing a higher degree of fibrosis reduction. For alcoholic hepatitis, complete abstinence from alcohol is essential. In many mild cases, avoiding alcohol can resolve clinical symptoms. For non-alcoholic hepatitis, lifestyle improvements through diet and exercise, as well as weight loss surgery, are the most effective methods [61]. For cholestatic hepatitis, Ursodeoxycholic Acid (UDCA) is widely used as a first-line treatment, and bile acid inhibitors, anti-fibrotic drugs, and anti-inflammatory drugs can also be utilized [62]. Continuous corticosteroid therapy can improve autoimmune hepatitis [63]. For patients with advanced liver disease, the limited availability of approved drugs necessitates liver transplantation as the final option for advanced fibrosis [64]. Liver fibrosis caused by schistosomiasis (*Schistosoma mansoni* and *Schistosoma japonicum*) is currently treated with praziquantel, while several urgently needed vaccines are in various stages of clinical development and are not yet approved for public use [65]. Etiological treatment is crucial for different causes of liver fibrosis to achieve optimal therapeutic outcomes.

Anti-fibrotic Treatment

For patients with advanced cirrhosis, etiological treatment is relatively slow in effectiveness, necessitating the development of anti-fibrotic treatments with fewer side effects. To date, many anti-fibrotic therapies have shown success in trials and experiments. These therapies include growth factors, cytokines, miRNA, monoclonal antibodies, stem cell-based methods, and other ECM-targeting approaches. The results of preclinical and clinical studies have laid the foundation for future alternatives to liver transplantation [66].

Activated HSCs, portal vein myofibroblasts, and the ECM they produce play pivotal roles in fibrosis and are thus the primary focus of anti-fibrotic therapies. These cells not only drive fibrotic processes but also interact with other cell types to influence fibrosis dynamics. For instance, they can promote fibrotic activation, induce quiescence or apoptosis, and contribute to ECM remodeling through the release of fibrinolytic enzymes and phagocytosis. This complexity highlights the potential for diverse therapeutic strategies. Therapies directly targeting HSCs, myofibroblasts, and ECM are categorized as “direct anti-fibrotic drugs,” while

those modulating other cellular interactions or pathways, such as immune cells or inflammatory mediators, are classified as “indirect anti-fibrotic drugs [67].”

Inhibition of the TGF- β 1/Smad Signaling Pathway

Extensive research has shown that dysregulation of the TGF- β 1/Smad pathway plays a crucial role in the development of tissue fibrosis. Smad2 and Smad3 serve as major downstream mediators that facilitate TGF- β 1-driven fibrosis, while Smad7 functions as a negative feedback regulator, providing protection against fibrosis induced by TGF- β 1.[68]. Therefore, inhibiting the TGF- β 1/Smad signalling pathway prevents HSC activation and proliferation, thereby improving liver fibrosis [69]. Experimented by Leonel et al., It is revealed that pirfenidone treatment in animal livers significantly downregulated TGF- β 1. It was concluded that pirfenidone might inhibit TGF- β 1 expression, leading to upregulation of the Smad-7 gene, which could benefit the treatment of human liver fibrosis [70]. Drugs such as praziquantel and fluorofenidone (Figure 5) also inhibit hepatic stellate cell activation by targeting the TGF- β 1/Smad pathway, serving as useful anti-fibrotic agents [71, 72].

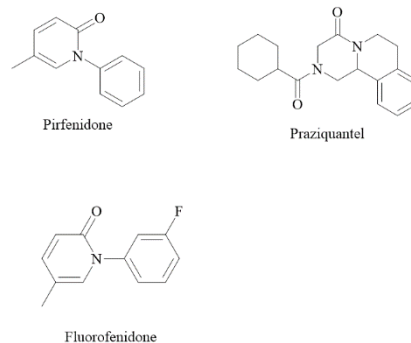


Figure 5: Chemical Structure of Drugs Inhibiting the TGF- β 1/Smad Signaling Pathway.

