

REVIEW

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Emerging advanced approaches for diagnosis and inhibition of liver fibrogenesis

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Abstract

Background Liver fibrosis results from chronic liver injury and is characterized by excessive deposition of extracellular matrix proteins including collagen. It can progress to cirrhosis and liver failure.

Main body of the abstract Multiple cellular signaling pathways drive hepatic stellate cell activation and fibrogenesis. Advances in biomarkers, imaging modalities, and omics platforms enable noninvasive diagnosis and staging of liver fibrosis. Emerging antifibrotic approaches include medications like pirfenidone, obeticholic acid, and monoclonal antibodies targeting pro-fibrotic mediators. Cell therapies using mesenchymal stem cells demonstrate antifibrotic potential through paracrine immunosuppression. Tissue-engineered liver grafts and biomaterial carriers for localized drug delivery are promising technologies. Microfluidic liver-on-a-chip platforms with patient-derived cells provide unprecedented models to study human liver fibrosis and test drug candidates.

Short conclusion Significant progress has elucidated mechanisms underlying liver fibrogenesis and uncovered novel therapeutic targets. Ongoing challenges include translating preclinical findings, improving antifibrotic efficacy, and enabling personalized precision medicine approaches. Further research into combinatorial therapies, biomarkers, and tissue engineering technologies will advance the treatment of liver fibrosis from all causes.

Keywords Liver fibrosis, Antifibrotic therapeutic targets, Inflammatory mediators, Liver-on-a-chip platforms, Cell-based therapies, Tissue engineering

Background

Liver fibrosis involves excessive deposition of extracellular matrix proteins like collagen. It occurs due to chronic liver injury and is an active wound-healing response that can progress to cirrhosis [1]. Fibrosis

develops from sustained inflammation caused by chronic viral hepatitis, fatty liver disease, alcohol, autoimmune disorders, metabolic disorders, and cholestatic diseases [2]. During the initiation of liver injury, the activation of Kupffer cells and recruitment of inflammatory immune cells result in the release of cytokines, chemokines, and growth factors that trigger a response from hepatic stellate cells. Hepatic stellate cells are the major effector cells that become activated, proliferate, and transform into myofibroblast-like cells that deposit excessive amounts of extracellular matrix [3]. Key cytokines and growth factors involved in this process include transforming growth factor beta, platelet-derived growth factor, connective tissue growth factor, and endothelin-1. Activated stellate cells increase their expression of alpha-smooth muscle actin and exhibit contractile, proinflammatory, and fibrogenic properties [4]. The pathological accumulation of extracellular

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matrix in liver fibrosis occurs primarily from increased synthesis and deposition of collagen type I and type III. This is mediated by activated stellate cells which upregulate tissue inhibitors of metalloproteinases, causing an imbalance in extracellular matrix turnover favoring excessive accumulation. In addition to activated stellate cells, other cell types contribute to the fibrogenic response including portal fibroblasts, bone marrow-derived fibroblasts, and hepatocytes undergoing epithelial to mesenchymal transition. Chronic inflammation also perpetuates liver fibrosis through paracrine signaling between macrophages, stellate cells, and damaged hepatocytes. Figure 1 illustrates the role of hepatic stellate cells (HSCs) in the progression and regression of liver fibrosis. In a healthy liver, HSCs are quiescent (qHSCs). Liver injury triggers qHSC activation mediated by factors like hepatocytes, liver sinusoidal endothelial cells (LSECs), macrophages, transforming growth factor-beta (TGF-β), and platelet-derived growth factor (PDGF). Activated HSCs (aHSCs) proliferate, secrete excessive extracellular matrix (ECM) proteins, and promote fibrosis. Fibrosis can be reversed by inhibiting aHSC proliferation, inducing aHSC death, or inactivating aHSCs (iHSCs) [5]. Liver fibrosis results from chronic damage from etiologies like viral hepatitis and can progress to cirrhosis with distortion of hepatic

architecture and regenerative nodules. Cirrhosis has complications like portal hypertension and liver failure. Accurate fibrosis staging is crucial for prognosis, guiding treatment, and monitoring progression. Liver fibrosis advances through stages of increasing collagen deposition and architectural distortion. Scoring systems categorize fibrosis stages based on patterns of fibrotic areas [6, 7]. Collagen first appears around vessels, then bridges portal and central areas causing distortion and nodules-defining cirrhosis [8]. Cirrhosis disrupts liver function and blood flow, increasing complications like cancer, bleeding, ascites, infections, and organ failure [9–13]. Genetic and epigenetic factors also influence fibrosis progression [14]. Biomarkers and imaging techniques are being developed for noninvasive diagnosis [15]. This review comprehensively summarizes recent advances across the spectrum of liver fibrosis research, from elucidating molecular pathways to emerging diagnostics, biomarker technologies, and novel therapeutic candidates. Notably, we highlight promising areas including stem cell, tissue engineering, nanotechnology, and microfluidic model approaches offering hope for personalized, precision antifibrotic therapies based on rapidly expanding knowledge of pathways underlying fibrogenesis. The background explains the focus is providing a comprehensive review of the multifaceted

