

Review Article

The role of autophagy in hepatic fibrosis

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Abstract: Hepatic fibrosis is a chronic liver injury process, and its continuous development can lead to cirrhosis, hepatic failure and even hepatocellular carcinoma (HCC). Autophagy has attracted much attention because of its controversial role in the course of hepatic fibrosis. In this review, we introduce the mechanism related to noncoding RNAs and some of the signaling pathways that promote or inhibit fibrosis by affecting autophagy. Finally, we list some targets related to autophagy that enable hepatic fibrosis therapy and forecast its prospect in hepatic fibrosis. This review will provide new ideas in diagnosing and treating hepatic fibrosis, which will be helpful to reduce the incidence of cirrhosis and its complications.

Keywords: Hepatic fibrosis, autophagy, hepatic stellate cell, noncoding RNA, exosomes

The development of hepatic fibrosis

Hepatic fibrosis is a chronic liver injury process with many causes, such as long-term alcoholism, viral infection, obesity, and familial hereditary diseases [1, 2]. Studies have shown that various cells participate in the progression of hepatic fibrosis, such as hepatocytes, cholangiocytes, bone marrow-derived cells, and especially hepatic stellate cells (HSCs) [3-5]. In this pathological process, pro-inflammatory and pro-fibrotic factors promote the activation and proliferation of these cells into myofibroblasts. These myofibroblasts can produce excessive extracellular matrix (ECM) [6, 7] and finally lead to the occurrence of hepatic fibrosis, and eventually cause liver cirrhosis, liver failure and even HCC (**Figure 1**).

Previous studies have suggested that hepatic fibrosis is a static process [8]. However, this finding is contrary to that of Dawood, who argues that hepatic fibrosis is a dynamic and reversible process [6]. Subsequently, a study analyzed the role of the retinoic acid signaling pathway in regulating the fibrogenic capacity of HSCs and found that the retinoic acid and PPAR- γ signaling pathways synergistically reverse hepatic fibrosis [9], suggesting that there is a reversible repair process occurring in he-

patic fibrosis. Whether hepatic fibrosis is irreversible or reversible, it is the main pathological process of the irreversible liver diseases, such as cirrhosis and HCC. Thus, it is vital to inhibit the emergence of hepatic fibrosis.

Hepatic fibrosis and autophagy

Previous investigations have shown that autophagy plays an important role in the development of hepatic fibrosis [10]. To date, the role of autophagy in promoting or inhibiting hepatic fibrosis is still controversial. It is believed that, on the one hand, autophagy could participate in the digestion of lipid droplets and provide energy for the activation of HSCs, which plays a direct impact on promoting fibrosis; on the other hand, autophagy could inhibit the emergence of hepatic fibrosis through anti-inflammatory effects [11]. An increasing number of studies have suggested that there is a complex relationship between autophagy and hepatic fibrosis.

Autophagy promotes hepatic fibrosis

TGF- β 1 is considered to be an essential factor in promoting HSC activation [12]. Ye et al. [13] showed that the expression of the autophagy-related protein LC3II/I was increased in HSCs