Inhibition of Fibroblast Growth Factors

There are 22 known members of the human Fibroblast Growth Factor (FGF) family. With several physiological roles, FGF21 is an endocrine hormone that is expressed in the liver, pancreas, white adipose tissue (WAT), and brown adipose tissue (BAT) [73]. FGF21 is a crucial metabolic regulator predominantly expressed in the liver. Cong evaluated the effects of the FGF21 analogue LY2405319 in a mouse model of liver fibrosis and concluded that FGF21, as an inhibitor of the succinate-GPR91 pathway, effectively controls liver fibrosis [74]. Thus, FGF21 presents a novel approach for treating fibrosis. Pegbelfermin (BMS-986036), a PEGylated FGF21 analogue, demonstrated in a multicenter, randomized, placebo-controlled phase 2a study that daily doses of 10 mg or weekly doses of 20 mg for 16 weeks were generally well tolerated and considerably reduced liver fat in patients with non-alcoholic steatohepatitis

(fibrosis stages 1-3). It also positively affected several parameters related to non-alcoholic steatohepatitis. Furthermore, Harrison et al. found that NGM282, an FGF19 analogue, inhibits classical bile acid synthesis and suppresses fatty acid synthesis and lipogenesis, effectively addressing non-alcoholic steatohepatitis and inhibiting liver fibrosis [75].

Inhibition of ECM

ECM buildup is a hallmark of liver fibrosis, a wound-healing reaction to persistent liver injury. Urokinase Plasminogen Activator (UPA) promotes ECM degradation by activating Matrix Metalloproteinases (MMPs). Increased MMP-9 activity contributes to liver fibrosis, while FRETTOH downregulates UPA, MMP-2, and MMP-9 activities in hepatocytes and reduces MMP-9 activity [76]. Thus, FRETTOH demonstrates a therapeutic effect on liver fibrosis.

Inhibition of LOX

A humanized IgG4 monoclonal antibody named simtuzumab specifically targets LOXL2, blocking its function in vitro and in mouse studies. In mouse models, simtuzumab reduces fibroblast activation and the production of cytokines and growth factors, such as transforming growth factor- β (TGF- β) [77]. However, further research is required to investigate Simtuzumab's effect on transaminases, as current experiments do not provide sufficient evidence of its efficacy.

MiRNA and SiRNA

Hepatic stellate cells (HSCs) and fibroblasts, the main source of fibrotic cells in liver tissue, have significant levels of MiR-122 expression. MiR-122 suppresses the expression of serum response factor (SRF), which in turn lowers the transcription of α -SMA and COL1A1, and directly inhibits FN1 expression by binding to its 3'-untranslated region. SRF is an essential transcription factor that mediates fibrotic cell activation [78]. Li demonstrated through loss-of-function analysis that the deletion of HDAC2 in TGF- β 1 activated

HSC-T6 cells promotes cell cycle arrest and blocks the expression of Col1 α 1 and α -SMA proteins [79]. Therefore, HDAC2 plays a crucial role in HSC activation and liver fibrosis and could be a target for liver fibrosis therapy, although its regulatory role requires further investigation.

Renin-Angiotensin Inhibitors

The renin-angiotensin system (RAS) is an essential regulator of liver cirrhosis and portal hypertension. As liver fibrosis progresses, the levels of RAS components, including angiotensin (Ang) II, Ang-(1-7), Angiotensin-Converting Enzyme (ACE), and Ang II Type 1 Receptor (AT1R), increase [80]. The fibrotic and inflammatory effects of Angiotensin II in the liver are primarily mediated by the Angiotensin Type 1 (AT1) receptor. AT1 receptor blockers have been proposed for treating liver fibrosis in patients with chronic liver diseases and have shown good tolerance. Losartan can reduce serum transaminase levels without adverse effects as a representative drug. Valsartan and Irbesartan (Figure 6) are also commonly used AT1 receptor blockers, showing potential for treating liver fibrosis.

