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Portal hypertensive enteropathy

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Abstract

Portal hypertensive enteropathy (PHE) is a condition that describes the pathologic changes and mucosal abnormalities observed in the small intestine of patients with portal hypertension. This entity is being increasingly recognized and better understood over the past decade due to increased accessibility of the small intestine made possible by the introduction of video capsule endoscopy and deep enteroscopy. Though challenged by its diverse endoscopic appearance, multiple scoring systems have

been proposed to classify the endoscopic presentation and grade its severity. Endoscopic findings can be broadly categorized into vascular and non-vascular lesions with many subtypes of both categories. Clinical manifestations of PHE can range from asymptomatic incidental findings to fatal gastrointestinal hemorrhage. Classic endoscopic findings in the setting of portal hypertension may lead to a prompt diagnosis. Occasionally histopathology and cross sectional imaging like computed tomography or magnetic resonance imaging may be helpful in establishing a diagnosis. Management of overt bleeding requires multidisciplinary approach involving hepatologists, endoscopists, surgeons, and interventional radiologists. Adequate resuscitation, reduction of portal pressure, and endoscopic therapeutic intervention remain the main principles of the initial treatment. This article reviews the existing evidence on PHE with emphasis on its classification, diagnosis, clinical manifestations, endoscopic appearance, pathological findings, and clinical management. A new schematic management of ectopic variceal bleed is also proposed.

Key words: Portal hypertension; Enteropathy; Intestinal vasculopathy; Ectopic varices; Gastrointestinal bleeding

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Core tip: Portal hypertensive enteropathy (PHE) is an under recognized complication of portal hypertension. It can present with a broad spectrum of clinical manifestations and endoscopic findings, making its diagnosis challenging. Video capsule endoscopy and deep enteroscopy are diagnostic tools of choice. PHE should be considered in patients with portal hypertension who present with occult or overt gastrointestinal bleeding, especially when portal hypertensive gastropathy and advanced cirrhosis are also present. Adequate resuscitation, reduction of portal pressure, and endoscopic therapeutic intervention remain the mainstay of initial treatment though definitive management may require a multidisciplinary approach involving hepatologists, endoscopists, surgeons, and interventional radiologists.

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INTRODUCTION

Portal hypertension is defined as increased pressure in portal circulation, as estimated with the measurement of hepatic venous pressure gradient (HVPG)^[1] which is the difference between wedge hepatic venous pressure (hepatic sinusoidal pressure) and free hepatic venous pressure. HVPG of more than 6 mmHg is considered abnormal^[1,2]. Clinically significant portal hypertension occurs when HVPG is more than 10 mmHg and it is considered severe portal hypertension when HVPG is more than 12 mmHg^[3,4]. Cirrhosis with consequent increased hepatic vascular resistance and portal venous inflow remains the most common etiology of portal hypertension, though it can also occasionally be seen due to other pre and post hepatic conditions such as congestive heart failure, Budd-Chiari syndrome, portal vein thrombosis, *etc.* The major clinical manifestations of portal hypertension were first recognized in the late nineteenth century and portal hypertension was described as a clinical syndrome of ascites, splenomegaly, and esophageal hemorrhage by Gilbert and Carnot in 1902^[5]. Rupture and consequent hemorrhage from esophageal varices remains the most lethal complication of portal hypertension and therefore guidelines now call for periodic endoscopy for surveillance of these varices. Upper and lower endoscopy have led to recognition of other mucosal changes of gastrointestinal tract which previously had been less well recognized.

Microcirculatory changes of gastric mucosa, as a result of portal hypertension was first described as congestive gastropathy in 1985 by McCormack *et al.*^[6]. The characteristic findings of mosaic-pattern mucosa (irregular, cleft-bordered polygonal reticulated area, or snake skin-like pattern), small flat red-point lesions, elevated large cherry red spots, and black-brown spots have been variously termed as inflammatory gastritis, mucosal vasculopathy, portal hypertensive mucosa, and portal hypertensive gastropathy^[6-10]. Subsequent studies suggested that these changes may be found in other areas of gastrointestinal tract as well^[11-14]. The term, portal hypertensive colopathy, was used to describe similar abnormalities found in the colon of patients with portal hypertension. Endoscopic findings of portal hypertensive colopathy include diffuse erythematous and edematous mucosa, inflammatory (colitis-like) lesions, angiodysplasia-like lesions, and ectopic or anorectal varices^[15-24].

Once the gastric and colonic mucosal changes associated with portal hypertension were recognized, jejunal and ileal mucosal changes were also noted. Occasional reports suggested small bowel bleeding in cirrhotic patients which was presumed to be a consequence of portal hypertension. The introduction of video capsule endoscopy and deep enteroscopy has increased our ability to evaluate the small bowel completely with good quality imaging and has led to better recognition and understanding of these portal hypertensive changes in the small bowel. This article reviews the existing evidence on portal hypertension-related changes in the small intestine, also known as portal hypertensive enteropathy (PHE) with emphasis on its classification, clinical manifestations, endoscopic appearance, and clinical management.

DEFINITION AND CLASSIFICATION

PHE previously termed portal hypertensive intestinal vasculopathy^[25], includes pathologic changes and mucosal abnormalities observed in the small intestine of patients with portal hypertension^[26]. Like other portal hypertensive related changes, PHE has also been reported in non-cirrhotic etiologies of portal hypertension^[27]. The definition and diagnosis of PHE have evolved over the past decade due to increased accessibility of the small intestine. Introduction of video capsule endoscopy (VCE) and deep enteroscopy has shed more light into this area of the gastrointestinal tract and redefined the disease. Multiple scoring systems have been proposed to classify PHE and to grade its severity^[28-30].

De Palma *et al.*^[29] were among the first groups to study small intestinal changes in cirrhotic patients with portal hypertension using VCE^[29]. They found that portal hypertensive gastropathy and portal hypertensive colopathy are significantly associated with PHE and suggested these changes across different parts of the gastrointestinal tract could be the regional manifestations of the same process in portal hypertension, rather than distinct entities. Other parameters that were associated with PHE in their study were grade 2+ or larger esophageal varices and Child-Pugh class C cirrhosis. They classified endoscopic findings of PHE into two categories: mucosal inflammatory-like abnormalities (edema, erythema, granularity, friability) and vascular lesions (cherry red spots, telangiectasias, or angiodysplasia-like lesions, varices)^[29].

Abdelaal *et al.*^[28] classified PHE lesions into 4 subtypes: inflammatory-like lesions, red spots, angioectasia, and small bowel varices. They created a scoring system based on VCE findings, giving each type one point with an additional point for multiple (more than two) lesions. Using this PHE score, they confirmed the findings by De Palma *et al.*^[29] by redemonstrating the association between PHE and

Table 1 Endoscopy based classification systems for portal hypertensive enteropathy

	De Palma <i>et al.</i> ^[29]	Abdelaal <i>et al.</i> ^[28]	Kodama <i>et al.</i> ^[30]
Classification	Inflammatory lesions Vascular lesions	Inflammatory-like lesions Red spots Angioectasia Small bowel varices	Villous abnormalities Edema Atrophy Erythema Vascular lesions angiodysplasia-like lesions dilated/proliferated vessels varices
Scoring system	None	One point for each type of lesion. An additional point for > 2 lesions	One point for each type of lesion
Clinical significance	Associated with PHG, PHC, large esophageal varices and Child-Pugh class C cirrhosis	Associated with PHG, large esophageal varices, Child-Pugh class C cirrhosis, and a history of EVL	Associated with ascites

PHG: Portal hypertensive gastropathy; PHC: Portal hypertensive colopathy; EVL: Endoscopic variceal ligation.

portal hypertensive gastropathy, large esophageal varices, Child-Pugh class C cirrhosis, and a history of prior endoscopic variceal sclerotherapy or banding ligation in a non-randomized, case-controlled, prospective study^[28].

Kodama *et al.*^[30] proposed a PHE scoring system based on double balloon enteroscopy findings. They classified PHE lesions into 2 categories: villous abnormalities and vascular lesions. They further subclassified each category into 3 subtypes: edema, atrophy and reddening for villous abnormalities, and angiodysplasia-like lesions, dilated/proliferated vessels, and varices for vascular lesions. A single point is given for each type, resulting in a scoring system with a maximum of 6 points. This scoring system, however, was associated with only the presence of ascites, making its clinical significance unclear^[30].

In summary, PHE lesions can be described based on VCE findings or optical endoscopic findings. They can be categorized into subtypes of vascular and non-vascular lesions. Despite multiple proposed scoring systems as described in Table 1, presently there is insufficient data to standardize or validate these systems. Besides ectopic varices and bleeding lesions, the clinical significance of other subtypes of mucosal changes remains unclear.

EPIDEMIOLOGY

There is great heterogeneity among reported prevalence of PHE ranging from 15% up to 82% in cirrhotic patients^[5,27,29,31]. We believe that this heterogeneity is a result of a wide spectrum of clinical severity of portal hypertension in cirrhotic patients, together with increase in reported prevalence due to recent advances in small bowel imaging in the past decade. The prevalence of PHE ranges only 15%-25% in studies where traditional endoscopic modalities like duodenoscopy, push enteroscopy or colonoscopy with ileal intubation were used^[14,20] but is

much higher (40%-82%) in other studies when VCE was used^[5,26,29,32-35].

Prevalence of each type of endoscopic finding also varies among studies. Red spots (22.2%-62.2%) and angiodysplasia-like lesions (24.3%-55.7%) seem to be more common than inflammatory-like lesions (5.6%-13%) or varices (8.1%-38.9%). Mixed lesions, especially multiple vascular lesions (varice, angiodysplasia-like, and red spots) can be seen in up to 22.3%^[26,29,36]. Small bowel varices account for 12%-35% of all ectopic varices^[37,38]. Portal hypertensive polypoid enteropathy is a rarer manifestation of PHE and polypoid lesions are less common in the small intestine (0.3%) compared to polypoid gastropathy (0.6%)^[39].

Actively bleeding lesions are not uncommon and can be seen in up to 17.8% of all patients with PHE, suggesting their clinical significance as a possible culprit source of obscure overt or obscure occult gastrointestinal bleeding. The bleeding lesions are commonly from angiodysplasia-like lesions and varices, though occasionally can be associated with polypoid enteropathy as well^[26,29].

CLINICAL PRESENTATION

PHE should be suspected when there is gastrointestinal bleeding or anemia not otherwise explained by more common etiologies, along with signs of portal hypertension such as ascites, splenomegaly, thrombocytopenia, or hepatic venous pressure gradient more than 8 mmHg^[39,40]. PHE can present as anemia, melena, hematochezia, hematemesis or may be asymptomatic. Fatal and life threatening ectopic variceal hemorrhage in small intestine has also been reported^[26,38-41]. Small intestinal variceal rupture can present with a classic triad of hematochezia (without hematemesis), portal hypertension, and previous intra-abdominal surgery^[38,42,43]. The most common indication for diagnostic work up is occult gastrointestinal bleeding.

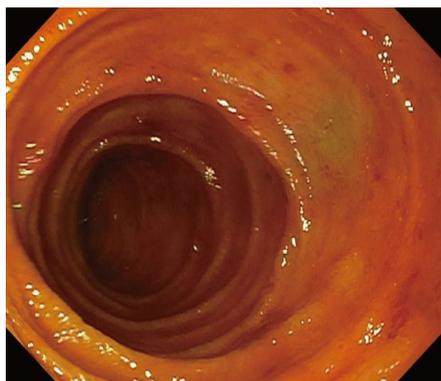


Figure 1 Mucosal red spots.

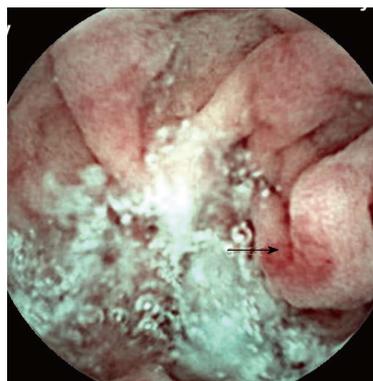


Figure 2 Angiodysplasia-like lesion.

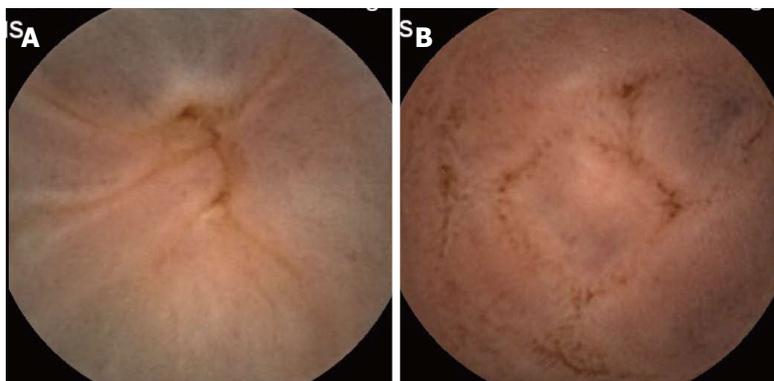


Figure 3 Ileal varices (A and B).

Certain clinical parameters have been shown to associate with PHE and can serve as clues to search for the disease. These parameters include large esophageal varices, history of endoscopic variceal injection sclerotherapy, history of endoscopic variceal ligation, portal hypertensive colopathy, portal hypertensive gastropathy, low hemoglobin, thrombocytopenia, splenomegaly, multiple signs of portal hypertension seen on CT scan, and Child-Pugh class C^[26,28,29,36,44,45]. However, not all of these associations were seen in every study. Only advanced cirrhosis (Child-Pugh class C) and the presence of portal hypertensive gastropathy were consistently associated with PHE in most studies^[28,29,36,44]. PHE has also been reported in non-cirrhotic etiologies of portal hypertension like polycystic liver disease, portal vein thrombosis, and Budd-Chiari syndrome^[27,46-48]. PHE is usually diagnosed by a VCE or a deep enteroscopy^[35].

ENDOSCOPIC FINDINGS

Endoscopically, PHE can be associated with a wide range of mucosal changes including the mucosal edema, congested rounded blunt villi giving a classic “herring-roe” appearance, loss of vascularization, friability, hyperemia, flat red spots, angiodysplasia-like lesions, pigmented black-brown spots, mucosal granularity, ulcers, reticulated mosaic-like pattern mucosa, protruding red bumps, inflammatory polyps, and varices^[11,26,39,44,47,49,50]. Recognizing these

findings through an optical endoscope or capsule endoscopy can lead to a prompt diagnosis and avoid further unnecessary and potentially harmful intervention. As mentioned above, these endoscopic findings can be classified into vascular and non-vascular lesions. They can be further categorized as inflammatory-like lesions, red spots, angioectasia, and small bowel varices^[28]. Special attention should be paid to vascular lesions such as red spots, angioectasia-like lesions, and varices, which are more likely to cause clinically significant bleeding and are amenable to endoscopic intervention^[28].

Red spots are small, symmetrical, uniformly erythematous, vascular areas on intestinal mucosa as shown in Figure 1. The lesions are usually flat. They are a very common manifestation of PHE and have been reported in up to 55% of cirrhotic patients with portal hypertension in one series^[28].

Angioectasia are aberrant submucosal vascular lesions, characterized as small red patches with arborizing ectatic vessels as shown in Figure 2. Variceal lesions in the small bowel are described as tortuously enlarged veins that usually have serpiginous or nodular shape with overlying mosaic-like shining mucosa with bluish discoloration as shown in Figure 3A and B^[26,27]. Endoscopic characters of these lesions should be promptly recognized to avoid potentially disastrous diagnostic biopsy attempts.

Duodenal varices are most commonly found in the duodenal bulb and the second portion of duo-

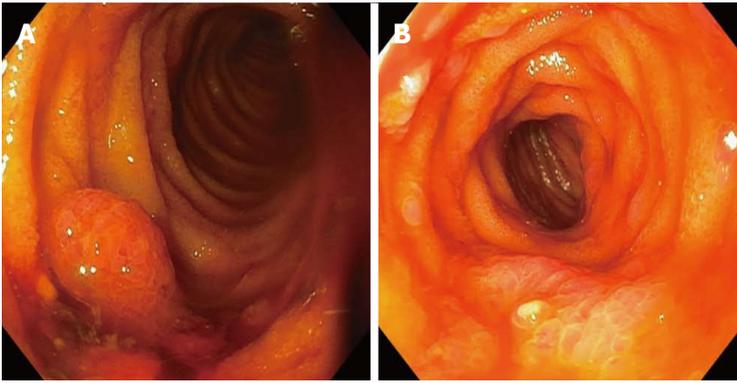


Figure 4 Portal hypertensive polypoid enteropathy (A and B).

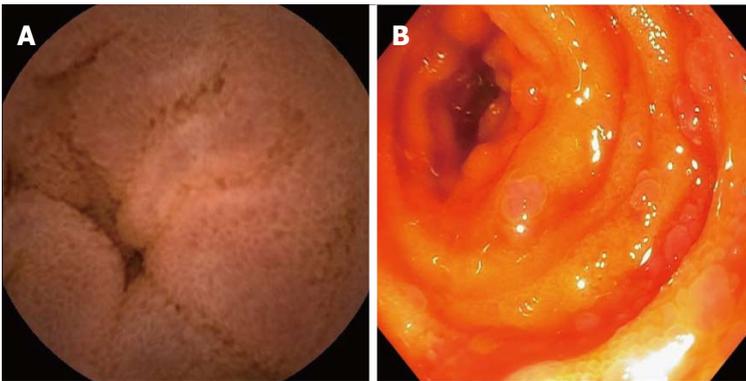


Figure 5 Herring roe appearance of small bowel mucosa (A and B).

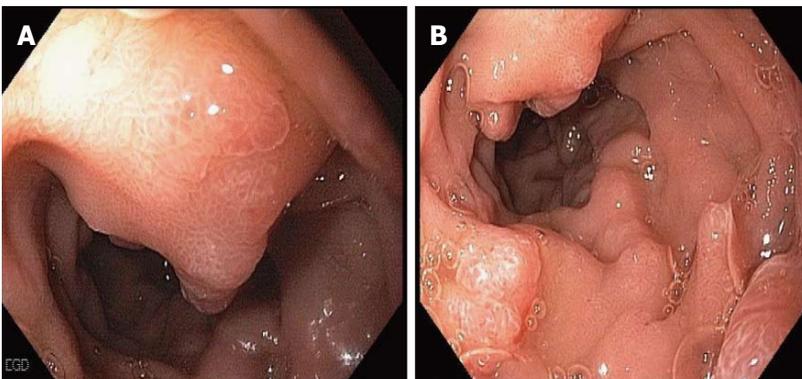


Figure 6 Mucosal edema with granularity of the small bowel (A and B).

denum^[51-53]. Jejunal and ileal varices are more commonly seen in patients with history of abdominal surgery due to post surgical portosystemic collaterals related to bowel anastomosis and are more difficult to diagnose due to their deep location in the intestinal tract^[38,42,43]. Small intestinal varices usually originate from portal venous trunk or superior mesenteric vein^[48,54,55]. Polypoid enteropathy is a rare manifestation of PHE and can present in any part of the small intestine. Polypoid lesions can have varied endoscopic manifestations. They can be single or multiple, sessile or pedunculated, small or large and may occasionally mimic adenomatous polyps as shown in Figure 4A and B. They usually arise in the background of inflamed mucosa with classic mosaic or herring-roe pattern and have been associated with occult or overt gastrointestinal bleeding^[39,49,50,56].

Other inflammatory lesions include mucosal ede-

ma, granularity, patchy erythematous mucosa, and herring roe appearance (rounded blunted villi on the background of congested mucosa with granularity) as shown in Figures 5 and 6. Clinical significance of these lesions is unclear but it is believed that they are less likely to cause overt gastrointestinal bleeding^[26,28,44].

Some of these endoscopic findings are non-specific and can pose a diagnostic dilemma especially in a patient whose portal hypertensive status is unknown. Differential diagnosis includes inflammatory bowel disease, celiac disease, arteriovenous malformations, and familial adenomatous polyposis. Biopsy of non-vascular lesions can be performed with caution to confirm the diagnosis.

In portal hypertension when splanchnic blood flow cannot effectively return to the systemic circulation, consequently splanchnic vasodilation ensues and mucosa of small intestine becomes congested.

Therefore, histopathologic changes of PHE show evidence of congested mucosa and vascular ectasia. These histologic findings include capillary dilation in the lamina propria (mean vascular diameter of 380 micrometers), increased capillary wall thickness, fibromuscular proliferation, a decreased villous/crypt ratio, neovascularization, vascular ectasia, vessels containing fibrin thrombi, inflammatory lymphoplasmacytic cells infiltration in the lamina propria, reactional nucleocytoplasmic atypia in the epithelial cells, and crenulated aspect of the glands^[13,14,39,57-59]. However, these histopathological changes are non-specific and can be seen in patients without portal hypertension and in patients with normal endoscopic findings^[12-14]. Therefore, the diagnosis of PHE should not be made on histopathology alone, but rather in conjunction with other clinical and endoscopic characteristics.

OTHER DIAGNOSTIC TESTS

Jeon *et al.*^[26] has evaluated the use of computerized tomography (CT) scan findings of portal hypertension such as esophageal varices, gastric varices, peri umbilical varices, portal hypertensive colopathy, portal hypertensive gastropathy, portal hypertensive cholecystopathy, splenomegaly, and ascites as the radiologic predictors of PHE. They have created a scoring system giving each CT finding one point with a maximum score of 6 points. A CT score of more than 3 was found to be significantly associated with PHE^[26]. Cross sectional imaging by computed tomography angiography and magnetic resonance angiography can also aid in evaluation of vascular origin of the ectopic varices^[55,60].

Abdelaal *et al.*^[28] have explored the use of a transient elastography (FibroScan®), a novel non-invasive ultrasound-based technology using pulse-echo ultrasound signals to measure liver stiffness as a surrogate marker of severity of portal hypertension in cirrhotic patients^[4,61]. They found that a high transient elastography score had a linear relation with a high PHE score ($r^2 = 0.314$, P -value 0.004), suggesting that higher severity of PHE may be associated with higher degree of liver disease and portal hypertension. Mean liver stiffness measurement in PHE group was 29 kPa, which is much higher than the portal hypertensive cut-off value of 13.6 kPa^[4,28]. They concluded that transient elastography may be a new non-invasive method for detecting the presence and severity of PHE in cirrhotic patients^[28].

MANAGEMENT

VCE is the preferred initial diagnostic modality in evaluating the small bowel due to its non-invasive nature^[5,27,33,35]. It also serves as a road map for subsequent interventions. A deep enteroscopy is warranted when a therapeutically amenable lesion is

found on VCE in patients with obscure gastrointestinal bleeding^[27]. Different endoscopic findings may require different therapeutic interventions.

Due to its rarity and insufficient evidence, there are no standardized therapeutic guidelines for symptomatic PHE^[40]. Clinical significance and the need for any intervention for inflammatory lesions and red spots remain unclear. Argon plasma coagulation is generally used for angioectasia^[27] while multiple approaches have been used for small bowel varices and polypoid enteropathy^[39,55,56,62,63].

Treatment of portal hypertensive polypoid enteropathy depends on number of polyps and endoscopic accessibility. A polypectomy can be safely performed if the polyp is accessible and amenable for endoscopic removal. Endoclip can be used at the stalk to achieve complete hemostasis^[56]. Argon plasma coagulation can be used on the inflamed surface of bleeding polyp to achieve hemostasis but recurrent bleeding has been reported^[39]. Non-selective beta blocker, transjugular intrahepatic portosystemic shunt (TIPS), surgical small bowel resection, and liver transplantation have all been reported anecdotally to be successful treatment for portal hypertensive polypoid enteropathy^[39,56].

Bleeding small bowel varices occurs in 0.4% of patients with portal hypertension and account for up to 5% of all variceal bleeding. Similar to esophagogastric variceal hemorrhage, it usually presents as massive life-threatening hemorrhage with mortality rate as high as 40%^[38,40,51,54]. Available therapeutic options are endoscopic treatment, surgical interventions, and interventional radiological approaches. Similar to polypoidal lesions, management of small bowel varices depends largely on endoscopic accessibility, patient's surgical risk, available therapy and local expertise.

Adequate resuscitation with intravenous fluids, blood product transfusion, close monitoring, and airway protection remain the main principles of initial management. The benefit of medical management has not been extensively studied in ectopic varices or other manifestations of PHE, but given its established role in esophageal and gastric variceal management^[64], it is reasonable to consider the use of vasoactive agents such as octreotide and non-selective beta blockers to reduce the splanchnic and portal pressure in both primary and secondary prophylaxis of ectopic varices^[40].

