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# **Exosomes in liver pathology**

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# Abstract

Exosomes are small (~100 nm) membrane-bound extracellular vesicles released by various types of cells into biological fluids. They contain proteins, mRNAs and miRNAs as cargo. Different cell types can take up exosomes by endocytosis and the cargo contained within them can be transferred horizontally to these recipient cells. Exosomal proteins and miRNAs can be functional and regulate physiological cell events modifying the microenvironment in target cells, a key event of liver pathology. Exosome-mediated cell-cell communication can alter tumor growth, cell migration, anti-viral infection and hepatocyte regeneration, indicating that exosomes have great potential development as diagnostic or therapeutic tools. Analyses of circulating total or exosomal miRNAs have identified a large number of candidate miRNAs that are regulated in liver diseases, and the diagnostic testing using single or multiple miRNAs shows good sensitivity and specificity. Some candidate miRNAs have been identified to play an important role in various liver disorders. This review summarizes recent findings on the role of extracellular vesicles in liver diseases and their diagnostic and therapeutic potential, mainly focusing on exosomes but also includes microvesicles in liver pathology.

# Keywords

exosomes; microvesicles; microRNAs; liver disease; cancer

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# Introduction

Exosomes are one of three types of extracellular vesicles (EVs) secreted in a wide variety of cells. EVs are spherical particles enclosed by a phospholipid bilayer. There are three types of vesicles according to their size: exosomes (40~100 nm), microvesicles or microparticles  $(0.1-1 \,\mu\text{m})$  and apoptotic bodies  $(1-4 \,\mu\text{m})$  [1]. Exosomes are derived from multivesicular bodies (MVBs), which are the type of endosomes containing membrane-bound vesicles. These vesicles form by budding into the lumen of the MVB, and some MVBs are degraded in lysosomes. Some MVBs, however, are fused with the plasma membrane releasing internal vesicles as exosomes via exocytosis. Microvesicles are released directly from plasma membrane of cells by outward blebbing without MVBs. EVs have been the topic of great interest in recent years in medical research. In particular, EVs are of great interest in liver pathology because they regulate cell-cell communication and a number of pathophysiological events in various types of cells via horizontal transfer of their cargo [2]. EVs contain a variety of cargoes including proteins, mRNAs and microRNAs (miRNAs) and they can be transferred from donor cells to recipient cells and can activate or regulate cell activities such as protein expression, cell proliferation and differentiation or antiviral responses.

miRNAs are small non-coding RNAs which regulate gene expression by binding to target mRNAs and interfering with their translation (Figure 1). Therefore, cargoes of EVs especially miRNAs have attracted research interests in recent years. It is important to study EVs and their functions to understand mechanisms of liver disorders. More importantly, EVs can be used as specific biomarkers to diagnose liver diseases. This review summarizes latest liver cell derived-EV research mainly focusing on exosomes but also includes microvesicles in liver pathology.

# Circulating EVs as biomarkers of liver diseases

Searching for a novel biomarker of liver diseases has been a challenge for researchers. The isolation of EVs, however, may be a potential approach due to the unique molecules such as proteins and miRNAs could be transported only in exosomes or microvesicles. Sugimachi et al has compared expression profile of miRNAs in serum exosomes between patients with and without hepatocellular carcinoma (HCC) recurrence after liver transplantation and found that miR-718 is significantly decreased in exosomes of patients with HCC recurrence [3]. A study comparing 30 patients with HCC and 30 patients with chronic hepatitis B virus (HBV) infection has demonstrated that serum exosomes from HCC patients contain a higher level of miR-21 [4]. Another study recruiting 20 HCC patients and 20 HBV patients has reported that miR-18a, -221, -222 and -224 are increased and miR-101, -106b, -122 and -195 are decreased in serum exosomes from HCC patients although no significant difference was observed for miR-21 [5]. Microvesicles may also have a potential for a novel biomarker. A study comparing patients with chronic hepatitis C virus (HCV) infection with healthy individuals has shown that patients with HCV secrete increased levels of microvesicles derived from CD4+ and CD8+ T cells [6]. Although further studies are required to identify a

unique biomarker that is specific in the certain liver disorder, secreted EVs in positive patients may be a good candidate for identification or diagnosis for liver diseases.