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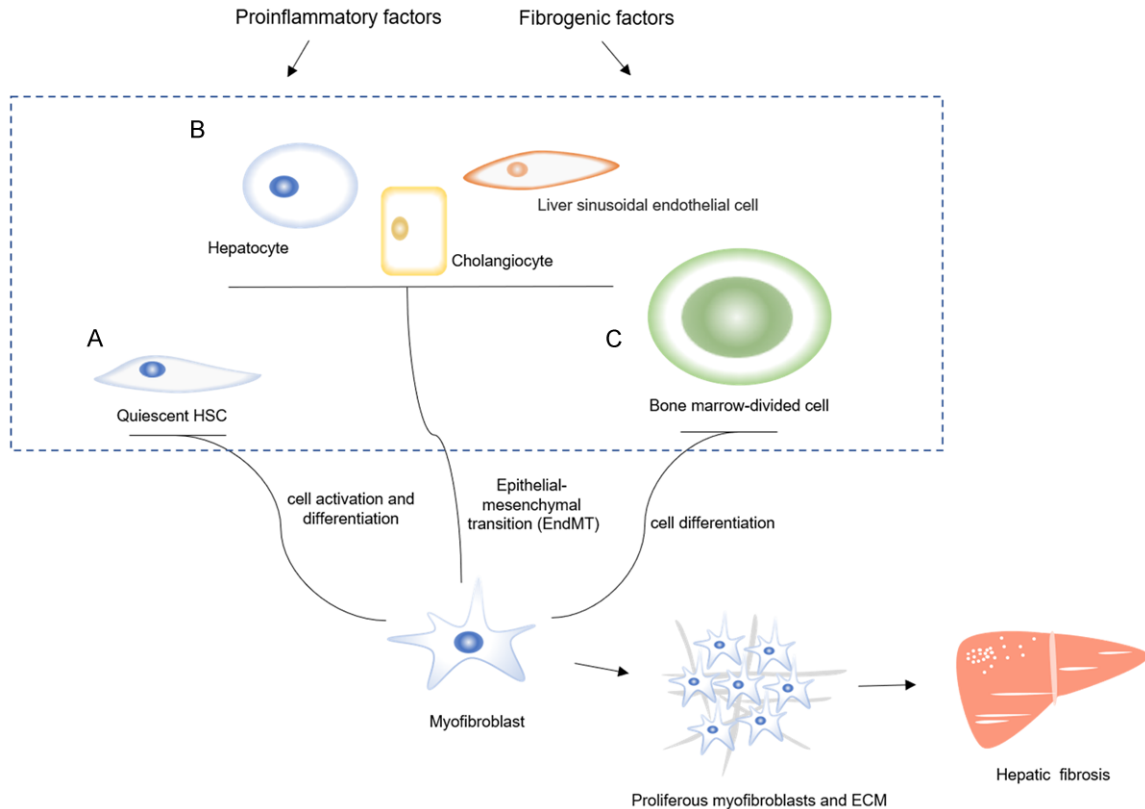


Figure 1. The development of hepatic fibrosis. The pro-inflammatory and pro-fibrotic factors could activate many of the different cell types, which include (A) the activation of quiescent HSCs, (B) the enhanced ability to acquiring myofibroblast characteristics (endothelial-mesenchymal transition, EndMT) in hepatocytes, cholangiocytes and liver sinusoidal endothelial cells, and (C) the differentiation of bone marrow-divided cells, which leads to these cells proliferation into myofibroblasts, and finally the myofibroblasts secrete a large quantity of ECM. This process promotes the hepatic fibrosis.

after TGF- β 1 treatment, which prompted HSC activation to induce autophagy and ultimately accelerate the process of hepatic fibrosis. In addition, a research compared the degree of hepatic fibrosis in mice with ATG5 deficiency and found that the mice without ATG5 deficiency had obvious hepatic fibrosis, which was not observed in the mice with ATG5 deficiency [14]. Therefore, it is speculated that autophagy can promote the emergence of hepatic fibrosis, while inhibiting autophagy may alleviate this disease.

Noncoding RNAs participate in autophagy and promote hepatic fibrosis

The lncRNA/miRNA/autophagy related-gene axis: Studies have shown that DNA methyltransferase 3A (DNMT3A) is involved in the regulation of autophagy and contains binding sites for miR-29b [15-17]. Xie et al. [18] found

that the knockout of lncRNA-SNHG7, which is highly expressed in HCC, downregulated the expression of miR-29b and upregulated DNMT3A, which promoted HSC activation and autophagy. This result suggested that the lncRNA-SNHG7/miR-29b/DNMT3A axis was involved in autophagy-induced hepatic fibrosis. Previous studies found that endothelial cells also had the ability to acquire myofibroblast characteristics (endothelial-mesenchymal transition, EndMT), which further promoted the occurrence of fibrosis [19]. Studies found that the downregulation of lncRNA-Tug1, which has high expression in hepatic fibrosis, increased the level of miR-142-3p and decreased the autophagy gene ATG5, which weakened EndMT and autophagy [20, 21]. This result indicated that the lncRNA-Tug1/miR-142-3p/ATG5 axis is involved in autophagy-induced hepatic fibrosis. According to the previous research, the levels of lncRNA-NEAT1 and the autophagy gene

ATG9a were increased, but miR-29b was decreased in IGFbPrP1-induced hepatic fibrosis [22]. Moreover, lncRNA-NEAT1 also increased HSC autophagy. These results indicate that the lncRNA-NEAT1/miR-29b/ATG9a regulatory axis is involved in IGFbPrP1-induced HSC autophagy and activation. Studies have indicated that lncRNA-XIST is highly expressed in HCC [23]. High-mobility group box-1 (HMGB1) was discovered to induce autophagy [24]. And another research found that the activation of the lncRNA-XIST/miR-29b/HMGB1 axis promoted alcoholic liver fibrosis (ALF) injury and autophagy in the ALF mouse model [25].