Despite the technical challenges, endoscopic band ligation and endoscopic variceal obturation with tissue glue monomer such as N-butyl-2-cyanoacrylate are endoscopic interventions of choice for hemostasis in ectopic variceal rupture. However, the achieved hemostasis is temporary and re-bleeding is a major concern^[52,53]. Endoscopic variceal obturation has shown better success and lower re-bleeding rate compared to endoscopic variceal banding^[38,65-68]. Endoscopic sclerotherapy and band

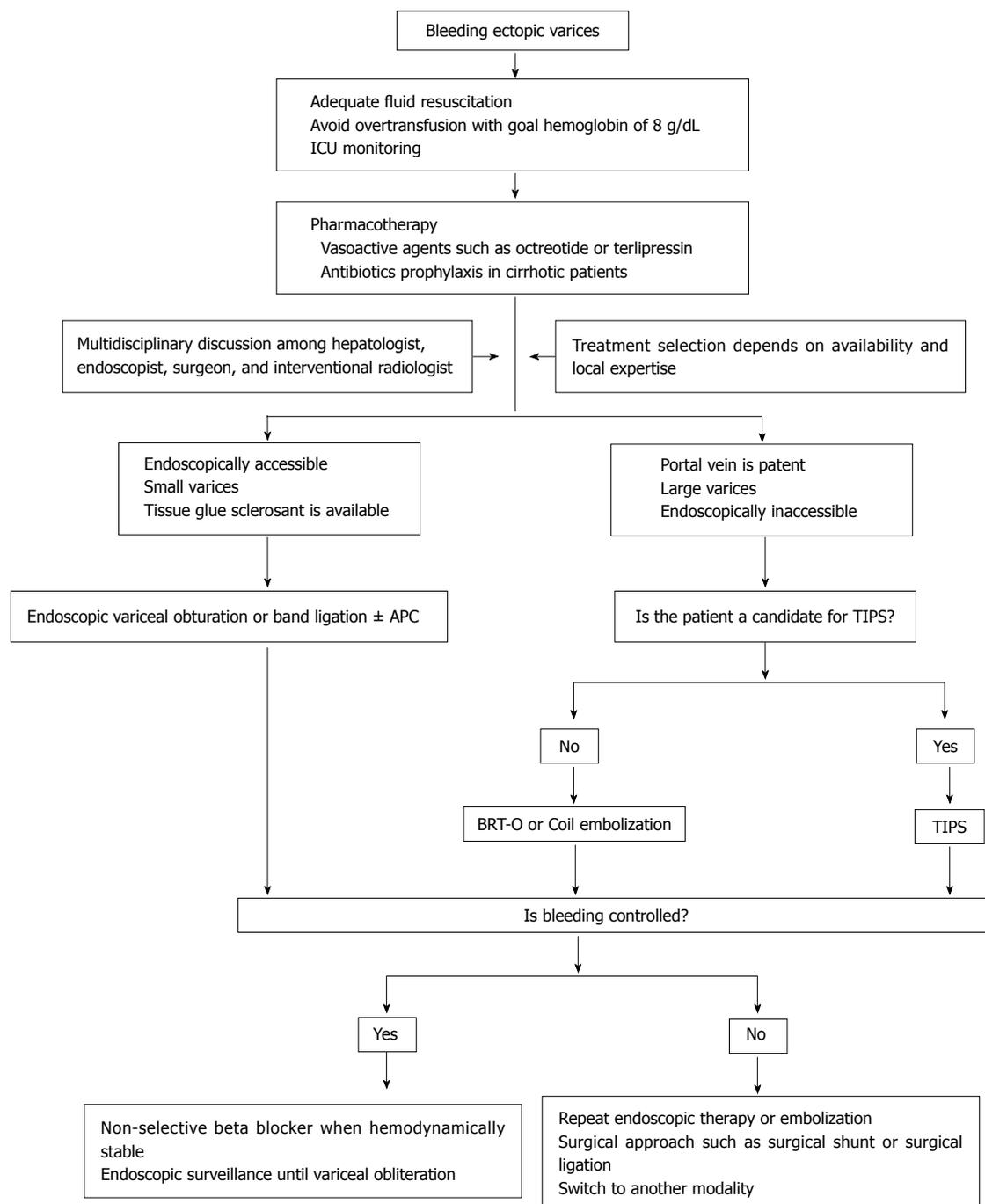


Figure 7 Management of bleeding ectopic varices. TIPS: Transjugular intrahepatic portosystemic shunt; APC: Argon plasma coagulation; BRT-O: Balloon-occluded retrograde transvenous obliteration.

ligation are not recommended for large varices, especially the varices with larger diameter than the endoscope itself. This is because incomplete banding can lead to mucosal defect in the remaining varix causing recurrent hemorrhage while excessive dilution of injected sclerosant in large varices decreases the success rate of the sclerotherapy^[40,63]. Limited evidence suggests supplemental use of argon plasma coagulation after variceal band ligation as a successful intervention in prevention of rebleeding in esophageal and ileal varices and can

be considered in endoscopic management of ectopic varices as well^[69-71].

Interventional radiology approaches are effective modalities especially for large varices, lesions that are not endoscopically accessible, and for patients with poor overall general condition who are poor surgical candidates^[38,63,72]. These interventions include TIPS, balloon-occluded retrograde transvenous obliteration (B-RTO), and percutaneous coil embolization^[38,40,55,62].

Even though TIPS is relatively safe procedure,

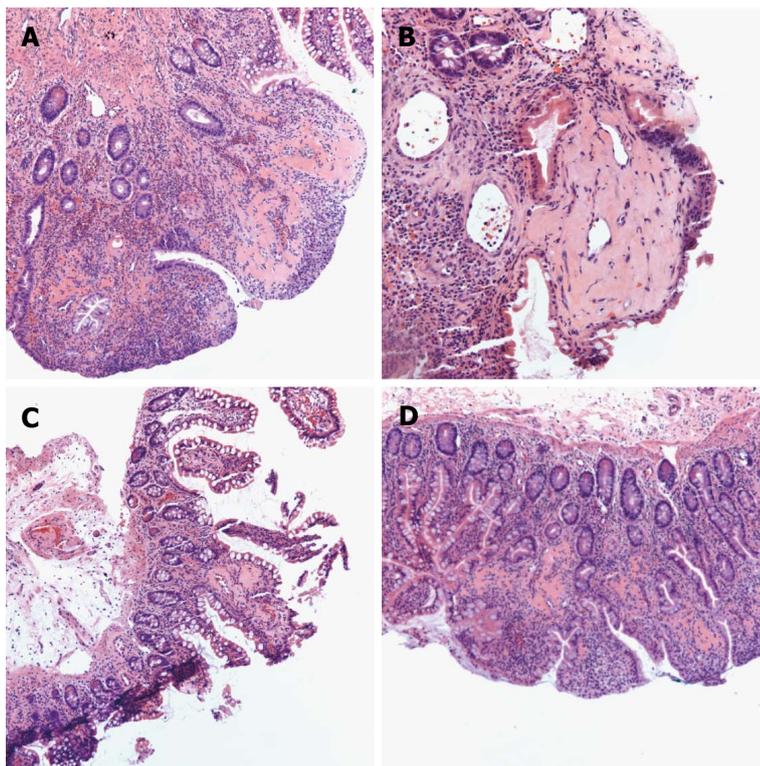


Figure 8 Histopathological changes of portal hypertensive enteropathy (A-D).

widely available, and with good success rate, it has limitations in patients with hepatic encephalopathy, high MELD score, severely decompensated cirrhosis, and severe hepatic atrophy^[73,74]. It, however, remains an effective intervention to control the variceal bleeding and prevent recurrence, especially in those who have failed endoscopic and medical management. The American Association for the Study of Liver Diseases (AASLD) has recommended TIPS as the preferred approach for the prevention of rebleeding of ectopic varices (including intestinal, stomal, and anorectal varices)^[75].

Percutaneous transhepatic and transjugular coil embolization are other radiological interventions that may offer a focal occlusion of the feeding vein of the varices. They can be safely performed even in a large varix, and have good short term results. However, as the portal system is not decompressed, they carry a very high recurrent bleeding rate^[76-81]. A retrospective study by Macedo *et al.*^[76] showed that even though an immediate bleeding control can be achieved in 75% of patients receiving coil embolization, the rebleeding rate was 67% with a mean bleeding-free interval of 7.8 mo. This result was similar to previous studies that demonstrated a rebleeding rate as high as 55% at 6 mo and 92% at 4 years^[76,82]. A combined approach of TIPS and percutaneous embolization in the setting of persistent bleeding after TIPS is preferred^[76].

B-RTO is a non-surgical therapy that can occlude not only the varices, but also the feeding afferent and efferent vessel. It is particularly useful in patients who bleed at lower portal pressure, have

non-patent portal vein, or are not candidates for TIPS. The success of B-RTO in treating small bowel ectopic varices has been reported, especially in the Japanese literature^[83-86]. However, B-RTO can cause significant elevation of porto-systemic pressure gradient and subsequent variceal formation has also been reported^[55,87].

Surgical approaches, such as portosystemic surgical shunt, segmental small bowel resection and surgical variceal ligation, can be performed but are usually reserved for patients who are refractory to other therapy^[38,41,88,89]. Surgical shunt such as distal splenorenal shunt has equal efficacy as TIPS with no difference in survival or hepatic encephalopathy rate but is less cost-effective approach compared to TIPS^[75]. Other surgical approaches such as duodenotomy with simple oversewing of the varix and duodenal dearterialization with stapling have been reported^[90,91].

Similar to what Helmy *et al.*^[40] have proposed, management of bleeding ectopic varices should focus on adequate initial resuscitation and use a multimodality approach depending on availability and local expertise of each institution. We suggest an algorithm for management of bleeding ectopic varices in Figure 7. As mentioned above, we also recommend individually tailored therapies for non variceal bleeding depending on the clinical situation and local expertise.

CONCLUSION

PHE-associated small bowel mucosal changes are

increasingly being recognized due to introduction of VCE and deep enteroscopy of VCE and deep enteroscopy which enables a more thorough small bowel evaluation than previously possible. Endoscopically, PHE can have myriad presentations ranging from mild mucosal inflammatory changes to angioectasias, inflammatory appearing polyps, and occasionally as large ectopic varices. Due to the non-specific endoscopic and histopathological findings, a high index of suspicion is required to recognize and accurately diagnose this condition. VCE and/or a deep enteroscopy are the current preferred modalities for establishing the diagnosis. PHE has been known to cause significant life-threatening overt gastrointestinal bleeding or be a source of occult gastrointestinal blood loss. However, due to low prevalence and lack of large studies, its prognostic value and clinical significance on morbidity and mortality remain unclear. Management of PHE has not yet been standardized and should be individualized based on acuity and severity of the hemorrhage, endoscopic accessibility of the lesion, surgical risk of the patient, patency of portal vein, available therapy and expertise of each institution.

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Current biomarkers for hepatocellular carcinoma: Surveillance, diagnosis and prediction of prognosis

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Abstract

Biomarkers for surveillance, diagnosis and prediction of prognosis in patients with hepatocellular carcinoma (HCC) are currently not ready for introduction into clinical practice because of limited sensitivity and specificity. Especially for the early detection of small HCC novel biomarkers are needed to improve the current effectiveness of screening performed by

ultrasound. The use of high-throughput technologies in hepatocellular research allows to identify molecules involved in the complex pathways in hepatocarcinogenesis. Several invasive and non-invasive biomarkers have been identified already and have been evaluated in different clinical settings. Gene signatures with prognostic potential have been identified by gene expression profiling from tumor tissue. However, a single "all-in-one" biomarker that fits all-surveillance, diagnosis, prediction of prognosis-has not been found so far. The future of biomarkers most probably lies in a combination of non-invasive biomarkers, imaging and clinical parameters in a surveillance setting. Molecular profiling of tumorous and non-tumorous liver tissue may allow a prediction of prognosis for the individual patient and hopefully clear the way for individual treatment approaches. This article gives an overview on current developments in biomarker research in HCC with a focus on currently available and novel biomarkers, in particular on microRNA.

Key words: Hepatocellular carcinoma; Biomarker; Diagnosis; Prognosis; MicroRNA

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Core tip: The aim of this review is to provide an overview on current invasive and non-invasive biomarkers in hepatocellular carcinoma (HCC) with respect to their use in surveillance, diagnosis and prediction of prognosis. We also give an outlook on the future development of HCC biomarker research with a focus on microRNA.

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INTRODUCTION

The incidence of hepatocellular carcinoma (HCC) is rising throughout the world as a consequence of a rising prevalence of hepatitis B virus (HBV) and hepatitis C virus (HCV) infection and an increase in prevalence of (non-alcoholic) fatty liver disease due to the metabolic syndrome^[1-4]. The number of patients that are at risk to develop HCC and need to undergo structured surveillance is therefore constantly rising in parallel.

Transabdominal ultrasound currently is the only recommended tool for surveillance in the Western world which has been shown to be cost-effective. Its sensitivity is dependent on many factors, including the quality of the ultrasound machine, the experience of the examiner and also the patient. Especially in obese patients with NASH cirrhosis liver ultrasonography may be difficult and therefore not always appropriate to rule out HCC. In patients with liver cirrhosis regenerative nodules may be hard to distinguish from HCC on ultrasound, and the sensitivity of ultrasound to detect early HCC lies in a range of 32% to 65%^[5,6]. On the other hand, surveillance by contrast-enhanced computed tomography or contrast-enhanced magnetic resonance tomography is rather expensive and not cost-effective. In addition, they are associated with the additional exposure to radiation with an increased risk of tumor development and/or contrast-media related deterioration of kidney function. In developing countries, where even the availability of ultrasound surveillance is quite low, serological markers for surveillance are of special interest^[7].

Biomarkers in blood, other body fluids or tissue for screening, prediction of prognosis and monitoring of response to a therapy would be an important contribution to the management of patients with HCC.

Early detection of HCC is the most important factor to offer the patient the chance of cure. α -fetoprotein (AFP) is the most widely used and broadly known biomarker for HCC, but the measurement of serum AFP levels has been dropped from current surveillance guidelines in Europe and the United States because of low sensitivity and specificity. This is based on the knowledge that almost 80% of small HCCs do not show increased levels of AFP, and the sensitivity decreases to 25% in tumors smaller than 3 cm^[8]. Nonetheless, serum AFP measurement is still combined with ultrasound by many physicians worldwide to reduce the risk of missing small lesions in the cirrhotic liver that have not been detected by ultrasound. Alternative or additional biomarkers may be useful tools for surveillance or as a decisional tool in clinical practice to identify patients that will benefit from advanced imaging methods in a surveillance setting to augment the proportion of patients with HCC diagnosed in an early tumor stage.

Apart from their role as a surveillance tool, bio-

markers may play a role as diagnostic tool once a suspicious lesion in a patient with liver cirrhosis has been detected. In the past, a significant concentration of AFP in the serum of a patient with liver cirrhosis and a suspicious mass in the liver larger than 2 cm was sufficient to diagnose HCC^[9]. However, diagnostic algorithms endorsed by the American Association for the Study of Liver Diseases (AASLD) and the European Association for the Study of the Liver (EASL) nowadays strictly rely on typical radiological hallmarks in dynamic contrast-enhanced imaging apart from biomarkers^[9-11].

Once the diagnosis of HCC is confirmed, molecular biomarkers could potentially be used for prediction of prognosis of the individual patient and also for the guidance of therapeutic decisions. Currently, early studies on predictive biomarkers are on their way to make a step towards a personalized and individualized therapy of patients with HCC.

This review gives a concise overview on current clinical-translational knowledge on biomarkers in surveillance, diagnosis and prediction of prognosis with a focus on miRNA.

ROLE OF BIOMARKERS IN SURVEILLANCE AND DIAGNOSIS

Geographical differences in tumor prevalence, tumor biology and resources have resulted in differences in current guidelines with respect to screening recommendations. The main pillar of surveillance in high risk populations is repeated transabdominal ultrasound with small differences with respect to the definition of target populations in various guidelines. Throughout the world three serum biomarkers are suggested as tools to determine the risk of liver cancer in high risk populations: AFP, the proportion of the fucosylated isoform of AFP, AFP-L3, and des-gamma-carboxy-prothrombin (DCP). These markers are FDA approved for this indication, but not a part of the surveillance guidelines published by the AASLD and the EASL^[10,11]. Current expert opinion from Western countries has been rather critical on these biomarkers regarding their clinical value^[12]. On the other hand, the Asia Pacific guideline recommends to combine ultrasound with the measurement of AFP levels in the serum and the Japanese society even recommends to apply all three mentioned biomarkers^[13,14] for surveillance.

However, most studies on the performance of biomarkers in HCC detection have not been performed in a surveillance setting but compared levels of predefined biomarkers in patients with HCC with a comparator group, in most cases in patients with chronic liver disease.

A randomized controlled study performed in a high-risk population in China showed that screening by AFP measurement led to earlier diagnosis of HCC but had no impact on mortality^[15]. On the

Table 1 Diagnostic performance of alpha-fetoprotein serum levels in selected studies

Ref.	Year	n	Comparator	Cut-off-level	Sensitivity	Specificity	AUC
Marrero <i>et al</i> ^[20]	2009	836 total (419 HCC)	Liver cirrhosis	20 ng/mL	59%	90%	0.8
Mao <i>et al</i> ^[21]	2010	4217 total (789 HCC)	Amongst others healthy controls, HBV carriers, liver cirrhosis	35 ng/mL	58.20%	85.30%	
Farinati <i>et al</i> ^[22]	2006	1158 HCC	No control	400 ng/mL	18%		0.59
Lok <i>et al</i> ^[23]	2010	39 HCC, 77 matched controls	Hepatitis C	20 ng/mL	61%	81%	0.79

AUC: Area under the curve; HCC: Hepatocellular carcinoma.

other hand, semiannual screening for HCC by AFP measurement in a population-based study in Alaska was effective in detecting HCC at early stages and significantly prolonged survival rates^[16].

As the discussion on the rise and fall of AFP as biomarker in HCC surveillance and diagnosis has been intense and sometimes even emotional during the last decade^[17-19], data on the most important diagnostic studies referred to in this discussion are summarized in Table 1.

A recent meta-analysis on the performance of AFP in diagnosis of HCC included seven studies and revealed a pooled sensitivity of 66% with a specificity of 86% and an area under the curve (AUC) of 0.87^[24]. In a further meta-analysis including ten studies the pooled sensitivity of AFP for the diagnosis of HCC was 51.9% at a specificity of 94% (AUC = 0.81)^[25]. It is a major drawback of AFP as surveillance tool that its serum levels are influenced by the activity of the underlying liver disease and therefore increased in patients with elevated ALT levels even in the absence of HCC as shown in the HALT-C trial^[26]. Additionally, only a proportion of patients with HCC exerts elevated AFP serum levels leading to low sensitivity of the marker. The heterogeneity of tumor biology in HCC therefore results in a necessity to find better or complementary markers to close this diagnostic gap.

The clinical utility of high-sensitivity AFP-L3 (hs-AFP-L3) in early prediction of HCC development in patients with chronic HBV or HCV infection was recently evaluated in a large Japanese study. Even at low AFP levels and in absence of suspicious ultrasound findings an elevation of hs-AFP-L3 was an early predictor of HCC development with an elevation in 34.3% of patients one year prior to diagnosis of HCC^[27]. In patients with low AFP levels (< 20 ng/mL), the diagnostic sensitivity for hs-AFP-L3 at a cut-off of 5% was 41.5% with a specificity of 85.1%^[28].

Numerous studies have investigated the performance of alternative markers or combinations of already established markers. New candidate markers include squamous cell carcinoma antigen-immunoglobulin M complex (SCCA-IGM), α -L-fucosidase^[29], glypican-3 (GPC-3), insulin-like growth factor (IGF)^[30], vascular endothelial growth factor (VEGF), or Dickkopf-1 (DKK1)^[31].

Three further biomarkers have intensively been studied for their potential use in screening for HCC, namely Golgi protein 73 (GP73), interleukin-6 (IL-6) and squamous cell carcinoma antigen (SCCA) and were addressed in a recent meta-analysis^[32]. The transmembrane glycoprotein GP73 has a sensitivity of 62% with a specificity of 88% at a cut-off of 10 relative units in a study comparing 144 patients with HCC to 152 patients with cirrhosis and 56 healthy controls^[33]. A further study including 4217 subjects of whom 789 were patients with HCC revealed a sensitivity of 74.6% with a specificity of 97.4% at a cut-off of 8.5 relative units^[21]. Two smaller studies were identified in the meta-analysis studying the cytokine IL-6. Using different cut-off-values, sensitivity for HCC ranged from 46% to 73% with a specificity of 87% to 95%^[32,34,35]. The largest study on the role of the serine protease inhibitor SCCA included 961 patients and resulted in a sensitivity of 42% at specificity of 83% using a cut-off of 3.8 ng/mL^[32,36].

Seven well-designed studies on the diagnostic performance of osteopontin, an integrin-binding glycoprophosphoprotein, were published and recently summarized in a meta-analysis^[24]. Osteopontin is expressed by transformed malignant cells and has been evaluated also in colon and pancreatic cancer. All of the reported studies were retrospective in design and included a range of 30 to 179 patients with HCC. The pooled sensitivity of osteopontin for HCC was 86% with a specificity of 86% resulting in a diagnostic accuracy comparable to that of AFP in the included studies. The authors of the meta-analysis conclude that further validation studies are needed before the marker could be suggested for the use in daily clinical routine.

By combining two or more biomarkers the diagnostic performance of a single non-invasive test can be optimized. This has been investigated for the three best established non-invasive biomarkers in HCC, AFP, AFP-L3 and DCP.

When comparing 164 European patients with HCC to 422 controls with chronic liver disease a significant increase in AFP serum levels was mainly shown in patients with advanced stages of HCC and in patients suffering from viral hepatitis while DCP was more frequently elevated in patients with early-stage and NASH associated HCC. Taken alone,

Table 2 Diagnostic performance of novel non-invasive biomarkers

Ref.	Year	Marker	n	Comparator	Cut-off-level	Sensitivity	Specificity	AUC
Toyoda <i>et al</i> ^[31]	2011	hs-AFP-L3%	666	Chronic liver disease and AFP < 20 ng/mL	5%	41.50%	85.10%	0.707
Ertle <i>et al</i> ^[37]	2013	DCP	586	Chronic liver disease	5 ng/mL	45.80%	95%	0.87
Wan <i>et al</i> ^[24]	2014	Osteopontin	Meta-analysis (7 studies)	mixed		Pooled: 86%	Pooled: 86%	0.92
Hsia <i>et al</i> ^[35]	2007	IL-6	128	Mixed, including chronic liver disease and healthy controls	3 pg/mL	46%	95%	
Mao <i>et al</i> ^[21]	2010	GP73	4217	Mixed, including chronic liver disease and healthy controls	8.5 rel. units	74.60%	97.40%	0.94
Giannelli <i>et al</i> ^[36]	2007	SCCA	961	Liver cirrhosis	3.8 ng/mL	41.90%	82.60%	0.656
Ertle <i>et al</i> ^[37]	2013	AFP combined with DCP	586	Chronic liver disease	DCP 5 ng/mL AFP 10 ng/mL	78%	89.30%	0.91
Johnson <i>et al</i> ^[40]	2014	GALAD-score	670	Chronic liver disease		93%	89%	

hs-AFP: High-sensitivity α -fetoprotein; DCP: Des-gamma-carboxy-prothrombin; SCCA: Squamous cell carcinoma antigen; GP73: Golgi protein 73; IL-6: Interleukin-6; AUC: Area under the curve.

neither of the two parameters could detect more than one third of HCC patients independently of stage or etiology but by combination of AFP with DCP a sensitivity of 55% for early stage HCC and 78% for all stages (cut-off for AFP 10 ng/mL and for DCP 5 ng/mL) was reached^[37]. The addition of AFP-L3% to this combination, led to a further gain in sensitivity (84%) in another European study^[38].

The incorporation of clinical variables like age and gender into models based on a combination of biomarkers for HCC detection further improve the predictive performance of these models^[39]. A model using a combination of age, gender, AFP, AFP-L3 and DCP estimates the probability to suffer from HCC in an individual patient with chronic liver disease with a sensitivity of 86% for HCC in BCLC stage 0 or A and a sensitivity of 94% for later tumor stages^[40]. The diagnostic performance of novel circulating biomarkers and scores is summarized in Table 2.

Although the complex process of hepatocarcinogenesis is still not fully understood, several signal transduction pathways have been identified as critical players in the pathophysiology of HCC, including the Wnt/ β -Catenin pathway, the p53 pathway, the tumor suppressor retinoblastoma protein pRb1 pathway, the mitogen-activated protein kinase pathway, the Ras pathway, JAK/STAT signaling, mechanisms of cellular stress response like heat shock proteins and epidermal growth factor receptor and transforming growth factor- β signaling^[41]. As a consequence of different risk factors causing HCC in the individual patient, the alterations in these pathways differ in different settings which is probably the cause of insufficient sensitivity of single biomarkers. Genetic and epigenetic alterations occur in these pathways and mediate cell proliferation. The possibility to perform proteomic profiling and whole genome sequencing in combination with systems biology has led to a new era in biomarker development that will hopefully help to understand the complex interactions

in hepatocarcinogenesis of multiple proteins, genes and transcription factors. First examples of this approach have successfully been evaluated in clinical studies, but none of the signatures has been validated in large prospective studies.