# **Exosomal alteration in liver diseases**

## **Drug-induced liver injury**

As EVs can be used as a novel biomarker, several groups have performed profiling exosomal cargoes secreted in liver diseases. A recent study has analyzed exosomal protein levels in urine of rats with drug-induced liver injury (DILI) and reported that exosomes in DILI rat urine contain reduced exosomal proteins, including CD26 and CD81 and candidate marker proteins compared with exosomes in control rat urine [7]. Another study has demonstrated, however, that exosomes isolated from blood serum of rats with DILI contain higher expression levels of exosomal proteins, including HSP70 and HSP90 and other candidate marker proteins although Alix, endosome-associated protein is decreased compared with exosomes from healthy rats [8]. This study also demonstrated that protein expression profiles differ between liver extracts and circulating exosomes isolated from blood serum. These observations indicate that a biomarker for liver diseases could be identified only in exosomal cargoes isolated from specific body fluids and may not be identified in liver tissue or serum. In fact, another study has shown that total miRNA concentrations are increased in plasma of porcine models of DILI compared with control pigs but this enhancement is not observed in isolated exosomes from plasma [9].

#### Alcoholic and non-alcoholic fatty liver disease

Alcohol-fed mice were found to secrete significantly increased amounts of exosomes in the blood, and miRNA array has shown increased levels of candidate miRNAs in the exosomes of alcohol-fed mice [10]. Furthermore, previous studies have demonstrated that human monocytes, precursors of macrophages, release significant amounts of exosomes when exposed to ethanol, and these exosomes can stimulate naïve monocytes to polarize and differentiate into M2-macrophages. Monocytes exposed to alcohol also secrete exosomes containing increased levels of miR-27a leading to increased cytokine secretion, followed by activation and polarization of other monocytes [11, 12]. Alcohol treatment also induces EV release from hepatocytes, which is associated with the activated caspase-3. Additionally alcohol-induced EVs contain CD40 ligand that promotes macrophage activation followed by inflammation in alcoholic liver disease [13].

Patients with non-alcoholic fatty liver disease (NAFLD) or non-alcoholic steatohepatitis (NASH) secrete increased levels of microvesicles derived from macrophages and natural killer T cells [6]. Another EV study has shown that the expression levels of various proteins within vesicles are enhanced in a mouse model of NAFLD, and that protein expression pattern differs between exosomes and microvesicles [14]. Another study has demonstrated that lipids induce EV release from primary hepatocytes and the Huh7 cells (well differentiated hepatocyte derived cellular carcinoma cell lines) and those secreted EVs activate an inflammatory response in macrophages leading to NASH [15]. These findings suggest that cells under disease conditions may secrete elevated amounts of EVs containing unique cargoes, and this could be a trigger for various pathophysiological events.

#### Cell-cell communication via exosomes

There is strong evidence that EVs can play an important role in cell-cell communication by transferring proteins and miRNAs from the source to the target cells (Figure 2). Such communication allows a cell to influence neighboring cells and alter the liver's microenvironment. A study has also demonstrated that exosomes secreted from mouse and human mast cell lines can be taken up by other mast cells, and their cargos, mRNAs and miRNAs are still functional in recipient cells *in vitro*, which means that transferred exosomal mRNAs can be translated and the new proteins could regulate physiological events in recipient cells [16].

#### Liver cancer

Recent studies have revealed that cancer cells can modulate surrounding hepatic environments to aid their growth, proliferation and invasion. Exosomes containing mRNAs, miRNAs and natural antisense RNAs secreted by colorectal cancer cells can be taken up by other cancer cells such as hepatoma and lung cancer cells in vitro [17]. Human HCC cells can secrete exosomes that contain a variety of miRNAs, which attenuate protein expression of the TGF- $\beta$  pathway in other HCC cells. As a result, growth of HCC cells is increased dramatically with HCC cell-derived exosomes [18]. Another study using an immortalized non-tumorigenic hepatocyte cell line, MIHA has demonstrated that non-motile MIHA cells show significantly higher cell migration and invasion after incubated with exosomes isolated from HCC cells [19]. Similar results are also observed in studies with cholangiocarcinoma (CCA). A study using CCA cells derived from human patients were shown to secrete exosomes that increased migration and invasion of human cholangiocyte cell line H69 [20]. The exposure of mesenchymal stem cells to CCA cell-derived exosomes enhance their migratory capability and the release of cytokines such as CXCL1, CCL2 and IL-6 leading to cell proliferation [21]. These results suggest that malignant cells can co-operate with other cancer cells or even "educate" non-cancer cells for their cell growth, migration and invasion by modifying microenvironment by secreting exosomes.