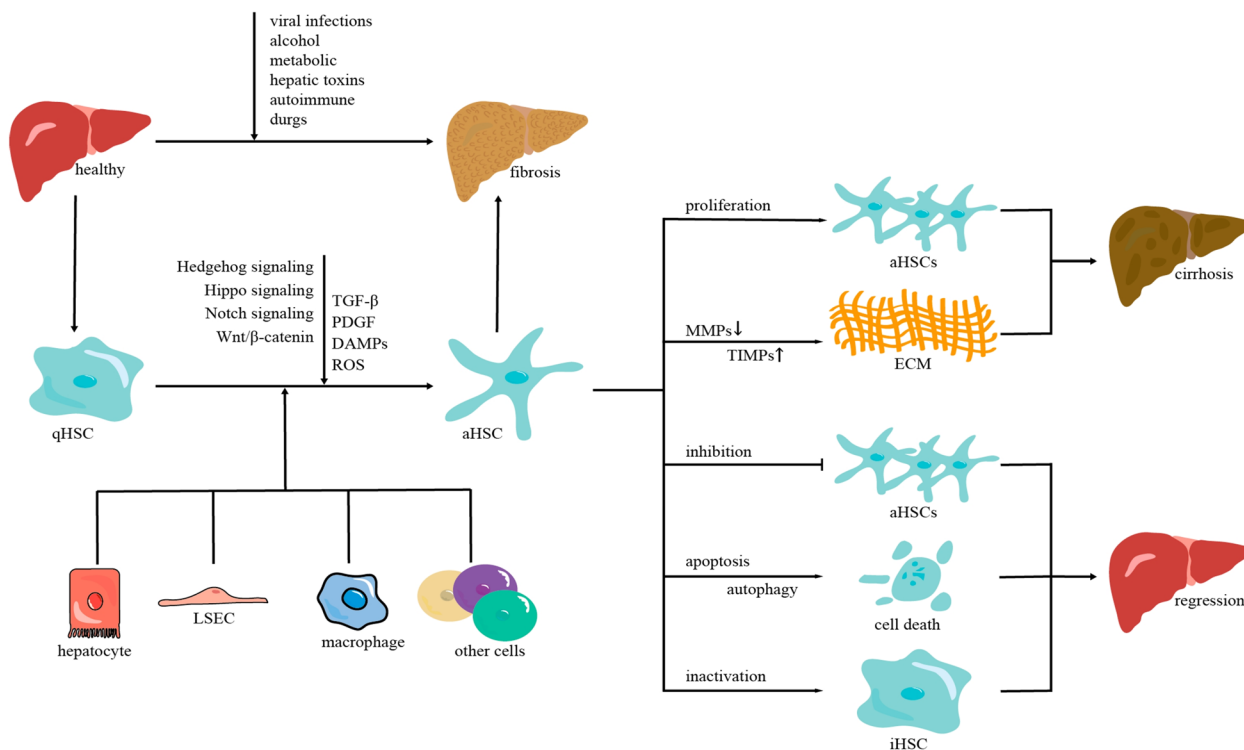


Fig. 1 Role of hepatic stellate cells in liver fibrosis progression and reversal [5]

progress in understanding and treating liver fibrosis. It emphasizes emerging tools enabling precision, and personalized approaches as a key aspect differentiating this review.

Cellular and molecular mechanisms

Liver fibrosis results from a complex interplay between multiple cell types and signaling pathways that together stimulate hepatic stellate cell activation and excessive extracellular matrix (ECM) production and remodeling. Key events include hepatic stellate cell (HSC) activation into proliferative, fibrogenic myofibroblasts under paracrine stimuli from damaged hepatocytes and cells like Kupffer cells releasing inflammatory cytokines like tumor necrosis factor- α (TNF- α) and interleukin-6 (IL-6) [16]. Activated HSCs upregulate ECM components like fibrillar collagens and tissue inhibitors of metalloproteinases (TIMPs). Chronic inflammation further drives fibrosis progression through recruited leukocytes secreting profibrogenic cytokines. Epithelial-mesenchymal transition (EMT) of hepatocytes into migratory, invasive fibroblast-like cells is triggered by stimuli like transforming growth factor- β (TGF- β) activating transcription factors like Snail and Zeb [17]. Strategies targeting HSC activation, inflammation, EMT, oxidative stress, and extracellular matrix balance show promise for antifibrotic therapies.

Genetic and epigenetic regulation

Liver fibrosis progression and severity exhibit interindividual variability influenced by genetic and epigenetic factors. Polymorphisms in genes encoding pro- and anti-fibrogenic mediators can regulate an individual's fibrotic response to injury. Epigenetic changes including DNA methylation, histone modifications, and non-coding RNAs control hepatic stellate cell activation and profibrogenic gene expression. Functional polymorphisms affecting cytokines, the renin-angiotensin system, collagen remodeling enzymes like matrix metalloproteinases (MMPs), and genes regulating oxidative stress and immunity modify fibrosis progression [18]. Genome-wide association studies have uncovered additional genetic loci correlated with fibrosis severity, like toll-like receptor 4 (TLR4), neurocan (NCAN), and patatin-like phospholipase domain containing 3 (PNPLA3) variants [19]. Key epigenetic mechanisms include DNA methylation, post-translational histone modifications, and non-coding RNAs such as microRNAs (miRNAs). Hypermethylation of antifibrotic genes promotes stellate cell activation. MiR-29 downregulation correlates with progression [20]. Histone modifications alter chromatin structure to promote profibrogenic gene transcription. Manipulating these changes using histone deacetylase inhibitors

or miR-29 restoration reduced fibrosis in models [21]. Hepatic stellate cells exhibit major DNA methylation reprogramming during activation warranting further epigenome analysis [22].