In patients with chronic HBV infection and liver cirrhosis, proteomic analyses in the plasma identified a cluster of 11 proteins that is able to identify patients at high risk for HCC development (OR = 4.83, 95%CI: 1.26-18.56)^[42].

Gene expression profiling of peripheral blood mononuclear cells in HCC patients using microarrays and bioinformatics-driven analysis of the data has identified a blood-based signature of three genes, namely Chemokine (C-X-C motif) receptor 2 (CXCR2), C-C chemokine receptor type 2 (CCR2) and E1A-Binding Protein P400 (EP400), that predicts HCC with an AUC of 0.96 yielding at a sensitivity of 93% with a specificity of 89%^[43].

High-throughput metabolomics technologies with the comprehensive analysis of small molecular metabolites may additionally identify serum metabolic profiles that can be used as diagnostic biomarkers. First steps into this direction have also already been taken^[44,45].

To distinguish dysplastic nodules from well-differentiated HCC is a challenge, not only for the radiologist on imaging, but also for the pathologist on tissue samples.

Molecular signatures derived from gene expression profiling have been identified that are helpful to answer this critical question that is decisive for the further management of the patient.

Characteristic genomic changes during hepatocarcinogenesis have been identified. Specific gene signatures accurately reflect the pathological progression of disease from cirrhosis to dysplasia to early and advanced HCC in patients with HCV infection in Asian and Western patients^[46,47].

A three gene set in the tissue including glypican

3 (GPC3; 18-fold increase in HCC, $P = 0.01$), LYVE1 (12-fold decrease in HCC, $P = 0.0001$), and survivin (2.2-fold increase in HCC, $P = 0.02$) has an accuracy of 94% to discriminate dysplastic nodules from early HCC in HCV cirrhosis. Especially immunostaining for GPC3 is highly discriminative^[48].

Heat shock protein 70 and cyclase-associated protein 2 are further examples for tissue biomarkers identified in comprehensive approaches that found their way into clinical testing and application^[49,50].

MICRORNAS AS NOVEL BIOMARKERS

With respect to novel potential biomarkers, non-coding RNA and specifically microRNA (miRNA) have received the greatest attention over the past years^[51]. MiRNAs are small non-coding and evolutionary conserved RNA molecules that serve as posttranscriptional regulators of mRNA expression and interfere with translation to protein. Following several common modifications steps, miRNA become a part of the so called RISC (RNA silencing complex) to be functionally active^[52]. MiRNAs can either preserve their function intracellularly by regulating the expression of a target population of molecules, or can be released from the cell bound to other proteins and also as a free molecule^[53,54]. As part of the released vesicles, specific miRNAs can further preserve their functional activity locally or be transported in blood or probably other specimens to other tissues or organs^[55].

The most exiting advantages of miRNA over various other molecules are their stability against degradation, cell-type specific miRNA expression patterns and detectability in all types of human specimens such as blood, feces, saliva, *etc.*^[56-59]. While mRNA or various proteins are relatively sensitive to extracellular enzymes, miRNA expression levels remain, as long as they are preserved in a natural milieu, relatively resistant to RNA digestions, heating, storage, drying, formalin fixation, *etc.* For detailed information regarding the biogenesis of miRNA as well as regarding the current knowledge on molecular function we refer to the several excellent reviews from the field^[51,60,61].

Shortly after definite recognition of miRNA, several groups have provided seminal evidence for differences in miRNA expression patterns between different tissues and malignant conditions including HCC^[62,63]. High quality analyses using deep sequencing have recently provided an important view in microRNAome in liver tissue and HCC^[64]. Interestingly, about 86% of the miRNA were expressed in very low concentrations and only about 1% were expressed abundantly. Three of those miRNAs, namely miR-122, miR-192 and miR-199a/b-3p, were responsible for 74% of all miRNA in normal liver tissue with miR-122 accounting for almost 52% suggesting that those miRNAs are the

most important ones in liver biology^[64]. Many recent reports have shown a broad spectrum of changes in microRNAome in HCC^[64-67]. Therefore, miRNAs may have the potential to become valid biomarkers in HCC.

MIRNAS AS NON-INVASIVE DIAGNOSTIC BIOMARKERS FOR HCC

The biggest effort from miRNA-based biomarker research has been made to improve the diagnostic utility in HCC. In parallel with the dominant etiological factors in HCC development, the largest body of data comes from Asian populations and virus-related HCC cohorts. In one of the first profiling studies, Li *et al.*^[68] performed deep sequencing in pooled samples from chronic HBV virus patients, HCC patients and controls with and without cancer. They identified a pattern of 21 miRNA that show differential expression in cHBV patients and 6 miRNA differentially expressed in HCC patients. Following subsequent testing and validation, 13 miRNA, including miR-122, miR-375, miR-92a, miR-10a and let-7c, were identified as a biomarker for patients with HBV (acute and chronic) and HCV virus infection. Furthermore, using only 3 miRNAs (miR-25, miR-375, let7f) the authors could reach an AUC of 99.7% with a 97.9% sensitivity and a 99.1% specificity to discriminate controls from HCC patients. Most interestingly for HCC diagnosis, the comparison of the two cohorts with chronic HBV and HBV-associated HCC lead to identification of two miRNAs (miR-10a and miR-125b) that could separate the HCC cohort with an AUC of 99.2% (sensitivity 98.5% and specificity 98.5)^[68]. Recently, another large scale study studied plasma samples from 934 patients with various conditions including healthy subjects, patients with chronic HBV, liver cirrhosis and HBV-related HCC^[69]. Following discovery and training phases, the authors identified a panel of 7 microRNAs (miR-122, miR-192, miR-21, miR-223, miR-26a, miR-27a and miR-801) that provided the highest diagnostic accuracy for the identification of HBV-related HCC. Using an independent validation cohort of 390 samples, the area under the curve value was comparable with the training data and reached 0.888 with a sensitivity of 81.8% and a specificity of 83.5%. Interestingly, the diagnostic accuracy was independent of disease stage and was comparable to healthy subjects, patients with chronic hepatitis or livers cirrhosis. At present, this study is one of the largest to evaluate the biomarker potential of miRNAs in cancer. Notably, the expression of selected miRNA was analyzed using RT-PCR which may be critical for clinical translation of the results. Whether miR-122 is the optimal normalizer needs further evaluation^[69].

Besides profiling studies, a candidate-based

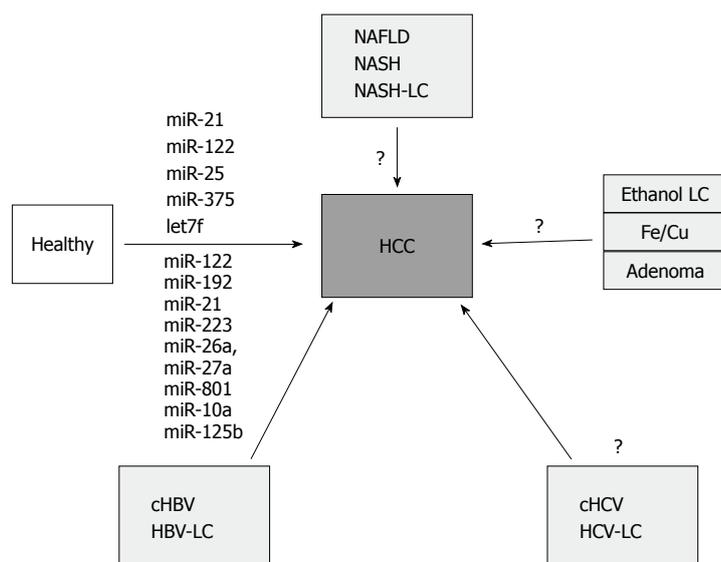


Figure 1 Schematic presentation of the potential microRNAs in diagnosis of hepatocellular carcinoma in relation to etiology. HBV-LC: Hepatitis B virus-liver cirrhosis; HCC: Hepatocellular carcinoma; NAFLD: Non-alcoholic fatty liver disease; NASH: Nonalcoholic steatohepatitis; cHCV: Chronic hepatitis C virus; cHBV: Chronic hepatitis B-virus; Fe: Hemochromatosis; Cu: M. Wilson; miR: microRNA.

approach has been also used to evaluate the expression differences of liver and tumor-related miRNA. In particular miR-21, which is most frequently deregulated miRNA in cancer, was found at higher level both in sera and plasma from HCC patients^[70,71], while others showed no significant expression differences^[72,73]. In similar fashion, miR-122, the most abundant miRNA of the liver, was also found at high level in sera from HCC patients^[71,72]. However, the level of circulating miR-122 may be strongly influenced by inflammation or apoptosis of hepatocytes in such conditions as acute or chronic hepatitis or nonalcoholic fatty liver disease, suggesting that background condition of the liver inflammation may strongly influence the miR-122 level^[74]. Nevertheless, the data from candidate-based studies correlate with the data from Link *et al.*^[54] at least for both miRNAs miR-21 and miR-122^[69]. There are several other studies that have identified additional miRNA, however, at present no independent validation has been performed (Figure 1). It is further more important to mention that the potential of miRNA as biomarker has not been equally analyzed in all HCC-related risk conditions. Systemic analyses for alcohol, NASH or HCV-related conditions are pending.

NON-INVASIVE PROGNOSTIC BIOMARKERS IN HCC

The molecular heterogeneity of HCC results in differences in outcome of affected patients. Clinically, main factors that have an impact on patient survival have been identified. These include tumor related factors like number and size of nodules, vascular invasion, existence of extrahepatic metastases, liver function and patient related factors. The Barcelona Classification summarizes these factors in a comprehensive algorithm and is endorsed by current guidelines^[10,11]. However, within the defined tumor stages

the survival of patients is still heterogeneous, and some patients that are treated in curative intent or even undergo liver transplantation show early recurrence of disease. Knowledge on high-risk profiles would therefore be important to guide individualized treatment.

Several of the non-invasive biomarkers that have been evaluated for their diagnostic power in HCC have also been studied for their prognostic significance.

High expression of AFP in serum correlates with high cell proliferation, high angiogenesis and low apoptosis and is associated with poor prognosis^[75,76]. The fraction of AFP-L3 is another prognostic biomarker for survival after resection of HCC^[77,78]. Patients that have undergone resection of HCC and had elevated levels of AFP, AFP-L3 and DCP at baseline had a worse prognosis than those patients that are positive for just one or two of the markers before surgery^[79]. The combination of AFP, the percentage of AFP-L3 and DCP combined with the concentrations of bilirubin and albumin, summarized in the BALAD score, is prognostic for survival of patients with HCC in an Asian population^[80]. Recently, a modification of this model, the BALAD-2 score, was validated in an international setting and confirmed to reliably predict the prognosis of patients with HCC^[81].

Other circulating biomarkers that mirror current knowledge on pathways involved in hepatocarcinogenesis and shown to be of prognostic value are, amongst others, IGF1^[82], DKK1^[83,84], GPC-3 and HSP 70^[85] although prospective validation studies are still to come. In patients with advanced HCC, baseline angiopoietin 2 (Ang2), and VEGF concentration in the plasma also independently predict survival^[76].

MIRNAS AS NON-INVASIVE PROGNOSTIC BIOMARKERS FOR HCC

In addition to their diagnostic potential, miRNAs may be helpful in prediction of the prognosis of HCC. Li

Table 3 Genetic signatures from tumor tissue and their prognostic significance

Ref.	Year	Correlation with	No. of genes in signature	AUC	P-value
Nault <i>et al</i> ^[95]	2013	Disease-free survival after resection	5 (tumor)	0.8	< 0.0001
Lim <i>et al</i> ^[96]	2013	Disease-free survival after resection	25 (tumor)		0.002
Kurokawa <i>et al</i> ^[97]	2004	Tumor recurrence after resection	20 (tumor)		0.001
Yoshioka <i>et al</i> ^[98]	2009	Tumor recurrence after resection	172 (tumor)		< 0.0001
Woo <i>et al</i> ^[99]	2008	Recurrence free survival	628 (tumor)		< 0.01

AUC: Area under the curve.

et al^[86] studied the expression of several miRNAs in sera from 46 HCC patients and 20 controls. Specifically, miR-221 was found in high concentration in HCC sera samples, which correlated with tumor size, cirrhosis and tumor stage. Kaplan-Meier survival analyses revealed an inverse correlation between miR-221 expression and survival rates. In another study, Tomimaru *et al*^[70] analyzed miR-21 expression in plasma from 126 HCC patients. MiR-21 expression was high in HCC and diminished after surgical treatment. Most importantly, high miR-21 expression level in plasma correlated with shorter cumulative survival following treatment. Köberle *et al*^[87] analyzed the performance of miR-1 and miR-122 in European HCC patients. Higher miR-1 and miR-122 serum levels were associated with longer overall survival compared to low expression of those miRNAs. However, miR-122, but not miR-1, showed a correlation with hepatic inflammation, liver function and synthetic capacity. The authors conclude that miR-1 may be a liver function independent predictive biomarker of HCC. There is also growing evidence that miRNA signature profiling can be useful in prognostic stratification^[88]. A signature of 31-miRNA correlates with stage of disease^[89]. A distinct 20-miRNA signature associated with metastases of HCC has also been identified^[90].

Despite of the promising potential, there are several pitfalls in utility and implementation of miRNA-based biomarkers in clinical practice. First, the majority of data comes from Asian populations with predominantly virus-related HCC (Figure 1)^[54]. However, in European or American populations the incidence of virus-related HCC is dropping and increases for NAFLD-related conditions therefore the data in these patients may probably be different. Second, the complexity of the miRNA alterations in the background of liver pathology (ex. chronic hepatitis with early fibrosis or cirrhosis) may impact the pattern of miRNA expression with increasing expression in one of the conditions and decreasing level in another. Furthermore, the ideal biomarker is probably the one that is expressed in HCC tissue with increasing concentration during progression of the disease. A combination of miRNA with the established-although not ideal-biomarker AFP may probably be beneficial^[70].

In the above section, we provided a brief insight into the growing field of miRNA-based biomarker research for HCC. Before this approach may be further utilized for clinical testing there are also critical technical questions that need to be answered. What is the best non-invasive specimen for the early diagnosis of HCC: plasma or serum? What normalizer is the best for the analyses? What is the best method for translational testing? Indeed, array-based analyses may be probably too expensive to apply, therefore, a candidate-based approach will need to be standardized for effective implementation. Those are only few reasons why the currently available data are so heterogeneous^[54,67]. Nevertheless, this miRNA-based approach may provide an additional value in personal-based management by prediction and application of new therapeutic targets^[91].

INVASIVE BIOMARKERS AS PROGNOSTIC TOOLS FOR HCC

After curative treatment of HCC the prognosis of the patient depends on the characteristics of the resected cancer but in addition on the risk of carcinogenesis due to the underlying etiology and inflammatory activity of chronic liver disease which persist after surgical resection or ablation. In a landmark study Hoshida *et al*^[92] demonstrated that gene-expression profiling can be performed in frozen as well as in formalin-fixed paraffin-embedded tissues and identified a gene-expression signature in liver tissue adjacent to tumor in patients who underwent resection of HCC that correlated with survival.

Since then, a large number of gene-expression profile studies has been performed in HCC with the aim to distinguish molecular subtypes. A validated and commonly accepted molecular classification has not been identified so far. Based on a meta-analysis of gene expression profiles from eight European cohorts of patients with HCC, a classification framework for HCC based on gene expression profiles was proposed that distinguishes three HCC subclasses, each correlated with clinical parameters such as tumor size, extent of cellular differentiation, and serum α -fetoprotein levels^[50,93]. The results of a

selection of recent studies with respect to prognostic gene expression profiles are summarized in Table 3. In a large validation study gene expression profiles of tumor and adjacent tissue were evaluated for their prognostic significance and a composite prognostic model was developed^[94]. A further validation of this model is pending before it may be considered for clinical use.

CONCLUSION

High-throughput technologies allow the identification of new molecules involved in complex pathways and their interaction in hepatocarcinogenesis. A single perfect biomarker has not been found so far to accomplish with the clinical demand for optimal HCC patient care. A combination of serological biomarkers may offer a better risk stratification of patients belonging to high-risk populations in the future. The combination of clinical characteristics and morphological signatures of tumor and the surrounding tissue will most likely be the best option for risk stratification and prediction of prognosis in patients with HCC in the future. Individualized treatment approaches that take into account the patient's own cancer genetic profile need to be addressed in further research^[12].

However, the critical step for translational research is to move the identified candidate signatures or single biomarkers from bench to bedside. There is a great hope that new molecular biomarkers can support clinicians in their daily routine and improve the care of patients with HCC. However, analyses tools need to be standardized and simplified in order to be useful, reliable and widely available.

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Gene therapeutic approaches to inhibit hepatitis B virus replication

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Abstract

Acute and chronic hepatitis B virus (HBV) infections remain to present a major global health problem. The infection can be associated with acute symptomatic or asymptomatic hepatitis which can cause chronic inflammation of the liver and over years this can lead to cirrhosis and the development of hepatocellular

carcinomas. Currently available therapeutics for chronically infected individuals aim at reducing viral replication and to slow down or stop the progression of the disease. Therefore, novel treatment options are needed to efficiently combat and eradicate this disease. Here we provide a state of the art overview of gene therapeutic approaches to inhibit HBV replication. We discuss non-viral and viral approaches which were explored to deliver therapeutic nucleic acids aiming at reducing HBV replication. Types of delivered therapeutic nucleic acids which were studied since many years include antisense oligodeoxynucleotides and antisense RNA, ribozymes and DNazymes, RNA interference, and external guide sequences. More recently designer nucleases gained increased attention and were exploited to destroy the HBV genome. In addition we mention other strategies to reduce HBV replication based on delivery of DNA encoding dominant negative mutants and DNA vaccination. In combination with available cell culture and animal models for HBV infection, *in vitro* and *in vivo* studies can be performed to test efficacy of gene therapeutic approaches. Recent progress but also challenges will be specified and future perspectives will be discussed. This is an exciting time to explore such approaches because recent successes of gene therapeutic strategies in the clinic to treat genetic diseases raise hope to find alternative treatment options for patients chronically infected with HBV.

Key words: Gene therapy; Hepatitis B virus; Antisense nucleic acid; RNA interference; Designer nuclease; Ribozyme; DNase; Dominant negative mutant; External guide sequence; DNA vaccination

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Core tip: With various successful clinical trials ongoing, gene therapeutic approaches gained increasing

attention in the community over the recent years. Here we introduce gene therapy as a versatile platform for treatment of hepatitis B (HBV) virus infection. Newest delivery methods based on non-viral and viral techniques combined with most advanced technologies for inhibition of HBV replication based on DNA, RNA and designer nucleases are discussed. An overview of various gene therapeutic systems which were explored *in vitro* and *in vivo* is provided. Advantages but also limitations of the different strategies to inhibit HBV replication are mentioned.

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INTRODUCTION

Hepatitis B virus

The hepatitis B virus (HBV) is an enveloped, partially double-stranded DNA virus which replicates through an RNA intermediate. Upon cell entry the DNA containing core particle is transported to the nucleus where the DNA is released. Next the partial double-stranded DNA is repaired by host enzymes to form the covalently closed circular DNA (cccDNA). The cccDNA serves as a template for transcription of viral proteins and reverse transcription of new viral genomes by the viral polymerase. The large 3.5 kb HBV transcript represents the pregenomic RNA serving as a template for virus replication. Furthermore, there are three additional mRNAs with a length of 2.4, 2.1, and 0.9 kb. The HBV genome is 3.2 kb in size and contains four overlapping major open reading frames tightly arranged that encode polymerase, surface (HBsAg), core (HBcAg) and X proteins (HBx)^[1,2]. In addition, especially early during infection the HBV early antigen (HBeAg) can be detected which is a proteolytic product of the pre-core protein.

HBV infection counts as a major global health problem since more than two billion people show evidence of a past or present infection with the virus. This hepatotropic virus can cause acute and chronic infection of the liver. Fortunately, for most people the infection proceeds nearly without symptoms when taking an acute course of disease and complete recovery is likely. However, 240 million people suffer from chronic HBV infection and more than 780000 people die every year because of hepatitis B related secondary diseases. Mostly newborns and infants are prone to develop the chronic type of the infection^[3] and so far no treatment is available that reliably cures those patients.

Current therapeutics for chronic HBV infection are

intended to reduce viral replication and slow down or stop the progression of the disease. To date there are seven Food and Drug Administration (FDA) approved compounds for the treatment of chronic hepatitis B. These include interferon alpha and pegylated interferon alpha, nucleoside analogues (lamivudine, entecavir and telbivudine) and nucleotide analogues (adefovir, dipivoxil and tenofovir)^[4].

Interferon alpha has an antiviral effect by inhibiting the synthesis of viral DNA and activating antiviral enzymes and additionally, acts in an immunomodulatory way by enhancing the cellular immune response against infected cells^[5]. It has to be administered daily or three times a week as unmodified version and once in a week in the pegylated form. The main disadvantages of interferon alpha are the parenteral administration causing discomfort to the patients and potential adverse effects such as flu-like symptoms in the beginning of treatment and later on for instance fatigue and low blood counts. It is only given to selected patients because under certain conditions administration of interferon alpha is contraindicated^[4]. In a long-term follow-up study of HBeAg-positive patients, 11% lost HBsAg after treatment with interferon alpha^[6] which is considered as a cure of the disease.

Nucleos(t)ide analogues interfere with the HBV replication primarily by targeting the HBV polymerase functions such as reverse transcriptase and DNA polymerase activity^[5]. These drugs are administered orally as a daily dose. The major limitation associated with nucleos(t)ide analogues is the emergence of antiviral drug resistance and that life-long treatment can be indicated in the presence of chronic infection. In this context failure of medication adherence is another problem, because viral relapse is common when ending the treatment. Another prognostic marker of HBV infection is the presence of HBeAg which correlates with high viral replication rates. However, HBeAg seroconversion can be achieved with nucleos(t)ide treatment. In addition, HBV DNA levels can be decreased to an undetectable level but at the same time HBsAg is not lost^[4]. These features demonstrate another peculiarity of the hepatitis B virus. After entry into a cell the cccDNA is maintained as an episomally maintained template in the nucleus. It is not attacked by nucleos(t)ide analogues nor by interferon in general, so that it is able to serve as a reservoir from which previously cleared or treated infections can recur^[4]. The clinical management of chronic hepatitis B infection is reviewed in detail by Santantonio and Fasano^[7].

The viral reservoir and potential reactivation of the virus represents a major problem when developing novel HBV treatments options and this is a challenge that could be faced by gene therapy. Researchers seek to inhibit viral replication in a long-lasting manner without the need for continuous drug administration through gene therapeutic approaches.

The more ambitious goal is to completely eradicate the viral cccDNA depot and hence find a true cure for chronic hepatitis B virus infection. Here we discuss gene therapeutic approaches as a versatile platform to combat HBV infection.

Gene therapy

Gene therapy is a strategy to transfer therapeutic nucleic acids into the desired target cell for treatment of a variety of different diseases. To efficiently deliver the genetic payload, multiple gene transfection techniques were explored which can be subdivided into two major groups: virus-based and non-viral vector systems for delivery of respective therapeutic nucleic acid. Both delivery techniques were also utilized in gene therapeutic approaches to treat chronic infectious diseases such as HBV infection. Since HBV infection resides in liver, the majority of gene transfer approaches were focused on targeting hepatocytes.

Non-viral vectors are based on delivery of naked RNA or DNA which in combination with chemical and physical means can result in efficient delivery of the nucleic acid into the respective target cell^[8,9]. Chemical methods in the context of non-viral vector delivery rely on various chemical formulations such as cationic lipids^[10] and polymers including polyamidoamine dendrimers and polyethylenimine (PEI)^[11]. All chemical reagents were explored in different approaches and there are several commercially available transfection reagents which are commonly used for transfection of DNA and RNA resulting in sufficient transfer efficiencies in many cell lines *in vitro*. Major constraints of these methods are transfection reagent-associated toxicity and the difficulty to cross the nuclear membrane. In addition to chemical transfer methods, physical transfer techniques were explored involving needle injection^[12], gene gun^[13], electroporation^[14], sonoporation^[15], magnetofection^[16], and hydrodynamic gene transfer^[17]. These methods directly deliver therapeutic nucleic acid into the cytosol of the target cell and compared to chemical methods these techniques harbor a reduced risk of transfection-mediated side effects due to dispersion of the transfection reagent. However, limitations of these methods are exposed by the difficulty to cross the nuclear membrane, potential cellular damage caused by the transfection method, and the requirement of costly instruments.