#### Viral hepatitis

Exosomes and intercellular communication may also be involved in immune regulation and antiviral response during viral infection. It is known that virus-infected cells can secrete exosomes, which contain virus-derived miRNAs leading to suppression of target genes in recipient cells [22]. Exosomes isolated from HCV-infected patients' serum contain HCV RNA and elevated levels of proteins Ago2 and HSP90 helping viral receptor-mediated transmission to hepatocytes [23]. Virus-infected cells release type I and III interferons (IFNs) that are a cytokine to stimulate immune system and antiviral responses [24]. Liver cells stimulated by type I IFN, IFN- $\alpha$  secrete exosomes which contain antiviral molecules and can attenuate hepatitis B virus (HBV) replication [25]. Exosomes isolated from liver endothelial cells stimulated by either type I or III IFNs can also suppress viral replication in HCV-infected liver cells [26]. These findings suggest that both viral infection and antiviral response are mediated by cell-cell communication through exosomes.

## Exosomes as therapeutic tools for liver diseases

Exosomes can modulate microenvironment not only to support cancer cell proliferation or viral transmission, but also to facilitate cell regeneration and recovery to maintain homeostasis. Mesenchymal stem cells (MSCs) provide important approach for the therapy of liver disease. For example, direct injection of exosomes isolated from human MSCs into mouse liver alleviates carbon tetrachloride-induced liver fibrosis by suppressing collagen and TGF-\beta1 expression *in vivo* [27]. Induction of cellular senescence in activated hepatic stellate cells can reduce the secretion of extracellular matrix components and subsequently inhibit liver fibrosis. We have demonstrated that human MSC derived extracellular vesicles induced cellular senescence in cultured human hepatic stellate cells (Figure 3). Intrasplenic injection of MSC-derived exosomes can enhance protein expression including PCNA and cyclin D1 leading to liver recovery, and alleviate liver damage caused by carbon tetrachloride administration in mice [28]. Primary hepatocytes as well as MSCs can also enhance liver cell proliferation. Hepatocyte-derived exosomes induce hepatocyte proliferation in vitro while exosomes isolated from Kupffer cells or endothelial cells have no effect [29]. Furthermore, the cargo of hepatocyte-derived exosomes induces generation of sphingosine-1-phosphate leading to hepatocyte regeneration and liver repair after ischemia/ reperfusion injury in vivo. Endothelial cell-derived exosomes can regulate hepatic stellate cell signaling and migration via sphingosine-1-phosphate [30]. These studies indicate that exosomes and their components are involved in mechanisms of liver cell regeneration and migration, and hence they can be a potential therapeutic tool for liver disease and damage.

Microvesicles may also be used as a therapeutic tool. Microvesicles derived from liver stem cells inhibit HCC cell growth and this is mediated by various miRNAs contained in those microvesicles [31]. An intravenous anesthetic propofol has antitumor effects, and it has been shown that mice with propofol administration secrete macrophage-derived microvesicles and those microvesicles and their cargo miR-142-3p significantly inhibits tumor growth *in vivo* [32]. In liver cancers such as HCC, malignant cells can not only modify microenvironment to help their proliferation and invasion, but also be controlled by other cells via cell-cell communication using exosomes and microvesicles (Figure 4).

# Diagnosis of liver disease using circulating proteins and miRNAs

Diagnosis of liver disease can be challenging, especially for chronic liver diseases, because of its progression through several stages: hepatitis  $\rightarrow$  fibrosis  $\rightarrow$  cirrhosis  $\rightarrow$  HCC, and this can take 5–50 years to attain the advanced disease stages or the progression may be arrested in the early stage with no obvious symptoms. Diagnosis of HCC, for example, is generally based on imaging including ultrasound, MRI or CT scan, or elevated serum  $\alpha$ -fetoprotein level although it can produce false diagnoses and it is not very feasible to discriminate the specific stages such as HCC from chronic HBV infection or cirrhosis. As early diagnosis of HCC can improve the survival rate, scientists have been making an effort to search a novel diagnostic method to discriminate liver disease stages, and circulating proteins or miRNAs have attracted great interests as a tool of diagnosis.

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Table 1 shows selected diagnostic studies with circulating miRNAs performed for human patients with liver disease. These latest studies have usually focused on total circulating miRNAs including exosomes, microvesicles and free miRNAs. Some studies have isolated exosomes before miRNA analyses but only limited number of studies is available although exosomes are a main component of EVs circulating in the serum [10]. Majority of recent studies has used a miRNA panel with ~10 miRNAs for diagnostic analyses to increase sensitivity and specificity. Although single miRNA can be useful to discriminate HCC from healthy individuals with good sensitivity and specificity, miRNA panels may be more reliable to discriminate more sensitive cases such as HCC patients from non-HCC individuals who are infected by HBV or HCV and showing chronic liver disease or cirrhosis but not HCC (Table 1). A meta-analysis studying 13 publications has shown that a miRNA panel analyzing multiple miRNAs can produce better accuracy than single miRNA diagnosis for those cases [33].