Molecular pathogenesis of liver fibrosis in major liver diseases

Liver fibrosis can occur as a consequence of virtually all chronic liver diseases. However, the molecular triggers for hepatic stellate cell activation and fibrogenesis show some variations between different disease etiologies. In chronic hepatitis C virus (HCV) infection, fibrosis results from hepatic inflammation triggered by the immune response against the virus. HCV proteins directly activate stellate cells through oxidative stress. Loss of antifibrotic miRNAs like miR-29 contributes to fibrosis. Hepatitis B virus (HBV) similarly elicits fibrosis indirectly through immune-mediated inflammation and directly through profibrogenic viral factors altering the epigenetic regulation of genes in stellate cells. Effective antiviral suppression can normalize fibrosis. In nonalcoholic fatty liver disease (NAFLD), fibrosis is driven by inflammation from fat accumulation, insulin resistance, oxidative stress, and lipotoxicity activating Kupffer cells and recruiting macrophages. Hepatocyte apoptosis from lipotoxicity provokes stellate cell matrix deposition [23]. Oxidative stress is a key driver in NAFLD and alcoholic liver disease, with alcohol metabolism generating reactive oxygen species inducing inflammatory and fibrogenic genes. In cholestatic diseases like primary biliary cholangitis, fibrosis occurs from bile acid-induced cholangiocyte and hepatocyte apoptosis recruiting immune cells to stimulate stellate cells. While shared pathways potentiate fibrosis across diseases, elucidating molecular nuances related to the initial injury is imperative for targeted therapies, like suppressing inflammation, inhibiting lipotoxicity, and oxidative stress, blocking stellate activation, and normalizing dysregulated epigenetic/metabolic pathways [24].

Emerging biomarkers for liver fibrosis assessment

The development of non-invasive biomarkers to diagnose and stage liver fibrosis is an area of intense research. Beyond traditional protein and metabolic biomarkers, newer classes include microRNAs (miRNAs), long non-coding RNAs (lncRNAs) and extracellular vesicles (EVs), and their molecular cargos. Multiple miRNA families show altered expression during fibrogenesis and directly target extracellular matrix component mRNAs for repression. Declining miR-29 levels correlate with the advancing fibrosis stage. A 5-miRNA panel accurately differentiated significant fibrosis in nonalcoholic fatty liver disease. lncRNAs (long non-coding RNAs) are RNA transcripts longer than 200 nucleotides that regulate

gene expression but do not encode proteins. In fibrotic diseases affecting the liver, lungs, kidneys, and heart, the expression levels of many lncRNAs become dysregulated. One prominent example is HOTAIR (HOX transcript antisense RNA), which shows increased expression across multiple fibrotic diseases. Higher circulating blood levels of HOTAIR correlate with more severe fibrosis and worse clinical outcomes, suggesting it could potentially serve as a biomarker for fibrosis severity. Mechanistically, HOTAIR interacts with chromatin remodeling complexes and can silence antifibrotic genes. Overexpression of HOTAIR promotes collagen deposition and myofibroblast differentiation. Other pro-fibrotic lncRNAs include MEG3, PVT1, and FENDRR, while GAS5 and lncRNA-ATB have antifibrotic effects. Modulating the levels of dysregulated lncRNAs impacts fibrosis progression in preclinical disease models. The functional roles and dysregulation of lncRNAs like HOTAIR in fibrosis point to their potential as biomarkers and therapeutic targets in fibrotic diseases across different organs [25]. lncRNA panels may enhance diagnostic accuracy. EVs provide an enriched liquid biopsy source of miRNAs, proteins, mRNAs, and lipids reflective of liver status. Exosome surface proteins and miRNAs distinguished mild from significant fibrosis in chronic hepatitis C. Better characterization of liver-specific EVs is still needed [26, 27].

Advanced imaging techniques for assessing liver fibrosis and cirrhosis

Liver fibrosis and cirrhosis represent advanced stages of chronic liver disease characterized by excessive extracellular matrix protein accumulation, including collagen. Liver biopsy has limitations including invasiveness, sampling errors, and qualitative scoring. Noninvasive quantitative methods for assessing fibrosis are increasingly of interest for safe, dynamic monitoring. Advanced imaging techniques show promise, including elastography methods, MRI-based techniques, and second harmonic generation microscopy [28, 29].

Elastography methods

Elastography methods assess the mechanical properties of tissues by measuring their response to applied stress. Tissues stiffen with fibrosis since collagen deposition increases elastic modulus. Methods used to assess liver fibrosis include transient elastography, shear wave elastography, and magnetic resonance elastography (MRE). Transient elastography involves inducing a shear wave using a vibrating probe on the skin to measure its velocity, which correlates with tissue stiffness. Higher velocity indicates more advanced fibrosis [30]. It is rapid and validated but can be limited by obesity. Shear wave elastography methods include point shear wave elastography

and 2D shear wave elastography. These induce localized shear waves in the liver to quantify stiffness. 2D methods generate stiffness maps across larger regions. Shear wave elastography accurately stages fibrosis and avoids limitations of ribs/vessels. MRE involves inducing shear waves using a driver on the body wall and imaging propagation using MRI. This generates quantitative maps of tissue stiffness. MRE is highly accurate for staging fibrosis due to sampling larger regions and quantitative measurements. However, MRE requires specialized equipment and expertise. Still, MRE is increasingly used in clinical trials as a noninvasive reference standard [31].