Virus-based transfection techniques were utilized in numerous pre-clinical and clinical gene therapeutic applications. Predominantly used viral vectors can be attributed to three viruses which were converted into viral vectors by deletion of essential viral genes: adenovirus, adeno-associated virus (AAV) and retrovirus. All viral vector systems display advantages and disadvantages which were discussed in more

detail in previous reviews^[18]. Adenoviruses combine a large transgene capacity of up to 36 kilo bases (kb), an episomal nature of the adenoviral genome reducing the risk of genotoxicity, the possibility to produce high viral titers and the ability to transduce dividing and non-dividing cells at high efficiencies *in vitro* and *in vivo*^[19]. However, one major obstacle for *in vivo* applications are the innate and the adaptive immune responses induced by the incoming adenoviral particle. AAV vectors were explored in clinical trials, are non-pathogenic, lead to a reduced immune response and predominantly exist as extrachromosomal vector genomes in the transduced cell^[20]. One major disadvantage, however, is the small transgene capacity which is below 5 kb. Lentiviral vectors^[21] were broadly explored in clinical trials to treat rare genetic diseases in *ex vivo* gene therapeutic approaches. Various generations of lentiviral vectors are available which carry a transgene capacity of up to 8 kb. Although commonly used lentiviral vectors integrate their genetic cargo into the host genome, newest versions of these vectors can circumvent side effects associated with somatic integration by changing their integration profile.

Various non-viral and viral transfer techniques were exploited to combat HBV infection *in vitro* and *in vivo* which will be discussed in the following paragraphs.

GENE THERAPEUTIC APPROACHES AGAINST HBV

Various gene therapeutic approaches to treat HBV infection were studied in cell culture models and in animal models for HBV infection. Within the viral life cycle in an infected cell there are various points of attack which can serve as targets in gene therapeutic approaches to inhibit HBV replication. Figure 1 schematically shows the life cycle of HBV infection and indicates points of attack when considering a gene therapeutic treatment.

As shown in Figure 1 the mechanisms of viral inhibition in gene therapeutic approaches can be on the level of RNA (HBV derived transcripts), DNA (cccDNA) and proteins. On the RNA level antisense oligodeoxynucleotides and antisense RNA, catalytic nucleic acids such as ribozymes and DNazymes, RNA interference and external guide sequences (EGS) can be considered. On the level of DNA (cccDNA) as a potential target designer nuclease such as zinc finger nuclease (ZFN), transcription activator-like effector nucleases (TALEN) and the *clustered regularly interspaced short palindromic repeats* (CRISPR)/Cas9 technology can be used. On the level of proteins, dominant negative HBV mutants and a strategy based on capsid-targeted viral inactivation (CTVI) were studied. Another technology

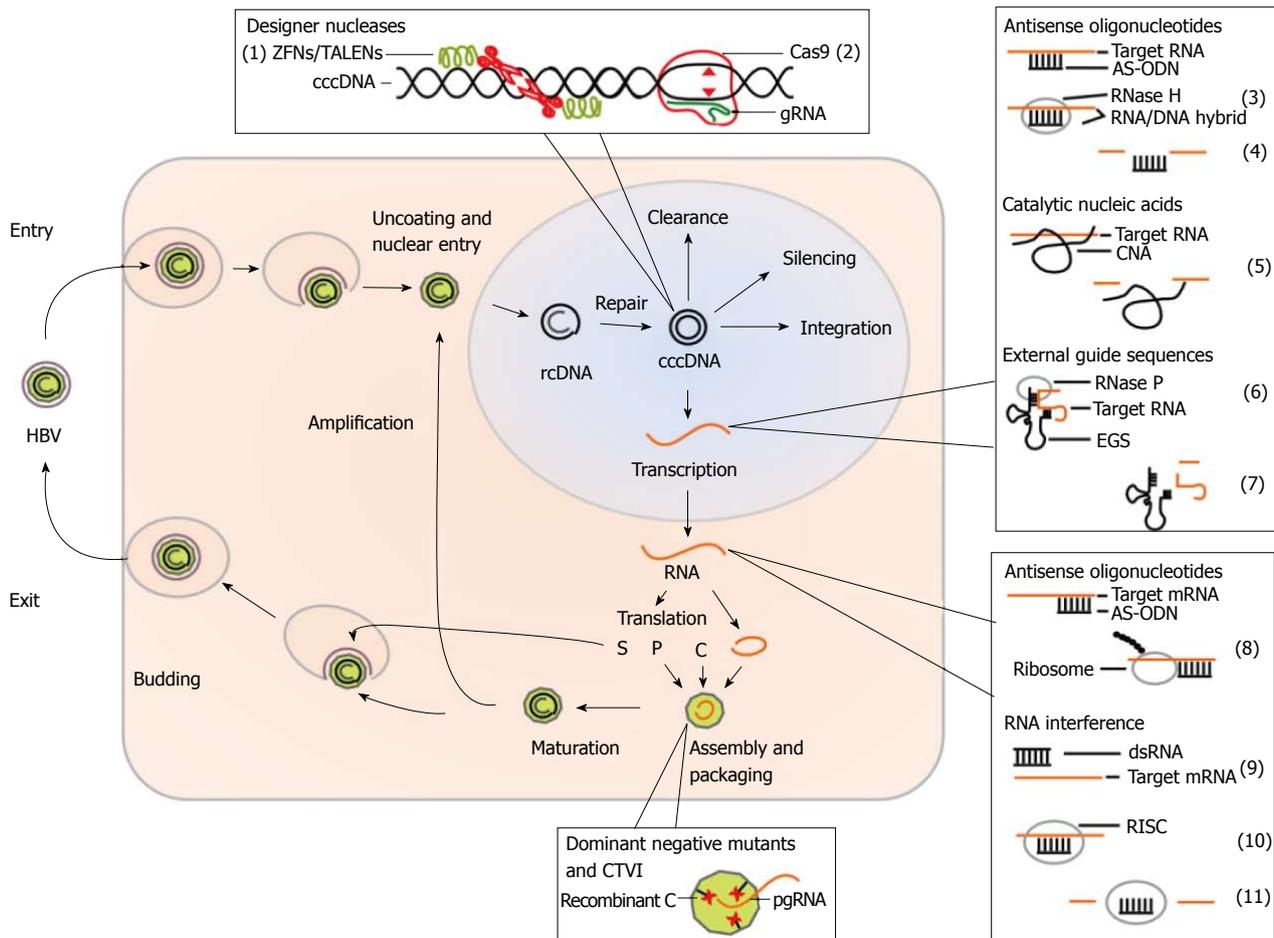


Figure 1 Hepatitis B virus replication cycle and gene therapeutic strategies. Enveloped virions of hepatitis B virus (HBV) infect liver cells *via* attachment to the cell membrane and endocytosis. The capsid with the relaxed circular (rc) DNA is released into the cytoplasm and the DNA is uncoated upon nuclear entry. In the nucleus the rcDNA is repaired to form the covalently closed circular (ccc) DNA. The cccDNA can be eventually cleared out, silenced or integrated into the host genome. Predominantly, it persists as an episome in the nucleus and is transcribed and translated by the host cell machinery. One of the transcripts forms the pregenomic (pg) RNA which is encapsulated together with the translated viral polymerase (P) by the translated viral capsid proteins (C). In the newly assembled nucleocapsid the pgRNA serves as a template for the viral polymerase which synthesizes the rcDNA. The nucleocapsid either migrates back to the nucleus to increase the pool of cccDNA or is internalized by the endoplasmic reticulum (ER). In the latter process it is enveloped with ER-membrane that harbors translated viral surface proteins (S) and finally released from the cell. Gene therapeutic strategies act on several steps of the viral replication cycle. Designer nucleases are intended to promote disruption of the cccDNA. Zinc finger nucleases (ZFNs) and transcription activator-like effector nucleases (TALENs) (1) use protein-based DNA-binding modules, while the CRISPR (clustered regularly interspaced short palindromic repeats)/Cas9 nuclease system (2) is directed by a guide RNA (gRNA) to the target site. Antisense oligonucleotides (AS-ODNs) act inhibitory on the viral replication in two ways. In the nucleus they recruit RNase H (3) after the formation of RNA/DNA hybrids, which cleaves the viral RNA (4). In the cytosol they bind to the viral RNA which leads to a steric blockade of subsequent processes (8). Catalytic nucleotides (CNA) as ribozymes and DNAzymes are able to cleave targeted RNA by themselves (5). External guide sequences (EGS) are designed in a way that they resemble precursor tRNAs (6) when they bind to their target RNA and trigger cleavage by RNase P (7). RNA interference can be induced by different dsRNA species (9). Longer dsRNAs can be delivered exogenously or expressed in the cells as shRNAs. The RNAs are processed to approximately 21 nucleotide long dsRNAs termed siRNAs which are used to form the large RNA-induced silencing complex (RISC) (10) which degrades target messenger RNAs (mRNAs) (11). Dominant negative mutants of capsid proteins form a hindrance for proper packaging of viral progenitor RNA. In contrast, capsid-targeted viral inactivation (CTVI) is an approach where the capsid proteins are additionally fused to a destructive compound.

for prevention of infection but also for a potential treatment option of chronically infected patients, DNA vaccination can be considered as an attractive alternative. Although the majority of the described strategies for inhibition of HBV replication were not translated into the clinic so far, we believe that gene therapy may represent a valuable alternative in the future. Studies describing milestones of the various gene therapeutic approaches are listed in Table 1 and are described in more detail in the following sections.

Antisense nucleic acids

The beginning of gene therapy against HBV can be linked to the first tests of antisense oligodeoxynucleotides (AS-ODNs) directed against the HBV genome^[22-24]. With respect to antisense nucleic acids post-transcriptional inhibition is achieved by blockade of ribosomal access, inhibition of ribosomal assembly and induction of RNase H cleavage^[25].

First *in vivo* studies showed applicability of this approach in duck-HBV-infected Peking ducklings. Infected animals were treated daily by intravenous

Table 1 Advances in hepatitis B virus infection gene therapy in chronological order

Year	Strategy	Milestone	Ref.
1990	AS-ODN	First <i>in vitro</i> application	[22]
1992	AS-ODN	First non-viral transfection (targeted polycation peptide complex)	[28]
	Ribozyme	First <i>in vitro</i> application	[44]
1993	AS-ODN	First <i>in vivo</i> application	[26]
	DNA vaccination		[75]
1994	Dominant negative mutants	First <i>in vitro</i> application	[65]
1997	AS-ODN	First viral transduction (retroviral)	[39]
	Ribozyme		[46]
1998	DNAzyme	First <i>in vitro</i> application	[60]
	EGS		[87]
2001	CTVI		[69]
2003	RNAi	First <i>in vitro</i> application	[93]
		First <i>in vivo</i> application	[94]
2004	Ribozyme	First <i>in vivo</i> application	[56]
	DNA vaccination	First clinical trial in chronic HBV carriers	[81]
2008	CTVI	First <i>in vivo</i> application	[72]
2010	ZFN	First <i>in vitro</i> application	[138]
2011	RNAi	First clinical trial in chronic HBV carriers	[117]
2013	EGS	First <i>in vivo</i> application	[89]
	TALEN	First <i>in vitro/vivo</i> application	[146]
2014	CRISPR/Cas9	First <i>in vitro/vivo</i> application	[148]

AS-ODN: Antisense oligonucleotides; HBV: Hepatitis B virus; EGS: External guide sequence; CTVI: Capsid-targeted viral inactivation; RNAi: RNA interference; ZFN: Zinc finger nuclease; TALEN: Transcription activator-like effector nuclease; CRISPR: Clustered regularly interspaced short palindromic repeats; Cas9: Cas9 nuclease.

injection of AS-ODNs for ten days. The treatment resulted in nearly complete inhibition of viral replication which was assessed by liver DNA analysis for DNA replicative intermediates and blockade of viral gene expression as demonstrated by disappearance of surface antigen in serum and core antigen in liver^[26]. AS-ODNs proved to be most effective when directed against the initiation site of the HBsAg-gene^[22] or at the encapsidation signal^[27]. Except for a study published by Wu *et al.*^[28] in which already targeted DNA complexes were used, delivery of AS-ODNs was initially restricted to simple cellular uptake and binding of the unmodified antisense DNA to its target sites. Soon the system was improved by enhancing stability of the respective nucleic acids and by increasing uptake of AS-ODNs by the chosen target cell^[29-32]. In other studies AS-ODNs were conjugated to ribonuclease H or manganese porphyrin which after binding to the target site can lead to cleavage of the desired target sequences^[33,34]. Furthermore, DNA carrier systems were used and also enhanced by making them targetable to hepatocytes^[28,35,36].

Other oligonucleotide based approaches include antisense RNA delivered by episomally replicating expression vectors^[37,38] or retroviral vectors^[39,40]. The advantage of these approaches is the fact that antisense RNAs can be expressed in the cells, whereas AS-ODNs have to be exogenously delivered. This allows for experimental settings with long-term effects. For instance efficacies lasting longer than ten months were observed after stable transfection of antisense RNA expression vectors^[37]. Antisense RNA-

mediated inhibition functions preferentially through destabilization of the sense RNA by targeting the antisense/sense-RNA duplex to dsRNase^[41].

A completely different idea unrelated to complementary antisense RNA was introduced by Hafkemeyer *et al.*^[42] and is based on so-called "antisense-toxin-RNA". In this approach the authors took advantage of the HBV reverse transcriptase in infected cells to selectively kill those cells through *Pseudomonas* exotoxin expression from reverse transcribed antisense-toxin-RNA.

Site-specifically cleaving nucleic acids

There are two types of catalytically active nucleic acids, ribozymes and DNAzymes. Ribozymes are naturally occurring RNA molecules that can execute enzymatic activity on itself in the absence of proteins (in *cis*) or on extrinsic targets (in *trans*). DNAzymes were generated by *in vitro* evolution. They resemble ribozymes and do not exist in nature. The substrate for both species is RNA. There are various types of ribozymes known, hammerhead and hairpin ribozymes being the most popular ones. All catalytic nucleic acids have in common that they consist of an antisense sequence recognizing the target site and a catalytic domain mediating cleavage^[43]. Scientists were able to manipulate the recognition of target sites, rendering such nucleic acids very attractive for enhanced transient knockdown of gene expression.

The first experiments with ribozymes targeting the HBV genome were performed in 1992 using a triple ribozyme construct. In this study three hammerhead ribozymes were encoded on a single

DNA template and it was shown that they were simultaneously active *in vitro*. However, cleavage kinetics were similar to single ribozyme constructs. Nonetheless, this approach may still be favorable because it enables facing high target variability and emergence of viral resistance^[44]. However, first studies performed in a cellular context disclosed a first drawback of the hammerhead ribozymes because they were active after *in vitro* transcription and in Mg²⁺-supplemented cell extracts, but not in intact cells. The authors suggested some non-viral block, or inappropriate target site selection being responsible for the lack of intracellular activity^[45].

However, to overcome these problems novel strategies were pursued using other types of ribozymes. Welch and colleagues^[46] designed several hairpin ribozymes against different conserved regions of the HBV genome and tested them in a human hepatoma cell line (Huh7) transfected with full-length HBV genomes. The ribozymes were transduced into target cells using a retroviral vector system. The HBV production could be inhibited to up to 83% assayed through an endogenous polymerase assay. In the following years further *in vitro* studies were performed using different types of modified ribozymes^[47-51]. These were predominantly delivered *via* transfection of an expression plasmid into various cell lines which were additionally transfected with a HBV genome-containing plasmid resulting in varying effects on inhibition of HBV replication^[52-55].

In 2004 the first *in vivo* experiment in a transgenic mouse model was conducted by Pan *et al.*^[56]. For this study a self-processing triple-ribozyme cassette was used, with ribozymes acting in *cis* and in *trans*. The constructs were packaged in liposomes that were targeted to hepatocytes in the presence of asialofetuin. Quantitative PCR analysis for quantification of HBV genome copy numbers in murine liver showed a more than 80% decrease of HBV genome copy numbers and immunohistochemistry revealed a robust reduction in the number of hepatocytes staining positive for HBV core antigen. This was the first proof of concept demonstrating *in vivo* feasibility of viral RNA degradation mediated by ribozymes.

Next, recombinant hepatitis D virus (HDV)^[57] as well as lentiviral vectors were utilized as delivery vehicles for respective ribozymes which achieved effective reduction of HBV mRNA levels over four months^[58]. Furthermore, HDV-derived ribozymes were also utilized which were delivered by a pseudotyped retroviral vector (Moloney murine leukemia virus)^[59]. According to the authors the main advantages of these ribozymes are the natural activity of HDV ribozymes in human cells even at physiological Mg²⁺-ion concentrations and the comparably highest cleavage rates among all known ribozymes. Their results revealed significant reduction in the intracellular HBV DNA concentration

in HepG2.2.15 cells, which secrete infectious HBV virions. Furthermore, decreased extracellular HBsAg and HBeAg levels were observed after treatment with HDV ribozymes in comparison to the negative control. The conclusion was that using this strategy, HBV can be effectively inhibited at post-transcription and replication levels.

DNAzymes cleave RNA substrates based on a similar mechanism also used by ribozymes. However, they may be superior to ribozymes because their production is comparably straight forward and DNAzymes are less sensitive to chemical and enzymatic degradation. The first DNAzyme directed against HBV mRNA was created by Asahina *et al.*^[60]. They targeted the direct repeat 1 (DR1) and polyadenylation signal regions of HBV. In this study the authors used stabilized forms of DNAzymes that on the one hand lost some degree of activity compared to unmodified versions but on the other hand the degradation level was less pronounced. The DNAzyme was tested in Huh7 cells on an HBV-luciferase fusion reporter system where it exhibited 48% suppression compared to untreated control groups. Further *in vitro* studies^[61-64], however, revealed a major disadvantage of this system because intracellular expression of these DNA species is not feasible. DNAzymes have to be transfected directly because they act on post-transcriptional level which is clearly disadvantageous if long-term administration for instance in clinical applications is required.

Dominant negative mutants and CTVI

Dominant negative mutants of viral proteins are able to inhibit viral replication by interfering with the function of the wild type protein. The first study on the molecular effects of dominant negative mutants on the HBV replication was conducted in 1994 by Scaglioni *et al.*^[65] They mutated the core protein and observed an inhibition of viral replication by 90%-95%^[65] and it was concluded that this was the result of the disruption of the viral nucleocapsid assembly process. In a follow-up study the authors provided a delivery system using retroviral and adenoviral expression vectors^[66]. von Weizsäcker *et al.*^[67] showed that carboxy-terminal, but not amino-terminal core mutants inhibit viral replication. Furthermore, it was discussed that rather the packaging of the viral pre-genome and the reverse transcription reaction within the particles than the nucleocapsid formation itself is restrained by dominant negative core mutants^[68]. However, it was also shown that some mixed particles retain replication competency suggesting a possible mechanism of viral escape.

Using the principle of CTVI may represent an interesting variant of capsid modification, as the goal is to introduce a destructive element into the virus. The first time this technology was pursued

was by fusing a Ca²⁺-dependent nuclease from *Staphylococcus aureus* to the HBV core protein. This led to inhibition of proper synthesis of viral DNA inside the capsid and rapid viral DNA degradation^[69]. Later human eosinophil-derived neurotoxin was fused to the HBV core protein by another group and thoroughly studied also *in vivo*^[70-72].

Therapeutic DNA vaccination

Initial considerations to use vaccination in a therapeutic manner against HBV infection were made in 1993 by a group around Christian Bréchet^[73,74]. This approach was based on classical vaccination compounds. Also in 1993 the first genetic immunization with expression vectors for HBsAg was introduced in mice^[75]. In a further step Mancini *et al.*^[76] proved that a combination of both strategies is feasible and they showed efficacy in a HBsAg transgenic mouse as a model for chronic HBV infection. A single intramuscular injection of HBV envelope encoding plasmid DNA resulted in the elimination of serum HBV antigen levels as early as 4 wk after administration of the therapeutic DNA in some mice. Notably no liver cell injury was detected. Soon genetic immunization employing retroviral delivery was tested first in rhesus monkeys^[77] and then in chimpanzees chronically infected with HBV^[78]. The first time a DNA vaccine encoding HBsAg delivered with a gene gun was tested in healthy humans was in 1999 and it turned out to be ineffective at the applied dosage (0.25 µg)^[79]. However, the vaccine was well tolerated. A subsequent study used higher dosages (1, 2, or 4 µg) and this time antigen-specific CD8⁺ T cells, T helper cells, and protective levels of antibodies could be induced^[80]. Therapeutic DNA vaccination for chronic HBV infection was first tested in 2004 and in other subsequent studies^[81-83]. It was shown that HBV DNA vaccination was safe and immunologically effective because T-cell responses were activated in some chronic HBV carriers who did not respond to current antiviral therapies. These results indicated that in combination with conventional therapy, DNA vaccination may have a synergic effect leading to complete recovery from HBV infection.

EGS and RNA interference

The recruitment of ribonuclease P (RNase P) by EGS represents another interesting gene interference strategy. RNase P is an enzyme that removes the leader sequence of tRNA precursors by recognizing a common structure shared amongst all tRNAs. Using a custom-designed EGS that hybridizes with an mRNA to form a tRNA-like structure, RNase P can be recruited to cleave the target mRNA^[84,85]. With respect to HBV infection, this strategy was first proposed by Werner *et al.*^[86] in 1997. They took advantage of EGS designed to target HBV sequences

as proof-of-concept for the use of EGS as anti-viral or anti-cancer therapeutics^[87,88]. The EGS technology gained more attraction in 2013, when EGS was considered as a sufficient method to antagonize HBV replication. One study was conducted in cell culture and in mice using *Salmonella*-mediated delivery. Oral inoculation of attenuated *Salmonella* carrying the EGS construct led to up to approximately 95% inhibition of HBV gene expression levels and a approximately 200000-fold reduction of viral DNA levels in the livers and sera of the treated mice transfected with a HBV plasmid^[89]. In a follow-up study efficacy could be improved about 50-fold by modifying the EGS derived from natural tRNA through *in vitro* selection^[90].

RNA interference (RNAi) is a natural intracellular antiviral immune response mechanism triggered by double-stranded RNA (dsRNA). After conversion of dsRNA into guide molecules (siRNA), dsRNA induces the degradation of RNA *via* the RNA-induced silencing complex (RISC). Synthetic siRNA can be delivered exogenously or produced endogenously in the form of precursor small hairpin RNAs (shRNA) from plasmid DNA or viral vectors^[91]. When RNAi was first discovered^[92] it released a wave of multiple studies and applications. Also in HBV therapy it gained attention. In 2003 several studies were published^[93-97] and marked the beginning of the most intense exploitation of a gene therapy method as a valuable treatment option for HBV infection.

Shlomai *et al.*^[93] were the first to publish the use of RNAi in an *in vitro* cell culture model of HBV replication. They found that HBV gene suppression was achieved with different efficacies for different gene targets. Analysis of HBV transcripts revealed a reduction of about 68% on the level of all viral transcripts for RNAi targeting HBx protein. However, RNAi against two different targets on the same HBcAg open reading frame, which is exclusively encoded by the large 3.5 kb transcript, resulted in about 13% and 50% reduction on the level of the 3.5 kb transcript, respectively. This led to the conclusion, that RNAi target sequences have to be evaluated carefully.

In the next study McCaffrey *et al.*^[94] tested the RNAi-based approach *in vivo* in immunocompetent and immunodeficient mice. DNAs containing a HBV expression plasmid and a shRNA expression plasmid were co-transfected into mouse liver by hydrodynamic plasmid delivery. The authors could show that RNAi can inhibit all the steps of HBV replication that occur in cell culture and in mice. This was indicated by reduced secreted HBsAg levels in the supernatant of transfected cells and in mouse serum, reduced HBV RNAs levels in mouse liver, reduction of HBV genomic DNA to undetectable levels in mouse liver, and decreased numbers of cells stained positive for HBcAg.

The first viral vectors used to transduce RNAi sequence expression cassettes were either based on a prototype foamy virus (PFV) or an adeno-associated virus (AAV)^[98]. The vectors expressing the respective RNAi molecule were assessed in 293T.HBs cells, a cell line stably expressing HBsAg and HepG2.2.15 cells. In 293T.HBs cells HBsAg was knocked down by approximately 90% if directly compared to controls cells. HBsAg expression was also inhibited in HepG2.2.15 cells even in the presence of HBV replication. Uprichard *et al.*^[99] introduced recombinant adenovirus vectors for the delivery of shRNAs. Additionally, they used for the first time HBV-transgenic mice to show that ongoing HBV replication *in vivo* can be cleared by RNAi-targeted suppression of viral RNA for at least 26 d. Other viral vectors used included retroviral vectors^[100,101], AAV-7, -8 and -9 pseudotyped vectors^[102-104], recombinant human foamy virus^[105], gene-deleted adenoviral vectors^[106,107], lentiviral vectors^[108,109], and recombinant baculovirus^[110].