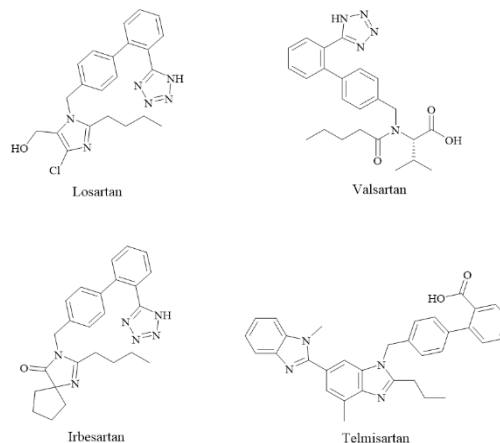


Figure 6: Chemical Structure of AT1 Receptor Blockers.

FXR Receptor Inhibitors

Obeticholic Acid (OCA) is an FXR receptor agonist with an efficacy 100 times greater than Chenodeoxycholic Acid (CDCA). FXR signalling protects hepatocytes from bile acid toxicity by inhibiting bile acid synthesis and stimulating bile secretion through the upregulation of bile acid transporters. Additionally, it regulates other pathways with direct anti-inflammatory and antifibrotic effects. In a clinical trial, OCA improved liver biochemistry [81]. Glycochenodeoxycholic Acid (GUDCA) and Tauroursodeoxycholic acid (TUDCA) (Figure 7) are FXR antagonists that significantly inhibit FXR transcriptional activity.

C-C Chemokine Antagonists

Geniciviroc (Figure 8), a C-C Chemokine Receptor Type 2 and

5 (CCR2/CCR5) antagonist, has been demonstrated to have anti-inflammatory and antifibrotic effects in animal models, which is primarily due to ceniciviroc's ability to block the inflammatory cascade of fibrosis. Compared to placebo, ceniciviroc significantly improved liver fibrosis by at least one stage after 48 weeks [82].

GR-MD-02

GR-MD-02 (Galactose arabinose-rhamnose cythiathate) has been shown to reduce liver fibrosis and decrease serum biomarkers of NASH fibrosis in animal studies. However, it was found that the drug only had a positive therapeutic effect on a small portion of liver disease patients due to insufficient dosage or lack of sensitivity to high fibrosis levels in experiments [83]. Further clinical trials are needed to determine its efficacy.

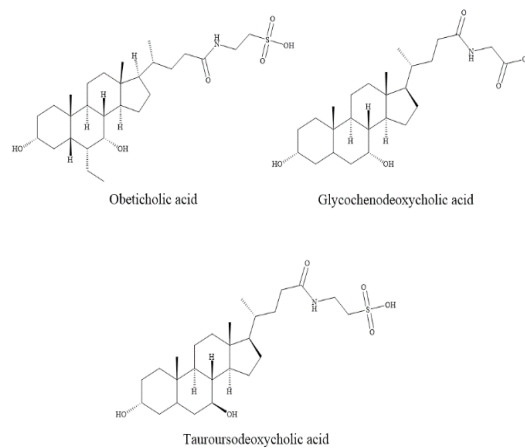


Figure 7: Chemical Structure of FXR Receptor Inhibitors.

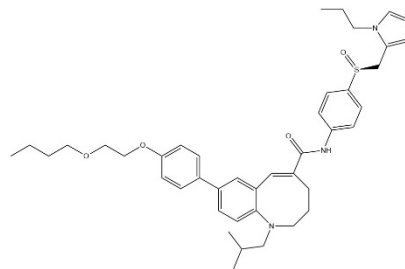


Figure 8: Chemical structure of cenicriviroc.

T2DM

In the general population, Type 2 Diabetes Mellitus (T2DM) has been found to be a risk factor for liver fibrosis, particularly in individuals with fatty liver disease. It has also been discovered that metformin, a first-line hypoglycemic medication, helps with liver fibrosis and steatosis [84]. According to liver biopsy results, 32%

of patients exhibited advanced fibrosis, and the risk of advanced fibrosis significantly decreased after metformin treatment. Studies have shown that metformin (Figure 9) inhibits liver fibrosis, reduces fibrosis severity, and curtails the progression of non-alcoholic fatty liver disease. Additionally, metformin intervention can inhibit the activation of HSCs, deplete lipid accumulation in hepatocytes, and prevent decompensated cirrhosis and liver fibrosis [85].