MRI-based techniques

Beyond magnetic resonance elastography, other MRI-based imaging techniques are being developed to non-invasively assess liver fibrosis, including gadoxetate MRI which provides functional information about liver function and fibrosis. Gadoxetate is a liver-specific contrast agent taken up by hepatocytes. Delayed phase gadoxetate MRI identifies alterations in liver function and contrast uptake kinetics associated with chronic liver disease that correlate with increased fibrosis stages [32–34].

Machine learning applied to multiphasic gadoxetate MRI can accurately stage fibrosis by detecting complex hepatic function loss patterns. Gadoxetate MRI also calculates the relative enhancement ratio that reflects hepatocyte uptake function, correlates with worsening fibrosis, and predicts clinical outcomes. Gadoxetate MRI provides concurrent quantitative functional information related to liver fibrosis severity [35].

Second harmonic generation microscopy

Second harmonic generation (SHG) microscopy allows visualization of tissue collagen fibers without exogenous stains by generating optical signals specifically from fibrillar collagen. SHG microscopy of unstained liver tissue can visualize collagen architecture, quantify collagen content, and determine the collagen-to-parenchyma ratio. Second harmonic generation microscopy distinguishes significant fibrosis from mild disease and normal livers. Since collagen remodeling is a key event during fibrogenesis, the second harmonic generation provides direct quantitative information on progression [36–38]. A major advantage is the ability to perform in vivo longitudinal monitoring of liver fibrosis to assess response to therapies. In animal models, SHG microscopy detected early regression of fibrosis reflected by decreasing collagen signals following injury resolution. Potential clinical uses include SHG endomicroscopy to quantify and monitor fibrosis. Limitations include limited depth penetration and sampling variability. Overall, second harmonic generation microscopy enables direct noninvasive

Table 1 Advanced noninvasive technologies for assessing liver fibrosis stage

Technology	Key features	Capabilities	Strengths	Limitations
Transient elastography	Quantifies liver stiffness	Stages fibrosis based on stiffness	Rapid, validated	Limited by obesity, ascites
Magnetic resonance elastography	MRI-based quantification of liver stiffness	Generates quantitative 3D liver stiffness maps	Accurate, samples larger regions	Requires specialized MRI equipment
Second harmonic generation microscopy	Visualizes collagen architecture without stains	Quantifies fibrosis progression/regression by collagen signals	In vivo monitoring capability	Limited depth penetration
miRNA panels	Measure dysregulated fibrosis-associated miRNAs	Distinguishes significant fibrosis, predicts outcomes	High stability in circulation	Variability, need validation
Extracellular vesicle biomarkers	Profile vesicles and cargo reflective of fibrosis stage	Assess functional proteomic/genetic information related to fibrosis	Enriched disease signals from liver-derived EVs	Isolation complexity, liver specificity
Multi-omics signatures	Integrate multidimensional molecular data	Identify prognostic indicators, subclasses	Gain insights not discernible from individual data types	Complex models, validation needs

visualization of liver collagen alterations occurring with fibrogenesis and fibrosis resolution [39]. Table 1 summarizes key parameters, strengths, and limitations of leading noninvasive technologies spanning imaging modalities, circulating biomarkers, and multi-omics data integration for assessing liver fibrosis stage. It provides an overview of evaluating these emerging tools.

Artificial intelligence and big data applications in clinical research

The advent of artificial intelligence (AI) and big data has opened new frontiers in clinical research. In chronic liver disease, AI and big data analytics are being applied in areas including image analysis, biomarker discovery, disease subgroups, and drug discovery [40–43]. A major application of AI is the automated analysis of medical images to detect liver pathology. AI image analysis algorithms can rapidly quantify specific features in a standardized, objective manner. For example, algorithms can evaluate liver collagen as a marker of the fibrosis stage from histopathology slides stained to enhance collagen visualization. Studies have shown AI collagen quantitation strongly correlates with stage and outcomes, outperforming qualitative scoring [44, 45]. Algorithms can also assess additional histological features like cell ballooning and inflammation. Integrating analysis of multiple domain-specific features allows AI assessment of biopsy samples rivaling or exceeding pathologists. Radiomic analysis extracts quantitative data from medical images by evaluating relationships between voxels. Algorithms can discern texture and shape patterns in MRI associated with fibrosis stages. Models identified imaging biomarkers related to the stage with high accuracy. Radiomics has the potential to identify prognostic imaging biomarkers for monitoring. Integrating multidimensional molecular

data is another promising application. Machine learning can integrate genomics, transcriptomics, proteomics, metabolomics, and microbiome data from patients. Analysis can identify molecular signatures differentiating disease subclasses and predict trajectories. For example, a meta-analysis combining serum metabolomics data from over 1000 nonalcoholic fatty liver disease patients revealed distinct subgroups, including increased severe fibrosis risk, and prognostic biomarkers [46]. Harmonizing big data uncovers insights not discernible when analyzing individual datasets. Big data also aids drug discovery through high-throughput screening and machine learning to identify antifibrotic candidates, refine hits in models, and facilitate in silico modeling of drug interactions. AI-enabled discovery expedites the search for promising new liver disease therapeutics [47].