MiRNAs/autophagy related-gene axis: MiR-29a is reduced in patients with liver cirrhosis [26]. A study found that overexpressed miR-29a is able to alleviate autophagy by lessening the levels of LC3BII and p-ULK, which leads to reduced liver damage and fibrosis in BDL mice [27]. MiR-30a is an important regulator in fibrosis and is downregulated in NASH [28-30]. There is research pointing out that the overexpression of miR-30a suppresses autophagy by inhibiting Beclin1 directly in TGF- β 1-induced activated HSCs, which leads to the restraint of HSC proliferation and remits the extent of fibrosis [31]. In addition, guanine nucleotide-binding α -subunit (G α) proteins promote G protein-coupled receptor (GPCR) activation, and the activation of autophagic processes requires many ligands of GPCRs that interact with G α 12 [32, 33]. The overexpression of G α 12 activated HSCs and promoted ATG12-5 conjugation, which accelerated autophagy, but this effect was reversed by upregulating miR-16 in HSCs [34]. A previous study showed that miR-223 participated in various types of liver diseases, and the autophagy gene ATG7 is the direct target of miR-223 [35]. In particular, Wang et al. [36] found that exosomes derived from natural killer (NK) cells (NK-Exos) with high miR-223 expression inhibited the activation of HSCs and restrained autophagy by decreasing the level of ATG7.

In conclusion, the expression of these non-coding RNAs is changed in unhealthy livers. Further, noncoding RNAs have the ability to affect the level of autophagy and promote the degree of fibrosis by the lncRNA/miRNA/autophagy-related gene axis or miRNA/autophagy-related gene axis. The discovery of these

noncoding RNAs provides more targets in diagnosing and treating hepatic fibrosis. Interestingly, we found that noncoding RNAs from exosomes mitigate fibrosis, which seems to provide a new method for clinical drug delivery in the treatment of hepatic fibrosis (**Figure 2**).

Signaling pathways participate in autophagy-induced hepatic fibrosis

The TGF- β 1/Smad signaling pathway: TGF- β 1 is a key factor in inflammation, and TGF- β 1/Smad signaling is a major pathway for fibrosis [12, 37, 38]. GNS561, an antifibrotic drug, inhibited LX-2 cell autophagy by destroying lysosomal function, and the phosphorylation of Smad2 (p-smad2) and Smad3 was reduced [39]. Another investigation [40] found that levo-tetrahydropalmatine (L-THP) reduced the formation of ECM and the autophagy biomarker levels of LC3 and Beclin1 by downregulating the TGF- β 1/Smad pathway in CCl4-induced hepatic fibrosis mouse models. Another study discovered that isorhamnetin decreased the deposition of collagen and the expression of Beclin1 in bile duct ligation (BDL)-induced hepatic fibrosis mouse models by inhibiting massive macrophage recruitment and downregulating the TGF- β 1/Smad signaling pathway [41]. Based on these studies, we conclude that inhibition of the TGF- β 1/Smad signaling pathway can inhibit autophagy and ameliorate liver fibrosis.

The NF- κ B signaling pathway: The NF- κ B signaling pathway is widely regarded to participate in inflammation, and there is research indicating that the NF- κ B signaling pathway is involved in the regulation of autophagy to reduce inflammation in the lung [42, 43]. However, studies on the role of NF- κ B in mediating autophagy to regulate hepatic fibrosis are limited. 3-Methyladenine (3-MA) is an autophagy inhibitor that inhibits NF- κ B translocation into the nucleus, decreasing the infiltration of inflammatory cells and the deposition of collagen in CCl4-induced hepatic fibrosis models, which ameliorates hepatic fibrosis [44]. Ghrelin is discovered to suppress the level of LC3 and maintain the balance between matrix metalloproteinases-2 (MMP2) and tissue inhibitor of matrix metalloproteinases (TIMP1) in HSCs by decreasing the expression of NF- κ B, which attenuates hepatic fibrosis [45]. Astaxanthin is an anti-inflammatory medicine and is found to inhibit autophagy by down-