# Strong candidate biomarkers for liver diseases

Exosomes contain diverse specific types of proteins/RNAs/miRNAs reflective of cell functions and conditions. The increasing number of exosomal proteins/RNAs/miRNAs have been found to be potential biomarkers for a variety of diseases including acute and chronic liver diseases. Studies for identification of candidate biomarkers in circulating total or exosomal miRNAs have reported a large number of miRNAs to date, but some miRNAs have been identified in several studies showing strong evidence that those miRNAs are involved in the specific liver disorders.

#### miR-122

miR-122 is the most frequently identified and well-studied miRNA in liver diseases. It is the major miRNA in liver, and decrease of exosomal miR-122 expression has been observed in HCC [5]. miR-122 can alleviate tumor growth and miR-122 knockdown upregulates HCC cell viability by inhibiting apoptosis in vitro suggesting that miR-122 may be a key for tumor proliferation and migration via exosomal intercommunication and hence could be a good therapeutic or diagnostic target for HCC. [34].

Whereas reduced levels of miR-122 are observed in HCC, they are, in contrast, elevated in alcoholic liver disease. In a rodent model of chronic alcohol feeding, elevated serum miR-122 levels were found to correlate with liver damage [35]. Alcohol drinking induces gut bacteria permeability and endotoxin (also known as lipopolysaccharide, LPS) elevation in blood. Hepatic cells exposed to alcohol secrete exosomes which contain elevated miR-122, and these exosomes are taken up by macrophages and sensitize them to LPS leading to enhanced cytokine secretion in vitro [36]. These current studies strongly support that this miRNA plays an important role in liver pathology but also suggest that even same miRNA can be expressed differently depending on liver diseases.

#### miR-21

miR-21 is another candidate exosomal biomarker identified in several liver disease studies. Previous studies indicate that miR-21 may have opposite functions to miR-122. In HCC,

miR-122 expression is lower in serum or circulating exosomes of positive patients compared with healthy individuals, but miR-21, on the other hand, is higher in HCC patients [37]. Although miR-122 knockdown inhibits apoptosis and increases HCC cell viability *in vitro* [34], miR-21 knockdown enhances apoptosis and decreases cell viability [38]. A study using human CCA specimens has revealed elevated miR-21 expression in the CCA tissue compared with noncancerous biliary epithelium. This study has also demonstrated that overexpression of miR-21 promotes CCA growth by targeting the key enzyme for prostaglandin conversion, 15-hydroxyprostaglandin dehydrogenase (15-PGDH) [39]. These studies support the notion that miR-21 may be another strong candidate biomarker in liver pathology.

#### Other miRNAs

miR-192 is often identified from patients with liver diseases as well as miR-122 and miR-21. Although only limited number of studies is available to date and functions of miR-192 in liver pathology are largely unknown, a study has revealed that miR-192 suppresses metastasis of HCC cells *in vivo* and *in vitro* by targeting an oncogene involved in different types of cancer, SLC39A6 [40].

Also, miR-155 is one of miRNAs which are associated with regulations of inflammation and its expression levels are elevated in the serum and peripheral monocytes of patients with chronic HCV infection [41]. A study using miR-155 knockout mice has shown that administration of exosomes containing miR-155 mimic leads to miR-155 expression in isolated hepatocytes of recipient knockout mice indicating its cellular uptake [42]. Another study using miR-155 knockout mice has demonstrated that miR-155 deficiency attenuates hepatic steatosis and fibrosis, but not liver injury and inflammation induced by methionine-choline-deficient diet [43]. These earlier studies suggest that some candidate miRNAs may be useful to understand mechanisms and to seek a novel biomarker or treatment in liver pathology.