Molecular targets for antifibrotic therapies in liver fibrosis

Extensive research into the cellular and molecular mechanisms driving liver fibrosis has uncovered numerous potential therapeutic targets to halt or reverse fibrosis progression. Key strategies include directly targeting activated hepatic stellate cells, inhibiting proinflammatory and profibrogenic signaling pathways, blocking epithelial-mesenchymal transition, and normalizing extracellular matrix remodeling. Hepatic stellate cells are the predominant collagen-producing myofibroblast-like cells in the fibrotic liver. Selectively inducing stellate cell apoptosis or reverting their activation state represents a direct antifibrotic approach [48]. Inflammation is a major driver of liver fibrosis, making inflammatory signaling pathways like TNF- α , TGF- β , IL-6, toll-like receptor 4, and JAK/STAT key targets. Anti-TNF biologics like infliximab reduce hepatitis and fibrosis in animal models. Blocking TGF- β signaling with neutralizing antibodies or TGF- β

receptor inhibitors effectively inhibits collagen deposition [49]. Oxidative stress generated during chronic liver injury directly activates stellate cells and increases collagen deposition. Antioxidants like vitamin E, superoxide dismutase (SOD) mimetics, and nuclear factor erythroid 2-related factor 2 (NRF2) activators inhibit stellate cell activation in preclinical studies. Imbalances between matrix metalloproteinases (MMPs) that degrade extracellular matrix and their inhibitors TIMPs drive net collagen accumulation in fibrosis. Recombinant MMPs like MMP-8 and MMP-9 degrade fibrotic matrix in preclinical models [50, 51].

Novel interventions for liver fibrosis

As understanding of the molecular mechanisms of liver fibrosis has advanced, numerous therapeutic approaches have emerged to target different aspects of the pathogenesis. These novel antifibrotic interventions include medications, stem cell therapies, RNA therapeutics, artificial liver technology, transplantation, and early-stage interventions undergoing preclinical investigation. Several antifibrotic medications are under clinical investigation after showing promise in animal models. Pirfenidone is an oral agent with anti-inflammatory, antioxidant, and antifibrotic properties. By inhibiting TGF- β and TNF- α , pirfenidone reduced fibrosis progression in idiopathic pulmonary fibrosis (IPF) patients and is undergoing trials for liver fibrosis. The farnesoid X nuclear receptor agonist obeticholic acid improved fibrosis in NASH clinical trials, likely by suppressing stellate cell activation and inflammation. Pentoxifylline, a TNF- α inhibitor, also exhibited antifibrotic effects in patients by reducing inflammation [52]. Cell therapies aim to repopulate the liver with healthy cells or deliver antifibrotic factors. Mesenchymal stem cells (MSCs) are administered intravenously home to injured tissues and secrete immunomodulatory and regenerative factors. MSC therapy alleviated fibrosis in animal models through paracrine effects on stellate cells. Early-stage trials found that MSCs were safe and improved liver function in cirrhosis patients. Co-administration with MSCs may enhance hepatocyte engraftment, while engineered MSCs overexpressing MMPs could boost collagen degradation. Hematopoietic stem cell therapy also improved fibrosis in early clinical studies potentially by promoting hepatic regeneration [53]. RNA interference therapeutics using small interfering RNAs (siRNAs) allow specific silencing of disease-causing genes. siRNAs targeting collagen, Timp1, integrin β 1, and heat shock proteins have shown antifibrotic effects in rodent models. CRISPR/Cas9 gene editing could correct disease-causing mutations or delete pro-fibrogenic genes. Induced pluripotent stem cell-derived hepatocytes offer another cell-based approach, although challenges exist in

maintaining function and integration [54]. Artificial liver support systems aim to bridge patients with acute liver failure to transplant or recovery. The molecular adsorbents recirculating system (MARS) removes circulating albumin-bound toxins using dialysis and adsorption columns. Bioartificial liver devices contain hepatocytes or stem cell-derived hepatocyte-like cells to provide metabolic and synthetic support. While promising, the clinical efficacy of these technologies requires further evaluation [55]. Liver transplantation remains the only curative option for end-stage cirrhosis. However, organ shortages and the need for lifelong immunosuppression limit its applicability. Emerging approaches like hepatocyte transplantation, bioengineered liver grafts, and 3D bio-printed liver tissues could eventually expand curative options [56]. Looking ahead, several preclinical stage interventions show promise for translating into antifibrotic therapies. These include engineered exosomes or nanoparticles for targeted siRNA/microRNA delivery, compounds targeting epigenetic modifiers, and modulating gut-liver axes implicated in fibrosis. Integrating cutting-edge techniques like nanotechnology, induced pluripotent stem cells, and gene editing could enable personalized cell and gene therapies. While significant challenges remain, the expanding antifibrotic pipeline offers hope for treating this major unmet need [57].

Preclinical studies targeting molecular pathways implicated in liver fibrosis

Preclinical research in cell and animal models has enabled extensive investigation into targeting the major molecular pathways that drive liver fibrosis progression. Strategies undergoing active investigation at the preclinical stage include blocking the profibrogenic cytokine TGF- β , suppressing inflammation via cytokines like TNF- α and IL-6, inhibiting oxidative stress through antioxidant activation, and preventing epithelial-mesenchymal transition. The TGF- β signaling pathway is a predominant target for antifibrotic drug development due to its critical role in stellate cell activation and fibrosis. Neutralizing antibodies against all three TGF- β isoforms (TGF- β 1, 2, and 3) demonstrate antifibrotic effects in animal models through reducing inflammation, matrix deposition, and stellate cell activation [58]. Multiple TGF- β receptor inhibitors such as galunisertib, SB525334, and SM305 are in preclinical testing. These compounds blunt liver, lung, kidney, and cardiac fibrosis by blocking receptor-mediated Smad signaling [5, 59–65]. Anti-TNF biologics like infliximab demonstrate antifibrotic efficacy in rodent models by suppressing inflammatory cell infiltration and stellate cell activation. IL-6 neutralization with monoclonal antibodies attenuates stellate cell activation and collagen production in mice. Oxidative stress is another key