Further studies of the RNAi system included lipid-encapsulation of chemically modified siRNAs for non-viral delivery^[111] and the assessment of efficacy and pharmacodynamic properties of different RNAi target sequences and constructs, including methyl-modified siRNAs and plasmid based DNA vectors^[112]. Moreover, high-throughput generation and screening of siRNAs was established^[113] and expression systems introducing multiple siRNAs were developed^[105,114-117]. Ely *et al.*^[116] found that a Pol II promoter may be advantageous compared to a Pol III promoter, which was traditionally used for shRNA expression. The Pol III promoter can result in shRNA overexpression and saturation of the endogenous microRNA pathway leading to serious toxic effect *in vivo*^[118]. This could be restricted with the Pol II promoter, which provides the possibility to control the production of RNAi activators^[119].

Another non-viral vector system which was used in shRNA approaches is the episomal replicating plasmid vector pEPI-1. Herein, the transcription unit is linked to a scaffold/matrix attachment region (S/MAR) which ensures that the vector is mitotically stable in transfected cells. It was shown that it provides long-term expression of shRNAs which resulted in suppression of HBV gene expression, intracellular HBV DNA replication and release of progeny HBV over 8 mo^[120].

Besides therapeutic DNA vaccination, siRNA is the only gene therapeutic approach that was translated into clinical trials. In 2006 a Phase Ib, first-in-human safety and tolerability study of an RNAi-based therapy (NUC B1000) in patients with mild to moderate chronic HBV infection was conducted^[117]. NUC B1000 is composed out of four expressed shRNAs on one plasmid carried on a nanoparticle (cholesteryl spermine complex) and administered through intravenous infusion. The results revealed elevated cytokines and no HBV DNA or HBsAg

decrease in the patients. However, the safety profile of RNAi therapy conducted among patients with HBV was considered as reasonable. A second compound, ARC-520, a liver-tropic cholesterol-conjugated siRNA (chol-siRNA), transported by the proprietary Dynamic Polyconjugate delivery system is just reaching a Phase II clinical study^[121,122].

In summary the RNAi system was thoroughly exploited for treatment of chronic HBV infection in the past and research will be ongoing on this topic to overcome major hurdles like evocation of immune responses and maintenance of long-term suppression. Long-term suppression is required because it was shown that the HBV cccDNA depot in the host cells is not affected by this approach^[110]. Interestingly, another limitation might be the manipulation of the host RNAi defense by the HBx protein that potentially functions as a RNA-silencing suppressor (RSS)^[123].

Designer nucleases

For sequence-specific DNA targeting designer nucleases such as zinc finger nucleases (ZFNs)^[124-126], transcription activator-like effector nucleases (TALENs)^[127,128] and the clustered, regularly interspaced, short palindromic repeats (CRISPR)-CRISPR-associated protein (Cas) system^[129-131] can be applied. They combine customizable DNA binding molecules for sequence-specific DNA-binding and a nuclease for introduction of double-strand DNA (dsDNA) breaks. The induced dsDNA breaks activate different cellular DNA repair pathways. The two most exploited pathways in gene therapy are homologous recombination (HR) and nonhomologous end-joining (NHEJ). In the presence of a respective homologous donor DNA, cells are able to repair the dsDNA break *via* HR by exchanging the respective sequence^[132]. Without any homologous donor DNA cells repair dsDNA breaks *via* NHEJ. This error-prone repair mechanism can lead to insertions or deletions of one or several base pairs and may cause specific knockout of a gene^[132]. Therefore, designer nucleases are valuable tools to specifically introduce knock out mutations at a desired DNA locus.

The DNA binding domains (DBD) of ZFNs commonly contains 3-4 zinc fingers. Each zinc finger consists of 30 amino acids and forms two β -sheets and one α -helix. Upon DNA-binding the α -helix is placed in the major groove of the dsDNA and directs contact with a certain base pair triplet. Depending on the amino acids within the α -helix and the number of zinc fingers, ZFNs can be designed specifically to target defined stretches of DNA triplets with high affinity^[133-135]. The DBD is connected to a sequence independent cleavage domain of the type II S restriction enzyme *FokI* which causes double strand breaks after dimerization^[136]. Therefore two ZFN monomers are necessary to create the desired dsDNA break in the spacer between the binding sites of

the ZFNs. The double strand breaks will be repaired *via* NHEJ, causing insertion and deletion mutations (indels) of several base pairs within the sequence. These indels can lead to frame shifts within the open reading frame or translation abortion by newly formed stop codons. Both options result in dysfunctional proteins.

There are some approaches using zinc fingers to target HBV cccDNA. Zimmerman *et al.*^[137] were the first to use zinc fingers in conjunction with HBV although they did not yet use ZFNs. Instead, they created several zinc finger proteins (ZFPs) targeting the enhancer region of duck HBV (DHBV), which probably form a steric hindrance for the RNA polymerase. After screening the candidates for binding efficiency, the two most efficient ZFPs were expressed in a special DHBV tissue culture system *via* transfection and both the transcription of viral genomic RNA and viral protein production was assessed *via* quantitative PCR and western blot analysis. They showed that both ZFPs significantly reduce transcription from cccDNA compared to controls. The authors concluded that ZFPs designed to target HBV DNA are able to substantially reduce viral transcription and interfere with viral replication of cccDNA.

In another study, Cradick *et al.*^[138] showed that ZFNs can mediate inhibition viral replication *in vitro*. They generated ZFN pairs targeting several conserved regions of HBV genomic DNA and chose the most robust pairs for further studies. Huh7 cells were transfected with plasmids encoding ZFNs under the control of a cytomegalovirus (CMV) immediate early promoter and the target plasmid pTHBV2. This target plasmid contains a 1.3-fold HBV genome which is capable of full transcription of the viral RNAs, translation of the proteins and production of infectious virus. Three days post transfection, Southern blotting analyses revealed about 26% linearized DNA and about 10% cleaved target plasmids being rejoined in a tail to tail orientation. Both DNA species indicate that ZFN-specific cleavage at the intended ZFN target site occurred. Linear genomes are the result of direct cleavage without any subsequent repair while concatamer formation is caused by NHEJ after ZFN cleavage. To investigate if NHEJ led to indels in recircularized genomes or head-to-tail concatamers, a *Xba*I site was inserted within the spacer region of the ZFN dimer. After NHEJ occurred this site should be destroyed. Target sites of treated samples were amplified by PCR, digested with *Xba*I and resistant amplicons were sequenced. 13 of 16 samples showed a frameshift which would lead to dysfunctional proteins. Furthermore this study indicated a 29% reduction of pregenomic RNA in northern blot analysis compared to controls. Besides these results it could also be demonstrated that the ZFN pairs caused only moderate toxicity. In conclusion, Cradick *et al.*^[138] revealed that

specifically designed ZFNs targeting regions which are conserved among many different HBV serotypes can significantly reduce viral replication which is associated with negligible toxicity.

A third study by Weber *et al.*^[139] deals with the delivery of functional ZFNs into target cells *via* a viral vector system. Here, in contrast to the latter study, obligate heterodimeric ZFNs were designed, which are not able to form homodimers. This measure minimizes off-target effects because ZFNs that are able to form homodimers might cleave at unintended genomic loci by tolerating some mismatches. The three designed ZFN pairs targeted the open reading frames of HBx, HBcAg and polymerase. For delivery a self-complementary AAV-vector (scAAV) was used, which shows higher transduction efficiencies compared to single stranded AAV vectors^[140]. Since scAAV vectors have a reduced transgene capacity each ZFN of a pair was delivered individually in co-transduction experiments. Transduction of each ZFN pair into HepAD38 cells, a model cell line for controllable HBV replication, revealed mutation rates ranging from 9.8% to 34%. Transduction of all three ZFN pairs simultaneously resulted in mutation rates of 8% to 20%. Off-target mutagenesis for seven potential off-target sites was investigated using single molecule real time (SMRT) sequencing which detected indels (> 1 nt) in four sequencing reads out of 9290 filtered reads. It is of note that indels of only one nucleotide were not taken into account because they could not be distinguished from sequencing artefacts. In order to test if ZFN treatment has an effect on HBV replication, ZFNs were transduced into HepAD38 cells in which HBV replication was shut down. After turning HBV replication on, the controls showed a 30-fold and 323-fold increase in HBV marker levels in cells and in the supernatant, respectively. In ZFNs treated cells no significant increase of cellular or supernatant HBV marker levels could be detected, leading to the conclusion that ZFN-mediated cleavage resulted in replication-deficient HBV. In summary, it was demonstrated that HBV specific ZFN can be delivered successfully *via* AAV vectors into target cells *in vitro*, resulting in an efficient inhibition of HBV replication.

TALENs are a promising new class of designer nucleases that can be specifically designed to bind DNA sequences of interest and to introduce dsDNA breaks. Comparable to ZFNs, TALENs are chimeric proteins consisting of a N-terminal nuclear localization signal, a central DNA binding domain and a C-terminal *Fok*I nuclease domain. The DBD originates from transcription activator like effectors (TALEs) of bacterial plant pathogens of the genus *Xanthomonas*, which secrete these proteins to regulate gene expression within their host cells^[141]. The DBD is comprised of a repeat region consisting of a number of incomplete tandem repeats containing repeat-variable diresidues (RVD)^[127]. The

RVD of each repeat is specific for binding a corresponding nucleotide in their contiguous target DNA sequence^[127,141]. Similar to ZFNs, the TALEN-DNA binding domain is combined with a non-specific endonuclease activity which is dependent on dimerization of two *FokI*-cleavage domains. Binding of the DBDs upstream and downstream of a DNA target brings the *FokI* nuclease domains of a TALEN pair in close proximity and dimerization of the *FokI* monomers introduces dsDNA breaks. Note that the spacer between the binding sites of a pair needs to be considered in the TALEN design to ensure optimal cleavage^[127,128]. Up to now several techniques are available to specifically design^[127,142] and to assemble TALENs for the desired application^[127,143-145]. Assembly can be performed without highly complex screening procedures or the need for special equipment, making it cheap, simple and fast and thereby with that respect superior to the ZFN technology.

Bloom *et al.*^[146] designed TALENs with target sequences in the HBsAg or HBcAg expressing region of the HBV genome. TALEN efficacy was determined after co-transfection of Huh7 cells with TALEN expression plasmids together with the pCH-9/3091 HBV replication-competent plasmid. They observed that the HBsAg production was diminished in cells expressing TALENs. In a more stringent experimental model of HBV replication, the HepG2.2.15 cell line was transfected with TALEN expression plasmids. After three subsequent transfections and culturing cells under hypothermic conditions, HBsAg-specific TALEN expression resulted in disruption of cccDNA molecules with efficiencies of approximately 31%. This was confirmed in a T7E1 mutation detection assay and correlated with a decrease of HBsAg secretion of these cells. Expression of the HBcAg-specific TALEN pair only mutated 12% of cccDNA molecules which was not sufficient to inhibit HBsAg secretion. After hydrodynamic tail vein injection of replication-competent HBV DNA together with HBsAg-specific TALEN expression plasmids, serum HBsAg levels were decreased by more than 90% and circulating viral particle equivalents (VPEs) were decreased by approximately 70%. The T7E1 assay demonstrated mutation rates of 58%-87% in HBV-DNA extracted from livers of TALEN treated mice.

Chen *et al.*^[147] used three TALEN pairs targeting regions of HBV genomic DNA conserved among HBV genotypes A-D. Huh7 cells were transfected with the monomeric linear full-length HBV DNA of subgroups A, B, C, or D, respectively and plasmids expressing one of the respective TALEN pairs. Cells containing HBV DNA simulated the complete HBV replication cycle, including the nuclear generation of cccDNA. Suppression of HBeAg and HBsAg production by the TALEN expression was observed for all four HBV genotypes. Additionally HBcAg-RNA as well as pregenomic HBV RNA levels were decreased in

TALEN expressing cells. Furthermore T7E1 mutation detection assay confirmed that mutations were successfully induced at the respective TALEN target site within the HBV genome leading to a 10%-50% decrease of cccDNA levels. In combination with interferon alpha treatment TALENs expression led to synergistic effects further increasing the inhibition of HBV transcription in Huh7 cells. After delivery of monomeric linear full-length HBV DNA and TALEN expression plasmids into C3H/HeN mice by hydrodynamic tail vein injection, serum levels of HBeAg, HBsAg and cccDNA as well as liver pregenomic HBV RNA significantly decreased compared to control animals. In summary, HBV-targeting TALENs were shown to be active in cell culture models as well as in *in vivo* models and are capable of introducing mutations at their target sites reducing HBV gene expression levels and cccDNA genome numbers.

The most recent approach exploited the CRISPR/Cas9 system for genome engineering^[131]. Because sequence specificity is achieved by a guide RNA (gRNA) which can be produced from an adaptable expression cassette, this system is easier to manipulate in comparison to ZFNs and TALENs where the DBD is based on a protein sequence. Lin *et al.*^[148] were the first to apply this system for achieving HBV genome degradation^[148]. They co-expressed the Cas9 nuclease together with eight HBV specific guide RNAs individually and in addition combined two of them on one expression vector. They tested all constructs in Huh7 cells which were transfected with a HBV-expression vector. Inhibition varied among the different guide sequences. A maximum inhibition of 70% of intracellular HBsAg expression was reached after using one guide RNA and Cas9 nuclease expressed from individual vectors and up to 96% inhibition was reached when they were combined on one expression vector. Next, they multiplexed the two most effective gRNAs, an approach which proved to be even more effective. Finally, the expression vectors that contained expression cassettes for one gRNA and the Cas9 nuclease were tested in the HBV-hydrodynamic mouse model. Serum HBsAg levels were significantly reduced two days post-injection but increased again on day seven. However, Southern blot analyses revealed a reduction of intrahepatic HBV-expression levels of 20% to 60%. In conclusion the RNA-guided Cas9 nuclease system may be a useful technique for achieving inhibition of HBV replication.

CONCLUSION

Numerous and highly diverse gene therapeutic approaches were pursued to combat chronic HBV infection. Although strategies that solely rely on nucleic acids like antisense oligonucleotides and catalytic nucleic acids have a great advantage in

their simplicity, these technologies are also limited due to their instability and imprecision. RNAi was most adopted and thoroughly investigated not only in the field of HBV therapy and this generated a deep knowledge regarding this technique. However, especially in the case of HBV, RNAi may be disadvantageous, because it does not affect the HBV cccDNA. It can rather be considered as an alternative to nucleos(t)ide analogues based therapy potentially associated with an improved ability to also respond to escape mutants. In addition, if long-term expression is required, RNAi-based approaches are superior compared to conventional therapy based on daily administration required for nucleos(t)ide analogues.

We believe that the most recent technology to combat HBV infection based on designer nucleases may be one of the most promising approaches to be explored in the future. This strategy bears the potential to actually eradicate cccDNA species in infected cells. Especially in combination with compounds that inhibit the HBV replication cycle this could be an attractive therapeutic option. However, major obstacles are the production and delivery of the designer nucleases and for translation of this approach into the clinic, this methodology needs to be further improved. The most promising results to date were obtained with genetic vaccines which were also pursued in the clinic. Also this strategy may hold great potential for eradicating chronic hepatitis B infection.

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Targeting the tumor stroma in hepatocellular carcinoma

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Abstract

Hepatocellular carcinoma (HCC) is one of the most common and deadly cancers worldwide. In ninety percent of the cases it develops as a result of chronic liver damage and it is thus a typical inflammation-related cancer characterized by the close relation between the tumor microenvironment and tumor cells. The stromal environment consists out of several cell types, including hepatic stellate cells, macrophages and endothelial cells. They are not just active bystanders in the pathogenesis of HCC, but play an important and active role in tumor initiation, progression and metastasis. Furthermore, the tumor itself influences these cells to create a background that is beneficial for sustaining tumor growth. One of the key players is the hepatic stellate cell, which is activated during liver damage and differentiates towards a myofibroblast-

like cell. Activated stellate cells are responsible for the deposition of extracellular matrix, increase the production of angiogenic factors and stimulate the recruitment of macrophages. The increase of angiogenic factors (which are secreted by macrophages, tumor cells and activated stellate cells) will induce the formation of new blood vessels, thereby supplying the tumor with more oxygen and nutrients, thus supporting tumor growth and offering a passageway in the circulatory system. In addition, the secretion of chemokines by the tumor cells leads to the recruitment of tumor associated macrophages. These tumor associated macrophages are key actors of cancer-related inflammation, being the main type of inflammatory cells infiltrating the tumor environment and exerting a tumor promoting effect by secreting growth factors, stimulating angiogenesis and influencing the activation of stellate cells. This complex interplay between the several cell types involved in liver cancer emphasizes the need for targeting the tumor stroma in HCC patients.

Key words: Hepatocellular carcinoma; Stellate cells; Cirrhosis; Angiogenesis; Macrophages; Inflammation

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Core tip: Hepatocellular carcinoma is a primary liver tumor that usually develops in a background of chronic liver disease and fibrosis. It is the underlying chronic inflammation that creates an environment that not only causes but also enhances the formation and growth of tumors. The stromal compartment-including hepatic stellate cells, macrophages and endothelial cells-actively contribute to tumorigenesis, while the tumor itself influences these cells to create a background that is beneficial for tumor growth. This review focuses on the interplay between stroma and tumor cells, as well as therapeutic strategies that aim to target these complex interactions.

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INTRODUCTION

Hepatocellular carcinoma (HCC) is a primary liver tumor that usually develops in a background of chronic liver disease. It is the underlying chronic inflammation that creates an environment that not only causes but also enhances the formation and growth of tumors. Firstly, the continuous state of inflammation as a result of sustained liver damage can lead to hepatocyte cell death as well as compensatory proliferation, which can generate an accumulation of genomic lesions in hepatocytes. Secondly, the initiated cells are surrounded by an inflammatory niche that facilitates their progression towards malignant tumors. For instance, the fibrotic liver is characterized by an increased formation of blood vessels^[1], which will benefit tumor cells for their blood supply as well as facilitating metastasis^[2]. In addition, several factors produced by macrophages and activated stellate cells are known to directly stimulate and enhance tumor growth. Once the cancer has been established, the microenvironment continues to regulate the tumor behavior, influencing the development, progression and even response to therapy. All players within the tumor stroma strongly interact with each other, creating an environment that supports tumor growth (Figure 1). It is therefore not unlikely that future therapies will more and more focus on targeting these complex interactions in the tumor stroma (Figure 2). Ongoing clinical trials are listed in Table 1.

HEPATIC STELLATE CELLS

One major player in the formation of the perfect tumor environment is the activated hepatic stellate cell (HSC)^[3,4]. During liver injury, the stellate cells undergo a transformation from quiescent cells that serve as the liver's resident vitamin-A storing cells, towards "activated" myofibroblast-like cells. These activated HSCs are characterized by increased proliferation and contractility, altered matrix protease activity and the secretion of extracellular matrix (ECM) proteins, as well as tumor growth factors and pro-angiogenic factors.

Several studies have shown that co-culturing HSC with different HCC cell lines induces phenotypic changes in the behavior of the tumor cells^[5,6]. *In vitro* studies show that HSCs can directly influence the tumor cells (through the secretion of growth factors^[7], matrix proteases^[8] and/or ECM proteins^[9]) and there is also evidence from *in vivo* studies that activated HSCs can create an immunosuppressive environment that promotes HCC growth^[10,11]. The

interaction between the tumor cells and HSCs is bidirectional, thereby allowing the tumor to alter the stellate cells (and the overall stromal environment) towards a more pro-tumoral phenotype^[8]. Consistent with these findings, several *in vivo* studies have shown that inducing stellate cell activation increases liver fibrosis and hepatocarcinogenesis^[12-15].

One of the key factors in this HSC-HCC cross talk is transforming growth factor (TGF)- β ^[14,16,17]. Activated HSC are the main source of TGF- β , however most liver cells (including malignant hepatocytes) have the ability to produce TGF- β as well. The TGF- β signaling pathway consists of three distinct ligands, TGF- β 1, TGF- β 2, and TGF- β 3 which all bind to a specific receptor by first engaging with the TGF- β R1, which then heterodimerizes with the TGF- β R2. This causes the phosphorylation of Smad2 and 3, initiating an activation cascade leading to the induction several nuclear transduction proteins. Alternative pathway activation is possible, including the activation of AKT and other intracellular activation proteins. Interestingly, Smad7 antagonizes TGF- β mediated activation of hepatic stellate cells and protects against liver damage^[18]. TGF- β signaling promotes HCC by several distinct mechanisms (reviewed more in detail by Dooley *et al.*^[17]): firstly, through functioning as a growth factor, by which it can act oncogenic or as tumor suppressor depending on the temporal and spatial availability of TGF- β in tumor and stromal cells^[19,20]. And secondly, by transforming HSC to activated myofibroblasts. Interestingly, Inhibitors of TGF- β signaling have been shown to block HCC in different experimental models^[21], leading to the clinical investigation of the TGF- β inhibitor LY2157299 (NCT01246986 and NCT02178358). LY2157299 is a small molecule kinase inhibitor that binds to TGF- β R1 and hence inhibits TGF- β signaling.

The connective tissue growth factor (CTGF) is an extracellular matrix-associated heparin binding protein that is overexpressed in fibrotic lesions, and the overexpression correlates with the severity of fibrosis and can be linked to malignant transformation in patients with chronic hepatitis B^[22]. CTGF is a downstream mediator of some TGF- β effects and it is induced by TGF- β in a SMAD2/3 and stat3 dependent way. Furthermore, IL-13 is able to induce CTGF expression in HSCs by activating TGF- β -independent Smad signaling *via* the Erk-MAPK pathway instead of the canonical JAK/Stat6 pathway^[23]. CTGF expression in HSC leads to increased migration, proliferation, and collagen expression of these cells. In addition, studies have shown that TGF- β can elicit a direct effect on hepatocytes *via* CTGF, thus making it an interesting therapeutic target for multiple cell types involved in the fibrogenesis^[18]. CTGF blocking antibodies have been tested in patients with idiopathic pulmonary fibrosis (NCT00074698) and animal studies have shown that CTGF-inhibition prevents liver fibrosis in

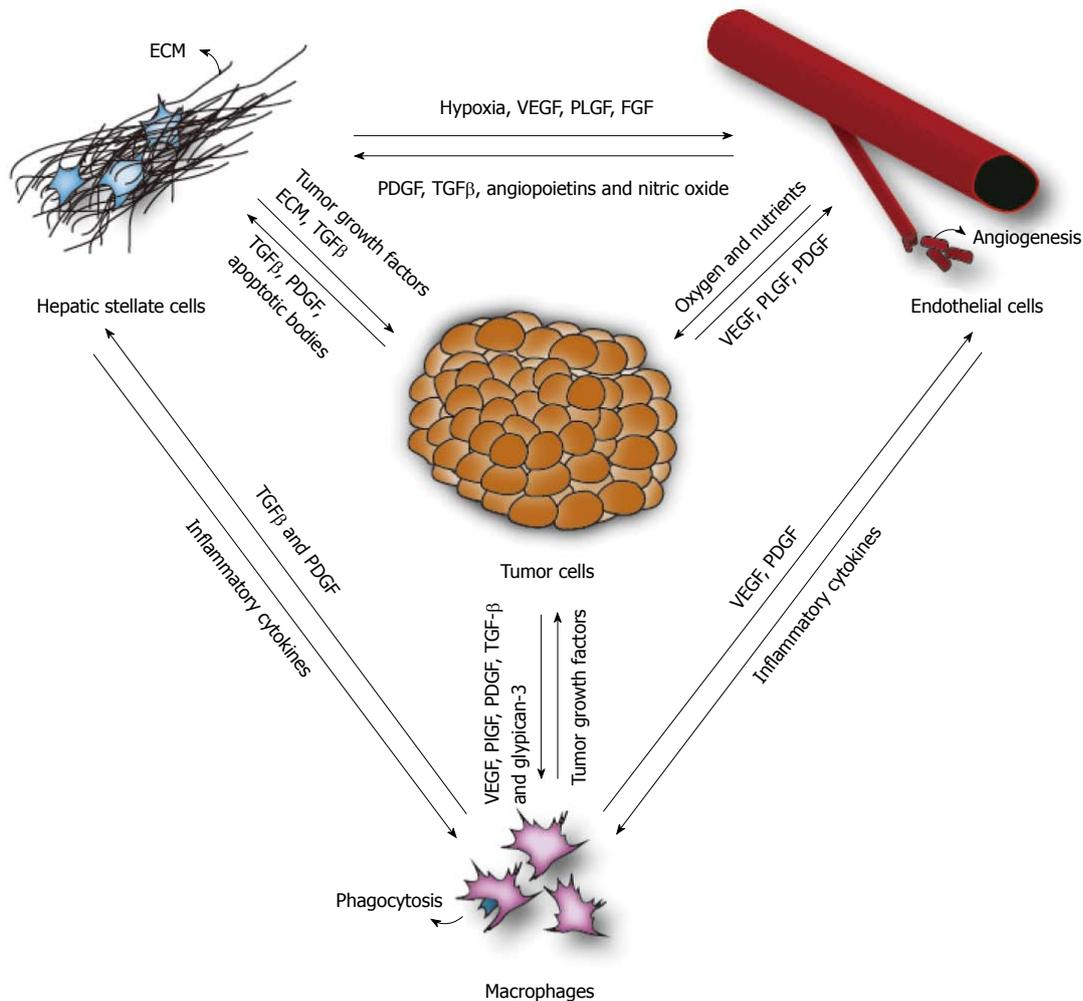


Figure 1 The interaction between tumor stroma and tumor cells in hepatocellular carcinoma. The several actors of the stromal compartment—including hepatic stellate cells, macrophages and endothelial cells—actively contribute to tumorigenesis, while the tumor itself influences these cells to create a background that is beneficial for tumor growth. Tumor cells activate the hepatic stellate cells, leading to the deposition of extracellular matrix (ECM), an increased production of angiogenic factors and the recruitment of macrophages. The increase of angiogenic factors (secreted by macrophages, tumor cells and activated stellate cells) will induce the formation of new blood vessels, thereby supplying the tumor with more oxygen and nutrients, thus supporting tumor growth. The increase of inflammatory cytokines, leads to the recruitment of macrophages, which can exert a pro-tumoral effect by secreting growth factor and influences the activation of stellate cells. This complex interplay between the several cell types involved in liver cancer emphasizes the need for a multi-targeted approach in hepatocellular carcinoma (HCC) patients. VEGF: Vascular endothelial growth factor; PDGF: Platelet-derived growth factor; FGF: Fibroblast growth factor.

rats^[24]. However, no clinical trials on the effect on liver fibrosis have been done.