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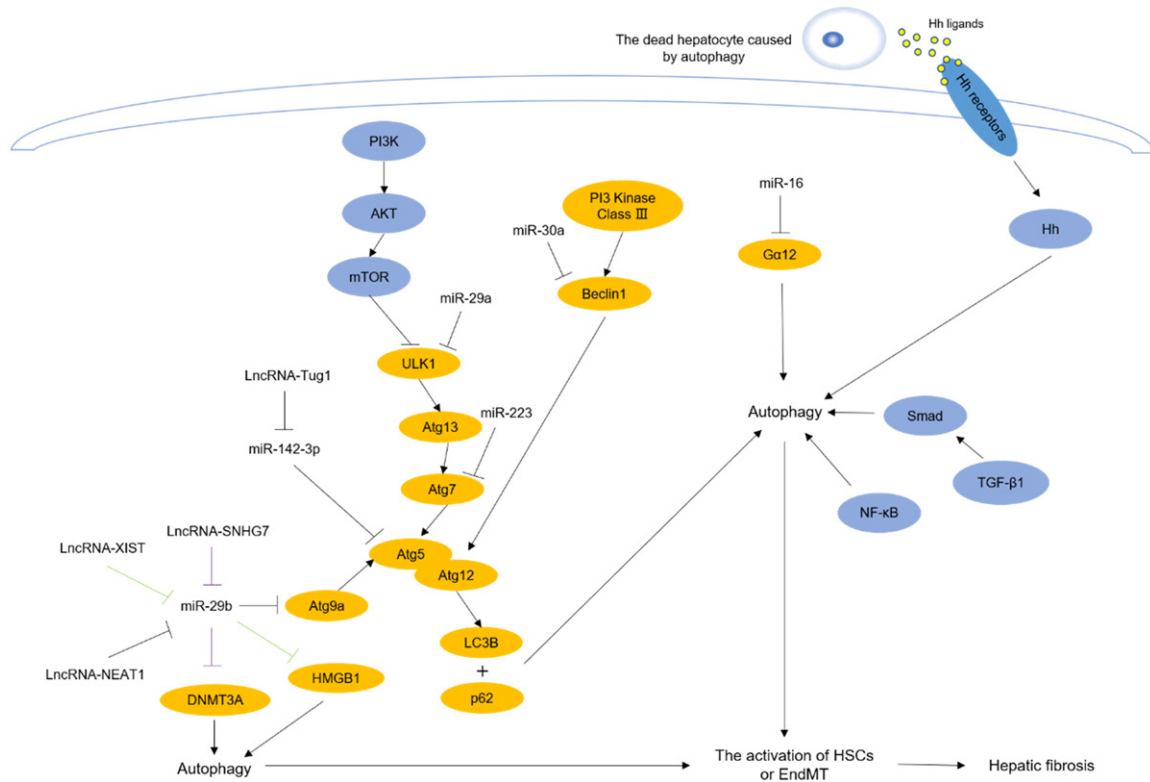


Figure 2. Non-coding RNAs and signaling pathways participate in autophagy-induced hepatic fibrosis. The expression of these non-coding RNAs and signaling pathways are altered in different liver diseases, and they could lead to the autophagy-related genes changed and promote the activation of HSCs or EndMT, which could induce hepatic fibrosis. The pathway marked in purple is LncRNA-SNHG7/miR-29b/DNMT3A, and the pathway marked in green is LncRNA-XIST/miR-29b/HMGB1.

regulating the expression and translation of NF- κ B, which restrains the provision of energy for HSCs, playing a protective effect against hepatic fibrosis [46]. According to these studies, we conclude that downregulation of the NF- κ B signaling pathway can inhibit autophagy and alleviate liver fibrosis.