driver of fibrogenesis [66, 67]. Antioxidants like vitamin E, SOD mimetics, and mitochondrial-targeted antioxidants reduce ROS production and associated inflammation and matrix deposition in animal studies. Inhibiting epithelial-mesenchymal transition represents another antifibrotic approach. Transcription factor inhibitors targeting Zeb, Snail, and Twist suppress epithelial-mesenchymal transition in vitro, reducing ECM production. Given the key role of inflammation in driving hepatic stellate cell activation and liver fibrosis progression, prophylactic vaccination approaches targeting high-risk inflammatory mediators may hold promise for halting or reversing fibrogenesis in susceptible individuals before significant matrix deposition and architectural damage ensues [68]. Overall, Table 2 concisely summarizes key details on preclinical studies targeting 5 major profibrogenic pathways using various animal models and intervention strategies. It aims to provide a snapshot of cutting-edge antifibrotic approaches under active investigation at the preclinical stage.

Clinical trials of antifibrotic drugs for liver fibrosis

As the complex cellular and molecular mechanisms of liver fibrosis have been unraveled, a growing number of targeted antifibrotic compounds have progressed into clinical testing. Key drug classes reaching clinical trials include pirfenidone, obeticholic acid, and monoclonal antibodies against profibrotic factors and pathways. Pirfenidone is a small-molecule drug with anti-inflammatory, antioxidant, and antifibrotic properties. In cell and animal models, pirfenidone reduced fibrosis by decreasing TNF- α , TGF- β 1, collagen I, and other pro-fibrogenic mediators. In a phase 2 trial for liver fibrosis, pirfenidone treatment for 1 year significantly reduced liver stiffness compared to placebo in chronic hepatitis C patients. Phase 3 trials FAILED-B and REGENERATE evaluating pirfenidone in non-alcoholic steatohepatitis (NASH) patients are ongoing [69]. Obeticholic acid is a synthetic bile acid agonist for the farnesoid X nuclear receptor

(FXR). FXR activation is antifibrotic by inhibiting stellate cell activation and reducing inflammatory cytokine production. In the phase 2 FLINT trial for NASH treatment, obeticholic acid improved histological features including fibrosis. However, adverse effects included pruritis and elevated LDL cholesterol. The phase 3 REGENERATE study confirmed its fibrosis improvement, leading to recent FDA approval for NASH fibrosis [70]. Monoclonal antibody therapies are also emerging as antifibrotics by targeting specific mediators. Simtuzumab is a humanized monoclonal antibody against lysyl oxidase homolog 2 (LOXL2), an enzyme involved in collagen crosslinking. In phase 2 studies, simtuzumab treatment yielded significant hepatic collagen reductions in primary sclerosing cholangitis and NASH fibrosis patients. Another recently evaluated monoclonal antibody is BMS-986036, which blocks lysophosphatidic acid 1 receptor (LPA1) signaling, implicated in myofibroblast activation. Treatment for 15 weeks showed antifibrotic activity by reducing hepatic collagen, TGF- β 1, and α SMA in a phase 2 study of patients with NASH and liver fibrosis [71]. Ongoing and planned trials are evaluating other compounds like emricasan, an apoptosis signal-regulating kinase 1 inhibitor, galectin inhibitors, and cytokine inhibitors. Despite the promise, challenges exist in replicating preclinical efficacy in humans and optimizing delivery specifically to the liver. Combination treatment regimens may augment the modest efficacy of single agents. Further research on bioactive phytoconstituents, targeted delivery, and clinical translation is needed to realize the full potential of nano-herbal antifibrotics [16, 72–76].

Tissue engineering and biomaterials for treating liver fibrosis

Tissue engineering and biomaterials offer new approaches to recreating functional liver tissue and deliver targeted antifibrotic therapies. Key technologies include decellularized liver scaffolds, biomaterial implants for localized drug delivery, and 3D bioprinting

Table 2 Summarizing key preclinical studies targeting major molecular pathways implicated in driving liver fibrosis progression

Target pathway/mediator	Model system	Intervention strategy	Outcomes on fibrosis markers
TGF- β signaling	CCl4 mouse model	Anti-TGF- β neutralizing antibody	↓ Hepatic collagen I/III, ↓ α -SMA, ↓ Fibrosis histology score
TGF- β signaling	Bile duct ligation rat model	TGF- β receptor kinase inhibitor (SB525334)	↓ Collagen, ↓ TIMP-1, ↑ MMPs
Inflammation (TNF- α)	CCl4 mouse model	Anti-TNF monoclonal antibody (Infliximab)	↓ Serum TGF- β 1, ↓ Fibrosis area
Oxidative stress	CCl4 mouse model	Mitochondrial-targeted antioxidant (MitoQ)	↓ Hydroxyproline, ↓ α -SMA immunostaining
Epithelial-mesenchymal transition	In vitro model	Transcription factor Zeb1 siRNA	↓ Collagen 1, ↓ Fibronectin