# Conclusions and perspectives

Despite the progress in research for liver diseases to date, diagnosis of liver pathology remains a difficult task. Because methods such as imaging are not completely reliable, invasive liver biopsy continues to be a gold standard for diagnosis. Thus the development of novel, reliable and noninvasive diagnostic method has been of major interests. Recent research indicates that EVs and their cargoes may have a great potential as biomarkers for the diagnosis of liver pathology. Current studies suggest that circulating miRNAs and proteins may be a useful tool for diagnostic testing and in fact, especially miRNA panels show good sensitivity and specificity against various liver disorders. Figure 5 summarizes potential achievements in EV research. However, the current miRNA-based diagnoses have some limitations. First, researchers have a large number of options for EV studies: disorder, circulating miRNAs or local miRNAs in specific tissues, body fluid, and exosomes, microvesicles or total miRNAs and experimental conditions. Biomarkers could differ depending on liver diseases and even same miRNA may be expressed differently in a different disorder. miR-122 is increased in exosomes of alcohol-fed mice but decreased in

exosomes of HCC patients. Chronic alcohol feeding increases miR-122 in exosomes but decreases in liver tissues. Exosomal cargoes may differ between serum exosomes and urine exosomes. Although exosomes constitute majority of extracellular vesicles, cargoes may differ between exosomes and microvesicles and specific biomarker could exist in specific EVs. Moreover, careful attention should be paid to experimental conditions involving EV research for reliable and reproducible results. The secretion levels of EVs and protein composition in their cargoes can vary depending on culture medium with or without serum in vitro [44]. Second, although numerous miRNAs have been identified in samples from liver diseases, their functions and roles in liver pathology are largely unknown. Functional analyses of exosomal cargoes may be important in understanding the mechanisms of liver pathology and/or to identify therapeutic targets. Identification of donor and recipient cells for cell-cell communication in the specific liver disease and underlying mechanisms also need to be further clarified.

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# Abbreviations

CCA	cholangiocarcinoma
DILI	drug-induced liver injury
EV	extracellular vesicle
MVBs	multivesicular bodies
HBV	hepatitis B virus
НСС	hepatocellular carcinoma
HCV	hepatitis C virus
IFN	interferon
LPS	lipopolysaccharide
miRNA	microRNA
MSC	mesenchymal stem cell
N-Heps	normal human hepatocytes
MVB	multivesicular bodies
NAFLD	non-alcoholic fatty liver disease
NASH	non-alcoholic steatohepatitis

**PGDH** 15-hydroxyprostaglandin dehydrogenase

Pre-miRNA miRNA precursor

Pri-miRNA primary transcript of miRNA

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Author names in bold designate shared co-first authorship.

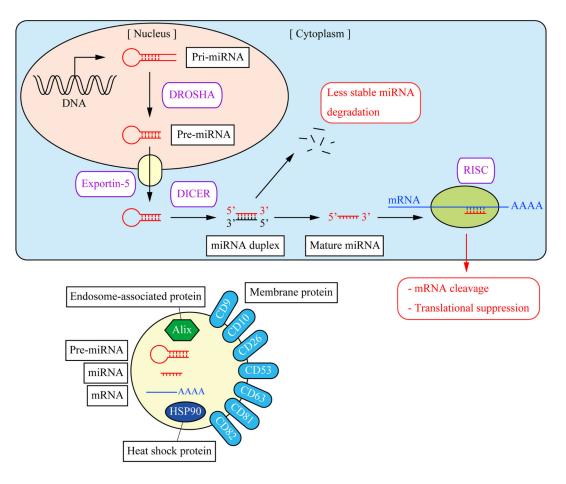
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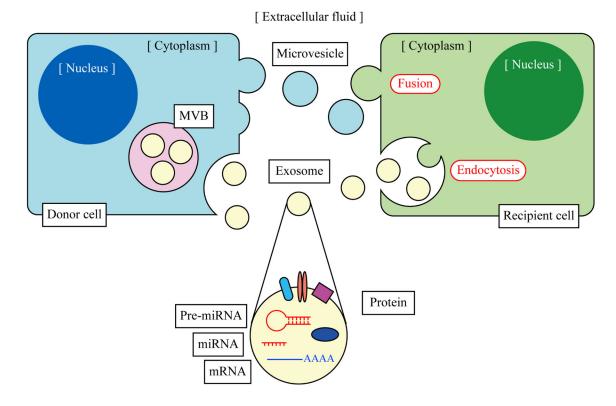
#### Key points

- Exosomes are small membrane-bound vesicles secreted by various types of cells
- Exosomes contain proteins, mRNAs and miRNAs as their cargoes and can transfer them horizontally to the other type of cells
- Exosomal cargo can be functional after being taken up by the recipient cell and regulate physiological events
- Exosomal miRNAs are associated with liver pathology and have a great potential as a therapeutic target or as a molecular biomarker which may be useful for diagnostic testing