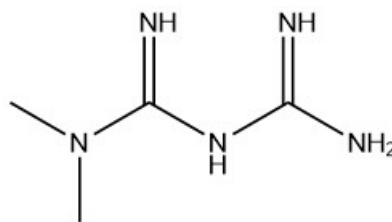


Figure 9: Chemical structure of metformin.

Targeting Reactive Oxygen Species (ROS) in Therapeutic Interventions

Research has identified ROS-generating enzymes in many molecules and subcellular organelles, which produce ROS during physiological processes. These ROS-producing enzymes are promising targets for treating oxidative stress-related diseases,

including liver fibrosis. For example, NOX and TLR in the liver can generate ROS through crosstalk, potentially adversely affecting liver cell function and leading to fibrosis. Drugs targeting ROS production have received significant attention in treating oxidative stress-related diseases, especially liver diseases, providing adequate means for current clinical therapies [86].

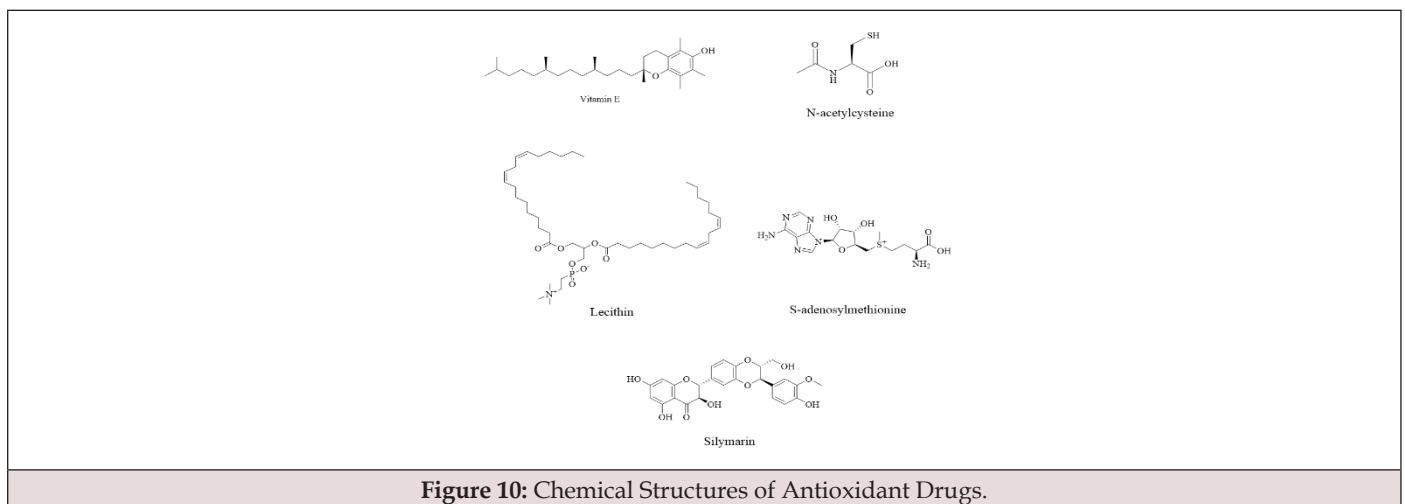
NOX Inhibitors

Reactive Oxygen Species (ROS) generated by NADPH Oxidase (NOX) play a critical role in liver injury and fibrosis. Therefore, inhibiting NOX1 and NOX4 is crucial for treating liver fibrosis. The dual NOX1/4 inhibitor GKT137831 has been shown to inhibit ROS production and inflammation and proliferation genes in primary mouse HSCs. Clinical trials have found that it effectively reduces the degree of liver fibrosis and exhibits good pharmacological properties and safety [87], making it a novel therapeutic approach. Losartan, another drug affecting NOX, is widely used in clinical

practice; however, its therapeutic effects on liver-related diseases are not yet significant [86]. NOX1 and NOX4 provide new directions for future research in liver fibrosis treatment.