Another driver of HSC activation are members of the platelet derived growth factor (PDGF) family. PDGFs are potent mitogens for mesenchymal cells and work synergistically with TGF to activate stellate cells. Specific hepatic over-expression of PDGF-C leads to an increase in fibrosis and enhances hepatocarcinogenesis^[12,15]. PDGF-B is also involved in different stages of liver cancer development and is an essential regulator in the development of liver fibrosis^[25]. Hepatic overexpression of PDGF-B accelerates liver cancer, possibly by up regulating TGF- β receptor and by increasing expression of β -catenin as well as VEGF, CD31 and FGF. Several protein tyrosine kinase inhibitors—such as sorafenib, orantinib, sunitinib and SU6668—target PDGFR amongst other targets including VEGFR and FGFR. The protein tyrosine kinase

inhibitor imatinib reduces stromal cell proliferation in this mouse model, which successfully inhibits tumor progression^[13]. Imatinib is currently used to treat gastrointestinal stromal tumors^[26] and could possibly benefit HCC patients.

The deposition of ECM proteins is one of the most characteristic hallmarks of the activated stellate cell. Several of the ECM components such as proteoglycans, laminins, collagens, and fibronectin interact directly and indirectly with HCC cells and the different stroma cell types. This not only changes the tumor phenotype, but also prepares a microenvironment that facilitates tumor growth. Since the ECM acts as a reservoir for growth factors and cytokines, it can rapidly release them to support the tumor's needs.

Heparan sulfate (HS) proteoglycans (PG) are expressed in the ECM and composed of a protein core to which HS is covalently attached as side

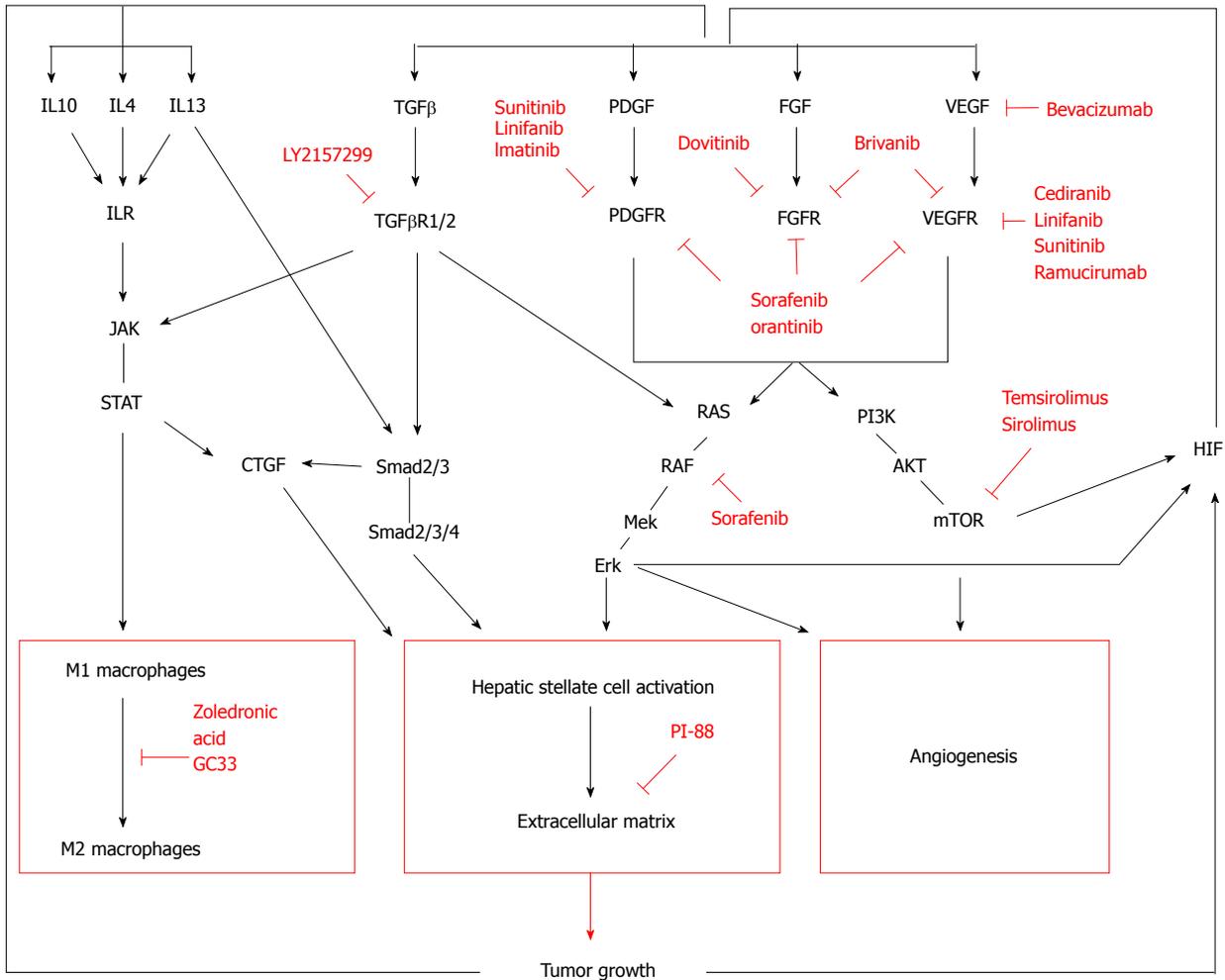


Figure 2 Schematic overview of (simplified) signaling pathways involved in the tumor-stroma interaction, and therapeutic targets that are currently tested in hepatocellular carcinoma. IL: Interleukin; TGF: Transforming growth factor; mTOR: Mammalian target of rapamycin.

chains. They maintain the structural framework of the tissue, store growth factors within the ECM or function as co-receptors. Desulfation of these co-receptor-PG's can abrogate growth factor signaling and inhibit tumor growth^[27,28]. Heparanase cleaves the HS side chains of HSPG, leading to the release of HS-bound proteins, such as growth factors. PI-88 is a heparin sulfate mimic that specifically targets heparanase in cancer, thus preventing the release of growth factors that otherwise would contribute to tumor growth, angiogenesis and metastasis^[29]. The safety and efficiency of PI-88 as an adjuvant therapy for post-operative HCC has been shown in a phase II trial^[30] and a recent follow up study revealed significant clinical benefits for patients with HCC^[31]. Phase III trials are currently ongoing (NCT01402908).

Another important glycoprotein is laminin-5. Laminin-5 is a member of the laminin family, which has been widely reported to be involved in the malignant phenotype of several cancers, including HCC^[32]. Laminin-5 is expressed higher in metastatic HCC and has been shown to stimulate HCC cell migration^[9,32].

Collagen is the major insoluble fibrous protein in the extracellular matrix. Besides its function as a supportive scaffold, collagens can also provoke a cellular response through the integrin family of transmembrane receptors. Several collagen types have been implicated in tumor growth and angiogenesis in different tumors^[33-35] and a recent study has shown that collagen matrix protects malignant hepatocytes from apoptosis^[36]. Antibodies targeting cleaved collagen epitopes have been clinically tested and show promising results in patients with solid tumors^[37,38].

This deposition of ECM leads to an increase in liver stiffness, an important hallmark of the cirrhotic liver, which is also used as a diagnostic tool for patients with CLD^[39]. This change in the mechanical properties of the tumor's surrounding has been associated with a higher risk of developing HCC^[40]. In addition, the increase of ECM and the capillarization of hepatic sinusoids cause a vascular resistance that leads to hypoxia, stimulating the production of pro-angiogenic factors and subsequently inducing angiogenesis^[1,41-43]. The activated HSCs also produce

Table 1 Overview of clinical trials that focus on the tumor environment of hepatocellular carcinoma

	Drug	Targets	Trial	Phase	Status	Ref.			
Tyrosine kinase inhibitors	Sorafenib	PDGFR VEGFR	NCT00105443	III	Completed ¹	[90]			
	Orantinib	RAF/MEK/ERK PDGFR, FGFR VEGFR	NCT02178358	I / II	Completed	[91]			
	Sunitinib	VEGFR	NCT00699374	III	Terminated	[71]			
		PDGFR	NCT00514228	II	Completed	[70]			
		RET	NCT00361309	II	Completed	[69]			
	Linifanib	CSF	NCT00428220	N/A	Ongoing				
		VEGF, PDGF, PDGFR- β , KDR	NCT01009593	III	Terminated				
	Brivanib	CSF	NCT00517920	II	Completed	[68]			
		VEGFR	NCT00858871	III	Completed	[59]			
				NCT00908752	III	Ongoing			
				NCT00825955	III	Ongoing			
				NCT01108705	III	Terminated			
				NCT00355238	II	Completed	[92]		
				NCT00437424	I	Completed	[93]		
				NCT00238394	II				
				NCT00427973	II	Terminated	[54]		
				NCT01232296	II	Ongoing			
				Antibodies	Bevacizumab	VEGFR, PDGFR, FGFR VEGF	NCT00335829	II	Completed
NCT00162669							II	Completed	[50]
Ramucirumab	VEGFR	NCT00605722	II		Completed	[51]			
		NCT00049322	II		Completed	[52]			
		NCT00280007	II		Terminated				
		NCT01180959	II		Ongoing				
		NCT00627042	II		Completed	[53]			
		NCT01140347	III		Ongoing				
GC33	Glypican-3	NCT01507168	II		Completed				
		NCT00746317	I		Completed	[84]			
Other kinase inhibitors	Temozolomide	mTOR	NCT00976170	I	Ongoing				
			NCT01008917	I	Ongoing				
	Everolimus	mTOR	NCT01687673	II	Recruiting				
			NCT01035229	III	Completed	[95]			
	LY2157299	TGF- β 1	NCT01488487	II	Ongoing				
			NCT00516165	I / II	Completed	[66]			
			NCT00828594	I	Terminated				
	PI-88	Heparanase	NCT01246986	II	Recruiting				
			NCT02178358	II	Recruiting				
	Zoledronic acid	Macrophages	NCT00568308	III	Terminated				
NCT01402908			III	Ongoing					
			NCT00247728	II	Completed	[30]			
			NCT01259193	II	Ongoing				

¹Sorafenib is currently used as the standard-of-care for advanced hepatocellular carcinoma. CSF: Colony stimulating factor-1-receptor; mTOR: Mammalian target of rapamycin.

angiogenic growth factors, thus enhancing neo-angiogenesis^[8]. This increased vasculature will allow small HCC lesions to progress and eventually metastasize.

ENDOTHELIAL CELLS

The prolonged fibrogenic process leads to an abnormal angioarchitecture distinctive for cirrhosis. Anatomical changes in the cirrhotic liver, such as fibrotic scar tissue compressing portal and central venules are responsible for an increased intrahepatic vascular resistance. In addition, the formation of fibrotic septa, as well as sinusoidal capillarisation, results in an increased resistance to blood flow

and oxygen delivery. This causes hypoxia and the transcription of hypoxia-sensitive pro-angiogenic genes, thus stimulating the formation of new vessels. These new vessels can contribute to the inflammatory response by expressing chemokines and adhesion molecules, thus promoting the recruitment of inflammatory cells, such as macrophages. In addition, hepatic stellate cells are recruited to the angiogenic areas (*via* a number of signaling pathways, including PDGF, TGF- β , angiopoietins and nitric oxide) to contribute in vascular remodeling and stabilization^[44]. Therefore, angiogenesis may contribute to the progression of liver cirrhosis and stimulate the growth of small dysplastic lesions to advanced solid tumors.

HCC is solid tumor that rapidly outgrows its blood

supply and therefore stimulates the formation of new blood vessels to fulfill its high needs in oxygen and nutrients. The malignant hepatocytes, as well as other actors in the microenvironment such as activated stellate cells and macrophages, secrete a number of angiogenic growth factors^[1]. This induces an “angiogenic switch”, which activates endothelial cells and basement membranes to remodel existing vessels, and form new vessels. These new vessels allow the tumor to rapidly expand and offer a passage in the circulatory system, thus facilitating metastasis. Therefore, targeting angiogenesis has become a common cancer therapy to treat solid tumors.

The vascular endothelial growth factor A (VEGF) is one of the key factors regulating angiogenesis. It is secreted by tumor cells, macrophages and stellate cells. VEGF binds to its receptors (VEGFR1 and VEGFR2) on the present endothelial cells, stimulating endothelial cell proliferation and migration into the tumor, which results in vascular sprouting. Elevated VEGF levels are associated with tumor vascularity, metastasis, chemoresistance and poor prognosis^[45-47].

Significant progress on the treatment of advanced HCC has been made possible by sorafenib. Sorafenib is a small molecular inhibitor targeting several tyrosine protein kinases in the Raf/MEK/ERK-pathway (anti-proliferative effect); and PDGF, VEGFR1 and VEGFR2 (anti-angiogenic effect). Sorafenib has become the standard-of-care for patients with advanced HCC and for those progressing after loco-regional therapies^[48]. The success of sorafenib has opened the door for several anti-angiogenic agents to enter clinical studies on HCC^[49]. At the moment, several multikinase inhibitors are being tested in clinical trials, including sunitinib, brivanib, linafinib, cediranib, pazopanib, lenvatinib and axitinib, as well as blocking-antibodies targeting angiogenic pathways.

Bevacizumab, a humanized monoclonal antibody that targets VEGF, has been approved for the treatment of various solid tumors and is currently being investigated as a treatment for HCC. Several phase II trials have been completed and show that bevacizumab is well tolerated in HCC-patients, and could be a promising therapy as a single-agent^[50], in combination with erlotinib^[51] or after loco-regional therapies^[52]. Ramucirumab is a monoclonal antibody targeting VEGFR2 which has been tested as a first line treatment (NCT00627042) for HCC-patients with promising results^[53] and is currently being investigated as second line treatment after sorafenib (NCT01140347)^[53]. Cediranib is a tyrosine kinase inhibitor that targets all VEGF receptors, which has been tested in two clinical trials (NCT00427973 and NCT00238394). Despite some anti-tumor effects, the high toxicity of cediranib makes it an unsuitable drug for HCC-patients HCC^[54,55].

However, targeting VEGF has been shown to

induce therapy escape mechanisms and many patients treated with VEGF-inhibitors or with sorafenib obtain a secondary resistance to therapy. Alternative angiogenic factors, such as the placental growth factor (PlGF), PDGF and fibroblast growth factor (FGF) have been implicated in this acquired tumor resistance and combination therapies could open the door for sustained treatment response^[56]. Additionally, combining sorafenib with conventional chemotherapy could improve outcome and is currently tested in several phase III trials (NCT01015833, NCT01214343)^[57].

Brivanib and dovitinib are tyrosine kinase inhibitors of VEGF and fibroblast growth factor (FGF) signaling pathways, hence anticipating FGF-mediated resistance to anti-VEGF therapy^[58]. Brivanib has been or is being investigated in several phase III trials, including first-line treatment with brivanib vs sorafenib (NCT00858871)^[59], second-line treatment with brivanib after progression on sorafenib treatment (NCT01108705), second-line treatment with brivanib after sorafenib (NCT00825955) and trans-arterial chemoembolization in combination with brivanib (NCT00908752). However, results from the study testing brivanib and sorafenib as first-line therapy in patients with HCC indicate there are no benefits of using brivanib over sorafenib^[59] and study NCT01108705-testing brivanib after sorafenib treatment-has been terminated before completing the trial. Dovitinib trials are still ongoing (NCT01232296).

A drawback of anti-angiogenic therapies is that they aim to deprive the tumor from oxygen, leading to a hypoxic environment that stimulates cancer cells towards a more aggressive phenotype^[60]. Therefore, long-term administration of anti-angiogenic treatment could trigger escape mechanisms and lead to increased metastasis^[61,62].

An interesting way to indirectly target VEGF signaling and the HIF-pathway, is through inhibitors of the mammalian target of rapamycin (mTOR) pathway. mTOR signaling increases VEGF expression by up-regulating hypoxia inducible factor 1 α ^[63]. Furthermore, mTOR-inhibitors can directly influence tumor growth by inhibiting the expression of anti-apoptotic proteins and by inducing autophagy^[64]. Everolimus binds the cyclophilin FKBP-12, which binds the serine-threonine (ST) kinase mTOR when it is associated with raptor and mLST8 to form a complex (mTORC1), and subsequently inhibits downstream signaling, which involves cell cycle regulators and transcription factors such as HIF. mTORC1 lies downstream of phosphatidylinositol 3' kinase (PI3K), which is frequently activated in human cancers. Everolimus has been used in several clinical trials^[65,66], but data from the latest phase III trial (NCT01035229) show no improvement in overall survival^[67]. Temsirolimus is a sirolimus ester, which binds the same receptors. Trials using

temsirolimus are currently ongoing (NCT01008917 and NCT01687673).

The activated stellate cells also play a pivotal role in vascular remodeling, by creating a hypoxic environment, by producing angiogenic factors and also by migrating to angiogenic sites to contribute in the stabilization and maturation of (tumor) blood vessels. Current anti-angiogenic strategies for cancer have mostly focused on endothelial cells. However, combining drugs that target endothelial cells and stellate cells (or pericytes) could work synergistically as a therapy.

Several receptor tyrosine kinase inhibitors target VEGF and PDGF. Linifanib is a potent inhibitor of VEGF, PDGF, PDGFR- β , KDR and colony stimulating factor-1-receptor (CSF). A phase II trial (NCT00517920) showed initial benefits for linifanib in HCC patients^[68], however, the subsequent phase III trial (NCT01009593) had to be terminated for unknown reasons. Sunitinib inhibits receptors for PDGF and VEGF, as well as other receptor tyrosine kinases such as CSF. While several phase II trials (NCT00514228, NCT00361309) have shown promising results^[69,70], it is inferior to sorafenib and the latest phase III trial had to be terminated for safety reasons^[71].

Orantinib is a receptor tyrosine kinase inhibitor that binds and inhibits the autophosphorylation of VEGFR2, PDGF-receptor and fibroblast growth factor receptor (FGFR), thereby inhibiting angiogenesis and cell proliferation. A phase I / II trial has shown a trend towards prolonged progression free survival in patients treated with orantinib after transarterial chemoembolization^[72], and a phase III trial is still ongoing (NCT01465464). Blocking PDGF signaling in mouse models of pancreatic carcinogenesis with orantinib caused regression of blood vessels, as a result of the detachment of pericytes from tumor vessels. The fact that tumor vessels lacking pericytes are more vulnerable suggests that they could be more responsive to other anti-angiogenic drugs^[73,74]. Combining receptor tyrosine kinase inhibitors targeting ECs and pericytes successfully diminished tumor angiogenesis and decreased tumor size compared to a monotherapeutic approach in colon cancer^[75]. Similar effects were seen when PDGF inhibitors were combined with anti-angiogenic treatments^[74]. Thus, targeting stellate cells and endothelial cells may destabilize the existing tumor vasculature more potently than targeting each cell type individually.

MACROPHAGES

After liver damage, the pool of the liver's resident macrophages-Kupffer cells-is rapidly expanded. A harmful incident causes the hepatic macrophages to secrete pro-inflammatory cytokines and chemokines such as IL-1 β , TNF, CCL2, and CCL5, resulting in the activation of protective or apoptotic signaling pathways of hepatocytes and the recruitment of

immune cells that support hepatic injury. There is increasing evidence suggesting that phagocytosis of apoptotic bodies by HSC and by macrophages may directly stimulate fibrogenesis through upregulation of TGF- β ^[76]. Furthermore, these repeated cycles of hepatocyte death and compensatory proliferation provide a mitogenic and mutagenic environment that fuels the development of HCC.

The location of Kupffer cells in the sinusoids allows close interactions with other non-parenchymal liver cells. Firstly, Kupffer cells interact with other immune cells by secreting inflammatory cytokines and chemokines. Secondly, they can activate HSC *via* paracrine mechanisms, likely involving TGF- β and PDGF. These profibrotic functions of Kupffer cells during chronic liver injury possibly contribute to a tumor-stimulating environment in the cirrhotic liver. *In vivo* studies have shown that depleting macrophages reduces angiogenesis and slows down tumor progression in mouse models, and enhances the response to sorafenib^[77].

Macrophages can be classified into two main classes depending on their phenotypic polarization: the M1-phenotype, triggering a Th1 immune response and exerting cytotoxic activity; and the M2-phenotype, which activates a Th2 immune response and promotes angiogenesis, tissue remodeling and tumor progression^[78]. Macrophages can adapt to signals from the microenvironment and change their functional phenotype accordingly^[79]. M1 macrophages are activated as a response to microbial stimuli and interferon gamma, while in a tumor environment the tumour-associated macrophages (TAMs) are mainly polarized towards a M2 phenotype. Increased numbers of M2-macrophages have been associated with angiogenesis, metastasis and poor prognosis.

Tumor associated macrophages are key actors of cancer-related inflammation, being the main type of inflammatory cells infiltrating the tumor environment^[80]. In HCC, tumor cells have been shown to recruit and activate TAMs by the secretion of VEGF, PlGF, PDGF, TGF- β and glypican-3. Glypican-3 is a member of the glypican family of heparin-sulfate proteoglycans linked to the cell surface. It is highly expressed in the majority of HCC cells and is known for its role in the regulation of cell proliferation and apoptosis^[81]. In addition, studies have suggested an involvement in the recruitment of M2-polarized TAM's in human HCC tissues^[82]. Possibly glypican-3 present on the cell surface of malignant cells, binds to CCL5 and CCL3, which are chemokines that attract TAMs. Glypican-3 antibodies could therefore block the recruitment of TAMs *via* CCL5 and CCL3. Antibodies targeting glypican-3 have been tested in several phase I trials for advanced HCC, with promising results. The antibody was well tolerated and preliminary antitumor activity show a threefold prolongation of the median time to progression in patients receiving glypican-3-antibodies compared to

untreated patients^[83,84].

Zoledronic acid (ZA) is a compound widely used to prevent skeletal complications associated with bone metastases. Recent studies have shown a possible direct role as an anti-tumor agent by targeting the TAMs. ZA is taken up by macrophages *via* phagocytosis and leads to apoptosis specifically in TAMs, thus causing a repolarization of the macrophage population^[85,86]. *In vivo* studies of ZA in combination with sorafenib have shown that the latter leads to an increase of M2-macrophages infiltrating the tumor stroma, which can be effectively depleted with ZA. This significantly inhibits angiogenesis, metastasis and tumor progression compared to sorafenib alone^[77]. A phase II study of sorafenib and ZA in advanced HCC has been conducted (NCT01259193), but no results have been published.

DISCUSSION

Several studies have shown that the stroma regulates the malignant transformation, survival, progression and metastasis of hepatocellular carcinoma. Factors derived from the tumor cells in their turn alter the tumor stroma to generate a tumor-permissive microenvironment. This complex interplay between the tumor and the different actors in the stroma establishes a promising axis for therapeutic targets (Figure 2).

VEGF targeting therapies have represented the first success in treating HCC patients in many years, reviving research in this field and leading to an explosion of clinical trials with anti-angiogenic therapies^[49]. However, the success of treatments such as sorafenib needs to be followed by better understanding of the mechanisms that underlie the intrinsic and acquired resistance to anti-angiogenic therapies. Perhaps targeting several actors of the stromal environment and the tumor cells at the same time could be the key for optimal treatment in future therapies.