#### Figure 1. miRNA synthesis and contents of exosomes

The pathway of miRNA synthesis and its function (top). In the nucleus, miRNAs are transcribed from DNA forming a primary transcript, pri-miRNA. The RNase III enzyme DROSHA cleaves pri-miRNA to produce a precursor, pre-miRNA that is transported to the cytoplasm through a nuclear export protein Exportin-5. In the cytoplasm, pre-miRNAs are cleaved by DICER to form a miRNA duplex. In general, only one strand of the miRNA duplex is used as the mature miRNA while the other strand is degraded because of less stability. Mature miRNA binds to the 3'UTR of target mRNA and is loaded into RNA-induced silencing complex (RISC). As a result, this complex inhibits the process of mRNA translation or enhances mRNA degradation leading to translational suppression. Contents of exosomes (bottom). Typical exosomes contain various cargoes including pre-miRNAs, mature miRNAs, and mRNAs as well as membrane marker proteins, endosome-associated (MVB-derived) proteins, and heat shock proteins (chaperones).



#### Figure 2. Cell-cell communication via exosomes and microvesicles

Exosomes are incorporated into vesicles in MVBs by budding into their lumen. MVBs are fused with the plasma membrane releasing internal exosomes to outside the cell by exocytosis. Microvesicles are formed and released directly from the plasma membrane by outward blebbing. These EVs are taken up by the other cell by fusion with the plasma membrane or endocytosis and EV cargoes are transferred horizontally from the donor cell to the recipient cell. EVs can contain various molecules including mRNAs, miRNAs, pre-miRNAs and proteins. These EV cargoes regulate physiological cell events in the recipient cell. Some EVs are circulating through the body in extracellular fluid and their cargoes can be useful as a biomarker to diagnose liver diseases.

# SA-B-Gal activity

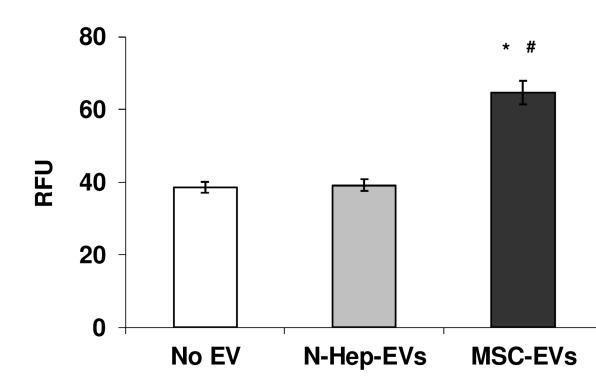
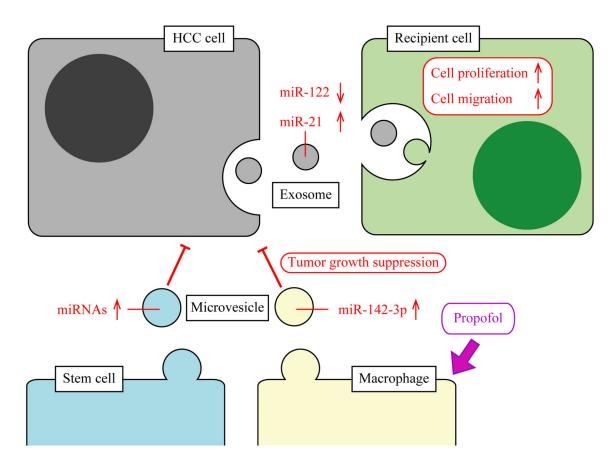


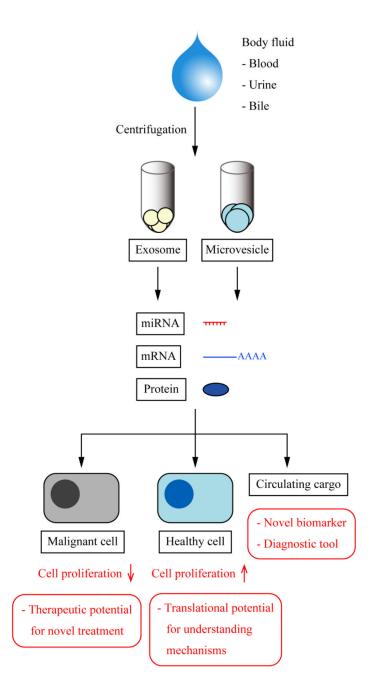
Figure 3. Enhanced Cellular Senescence in Cultured Human Hepatic Stellate Cells treated with MSC derived extracellular vesicles (EVs)