of liver constructs. Decellularized liver scaffolds leverage the innate extracellular matrix environment of the liver to promote implanted cell survival, differentiation, and tissue remodeling. In rodent models with liver injury, recellularized liver scaffolds improved survival compared to hepatocyte-only transplantation. Implanted cells proliferated, displayed metabolic activity, reduced fibrosis, and integrated with the host tissue. Challenges remain in recapitulating the intricate liver cell populations, perfusable vasculature, and biliary drainage system [77]. Controlled delivery of antifibrotics using biomaterial carriers is another promising approach. Both natural polymers like alginate, chitosan, and collagen and synthetic polymers like PLA, PLGA, and PEG have been formulated into nanoparticles, hydrogels, and implants for sustained drug release in the liver. Targeted delivery using vessels or molecules that are home to liver myofibroblasts can increase bioavailability and reduce off-target effects. Three-dimensional bioprinting of liver constructs offers precise control over the microarchitecture. Bioprinted liver mimics aim to replicate the lobule structure, hepatic zones, and vascular channels. However, challenges exist in printing multiple cell types, vascular integration, and scalable tissue generation. Further advances in bioprinting technology, bioinks, and tissue maturation will help realize implantable engineered livers [78].

Cell therapies for liver fibrosis

Cell-based therapies aim to repair or regenerate damaged tissue and provide beneficial paracrine signals that modulate liver inflammation and fibrosis. Key cell types under investigation include mesenchymal stem cells, fibrocytes, and hepatocytes. Mesenchymal stem cells (MSCs) exert antifibrotic effects through immunomodulation, inflammation resolution, and matrix remodeling. MSCs secrete hepatocyte growth factor, interleukin-10, and other anti-inflammatory mediators that reduce liver injury. In rodent models, MSC transplantation attenuated liver fibrosis by downregulating collagen, smoothing reticular fiber density, and inhibiting TGF- β 1 [58]. Several clinical trials have evaluated autologous MSC infusion in cirrhosis patients. A study found umbilical cord-derived MSC infusion was safe and improved liver function and model for end-stage liver disease (MELD) scores. While MSCs suppress fibrosis, fibrocytes originating from bone marrow appear to contribute to fibrosis progression. Fibrocytes infiltrate injured livers and deposit type I collagen. In mice, antibody-mediated fibrocyte depletion or chemokine receptor blockade ameliorated injury and fibrosis in the liver, pancreas, and other organs. Hepatocyte transplantation aims to restore metabolic and synthetic functions in liver failure as a bridge to recovery or transplantation. Limitations include the availability

of donor cells and poor long-term engraftment. Co-transplanting mesenchymal stem cells improves hepatocyte engraftment in animal models. Bioengineered liver grafts and primary human hepatocyte 3D spheroids show improved engraftment over isolated hepatocytes. Challenges remain in scaling up technologies for widespread application [79].

Liver-on-a-chip platforms for modeling liver disease

Liver-on-a-chip platforms are emerging as advanced *in vitro* models that integrate microfluidics, tissue engineering, and microfabrication to create biochips that recapitulate functional units of the human liver. These miniaturized devices allow precise control of the cellular microenvironment and application of physiologically relevant mechanical forces. Liver-on-a-chip technology enables researchers to investigate liver biology and model liver diseases such as fibrosis with unprecedented ability compared to conventional 2D culture systems [80].

Primary human hepatocytes are considered the gold standard cell type for liver-on-a-chip platforms. Isolation of hepatocytes from donor livers that are unsuitable for transplantation provides a source of highly functional human liver cells. Recent advances have enabled the use of stem cell-derived hepatocytes cultured on chips. Human induced pluripotent stem cells (iPSCs) can be differentiated into hepatocyte-like cells and integrated into liver-on-a-chip devices. Additionally, self-organizing 3D hepatic organoids generated from stem cells represent an exciting option for cellularizing liver chips. These new cell sources provide a scalable, renewable supply of human liver cells [81].

A major application of liver-on-a-chip platforms is modeling the development and progression of chronic liver diseases. The stepwise accumulation of extracellular matrix proteins and scar formation that occurs during liver fibrosis and cirrhosis can be recapitulated on-chip. Researchers developed a fibrosis-on-a-chip platform consisting of iPSC-derived hepatocytes cultured in a collagen hydrogel. By modulating cell–matrix interactions and culture conditions, they induced activation of stellate cells, focal matrix deposition, and progressive scarring reminiscent of early fibrotic remodeling. The biomimetic fibrosis model responded to profibrotic stimuli and known antifibrotic drugs similar to human liver. Such micro-engineered fibrotic tissues open new avenues for investigating mechanisms of fibrogenesis and testing potential treatments [82]. The schematic in Fig. 2A illustrates NAFLD progression mediated by free fatty acids (FFAs) through the gut-liver axis (GLA). The photograph in Fig. 2B shows an integrated gut-liver chip (iGLC) platform with separate perfusion and control layers colored pink and blue. Figure 2C depicts the iGLC