The PI3K/AKT/mTOR signaling pathway: The PI3K/AKT signaling pathway is associated with various inflammatory diseases [47-49]. As a downstream signaling molecule of this pathway, mammalian target of rapamycin (mTOR) directly and negatively regulates cell autophagy [50]. A growing number of scholars are devoted to exploring the role of PI3K/AKT/mTOR signaling in the relationship between autophagy and hepatic fibrosis. Ginsenoside Rg3 (G-Rg3) reduces the level of p62 and the conversion from LC3a to LC3b in inflammatory inducer lipopolysaccharide (LPS)-induced HSC-T6 cells by enhancing the phosphorylation of PI3K and AKT, which promotes regression from hepatic fibro-

sis through inhibiting the survival of activated HSC-T6 [51]. Quercetin is an anti-inflammatory drug that can inhibit the number of autophagosomes in CCl₄-induced hepatic fibrosis mouse models by activating the PI3K/AKT/mTOR signaling pathway, decreasing the formation of ECM, and preventing hepatic fibrosis [52]. Insulin-like growth factor binding protein-related protein 1 (IGFBP1) is a new profibrotic factor that interacts with TGF- β 1, and it increased the expression of LC3B and Beclin1 in HSCs by downregulating the PI3K/AKT/mTOR signaling pathway [53]. As we have described, the enhancement of the PI3K/AKT/mTOR signaling pathway can inhibit cell autophagy, protect the liver and ameliorate liver fibrosis.

The Hedgehog signaling pathway: The Hedgehog (Hh) signaling pathway is reported to be involved in the activation of HSCs in nonalcoholic fatty liver disease (NAFLD) [54]. There is crosstalk between Hh and mTORC1, which indicates that the Hh signaling pathway is associ-

ated with cell autophagy [55]. Tao et al. [56] proposed that because of autophagy, dead hepatocytes can release a large number of Hh ligands, and those ligands bind to the relevant receptors on HSCs, ultimately activating the Hh signaling pathway and HSCs, leading to the development of hepatic fibrosis. This shows that the inhibition of the Hh signaling pathway could be a new target in the treatment of hepatic fibrosis; however, the study of the Hh signaling pathway in autophagy and hepatic fibrosis is limited.

We found that the aforementioned pathways mostly involve inflammation, and the activation of these pathways is able to enhance cell autophagy, thus inducing hepatic fibrosis. However, the role of the PI3K/AKT/mTOR signaling pathway is different from the others we introduced. This suggests that we could reduce the inflammatory response and inhibit autophagy by regulating these signaling pathways, which could be helpful in relieving hepatic fibrosis (**Figure 2**).

Autophagy inhibits hepatic fibrosis

Although many investigations have shown that autophagy promotes the development of hepatic fibrosis, some studies have pointed out that it could also inhibit hepatic fibrosis. Autophagic flux was used to measure the level of autophagy. An investigation observed the relationship of oroxylin A with hepatic fibrosis, and it was found that this drug increased autophagic flux, inhibited ECM deposition in HSCs, and alleviated hepatic fibrosis in rats induced by CCl₄ [57]. In addition, increased autophagic flux could reduce the IL-1 secreted by Kupffer cells and further inhibited the activation of HSCs [58]. Thus, it is likely that autophagy also participates in the inhibition of hepatic fibrosis.

Noncoding RNA participates in autophagy-inhibited hepatic fibrosis

Contrary to the noncoding RNA that we described above, there are others involved in autophagy-inhibited hepatic fibrosis.

MiR-125a expression is higher in patients with liver cirrhosis than in those with fibrosis [59, 60]. As its target, vitamin D receptor (VDR) has been proven to bind with its ligand 1, 25-

(OH)₂D₃ and adjust autophagy by accommodating Beclin1 expression [61]. He et al. [62] observed that the silencing of miR-125a decreased α -SMA by restoring the level of VDR and autophagic flux in CCl₄-induced fibrotic mice. Deficiency of phosphatase and tensin homolog (PTEN) aggravates oxidative damage in hepatic fibrosis by increasing p62, which indicates that PTEN is associated with autophagy [63]. In addition, miR-20a is boosted in hepatic fibrosis and has a binding site for PTEN [64, 65]. A research observed that decreased miR-20a increases the expression of PTEN and promotes the levels of Beclin1 and ATG7, which reduces the levels of AST, ALT and collagen in hepatic fibrosis mice [66]. Furthermore, exosomes from adipose-derived mesenchymal stem cells (ADSCs) with high expression of miR-181-5P have been found to downregulate the levels of collagen I, vimentin, α -SMA and fibronectin in the liver by enhancing HSC-T6 cell autophagy [67].