Antioxidants

Antioxidants can reduce ROS production, thereby improving liver fibrosis. Antioxidants such as S-Adenosylmethionine (SAME), Silymarin, Lecithin, N-Acetylcysteine (NAC), and Vitamin E (Figure 10) have been proven to have beneficial effects in clinical trials [88].



Traditional Chinese Medicine (TCM) in Treatment

In recent years, TCM has continuously developed, and its efficacy and safety in the clinical treatment of liver fibrosis have been proven to surpass those of Western medicine, demonstrating a promising application prospect. Unlike Western medicine, TCM emphasizes unity, considering the patient and environmental factors together and treating the patient as an organic whole. TCM exhibits significant biological advantages [89]. The metabolism of intestinal microorganisms affects the therapeutic effect of TCM and is an essential factor to consider.

The liver and intestines are inseparable in the human body. Endotoxemia in liver cirrhosis is a phenomenon of bacterial translocation caused by bacterial overgrowth. Due to the interaction between the intestines and the liver, liver fibrosis affects the intestinal flora [90]. Similarly, the disruption of the gut microbiota can lead to the development of liver fibrosis. It has been found that TCM can effectively regulate the gut microbiota, playing a therapeutic role [91].

Scutellaria Baicalensis

Probiotics can improve the intestinal barrier and protect it from damage. Studies have found that probiotics such as *Lactobacillus*

casei GKC1 (GKC1), *Bifidobacterium lactis* GKK2 (GKK2), *Lactobacillus rhamnosus* GKLC1 (GKLC1), and *Lactobacillus paracasei* GKS6 (GKS6) have potential benefits in alleviating liver fibrosis [92]. *Scutellaria baicalensis*, a traditional Chinese herb widely used in China for its anti-inflammatory, antibacterial, and antioxidant properties, can effectively promote the healthy growth of intestinal bacteria, making it helpful in treating liver fibrosis [91]. Oroxylin A in *Scutellaria baicalensis* can induce autophagy of ferritin, thereby inducing HSC senescence and significantly inhibiting the expression of senescence marker SA- β -gal in HSCs, reducing liver fibrosis [93].

Astragalus

Astragalus polysaccharide, an important active component extracted from *Astragalus*, has antioxidant, antitumor, antiviral, and cardiovascular protective effects and is widely used in the clinical treatment of colitis. IL-17 has strong pro-inflammatory effects, leading to inflammation, while *Astragalus* polysaccharide can inhibit IL-17 and TNF- α , acting as an anti-inflammatory agent [92]. *Astragalus* polysaccharide also increases the number of *Lactobacillus* and *Bifidobacterium*, further inhibiting inflammation. Serum ALT and AST levels are essential indicators of liver function; *Astragalus* polysaccharide can significantly reduce serum ALT and AST levels, further indicating its potential to reduce liver disease and protect the liver [94].

Atractylodes

Atractylodes is a traditional Chinese medicine with spleen-strengthening and antidiarrheal effects. Treatment with Atractylodes significantly reduces the number of pathogenic bacteria, such as *Helicobacter pylori* and *Clostridium*, while increasing beneficial bacteria, such as *Lactobacillus*. This indicates that Atractylodes can promote the growth of beneficial bacteria, restore intestinal microecology, and improve intestinal structure, having a therapeutic effect on liver fibrosis [95].

Pueraria

Pueraria, the dried root of the leguminous plant Pueraria, contains many bioactive components with antioxidant, anti-inflammatory, and immunomodulatory activities [96]. Research has found that Pueraria extract (RPE) can reduce pathological changes and lipid deposition in rat liver tissue and decrease serum ALT, AST, and liver GGT activity, effectively mitigating liver damage caused by chronic alcohol intake [97]. Animal experiments have shown that puerarin, an active ingredient in Pueraria, activates endogenous PPAR- γ expression in rats, inhibiting the proliferation and activation of ECM-driven liver cells and strongly inhibiting fibrosis [98]. Pueraria and silymarin have therapeutic effects on alcoholic fatty liver and liver inflammation, showing potential as effective drugs for treating liver fibrosis [99].