Sorafenib does not only inhibit angiogenesis, but also alters the inflammatory environment. Sorafenib has been shown to suppress natural killer cells and facilitate tumor growth and metastasis^[87]. Furthermore, multi-tyrosine kinase inhibitors have been shown to increase infiltration of tumor-associated macrophages in the tumor environment which could contribute to the resistance or escape to anti-angiogenic treatment^[77] (although it is important to note that some studies have shown the opposite effect^[88]). Hence the solution could be the use of adjuvant immunotherapy along with tyrosine kinase inhibitors for patients with unresectable HCC in order to obtain long-term response. In fact, one of the first trials to confirm the efficacy of sorafenib in advanced, metastasized renal cell carcinoma, was performed in combination with immunotherapy with IL-2 and interferon-alpha^[89].

Tumor associated macrophages are important actors of cancer-related inflammation, being the main type of inflammatory cells infiltrating the tumor environment. Targeting macrophages as a therapeutic strategy could be done by depleting the overall population of macrophages, or by altering their phenotype from a M2 towards an M1 orientation. Again, macrophages are known to not only interact with the tumor cells and stimulate their growth, they also influence stellate cell activation and angiogenesis.

The stellate cells are one of the key players in the formation of the perfect tumor environment. Not only do they directly affect tumor growth by secreting growth factors^[7], matrix proteases^[8] and/or ECM proteins^[9], they also alter the mechanical properties of the tumor's surrounding. Activated stellate cells are known to stimulate angiogenesis, which allows the tumor cells to grow rapidly and invade in the circulatory system. The deposition of ECM proteins, such as collagens and proteoglycans, serve as a reservoir for growth factors, but also directly provoke a pro-tumoral cellular response. Indeed, the thick layer of ECM in the cirrhotic liver could impair drug delivery and hence decrease response to therapy. Therefore, preventing or reversing the activation of stellate cells could inhibit HCC growth, decrease angiogenesis and increase response to other therapies, such as classic chemotherapy or sorafenib.

As our understanding of the complex interplay between tumor and stroma evolves, the next-generation cancer drugs could target several actors in the tumor-stroma axis and offer a durable treatment for advanced HCC.

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Dysregulation of iron and copper homeostasis in nonalcoholic fatty liver

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with normal or mildly elevated transferrin saturation and mostly mild hepatic iron deposition are a characteristic finding in subjects with non-alcoholic fatty liver disease (NAFLD). Excess iron is observed in approximately one third of NAFLD patients and is commonly referred to as the "dysmetabolic iron overload syndrome". Clinical evidence suggests that elevated body iron stores aggravate the clinical course of NAFLD with regard to liver-related and extrahepatic disease complications which relates to the fact that excess iron catalyses the formation of toxic hydroxyl-radicals subsequently resulting in cellular damage. Iron removal improves insulin sensitivity, delays the onset of type 2 diabetes mellitus, improves pathologic liver function tests and likewise ameliorates NAFLD histology. Several mechanisms contribute to pathologic iron accumulation in NAFLD. These include impaired iron export from hepatocytes and mesenchymal Kupffer cells as a consequence of imbalances in the concentrations of iron regulatory factors, such as hepcidin, cytokines, copper or other dietary factors. This review summarizes the knowledge about iron homeostasis in NAFLD and the rationale for its therapeutic implications.

Key words: Dysmetabolic iron overload syndrome; Hepcidin; Iron overload; Metabolic syndrome; Non-alcoholic fatty liver disease; Nonalcoholic steatohepatitis

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Core tip: Hyperferritinemia with normal transferrin saturation and mostly mild hepatic iron deposition is a frequent finding in subjects with non-alcoholic fatty liver disease. Excess iron in non-alcoholic fatty liver disease (NAFLD) patients is referred to as the "dysmetabolic iron overload syndrome". Clinical evidence suggests that elevated body iron stores aggravate the clinical course of NAFLD with regard to liver-related and extrahepatic disease complications. Iron removal improves insulin sensitivity, delays the onset of type 2 diabetes mellitus,

Abstract

Elevated iron stores as indicated by hyperferritinemia

improves pathologic liver function tests and ameliorates NAFLD histology. The mechanisms contributing to iron excess in fatty liver include impaired iron export from hepatocytes and mesenchymal Kupffer.

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INTRODUCTION

Physiological regulation of iron homeostasis

Iron is essential for life of mammalian organisms due to its paradigmatic role in oxygen transport and also in being a central component of many enzymes and proteins involved in mitochondrial respiration, DNA biosynthesis and the citric acid cycle, among others. However, excess iron is detrimental and may lead to severe organ damage as it facilitates the formation of reactive oxygen species (ROS) *via* the Fenton reaction. On the other hand, iron deficiency can lead to anemia and fatigue which are among the most common disorders in the world. In order to provide enough iron for biological function and at the same time avoid iron overload and toxicity, iron trafficking and storage are diligently balanced by a mechanisms involving bone marrow, intestine, liver and the reticuloendothelial system (RES)^[1,2].

Many aspects of iron metabolism have been unravelled in recent years. Dietary iron is taken up as Fe²⁺ in the duodenum by the cation transporter divalent metal transporter 1^[3,4]. After transfer through the duodenal baso-lateral membrane *via* the iron exporter ferroportin (FPN)^[5,6], iron is oxidized by the copper containing ferroxidase hephaestin and loaded onto transferrin for systemic distribution^[7]. Most cells facilitate iron uptake by transferrin bound Fe³⁺ *via* the transferrin-receptor (TfR1). Most iron is required for erythropoiesis and the biosynthesis of other heme enzymes like cytochromes, and excess iron is stored in hepatocytes^[5,8]. Most iron for physiological requirements, mainly erythropoiesis, is obtained from re-utilisation of senescent erythrocytes which are taken up and degraded in splenic macrophages. Only approximately 1-2 mg of daily body iron requirements which are used for compensation of iron losses *via* bleeding, enteric and cutaneous cell desquamation are replenished *via* duodenal iron absorption. Iron export is facilitated by FPN from hepatocytes, macrophages and all other cells^[9].

Systemic iron homeostasis is equilibrated by the peptide hepcidin (hepatic bactericidal protein) mainly derived from hepatocytes and regulated by iron status, hypoxia, anemia and inflammation^[10-12].

Hepcidin impacts on iron trafficking by attaching to FPN which leads to the degradation of FPN and thereby to down-regulation of iron export inducing a decline in serum iron concentrations^[13]. Quantitatively hepatocytes are the most important source for hepcidin, however, expression has also been reported in adipose tissue, pancreatic islets, macrophages, and even cardiac myocytes. Hence, iron homeostasis *via* FPN mediated iron export may be regulated in an autocrine fashion in these cells^[14-16].

Perturbations of iron homeostasis are frequently observed in patients suffering from non-alcoholic fatty liver disease (NAFLD)^[17,18]. As the prevalence of obesity rises, NAFLD with or without associated metabolic syndrome (MetS), has become the most frequent cause of hyperferritinemia. The first report of non-hemochromatotic iron overload linked to metabolic characteristics such as insulin resistance and overweight in a French study subsequently stimulated extensive research on the potential mechanisms underlying iron accumulation in NAFLD^[19]. The dysmetabolic iron overload syndrome (DIOS) commonly refers to the characteristic association of fatty liver with moderate histological iron deposition (hemosiderosis) and increased serum ferritin^[17,20].

WHAT IS THE IRON PHENOTYPE OF NAFLD?

An increase in ferritin concentrations is the key feature of iron dysregulation in subjects with NAFLD. It is found in one third to half of patients with NAFLD and ranges from mild elevations to rarely 1000-1500 ng/mL^[17]. Serum ferritin concentrations increase with the number of features of the MetS^[21]. Transferrin saturation (TfS) is typically in the upper range of normal or mildly elevated (45%-50%) which is distinct from hereditary hemochromatosis, where hyperferritinemia is accompanied by markedly elevated TfS and usually TfS is elevated before the development of hyperferritinemia in early stages of hemochromatosis^[22].

Iron deposits in NAFLD are found in Kupffer cells which are the resident liver macrophages as well as in hepatocytes^[20]. Mesenchymal iron deposition is more frequent than hepatocellular iron accumulation but mostly both compartments are affected^[23]. This is different from tissue iron deposition in primary genetic iron overload, hemochromatosis, where the metal is almost exclusively found in the hepatocellular compartment (with the exception of ferroportin disease) and macrophages are iron deficient as a result of uninhibited iron export from these cells^[24,25]. The extent of hyperferritinemia in subjects with NAFLD and/or the MetS overestimates the degree of iron overload compared to hemochromatosis. Phlebotomy studies demonstrated that in DIOS patients the amount of iron need to be removed

for normalisation of circulating iron parameters is usually significantly less than in hemochromatosis, indicating only mild body iron excess^[26,27]. Few studies have performed liver iron quantification in NAFLD subjects and these results confirm the mild degree of tissue iron excess compared to genetic iron overload disorders^[19,28]. The mild degree of body iron excess compared to markedly raised serum ferritin concentrations suggests that iron overload in NAFLD subjects results from a combination of alimentary and inflammatory driven iron loading and retention^[20,29,30]. This is in line with the current evidence that NAFLD is both a metabolic and an inflammatory disease^[31].

WHAT IS THE CLINICAL RELEVANCE OF ELEVATED IRON STORES?

IR and associated metabolic conditions

In 1981 Sullivan^[32] suggested that the postponed occurrence of cardiovascular diseases in women compared to men and the subsequent postmenopausal increase could be caused by low premenopausal iron stores. This report likely is the first report of an impact of iron stores in non-hemochromatotic metabolic disorders. An association of iron stores with type 2 diabetes mellitus (T2DM) and various manifestations of IR has been repeatedly confirmed and a detailed discussion thereof is beyond the scope of this review^[33]. However, glucose metabolism and iron homeostasis appear to be functionally interconnected, due to the fact that gluconeogenic signals regulate iron homeostasis *via* hepcidin^[34] while iron loading or deficiency directly affect circulating glucose concentrations in mammals most likely *via* its effects on citric acid cycle enzyme activities^[35,36], thereby also affecting lipid profiles^[37]. Ferritin concentrations were associated with an increased rate of diabetes and gestational diabetes^[38-43], with BMI^[44], visceral fat mass^[45], serum glucose levels and insulin sensitivity^[46], blood pressure^[47], the MetS^[21,48], the polycystic ovary syndrome (PCOS)^[49] and cholesterol^[50]. Higher parameters of iron storage clustered with metabolic risk markers in a study of obese^[51] and healthy lean adolescents^[52]. These observations are epidemiologically important as patients with IR have a higher risk of developing cerebrovascular or cardiovascular disease^[53,54]. However, the most convincing argument for causative involvement of iron in obesity-related conditions is derived from iron removal studies mentioned in detail below. In summary, available studies convincingly suggest a direct impact of body iron on manifestations of IR or the MetS.

NAFLD

NAFLD has been firmly established as the hepatic manifestation of the MetS/IR^[55]. The disease spectrum of NAFLD ranges from simple steatosis

which is generally considered benign to steatosis with various stages of inflammation, hepatocellular ballooning and fibrosis called non-alcoholic steatohepatitis (NASH). NASH is the potentially progressive manifestation leading to cirrhosis, end-stage liver disease and hepatocellular carcinoma in a minority of patients^[56]. To our knowledge, there is no data available suggesting that excess iron is linked to the extent of hepatic steatosis. Although multiple associations between iron homeostasis and lipid metabolism have been reported^[57], no characteristic lipid phenotype has been documented to distinguish NAFLD with iron overload from NAFLD without iron. Underlying NAFLD may explain the link between MetS features and ferritin on the population level^[58].

Several studies provide evidence that iron may contribute to more advanced fibrosis and thus to progression of NAFLD^[18,59-63], however, this association was not confirmed in all studies^[64-66]. The to date largest study reported that iron in NAFLD liver biopsies, particularly in Kupffer cells, was linked to more fibrosis and disease severity^[67]. Iron deposition particularly in the Kupffer cell compartment was associated with higher markers of hepatocellular apoptosis and oxidative stress^[68]. Some studies also suggested that an increased rate of HFE mutations could account for more progressed stages of NAFLD, but this was not reported in all studies^[65,69-72]. Additionally the beta-globin trait^[73], TMPRSS6^[74], and the alpha-1-antitrypsin genotype^[75] may modify the iron phenotype of NAFLD. It appears reasonable to conclude that the contribution of the genetic background may vary according to the geographic region. Data evaluating causality of iron in disease progression is limited by the feasibility of a prospective study with serial liver biopsies in enough patients to adjust for known co-factors of disease progression^[64,76]. Retrospective studies demonstrated that, hyperferritinemia was linked to mortality of patients on the transplantation waiting list and it also had an impact on post-transplant mortality^[77,78]. It is important to note that particularly sinusoidal iron deposition may be linked to the development of HCC in NASH^[79].

In summary, the prevailing body of evidence suggests that excess iron is a contributing factor for the progression of steatosis to NASH, liver cirrhosis and also hepatocellular carcinoma. It remains to be established to what extent different patterns of iron deposition affect outcomes such as cirrhosis, HCC or cardiovascular diseases. The data mentioned above suggest that the pattern of iron deposition may have distinct effects.

HOW DOES IRON LEAD TO DISEASE PROGRESSION IN NAFLD?

It has been well recognized that iron overload leads to diabetes in patients with hemochromatosis where

IR increases and insulin secretion decreases with the rise of body iron stores^[25,80-82]. Hepatic insulin sensitivity and insulin secretion are re-established in the majority once iron is removed^[83,84]. However, the prediabetic stage in hemochromatotic mice and humans displays impaired β -cell function along with increased insulin sensitivity, whereas dietary iron overload similar to the prediabetic state in humans are characterized by peripheral IR^[85]. Hence, lessons drawn from hemochromatosis models are likely not fully applicable to the role of iron in human IR and NAFLD.

Iron is well-recognized as a catalyst for the production of reactive oxygen intermediates *via* the Fenton reaction, and it is generally held that an increase of oxidative stress is a central mechanism for IR although direct proof for this hypothesis has not been obtained so far. Oxidative stress is a central pathogenic factor in NAFLD, T2DM and obesity^[86-88] and markers of oxidative stress were increased in NAFLD with iron loading as compared to NAFLD without iron excess^[68,89,90]. Generation of ROS may induce lipid peroxidation and cellular damage which may contribute to the progression of NAFLD. Importantly, oxidative stress induced molecules such as malonyldialdehyde and 4-hydroxynonenal may induce the formation of *de-novo* antigens with subsequent activation of T-lymphocytes and development of immunoglobulin G reactive against these antigens. This response was further enhanced by previous immunization against these antigens with a stimulated M1 macrophage response^[91]. Although no studies have been performed, iron may contribute to this process by further augmentation of oxidative stress.

In cell culture, iron chelation re-established insulin receptor signalling and iron inhibited insulin receptor activity^[92]. Desferoxamine increased the phosphorylation of Akt/protein kinase B (Akt/PKB), forkhead transcription factor O1 (FoxO1) and glycogen synthase kinase 3 β (GSK3 β) reflecting insulin effects on gluconeogenesis and glycogen synthesis. Likewise, genes playing a role in glucose utilization such as GLUT1 or hypoxia-inducible factor 1 α (HIF1 α) were up-regulated in hepatoma cells resulting in enhanced glucose removal^[92]. In summary, these molecular observations indicate that iron affects IR by modulating insulin receptor signalling as has been recently reviewed^[93].

Importantly, dietary iron intake may impact on glucose metabolism by affecting circadian rhythm *via* heme mediated effects on RevErb- α . Disruption of circadian rhythms, *e.g.*, through night-shift work is an established risk factor for metabolic and cardiovascular diseases^[94,95].

In cultured fat cells, iron favored an IR, characterised by impaired glucose uptake and suppression of lipolysis in response to insulin^[96,97]. Ferritin was

inversely associated with adiponectin concentrations in insulin resistant and sensitive patients^[98,99]. Knockout of FPN1 in adipocytes increased intracellular iron and subsequently reduced adiponectin biosynthesis, thus establishing a molecular link between adipocyte iron concentration and insulin resistance^[100]. Furthermore, excess iron the diet may be routed to visceral adipose tissue and change the expression of adipokines, as demonstrated for resistin^[101]. Adipokines represent a diverse group of hormones which mediate the metabolic effects of diseased adipose tissue to organs and tissues. Associations have been observed between retinol-binding protein 4 (RBP4) and visfatin serum concentrations and parameters of iron metabolism^[102,103]. However, these reports may reflect the co-incidence of elevated iron stores with surrogate markers of IR and do not prove causality^[93].

Liver macrophages named Kupffer cells, which are an important site of iron storage in NAFLD, are tightly involved in the initiation of the hepatic inflammatory cascade in response to the uptake of oxidized lipoproteins^[104] or oxidized phosphatidylcholines^[105]. It is well known that macrophage iron status affects their inflammatory response pattern and polarization towards a pro-inflammatory phenotype^[106], however, the particular role of these potential interactions have to our knowledge not been investigated in NAFLD.

Thus, the potential mechanisms of iron-induced NAFLD disease progression are complex and involve protean effects of iron in extrahepatic tissues as well direct liver damage.

WHAT ARE THE MECHANISMS UNDERLYING IRON ACCUMULATION IN NAFLD?

Hepcidin is the key regulator of systemic iron homeostasis and plays a role for the hemochromatotic and the inflammatory driven misdistribution of iron. Whereas the lack of hepcidin in hemochromatosis leads to uncoordinated duodenal iron absorption and iron accumulation in parenchymal tissues such as the liver^[107], the inflammation driven iron retention occurs mainly in monocytes/macrophages as a consequence of increased iron accumulation and reduced FPN mediated iron export from these cells, the latter being due to increased circulating hepcidin levels along with negative effects of certain cytokines on FPN expression^[108]. The histological hallmarks of hemochromatosis, *i.e.*, hepatocellular iron, and also the inflammatory phenotype iron deposition in macrophages are both observed concurrently, suggesting that iron dysregulation is multifaceted in NAFLD. Several stimuli of hepcidin regulation have been reported which may be of particular relevance in NAFLD and also be related to different iron phenotypes. These stimuli and their relation

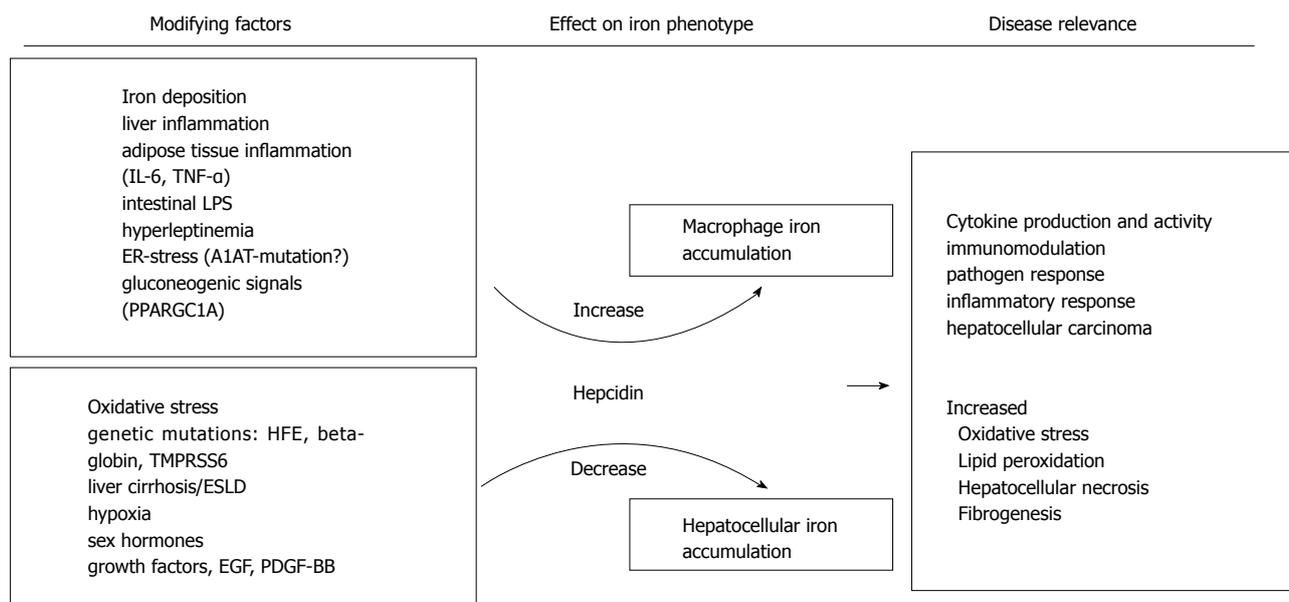


Figure 1 Summary of the potential stimuli that may affect iron homeostasis in non-alcoholic fatty liver disease. Both, increasing and decreasing stimuli have been reported in non-alcoholic fatty liver disease and it appears likely that the net balance of these frequently counteracting forces finally determines the iron phenotype in the individual. Patterns of iron deposition may also be linked to distinct clinical consequences. IL-6: Interleukin 6; TNF- α : Tumor necrosis factor- α ; LPS: Lipopolysaccharide; ER: Endoplasmic reticulum; A1AT: α -1-antitrypsin; PPARGC1A: Peroxisome proliferator-activated receptor gamma coactivator 1- α ; TMPRSS6: Transmembrane protease, serine 6; ESLD: End-stage liver disease; EGF: Epidermal growth factor; PDGF-BB: Platelet derived growth factor BB.

to NAFLD iron accumulation are summarized in Figure 1. For several of these, like sex hormones, growth factors and hypoxia-induced circulating factors, the contribution to the dys-regulation of iron homeostasis in NAFLD has not been directly demonstrated but is physiologically plausible and these have therefore been included in the summary figure^[109-112]. Additionally, alcohol consumption may decrease hepcidin expression and thus modify iron accumulation in NAFLD subjects^[113], and although relevant alcohol consumption should be excluded in NAFLD subjects both conditions frequently co-exist. Thus, in NAFLD multiple, potentially counteracting signals impacting on hepcidin expression may be present at the same time. It is likely that the net balance of these signals finally determines the pattern of iron accumulation in the fatty liver of the individual patient.

Hepcidin levels in urin, serum and liver were elevated in NAFLD patients with iron excess compared to healthy subjects, hemochromatosis patients and NAFLD subjects without excess iron^[28,114-116]. Hepcidin expression correlated directly with liver iron indicating an intact physiological response of hepcidin biosynthesis to iron in the liver^[28,114]. Additionally, hepcidin is expressed in adipocytes of morbidly obese subjects^[15]. Moreover, obesity is characterised by a chronic subclinical inflammation and in humans hepcidin concentrations and TNF- α were directly related, suggesting that both iron and inflammation contribute to hepcidin biosynthesis in NAFLD^[28]. Furthermore, hepcidin and cytokines may be derived from both, the inflamed adipose and the liver^[117,118].

Activation of gluconeogenesis *via* starvation, namely activation of peroxisome proliferator activated receptor gamma co-activator-1 α (PGC1 α) increased hepcidin expression in a mouse model^[34]. Likewise, iron fortification decreased gluconeogenesis *via* PGC1 α in a murine model^[119]. Hence, although PGC1 α offers an intriguing cellular link between glucose and iron homeostasis, its relevance to human NAFLD remains to be elucidated. Leptin, was demonstrated to up-regulate hepcidin in hepatocytes *in vitro* by activation of the JAK2/STAT3 pathway. Hence, hyperleptinemia may directly contribute to higher hepcidin and thereby to iron deposition in NAFLD^[120,121].

In NAFLD with iron overload the iron exporter FPN is lower than in controls and hemochromatosis patients in the liver and in the duodenum^[28,114,122,123]. In NAFLD without liver iron accumulation, FPN levels were comparable to control subjects, but were significantly lower in NAFLD with hepatic iron on histology^[28]. Along the same line of the observations, duodenal iron absorption was decreased in DIOS patients^[124]. Obesity also represents a risk factor for an inadequate dietary iron fortification, linked to high hepcidin and low FPN expression^[125]. Along this line mice feed a high fat diet presented with significantly reduced iron absorption which could be traced back diminished intestinal iron uptake. Mechanistically, the impaired iron absorption was independent of hepcidin but resulted from reduced metal uptake into the mucosa and transfer of iron across enterocyte membranes as a consequence of dietary induced discordant membrane-bound oxidoreductase expression^[126].

An additional mechanism may be the phagocytosis of fragile erythrocytes by liver Kupffer cells. This was documented in rabbits on a high-fat diet and the phagocytosis of fragile erythrocytes was observed *in vitro*. Accumulation of erythrocytes was microscopically detected in inflamed regions in human NAFLD^[127] suggesting that uptake of heme-iron *via* erythrophagocytosis may contribute to NAFLD iron accumulation, then promoting oxidative stress and inflammation.

Although cellular iron uptake *via* TfR1 is the most important route of iron uptake under physiological circumstances TfR1 appears not to be involved in excess iron uptake in NAFLD^[128,129]. Hepatic TfR1 expression in NAFLD patients with low iron was increased compared to NAFLD and iron accumulation or patients with hemochromatosis suggesting physiologically intact TfR1 expression in response to iron stimuli^[128] (Figure 2).

WHAT IS THE ROLE OF COPPER IN NAFLD?