In conclusion, we summarized numerous non-coding RNAs that are related to autophagy-inhibited hepatic fibrosis. Similarly, they have an abnormal change in the liver with fibrosis and target these noncoding RNAs by reversing their aberrant expression in the liver might accelerate autophagy and cure hepatic fibrosis (**Figure 3**).

Signaling pathways participate in autophagy-inhibited hepatic fibrosis

The ROS signaling pathways: Because ROS are important signaling molecules that regulate metabolism and inflammation, several studies have suggested that ROS participate in many kinds of inflammation [68-71]. Furthermore, ROS are connected with the early stage of autophagy [72]. There is evidence indicating that dihydroartemisinin (DHA) induces the cellular microenvironment with high ROS, enhances the phosphorylation of JNK1/2, which is the downstream target of ROS, increases autophagic flux in activated HSCs, decreases the secretion of inflammatory factors in the cellular supernatant, such as IL-4 and IL-6, and further inhibits inflammation-induced activation of HSCs [73]. However, in L02 human hepatocytes, glycochenodeoxycholic acid (GCDCA) promoted fibrosis by increasing ROS and inhibiting autophagic flux, which indicates that the low level of ROS

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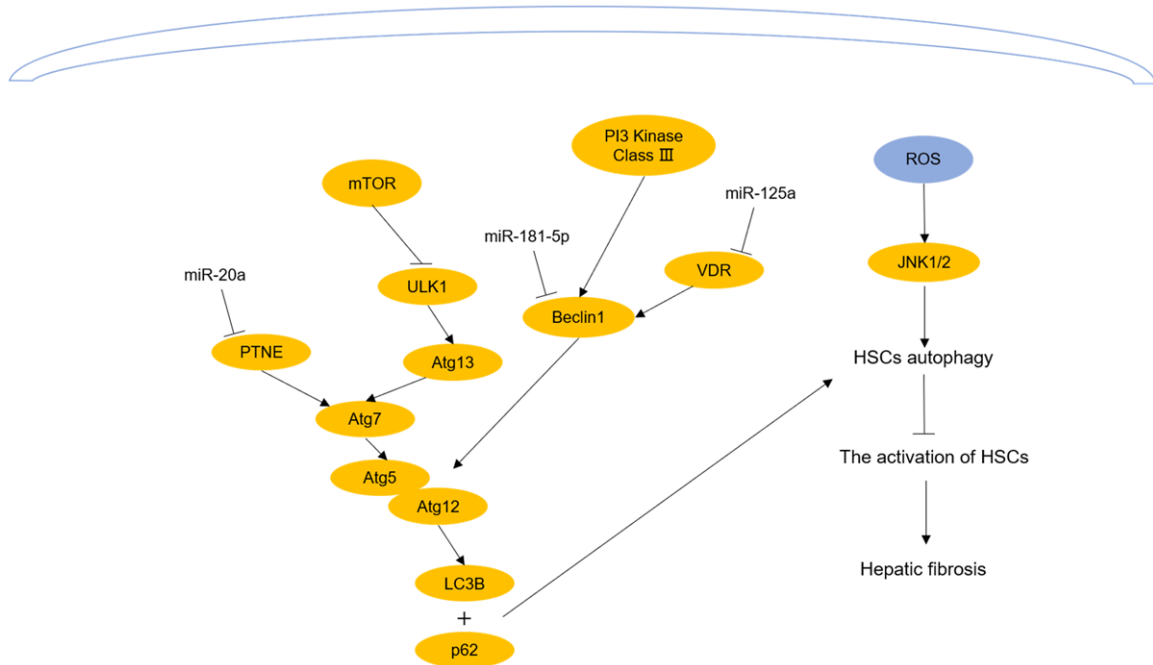


Figure 3. The non-coding RNAs and signaling pathways participate in autophagy-inhibited activation of HSCs. Reversing the expression of no-ncoding RNAs which are altered in different liver diseases or enhancing ROS could promote autophagy and inhibit the activation of HSCs, which could alleviate hepatic fibrosis.

in L02 human hepatocytes relieves hepatic fibrosis by increasing autophagy [74]. It is interesting that the role of autophagy in hepatic fibrosis is related not only to ROS concentration but also to cell type (**Figures 3 and 4**).