Schisandra

Schisandra regulates the gut microbiota, reversing the intestinal bile acid spectrum and microbiome to protect the liver and block fibrosis [100]. Schisandra acidic polysaccharides, one of the main components of Schisandra, improve damaged liver tissues pathologically and have an excellent protective effect on acute ethanol-induced liver injury. The mechanism may be related to reducing oxidative stress and regulating the expression of CYP2E1 [101]. Schisandra is a natural product with promising future research potential for developing liver-protective drugs.

Sophora Flavescens and Ligustrum Lucidum

Sophora root is an essential herb in treating hepatitis B in China, with its alkaloids showing significant liver-protective effects and polysaccharides demonstrating good immunomodulatory activity [102]. Ligustrum lucidum, the mature fruit of the Oleaceae family, has multiple pharmacological effects, including anticancer, liver protection, anti-inflammatory, anti-osteoporosis, and antioxidant properties [103]. The combination of Sophora and Ligustrum has been documented in ancient medical texts. Studies have found that when the ratio of these two drugs is less than 1:1, the synergistic therapeutic effect on liver fibrosis is significantly enhanced [104]. In the treatment of liver fibrosis, traditional therapies and herbal medicine demonstrate distinct advantages and characteristics. Traditional therapies typically target HSC activation, regulate inflammatory responses, and

degrade ECM through mechanisms such as those employed by drugs like lovastatin, which exhibit significant efficacy in inhibiting fibrotic signaling pathways. However, traditional methods also have limitations, including drug resistance, immunosuppression, and potential side effects. In contrast, herbal medicine stands out with its multi-targeted mechanisms, not only modulating HSC activation but also enhancing liver repair through anti-inflammatory, antioxidant, and gut microbiota-regulating pathways. For instance, *Scutellaria baicalensis* contains flavonoid compounds that exert anti-inflammatory and anti-fibrotic effects by inhibiting the TGF- β signaling pathway; Atractylodes macrocephaly has been shown to improve liver function, modulate the immune system, and reduce ECM deposition; while Pueraria protects hepatocytes through its antioxidant properties, alleviating oxidative stress and inflammation associated with fibrosis. Although herbal medicines generally pose fewer side effects, potential hepatotoxicity of certain herbs warrants caution. Overall, integrated traditional Chinese and Western medical approaches combining the multi-targeted mechanisms of herbal medicine with the precise efficacy of traditional therapies hold promise for achieving synergistic effects and advancing liver fibrosis treatment. Nonetheless, clinical evidence of the efficacy of herbal medicine requires further validation through large-scale randomized controlled trials.

Prevention of Liver Fibrosis

Intestinal Alkaline Phosphatase (IAP)

Research has shown that TLR4 drives myofibroblast activation and fibrogenesis in the liver, linking pro-inflammatory and pro-fibrotic signals through TLR4-dependent TGF- β signalling. The Lipopolysaccharide (LPS)-TLR4 pathway can deplete gut bacteria, which helps mitigate liver fibrosis [42]. IAP is expressed throughout the intestine, with the highest expression in the duodenum and the highest phosphatase activity at the terminal ileum [105]. IAP reduces the amount of active LPS transported from the intestine to the liver, effectively alleviating liver fibrosis and serving as a preventive measure against liver fibrosis [106].

Dietary Considerations

Fructose from carbohydrates, trans fatty acids, and omega-6 fatty acids can contribute to developing Non-Alcoholic Fatty Liver Disease (NAFLD). However, fibre from carbohydrates and omega-3 fatty acids, along with micronutrients such as vitamins C, E, D, and polyphenols, can prevent NAFLD onset [107]. Reducing the intake of fried foods and increasing the consumption of foods rich in monounsaturated and polyunsaturated fatty acids is essential for liver fibrosis patients. It is recommended that these patients consume more walnuts and wild cherries [108].