Human hepatic stellate cells (ScienCell Research Laboratories, Carlsbad, CA) were incubated with 30 µg/ml MSC derived EVs or normal human hepatocytes (N-Heps) derived EVs for 48 hours and compared to no EV treatment controls. Lysates were allowed to incubate with SA- $\beta$ -Gal Substrate for 1 hr at 37°C. SA- $\beta$ -Gal activities were measured using 96-Well Cellular Senescence Assay Kit from Cell Biolabs, Inc (San Diego, CA). Human MSCs and N-Heps were also purchased from ScienCell Research Laboratories. Significantly increased SA- $\beta$ -Gal activities were observed in cultured human hepatic stellate cells incubated with MSC-EVs when compared to no EV or N-Heps-EVs controls. \* p<0.05 relative to no EV group; # p<0.05 relative to N-Heps-EVs group.



#### Figure 4. Exosomes and microvesicles in hepatocellular carcinoma

HCC cells secrete exosomes that can be taken up by other HCC cells leading to enhancement of cell proliferation, migration and invasion in the recipient cell. Previous studies have shown that exosomes secreted from HCC cells contain decreased and increased levels of miR-122 and miR-21, respectively. Liver stem cells secrete microvesicles which contain elevated levels of various miRNAs. An anesthetic propofol induces macrophages to secrete microvesicles which contain elevated miR-142-3p. These microvesicles can inhibit HCC cell growth suggesting that microvesicles could be used as a therapeutic tool for liver cancer.



#### Figure 5. Pathological potentials in EV research

EV research could contribute to liver pathology. Major EVs, exosomes and microvesicles can be isolated by centrifugation from various body fluids. Circulating EV cargoes could be analyzed as disease-specific biomarkers leading to a novel non-invasive diagnostic method. EVs can be taken up by other liver cells and modify microenvironment by enhancing cell proliferation and migration, and this could lead to better understanding of pathophysiology of liver disorders. EVs could also be used as a tumor suppressor suggesting that EVs have a therapeutic potential to treat liver cancer.