The PI3K/AKT/mTOR signaling pathway: Moreover, there is other evidence confirming that the PI3K/AKT/mTOR signaling pathway participates in autophagy-inhibited fibrosis. Kong et al. [75] revealed that curcumin increased autophagic vacuoles in hepatocytes by down-regulating the PI3K/AKT/mTOR signaling pathway, which led to the inhibition of the EndMT in hepatocytes and alleviation of hepatic fibrosis. Another investigation substantiated that the strengthening of hepatocellular autophagy was able to reduce the level of AST in fibrotic mice by inhibiting the PI3K/AKT/mTOR signaling pathway [66]. Unlike studies showing that the PI3K/AKT/mTOR signaling pathway mediates autophagy and promotes the activation of HSCs, this pathway may remit fibrosis by regulating autophagy in hepatocytes (**Figure 4**).

Targets in hepatic fibrosis therapy

At present, there is no effective treatment for hepatic fibrosis, but autophagy can be a new

target. According to previous studies, we can identify some targets related to autophagy in hepatic fibrosis therapy.

Targets in noncoding RNAs

Many noncoding RNAs related to autophagy could be targets in the treatment of hepatic fibrosis.

Downregulation of the lncRNA/miRNA/autophagy-related gene axis, such as the lncRNA-SNHG7/miR-29b/DNMT3A axis, lncRNA-Tug1/miR-142-3p/ATG5 axis, lncRNA-NEAT1/miR-29b/ATG9a axis, and lncRNA-XIST/miR-29b/HMGB1 axis, and upregulation of the miR-30a/Beclin1 axis, miR-16/Gα12, and miR-29a/LC3B are capable of relieving hepatic fibrosis by inhibiting autophagy. The overexpressed miR-223 derived from the exosomes of NK cells also has the ability to treat hepatic fibrosis by inhibiting HSC autophagy.

Moreover, the downregulation of the miR-125a/VDR axis and miR-20a/PTNE axis could play a role in remitting hepatic fibrosis by promoting autophagy. However, the high expression of miR-181-5P in adipose-derived mesen-

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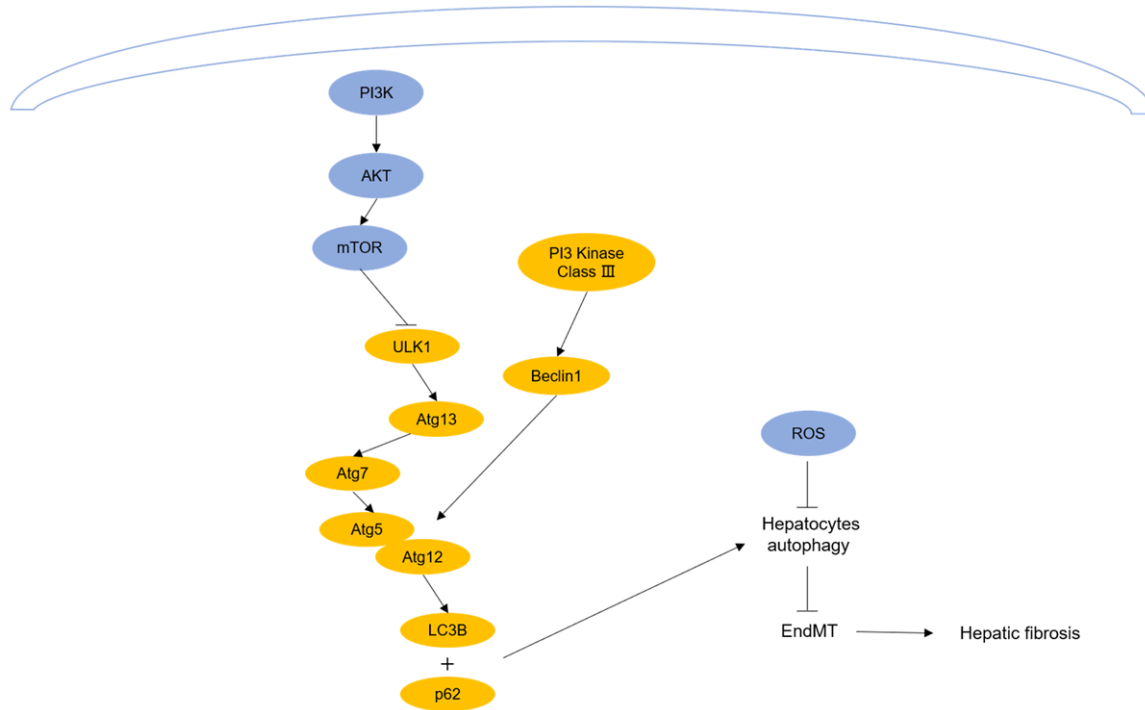