Exercise

Exercise is also an effective method to prevent liver fibrosis. Physical activity can limit liver inflammation and avoid progression

to advanced liver damage, such as fibrosis and cirrhosis [109]. Some studies suggest moderate to vigorous exercise is associated with improved liver indices and reduced severe liver fibrosis. In mice models, those subjected to exercise showed significantly reduced liver fibrosis [110]. Therefore, engaging in physical activity, including moderate-intensity aerobic exercise for at least 20-60 minutes five days a week and resistance training three times a week, has become a practical recommendation for preventing and treating liver fibrosis. Additionally, promoting weight loss and preventing overnutrition has been proven to alleviate liver fibrosis [111].

Omics Studies

A comprehensive analysis of liver fibrosis and omics technologies has been conducted using multi-omics approaches to study the relationship between liver fibrosis and omics technologies. Liver fibrosis is a pathological process caused by chronic liver injury, characterized by excessive extracellular matrix deposition, which can eventually lead to cirrhosis. Various omics technologies, including transcriptomics, genomics, and proteomics, provide a comprehensive understanding of the cell types involved in liver fibrosis.

Transcriptomics

The role of transcriptomics in liver fibrosis is primarily to elucidate changes in gene expression by analyzing mRNA expression patterns during liver fibrosis. Studies have found that therapeutic drugs that are effective in mouse liver injury models also show efficacy in zebrafish. Consequently, zebrafish are widely used in liver development and injury research [112]. Despite significant progress in understanding the molecular mechanisms of liver fibrosis, little is known about the specific gene regulatory networks of different cell types during the early stages of liver injury. When Thioacetamide (TAA) was used to induce early liver injury in adult zebrafish, Migdal identified changes in chromatin accessibility and transcription in three major liver cell types: hepatocytes, endothelial cells, and hepatic stellate cells. The study revealed significant transcriptional alterations in genes related to fatty acid and carbohydrate metabolism, immune response, and vascular-specific processes following TAA treatment. Notably, liver endothelial cells exhibited the most significant response at transcriptomic and chromatin levels, characterized by a loss of angiogenic phenotype. Multi-omics analysis provided valuable insights into cell type-specific transcriptomic and epigenomic responses to early liver injury, enhancing our understanding of the molecular mechanisms underlying liver fibrosis [113].

Genomics

Genomic studies have revealed the relevant gene mutations and variations associated with liver fibrosis. The systemic genetic analysis of the recombinant inbred BXD mouse population

identified core risk genes and gene interaction networks for liver fibrosis. A Carbon Tetrachloride (CCl₄)-induced liver fibrosis model combined with whole-genome QTL analysis identified seven genomic loci significantly affecting the fibrotic phenotype. Further expression QTL analysis reduced the 1,351 candidate genes to 11 core candidate genes. These findings demonstrated that the BXD reference population is a powerful experimental resource for identifying gene regulatory networks controlling susceptibility to chronic liver injury. The results provide new insights into the genetic basis of liver fibrosis and highlight the potential of systems genetics in studying complex traits [114].

Proteomics

Proteomics research focuses on proteins' expression, modification, and interaction, revealing changes in cellular functions and signalling pathways. For example, mass spectrometry can identify proteins that significantly change during liver fibrosis, such as collagen and extracellular matrix proteins. These studies help understand proteins' regulatory mechanisms and functional changes, revealing potential therapeutic targets. In one study, mice were subjected to a six-week CCl₄ injection to induce chronic liver injury, and proteomic changes were analyzed through RNA sequencing. The expression of various proteins in the mouse liver significantly changed after CCl₄ treatment. ECM and wound healing-related proteins were significantly upregulated, while proteins related to hepatocyte metabolism and drug metabolism were downregulated. These protein changes reflected alterations in cell composition and function during fibrosis. The study found that the expression of the Stard3nl gene was significantly associated with fibrosis. The protein encoded by this gene may play a crucial role in ECM organization and liver fibrosis. Other candidate genes, such as Clca3a2, also showed potential associations with liver fibrosis. Liver fibrosis involves complex gene and protein networks, particularly proteins related to ECM remodelling and immune response, which play critical roles in the fibrosis process [115].