Table 1

Selected studies for diagnosis of liver disease using human circulating miRNAs

HCCA field thy73910 $mR.143$ Seum95127Healthy80910816Up $mR.143$ Seum95127Healthy80910816Up $mR.215$ Seum3060Non-HCC87400.653Up $mR.124$ Seum11242Healthy82960.752Up $mR.121$ Seum6761Cirthosis96900.752Up $mR.Napanel2$ Seum10378Cirthosis82850.892Up $mR.Napanel3$ Seum10378Cirthosis82850.803Up $mR.Napanel3$ Bile4650Healthy86640.803Up $mR.Napanel3$ Bile4650Healthy74720.803Up $mRNApanel6$ Bile40100Cirthosis80740.93Up $mRNApanel6$ Bile40100Cirthosis80740.93Up $mRNApanel6$ Bile23Healthy82710.803Up $mRNApanel6$ Up100Cirthosis80700.93Up $mRNApanel6$ Up10101010101010 $mRNApanel6Pisana100100.94Up10mRNApan$	Biomarker	Sample	Cases	Controls	Control condition	Sensitivity	Specificity	AUC	Regulation	[Ref.]
73 83 0.795   80 91 0.816   82 98 0.949   87 40 0.655   87 70 80 0.752   87 96 90 0.752   88 82 85 0.892   89 82 85 0.802   80 74 72 0.803   81 82 71 0.803   82 82 71 0.803   83 82 71 0.803   849 72 0.803   85 71 0.849   86 70 0.803   80 70 0.858   81 90 90   82 80 70   83 90 90	HCC									
80 91 0.816   82 98 0.949   87 87 40 0.655   87 80 0.752 0.752   80 96 90 0.976   96 90 0.752 0.802   98 64 0.802   91 74 72 0.803   92 74 72 0.803   93 82 71 0.849   94 72 0.803 0.816   95 71 0.819 0.816   96 70 0.819 0.816   91 92 71 0.819   92 80 70 0.828   93 90 90 0.96	miR-143	Serum	95	127	Healthy	73	83	0.795	Up	[45]
82 98 0.949   CC 87 40 0.655   CC 70 80 0.752   CC 96 90 0.976   is 96 90 0.976   is 96 90 0.976   is 82 85 0.892   is 96 64 0.802   is 82 71 0.803   is 82 71 0.849   is 80 70 0.858   LI 90 90 0.96	miR-215	Serum	95	127	Healthy	80	91	0.816	Up	[45]
C 87 40 0.655   C 70 80 0.752   is 96 90 0.976   is 82 85 0.892   is 82 64 0.802   is 74 72 0.803   is 82 71 0.849   is 80 70 0.858   is 80 70 0.858   LI 90 90 0.96	miRNA panel <sup>1</sup>	Serum	112	42	Healthy	82	86	0.949	Up/Down	[46]
CC 70 80 0.752   is 96 90 0.976   is 82 85 0.892   is 86 64 0.802   i 74 72 0.803   i 74 72 0.803   is 82 71 0.849   is 80 70 0.858   is 80 70 0.858   L1 90 90 0.96	miR-221	Serum	30	60	Non-HCC	87	40	0.655	Up	[47]
is 96 90 0.976   is 82 85 0.892   86 64 0.802   67 96 N/A   74 72 0.803   82 71 0.849   93 80 70 0.858   11 90 90 0.96	miRNA panel <sup>2</sup>	Serum	27	135	Non-HCC	10	80	0.752	Up	[48]
is 82 85 0.892   86 64 0.802   7 96 N/A   7 74 72 0.803   7 74 72 0.803   82 71 0.849   19 90 0.658	miR-101	Serum	67	61	Cirrhosis	96	90	0.976	Down	[49]
86 64 0.802   67 96 N/A   71 72 0.803   72 82 71 0.849   82 71 0.849   1 82 71 0.849   1 82 71 0.849   1 82 71 0.849   1 82 71 0.849   1 90 70 0.858   1 90 90 0.96	miRNA panel $^{\mathcal{3}}$	Serum	103	78	Cirrhosis	82	85	0.892	Up/Down	[50]
67 96 N/A   74 72 0.803   82 71 0.849   1 82 71   1 90 70   1 90 90	miRNA panel <sup>4</sup>	Plasma	67	82	HBV	86	64	0.802	Up	[51]
67 96 N/A   7 72 0.803   82 71 0.849   80 70 0.858   11 90 90	CCA									
74 72 0.803   82 71 0.849   1 82 70   1 90 90   1 90 90	miRNA panel $5$	Bile	46	50	Healthy	67	96	N/A	Up	[52]
82 71 0.849   is 80 70 0.858   L1 90 90 0.96	miR-192	Serum	51	32	Healthy	74	72	0.803	Up	[53]
is 80 70 0.858 LI 90 90 0.96	miRNA panel $\delta$	Urine	22	21	Healthy	82	71	0.849	Up	[54]
is 80 70 0.858 LI 90 90 0.96	HBV									
LI 90 90 0.96	miRNA panel <sup>7</sup>	Plasma	100	100	Cirrhosis	80	70	0.858	Up	[55]
LI 90 0.96	DILI									
Notes: / miR-19a, -195, -192, -146a; 2 <sup>2</sup> miR-29a, -29c, -133a, -143, -145, -192, -505; 3 <sup>3</sup> miR-122, -1228, -141-3p, -192, -199a, -206, -26a, -433-3p;	miRNA panel <sup>8</sup>	Plasma	41	40	Non-DILI	06	06	0.96	Up/Down	[56]
/ miR-19a, -195, -192, -146a; 2 miR-29a, -29c, -133a, -143, -145, -192, -505; 3 miR-122, -1228, -141-3p, -192, -199a, -206, -26a, -433-3p;	Notes:									
$\frac{2}{3}$ miR-29a, -29c, -133a, -143, -145, -192, -505; $\frac{3}{3}$ miR-122, -1228, -141-3p, -192, -199a, -206, -26a, -433-3p;	<i>I</i> miR-19a, -195, -	192, -146a;								
3 miR-122, -1228, -141-3p, -192, -199a, -206, -26a, -433-3p;	2 miR-29a, -29c, -	133a, -143,	-145, -19	2, -505;						
	3 miR-122, -1228,	-141-3p, -1	92, -199a	, -206, -26a,	-433-3p;					

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<sup>4</sup> miR-20a, 25-3p, -30a, -92a-3p, -132-3p, -185, -320a, -324-3p;

5 miR-191, -486-3p, -1274b, -16, -484;

6 miR-192, -21; 7 miR-18a, -21, -29c-3p, -106b, -122, -185;

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185;

8 miR-382, -483-3p, -125a, -1290, -378i, -449b-3p, -151a-3p, -1915, -652, -27a-3p, -122, -151a, -151b, -885, -26a, -4783-3p, -2861