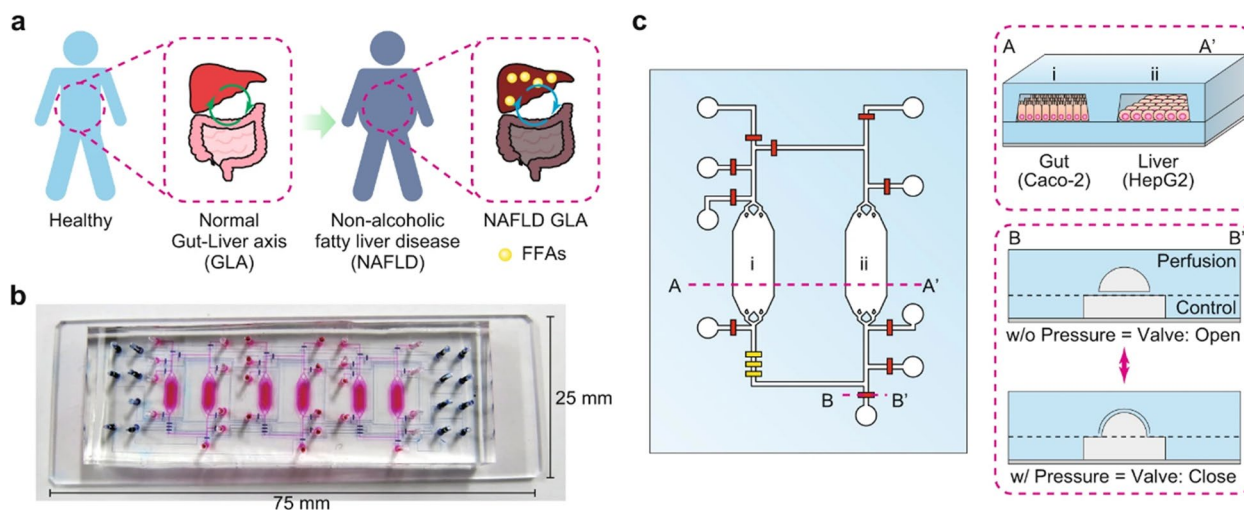


Fig. 2 Microfluidic iGLC platform design to model NAFLD [83]

system used to model NAFLD with Caco-2 gut cells and HepG2 hepatocytes in separate chambers connected by a microfluidic channel and micropump for closed FFA circulation. Integrated microvalves within the chip allow individual cell chamber access without cross-contamination. This iGLC platform enables the investigation of inter-tissue interactions between the gut and liver. Cross-section A-A' shows the Caco-2 and HepG2 cell culture chambers. Cross-section B-B' illustrates open and closed microvalves using an elastic PDMS membrane and differential pressure control [83].

In the future, personalized liver-on-a-chip models derived from patient-specific iPSCs could be used for therapeutic testing to identify optimal treatments for that individual. Integrated sensors could enable continuous monitoring of drug responses. Along with clinical information, data from patient-specific organ chip models could provide an invaluable tool for precision medicine. However, significant research is still required to fully realize the potential of liver-on-a-chip systems. Current challenges include recapitulating the full complexity of human liver physiology, the inclusion of non-parenchymal cells, vascularization, reproducibility, and validation of observations against the human liver [84].

Conclusions

Recent advances have uncovered critical pathways driving liver fibrosis, revealing numerous therapeutic targets. Approaches like pifrenidone, obeticholic acid, and monoclonal antibodies have shown promise in early clinical trials after demonstrating preclinical efficacy. Mesenchymal stem cell therapies also exhibit antifibrotic potential. Non-invasive biomarkers and advanced imaging enable accurate fibrosis diagnosis, staging, and monitoring.

Omics data integration facilitates disease subclassification and prognosis. Liver tissue engineering platforms provide unprecedented human models to study fibrosis and screen therapies. Looking ahead, combination therapies targeting multiple pathways may yield greater efficacy. Further research into biomarkers, tissue engineering, and AI will help translate discoveries into effective antifibrotics. A multifaceted approach harnessing expanding knowledge offers hope for managing fibrosis from all causes.

Recommendations

(1) Evaluate combination regimens in clinical trials; (2) develop additional non-invasive biomarkers; (3) leverage AI for data integration and analysis; (4) engineer biomimetic liver tissues for drug screening; (5) elucidate genetics and epigenetics of fibrosis subtypes. Implementation of these recommendations, alongside continuing research, will accelerate the development of safe, efficacious antifibrotic therapies.

Abbreviations

ECM	Extracellular matrix
TGF- β	Transforming growth factor beta
PDGF	Platelet-derived growth factor
CTGF	Connective tissue growth factor
α -SMA	Alpha smooth muscle actin
HOTAIR	HOX transcript antisense RNA
TIMPs	Tissue inhibitors of metalloproteinases
MMPs	Matrix metalloproteinases
TNF- α	Tumor necrosis factor alpha
IL-6	Interleukin 6
SOD	Superoxide dismutase
NRF2	Nuclear factor erythroid 2-related factor 2
EMT	Epithelial-mesenchymal transition
miRNAs	MicroRNAs
lncRNAs	Long non-coding RNAs
EVs	Extracellular vesicles

MRE	Magnetic resonance elastography
CPA	Collagen proportional area
NAFLD	Nonalcoholic fatty liver disease
HBV	Hepatitis B virus
HCV	Hepatitis C virus
FXR	Farnesoid X nuclear receptor
NASH	Non-alcoholic steatohepatitis
LOXL2	Lysyl oxidase homolog 2
LPA1	Lysophosphatidic acid 1 receptor
MSCs	Mesenchymal stem cells
IPF	Idiopathic pulmonary fibrosis

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Authors' contributions

The authors completed the study protocol and were the primary organizers of data collection and the manuscript's draft and revision process. Tamer A. Addissouky wrote the article and ensured its accuracy. All authors contributed to the discussion, assisted in designing the study and protocol, and engaged in critical discussions of the draft manuscript. Lastly, the authors (TA, MA, IE, YW) reviewed and confirmed the final version of the manuscript.

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Availability of data and materials

All data are available and sharing is available as well as publication.

Declarations

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The authors declare that they have no competing interests.

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