Figure 4. The non-coding RNAs and signaling pathways regulate autophagy and inhibit the EndMT in hepatocytes. Inhibiting PI3K/AKT/mTOR and ROS pathway could promote autophagy and inhibit the activation of HSCs, which could alleviate hepatic fibrosis.

chymal stem cell exosomes could suppress hepatic fibrosis by enhancing autophagy.

Targets in pathways

We found that the downregulation of the TGF- β 1/Smad signaling pathway, NF- κ B signaling pathway and Hedgehog signaling pathway can reduce liver inflammation, inhibit HSC autophagy and further improve hepatic fibrosis. On the basis of the distinction in cell types, we have also discovered that the inhibition of autophagy by upregulating the PI3K/AKT/mTOR signaling pathway is able to restrain the activation of HSCs, but the suppression of this pathway in hepatocytes could inhibit the EndMT or inflammation by enhancing autophagy. In addition, accelerating the ROS pathway in HSCs could facilitate autophagy and relieve hepatic fibrosis, which is contrary to the ROS pathway in L02 cells.

Expectations

Approximately 2,000,000 people die from liver disease each year worldwide, of which 1 million die from cirrhosis and its complications [1]. As

the early stage of liver cirrhosis, hepatic fibrosis is reversible. Therefore, inhibiting hepatic fibrosis has become a way to prevent serious liver disease. However, many investigations indicate that there is a compact relationship between autophagy and hepatic fibrosis, which indicates that autophagy could be a target in curing hepatic fibrosis.

From the aforementioned studies, we found that autophagy plays contradictory roles in the progression of hepatic fibrosis, and it is generally recognized that autophagy not only promotes fibrosis by digesting lipids and providing energy for the activation of HSCs but also inhibits fibrosis by resisting inflammation. Furthermore, we found that the activation of some inflammatory pathways, such as the TGF- β 1/Smad and NF- κ B signaling pathways, increases the partial inflammatory response, activates autophagy, induces the aggravation of liver injury and promotes the occurrence of hepatic fibrosis. However, there are few investigations on signaling pathways related to autophagy inhibiting fibrosis, but these pathways may still be related to inflammation. ROS

are closely related to inflammation, and autophagy plays an antifibrotic role in HSCs with high ROS, whereas it plays an antifibrotic effect in L02 cells with low ROS. Interestingly, the inhibition of the PI3K/AKT/mTOR signaling pathway could activate HSC autophagy and accelerate hepatic fibrosis, yet this pathway also has the ability to facilitate hepatocyte autophagy, reduce the levels of serum AST and ALT, alleviate the degree of liver injury or the EndMT, and ease fibrosis. This suggests that the dual role of autophagy in hepatic fibrosis is not only associated with different signaling pathways but also depends on the type of effector cells.

Moreover, various liver diseases contribute to abnormal changes in different noncoding RNAs, and these noncoding RNAs could regulate the expression of autophagy-related genes directly, alter the level of cell autophagy, and subsequently affect the progression of hepatic fibrosis, especially the noncoding RNAs in exosomes. Exosomes secreted from cells are membranous structures that contain the specific proteins and RNAs from parent cells and transmit intercellular information [40]. In the latest investigations, exosomes have been found to be involved in adjusting the process of hepatic fibrosis. Because exosomes are a part of the human body itself, it will be a new mode of administration in antifibrosis, which could increase the bioavailability of the drug.

Liver biopsy is the gold standard for the diagnosis of hepatic fibrosis, but it does not have extensive application due to the invasiveness and cost, and there is still a lack of effective drugs to treat hepatic fibrosis, which creates a serious challenge in diagnosing and curing hepatic fibrosis. Although autophagy may have two distinct effects on hepatic fibrosis, an increasing number of studies indicate that autophagy could be a new target in the diagnosis and treatment of hepatic fibrosis, which will improve grievous clinical problems in the liver.

Disclosure of conflict of interest

None.

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- hepatic stellate cell activation by suppressing autophagy. *Mol Med* 2020; 26: 81.
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