In summary, multi-omics research approaches provide a comprehensive perspective for revealing the molecular mechanisms of liver fibrosis, helping scientists understand the interactions between different molecular levels. These studies provide essential information for basic research on liver fibrosis and offer potential targets and resources for developing new therapeutic strategies and diagnostic tools.

Summary and Prospects

This paper comprehensively reviews the causes, pathogenesis, small molecule drugs, traditional Chinese medicine (TCM) treatments for liver fibrosis over the past decade, and corresponding preventive measures. The continuous progression of liver fibrosis can irreversibly lead to cirrhosis and liver cancer, posing a severe threat to liver tissue and affected populations. Addressing liver fibrosis is currently a critical issue that requires urgent solutions.

Liver fibrosis can result from chronic diseases, metabolic disorders, cholestasis, and schistosomiasis infection. Once it occurs, rapid intervention and treatment are essential to prevent dangerous outcomes. The accumulation of the Extracellular Matrix (ECM) can lead to complications, such as cardiovascular complications associated with cirrhosis, which have clinical latency and require physiological or pathological changes for detection. These pathological and physiological impacts are irreversible, making the prevention and treatment of liver fibrosis a key research focus.

Both parenchymal and non-parenchymal cells in the liver can cause fibrosis, and the mechanisms are complex. Identifying the underlying causes is crucial for treatment or alleviation. A thorough understanding of the pathogenesis is essential for targeted therapy. Western medicine and TCM have different focuses: Western medicine targets the causes of liver fibrosis for specific treatments, while TCM emphasizes holistic body regulation and necessary medication. Potential preventive measures, such as improved diet and rational exercise, can alleviate liver fibrosis. Early detection and intervention are the preferred methods for treating liver fibrosis. Numerous studies have demonstrated that small-molecule drug treatments and TCM preventive measures have significant potential in treating liver fibrosis. This paper explores liver fibrosis through the application of multi-omics technologies, including transcriptomics, proteomics, and genomics, to achieve a comprehensive understanding of the molecular mechanisms underlying fibrosis. Transcriptomics helps identify cell-specific gene expression changes in hepatocytes, HSCs, and endothelial cells, offering insights into cellular contributions to fibrosis. Proteomics complements these findings by examining protein expression, modification, and interaction, highlighting changes in cellular functions and signaling pathways. The integration of these multi-omics approaches provides a holistic view of the complex gene and protein networks.

Future research can focus on the following areas from both Western and TCM perspectives:

Early Detection Biomarkers: There is an urgent need for highly sensitive and specific biomarkers for early-stage liver fibrosis. Identifying novel molecules, such as non-coding RNAs or specific proteins, could improve early diagnosis and intervention.

Innovative Anti-Fibrotic Drugs: Developing new drugs with better efficacy and lower toxicity, such as inhibitors targeting TGF- β /Smad signaling or hepatic stellate cell activation, is critical. Drug repurposing and high-throughput screening offer promising approaches.

Modernization of TCM: Rediscovering and validating classical TCM formulas through active compound isolation and clinical trials could enhance their application. Technologies like metabolomics and proteomics can provide molecular insights.

Integrated Therapies: Combining Western anti-fibrotic drugs with TCM for synergistic effects requires further exploration. Studies should focus on identifying effective combinations and ensuring safety through comprehensive trials.

Personalized Treatment Strategies: Research on patient-specific genetic and environmental factors, as well as advanced technologies like RNA therapies and gene editing, could enable tailored interventions for liver fibrosis. Although liver fibrosis is an early stage of various liver diseases, without intervention, it can lead to irreversible effects. Therefore, early detection, intervention, and treatment are crucial for managing liver fibrosis. Future biological research should aim to identify more specific molecules reflecting liver fibrosis indicators as practical clinical diagnostic markers, enabling more efficient early detection, intervention, and treatment to safeguard human health.

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Declaration of competing interest

Authors have no conflict of interest to declare.

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