Similar to iron, an adequate supply of copper is essential for proper biological function. Chronic copper deficiency can elicit anemia, leucopenia, myelopathy or skin abnormalities and excess copper may also facilitate the formation of ROS.

Copper affects lipid and glucose metabolism

There are several ways in which inadequate copper supply may be involved in the pathogenesis of NAFLD. Epidemiological studies found that copper deficiency is linked to atherogenic dyslipidemia and dietary copper supplementation improved cardiovascular risk markers in healthy adults^[130]. Investigations in rodent models demonstrated that dietary copper restriction induces hypertension or cardiac dysfunction, hypertriglyceridemia, hypercholesterolemia and modifies LDL and VLDL composition^[131,132]. We recently reported low intrahepatic copper concentrations in human NAFLD compared to other liver diseases and that rats on a copper depleted diet developed IR and liver steatosis^[133]. Increased oxidative stress is considered a key trigger in the pathogenesis of human NAFLD and one of the enzymes counteracting oxidative stress, Cu/Zn superoxide dismutase (SOD) depends on adequate copper availability, suggesting a potential link between copper availability and impaired antioxidant defense in NAFLD^[134]. Sprague-Dawley rats exhibited an increased activity of the pro-inflammatory protein cyclo-oxygenase-2, when fed a diet with a low copper content^[135]. Systemic copper deficiency causes mitochondrial dysfunction in mice and similar morphological and functional alterations have also been described in human NAFLD^[136]. Recently, a detailed examination revealed an interaction of a high-fructose diet (which is also a

culprit in the rise of obesity-related conditions) with low copper intake in triggering liver steatosis and damage as well as iron overload. Fructose acts as an inhibitor of duodenal copper absorption thereby leading to impaired oxidant defense and augmented lipid peroxidation^[137]. As dietary copper content of the Western diet is rather low whereas iron and fructose are consumed in excess, this model offers attractive data to speculate that a dysbalance in micronutrient intake may have a significant role in NAFLD beyond calorie excess. Hence, animal and human data suggest that the therapeutic effect of dietary copper supplementation should be investigated as a subset of patients may potentially benefit.

Copper affects NAFLD iron homeostasis

Copper modulates iron homeostasis and is also linked to the iron perturbations of NAFLD. Hephaestin ferroxidase activity in duodenal enterocytes is critically dependent on copper as it oxidizes ferrous to ferric iron which is subsequently loaded onto Tf^[7]. Similarly, copper is necessary for ceruloplasmin function to export iron from the liver or the RES and also for FPN expression^[138]. Expression of a membrane-bound form of ceruloplasmin is mandatory for stable FPN expression^[139,140]. Accordingly, a lack of ceruloplasmin as found in the heritable disease aceruloplasminemia leads to tissue iron accumulation and damage most notably in the brain^[141].

Low liver and serum copper concentrations were reported in iron overloaded NAFLD and were linked to decreased ferroxidase activity of ceruloplasmin^[122]. The expression of FPN was found to be decreased in livers of rats on a copper deficient diet. These observations provide evidence that in addition to decreased FPN expression due to low-grade systemic inflammation, low copper bioavailability contributes to iron retention in NAFLD.

WHAT IS THE THERAPEUTIC POTENTIAL OF MODULATING IRON STORES IN NAFLD?

Elimination of iron may confer a beneficial effect on IR-associated conditions. Removal of iron using phlebotomies is usually well tolerated, with the caution that DIO patients frequently show a fast decline in TfS^[142]. These clinical observations are expected due to the underlying molecular mechanisms of impaired iron export. The incidence of diabetes, postprandial serum insulin and pancreatic insulin sensitivity, *i.e.*, beta cell function were all improved in subjects with previous phlebotomy treatment^[143]. Iron removal also improved coronary vascular dysfunction in patients with T2DM^[144] and endothelial function in patients with known coronary artery disease and in subjects with primary iron

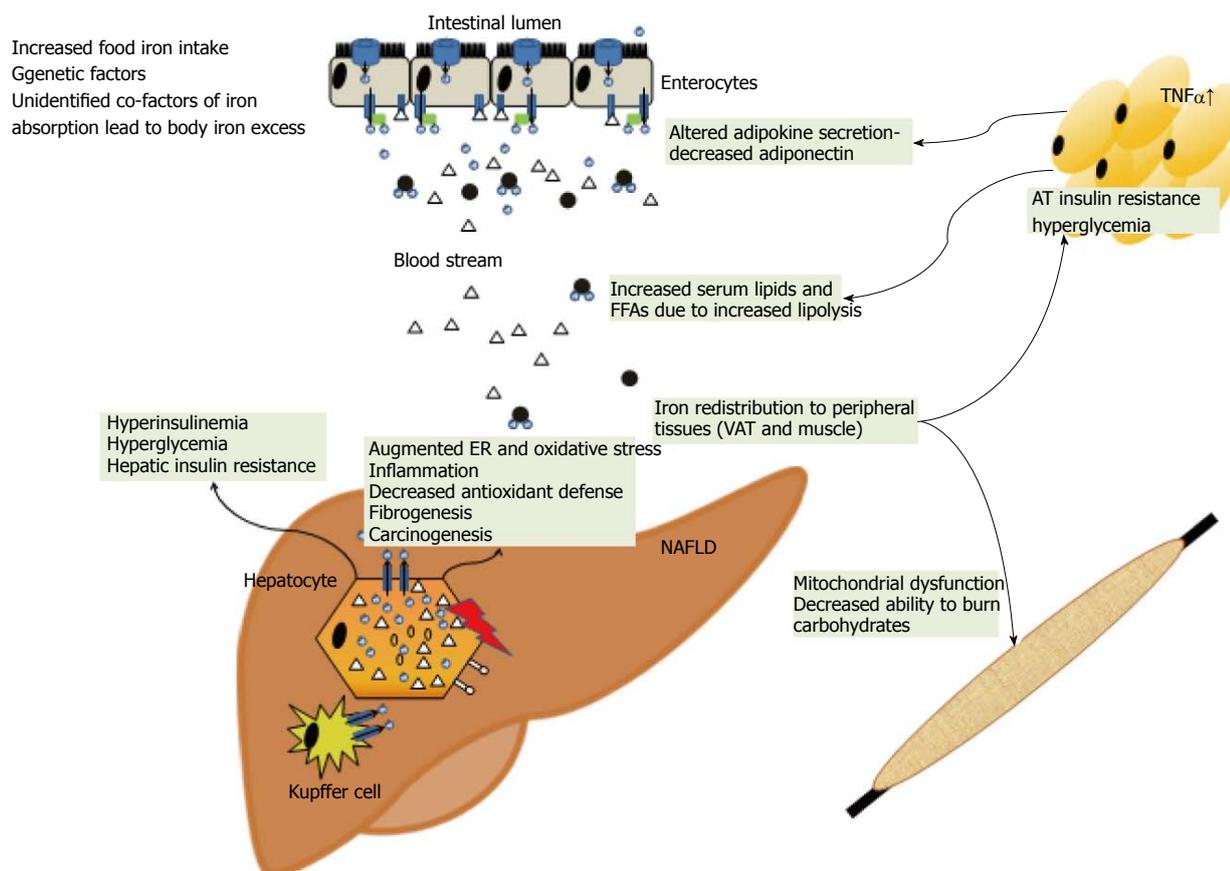


Figure 2 Summary of how iron excess and low copper availability may affect whole body glucose and lipid homeostasis. Iron excess may promote insulin resistance in the liver, muscle and adipose tissue. Iron may increase ER and oxidative stress whereas low copper is potentially associated with an impaired antioxidant defence. These factors may result in the propagation of inflammation, fibrogenesis and hepatocarcinogenesis. TNF- α : Tumor necrosis factor- α ; ER: Endoplasmatic reticulum; FFA: Free fatty acid; VAT: Visceral adipose tissue; AT: Adipose tissue.

overload^[145,146]. Blood donations were linked to insulin sensitivity even in healthy subjects^[46]. Studies on iron depletion in NAFLD in humans have demonstrated benefits regarding systemic or hepatic insulin resistance and pancreatic insulin sensitivity^[142,147,148]. A randomized trial demonstrated improved HbA1c, insulin sensitivity and secretion subjects who received phlebotomy treatment^[149]. The effects of iron depletion were additive to successful lifestyle modifications^[150]. Similar observations were reported the effect of iron depletion on other cardiovascular risk factors^[151] and iron removal may prevent development and progression of malignancies^[152].

As far as practical treatment of iron excess in NAFLD patients with elevated ferritin is concerned, available data suggest that iron removal may thus be beneficial in addition to weight loss, diet and lifestyle modification or antidiabetic medication as indicated in an individual patient. We have adopted the practice to perform biweekly phlebotomies in these subjects until serum ferritin concentrations are between 50 and 100 ng/L, however, no evidence-based recommendation for this is currently available. In contrast to hemochromatosis patients, NAFLD subjects have impaired

iron mobilisation from storage sites and may therefore develop anemia in response to phlebotomy treatment. We therefore recommend close monitoring of serum ferritin, TfS and hemoglobin at each visit for the period of time while these patients are on phlebotomy treatment^[26,153].

CONCLUSION

Elevated serum ferritin concentrations are a frequent finding in NAFLD. Excess iron is linked to IR, accelerated disease progression and adverse outcomes. Removing excess iron *via* phlebotomies is safe and has clinical benefits. We suggest that on the basis of available evidence it can be offered to NAFLD patients as it is linked to improvement of IR and inflammation. The mechanisms underlying iron accumulation in NAFLD are tightly linked to impaired iron export from liver cells as a consequence of low expression of the iron export molecule FPN and elevated hepcidin concentrations. Inflammation of adipose tissue as indicated by TNF- α and IL-6 and altered adipokine secretion (leptin, resistin) or hepcidin represent potent signals from diseased

adipose tissue to dysregulate iron as well as glucose or lipid homeostasis.

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Antiviral treatment for chronic hepatitis B in renal transplant patients

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showing a favorable safety and efficacy profile of nucleos(t)ide analogue (NUC) treatment in the renal transplant setting. Entecavir, a drug without major signs of nephrotoxicity, appears to be the first option for NUC naïve patients and tenofovir remains the preferred choice for patients with previous resistance to lamivudine or any other NUC. Renal transplant recipients under antiHBV therapy should be monitored for its efficacy against HBV but also for its safety with a close renal monitoring. Studies including a large number of patients with long term treatment and follow up are still needed to better demonstrate the safety and efficacy of newer NUCs in this population.

Key words: Tenofovir; Long term outcome; Hepatitis B; Renal transplantation; Entecavir

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Core tip: Nucleos(t)ide analogue treatment is safe and effective in renal transplant patients. It improves long term patients and graft survival.

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Abstract

Chronic hepatitis B infection is frequent in renal transplant patients. It negatively impacts long term outcomes reducing graft and patient survival. Current guidelines clearly define who needs treatment, when to start, what is the first line therapy, how to monitor treatment response, when to stop, and how patients must be controlled for its safety. There is some data

INTRODUCTION

Renal transplantation (RT) is the preferred treatment for patients with end-stage renal disease (ESRD) undergoing renal replacement therapy. Moreover, RT improves quality of life and survival when compared with remaining on the waiting list^[1]. In the last twenty years, improvement in immunosuppressive therapy resulted in a decline in acute rejection

prevalence and in 1-year improvement in graft survival. In spite of short-term survival, both long-term patients and graft survival has not improved as expected^[2,3]. Cardiovascular diseases, malignancy and infections are the most frequent determinants of death in RT recipients. Liver failure appears as the fourth cause of death in long-term survivors after RT^[3-6]. In these patients liver failure is mostly related to chronic viral hepatitis B (HBV) and C (HCV). Both HBV and HCV negatively impact on renal transplantation outcomes by reducing long term graft and patient survival; the magnitude of this impact may vary between these viruses and may differ from different series. Treatment strategies of these viruses are clearly different in patients with ESRD. Nowadays HCV must be treated with Peg-interferon combined with low dose ribavirin before transplantation; in the near future new antivirals will allow HCV treatment after renal transplantation. Conversely, HBV can be treated with the same drugs across all stages of chronic renal disease: before and in dialysis, and after RT.

In the last decades HBV prevalence has decreased in dialysis units due to the implementation vaccination programs and infection control procedures. Today it varies between 0% to 20% according to different sources^[7,8]. But prevalence in RT patients tends to vary and can be higher since some of these patients were transplanted before these programs were widely available. The present review focuses on the current management of patients with HBV after renal transplantation.

PRE-TRANSPLANT EVALUATION

Chronic HBV infected patients with ESRD must be adequately evaluated before being transplanted. Two key aspects must be taken into account: evaluation of HBV status and the severity of liver disease. Regarding HBV status, all HBsAg (hepatitis B surface antigen) positive patients and all patients with previous known history of acute or chronic hepatitis B or the presence of antiHBc (hepatitis B core antibody) with/without antiHBs (hepatitis B surface antibody) require a full virological evaluation including HBeAg (hepatitis B envelope antigen) and antiHBe (hepatitis B envelope antibody) determination and HBV DNA levels measurement. This evaluation will allow classifying these patients into different clinical situations^[9,10].

Chronic hepatitis B: (1) HBsAg positive for more than 6 mo; (2) Serum HBV DNA \geq 2000 (EASL)-20000 (AASLD) IU/mL (10^4 - 10^5 copies/mL), lower values 2000-20000 IU/mL (10^4 - 10^5 copies/mL) are often seen in HBeAg-negative chronic hepatitis B; (3) Persistent or intermittent elevation in ALT/AST levels; (4) Liver biopsy showing chronic hepatitis with moderate or severe necroinflammation; and (5)

Chronic hepatitis B can be subdivided into HBeAg positive and HBeAg negative chronic hepatitis B.

Inactive HBsAg carrier state: (1) HBsAg positive for more than 6 mo; (2) HBeAg negative, antiHBe positive; (3) Serum HBV DNA < 2000 IU/mL; (4) Persistently normal ALT/AST levels; and (5) Liver biopsy confirms absence of significant hepatitis.

Resolved hepatitis B: (1) Previous known history of acute or chronic hepatitis B or the presence of antiHBc with/without antiHBs; (2) HBsAg negative; (3) Undetectable serum HBV DNA; and (4) Normal ALT levels.

Concomitantly, the severity of liver disease must be evaluated before RT usually by obtaining a liver biopsy. There is some debate about the better route to perform the liver biopsy given that patients with ESRD undergoing hemodialysis have an increased risk of bleeding associated with clotting diseases, uremia-associated platelet dysfunction and intradialysis antiaggregants and anticoagulant treatments^[11]. Once transplanted this risk disappears with the restoration of normal renal function. In some centres the transjugular route is the preferred one since is associated with less a reduced risk of bleeding and pain, and may allow measuring the hepatic venous pressure gradient (HVPG) for confirming and grading portal hypertension^[12,13]. However in many centres the percutaneous trans-thoracic route is still widely and safely used^[14].

There are some noninvasive tests to evaluate the severity of liver fibrosis but they have not been widely evaluated in dialysis and RT patients^[11,15]. FibroTest (FT) and liver stiffness measurement (LSM) for noninvasive assessment of liver fibrosis had been evaluated in RT patients with chronic HBV or HCV infection^[16]. It had been reported that FT and LSM are sufficiently accurate for diagnosing mild liver fibrosis (\leq F2), but differed by 38.4% from the histological data in patients with more severe fibrosis (\geq F3); their predictive value for diagnosing severe liver disease needs to be confirmed. More information is needed in HBV infection to recommend its use.

Once the HBV clinical situation and the severity of liver diseases have been established, treatment indication and possibility of RT has to be defined. HBsAg positive patients receiving immunosuppressive therapy after renal transplantation must antiHBV prophylaxis or treatment (based on HBV DNA levels) with a NUC.

In the general population HBV inactive carriers do not need to be treated^[9,10]. But RT candidates who are inactive carriers have a higher risk of reactivation after transplantation. In HBsAg positive inactive carriers, treatment can be used as prophylactic (HBV DNA undetectable, no hepatocellular injury), preemptive (HBV DNA < 2000 IU/mL, no hepatocellular injury), and salvage

Table 1 Dosage adjustment of nucleos(t)ide analogue for patients with reduced creatinine clearance

	Recommended dosage		Dosage forms	
Creatinine clearance (mL/min)	Lamivudine ^[25]		Lamivudine ^[25]	
	> 50	100 mg once daily	Tablets: 100 mg	
	30-49	100 mg first dose, then 50 mg once daily	Oral solution: 10 mg/mL	
	15-29	100 mg first dose, then 25 mg once daily		
	5-14	35 mg first dose, then 15 mg once daily		
< 5	35 mg first dose, then 10 mg once daily			
> 50	Adefovir ^[26]		Adefovir ^[26]	
	10 mg every 24 h		Tablets: 10 mg	
	30-49	10 mg every 48 h	Oral solution: not available	
	10-29	10 mg every 72 h		
	Hemodialysis	10 mg every 7 d following dialysis		
> 50	Telbivudine ^[27]		Telbivudine ^[27]	
	600 mg every 24 h		Tablets: 600 mg	
	30-49	600 mg every 48 h	Oral solution: 100 mg/5 mL	
	10-29	600 mg every 72 h		
	Hemodialysis	600 mg every 96 h following dialysis		
> 50	Entecavir ^[28]	Entecavir in Lamivudine-Refractory ^[28]	Entecavir ^[28]	
	0.5 mg once daily	1 mg once daily	Tablets: 0.5 mg and 1 mg	
	30-49	0.25 mg once daily OR	0.5 mg once daily OR	Oral solution: 0.05 mg/mL
		0.5 mg every 48 h	1 mg every 48 h	
	10-29	0.15 mg once daily OR	0.3 mg once daily OR	
		0.5 mg every 72 h	1 mg every 72 h	
	Hemodialysis	0.05 mg once daily OR	0.1 mg once daily OR	
		0.5 mg every 7 d following dialysis	1 mg every 7 d following dialysis	
		Tenofovir ^[29]		Tenofovir ^[29]
	> 50	300 mg every 24 h		Tablets: 300 mg
30-49	300 mg every 48 h		Oral powder: 40 mg per 1 g of oral powder	
10-29	300 mg every 72 to 96 h			
Hemodialysis	300 mg every 7 d or after approximately 12 h of dialysis			

therapy after reactivation (HBV DNA > 2000 IU/mL, with hepatocellular injury). Even if the prophylactic/preemptive initiation is the generally accepted treatment, the data comparing these treatments are few^[17]. All RT candidates with *chronic HBV* need to be treated before transplantation with NUCs^[9,10]. Patients with *resolved HBV* have a low reactivation risk in the RT setting varying between 0.6% to 6%^[18-20]. Since there is a low reactivation risk in HBsAg negative patients, universal prophylaxis is not recommended in them. Among antiHbc positive patients, those having low antiHBs titers (< 100 IU/mL) have the higher risk of reactivation. Even though there is limited evidence, repeat vaccination may be considered for this group. Current HBV DNA tests allows to diagnose true occult infection in patients with isolated antiHbc positive serology. There is not enough information about the absolute risk of reactivation in this sub-population, so it is unclear whether prophylaxis is beneficial^[21].

The severity of liver disease will determine if the patient is a good candidate for RT or not. Evidence of decompensated liver disease (ascites, encephalopathy, variceal bleeding, etc.) precludes RT and is a clear indication of combined liver-renal transplantation (LRT). The presence of compensated cirrhosis with signs of portal hypertension is also an indication for LRT. Cirrhotic patients without portal hypertension must be carefully evaluated

for RT since cirrhosis is correlated with an higher mortality risk^[22]. Non cirrhotic patients are adequate candidates for RT.

TREATMENT

There are many guidelines regarding HBV treatment. Patients with ESRD and RT can be considered a special population and there are particular recommendations for them that may vary from those implemented in the general population^[9,10,15,23]. Patients with chronic hepatitis B are candidates for treatment and those inactive hepatitis B carriers are candidates for prophylactic or preemptive therapy.

There are two main treatment options in hepatitis B: interferon and NUCs. Interferon therapy has many disadvantages when compared with NUCs: poorly tolerated due to side effects, limited efficacy in this populations, subcutaneous administration and there is certain risk of graft rejection^[24]. So, there is agreement that in RT patients with chronic HBV infection interferon based therapy should be avoided^[15,21,23]. On the contrary, NUCs have a high antiviral potency, have a good safety and tolerability profile and can be orally administrated. These drugs can be easily used in RT and doses can be adjusted according to creatinine clearance^[25-29] (Table 1). The main limitations of NUCs include the need for long-

term therapy, which may be for indefinite time in HBeAg negative patients; the risk of development of NUC's resistant viral strains; and the unknown safety profile with long-term treatment^[11].

Regarding NUCs, there are five drugs currently approved for HBV treatment: lamivudine (LAM), adefovir (ADV), telbivudine (LdT), entecavir (ETV) and tenofovir disoproxil fumarate (TDF). Treatment with TDF or ETV is preferable to LAM in NUC naïve patients, since they are more effective due to a high antiviral potency and have a high barrier to resistance reducing the risk of drug resistance and treatment failure^[9,10,15,21,23].

Since LAM was the first NUC approved for clinical use, it has yielded the majority of data on the management of HBsAg-positive renal transplant recipients. Several observational studies have shown that LAM can improve liver function^[15]. A meta-analysis including 14 prospective cohort studies (184 patients) showed that LAM normalized ALT levels in 81% (95%CI: 70%-92%), cleared HBV-DNA in 91% (95%CI: 86%-96%) cleared HBeAg in 27% (95%CI: 16%-39%) of the patients. In most studies (11 of 14) LAM was administered for 6 to 12 mo^[30]. Even though LAM was associated with significantly improved patient survival^[31], prolonged treatment is associated with progressive increase in drug resistance and the cumulative probability of developing LAM resistance (LAM-R) was approximately 60% after 69 mo^[30,32,33]. LAM-R leads to treatment failure and can be associated with progressive liver disease and a negative impact in patient and graft survival. Fortunately today there are good treatment options for LAM-R. Given that there are better options for HBV treatment, LAM cannot be consider within the first treatment choices for these patients^[9,10,15,21,23].

Adefovir was the second available oral drug for HBV treatment infection. It has similar antiviral activity against both LAM-R and wild-type HBV, but it may be nephrotoxic (especially in high doses). Currently its major clinical application is as add on therapy for the management of lamivudine-resistance since it has lower antiviral activity than ETV and TDF for naïve patients^[9,10,15].

There have been reports on ADV short-term efficacy either as mono- or add-on therapy in LAM-R RT patients^[34-39]. One year ADV monotherapy showed a significant viral response in 11 patients with a median HBV DNA decline of 5.5 log₁₀. Only one patient cleared HBV DNA, one of the six HBeAg positive patients cleared HBeAg but without antiHBe seroconversion; none cleared HBsAg. Importantly, there were no significant clinical and laboratory adverse events^[34]. ADV as add-on therapy to LAM resulted in significant HBV suppression LAM-R RT recipients^[38,39]. In 11 ADV add on treated patients, HBV DNA was undetectable in 80%-83% after 36 to 42 mo^[38]. However, six patients (54%) had to lower

ADV dose due to a decline in glomerular filtration rate after a median of 11 mo (range: 9-42)^[38]. After 12, 24 and 36 mo of ADV treatment treatment 35.7%, 42.8% and 88.0% of treated patients cleared HBV DNA; there was no virological breakthrough and 92.8% of patients achieved normal ALT levels after 12 mo of treatment^[39]. Patients treated with add-on ADV therapy tended to normalize ALT levels and to reduce HBV DNA levels more effectively than those treated with ADV monotherapy^[39]. In this study 29% of the participants developed moderate to severe renal failure^[39].

However, when compared with treatment-naïve the virological response could be fluctuating and relatively slow in LAM-R patients^[40]. Nevertheless, rescue therapy with ADV resulted in significantly better viral suppression and liver biochemistry compared with continuation of LAM (75% vs 14.3% had persistent normalization of ALT), and the clinical response was sustained for at least 24 mo^[31]. Evidence of nephrotoxicity in the absence of proximal tubulopathy, despite dosage adjustment, was frequently observed, and could necessitate treatment discontinuation^[38,39]. ADV has a low antiviral potency at the currently approved dose and its efficacy could be further reduced with dose adjustment according to renal dysfunction. For these reasons ADV is not a first line option for naïve patients and its benefits for LAM-R may be less when compared with TDF.

There are currently no results about telbivudine treatment in RT recipients but it would be worthwhile to explore the use of this agent in treatment-naïve kidney allograft recipients given its relatively low resistance rate, lack of nephrotoxicity, and the relatively lower cost compared with other nucleoside/tide analogues^[40].

Entecavir is one of the first line treatment options for HBV^[9,10]. This drug has a high antiviral potency, a high genetic barrier for resistance and a good safety profile. It is very effective for treatment naïve patients but has a lower efficacy for LAM-R patients, and it is not the first option for this latter population^[9,10,15]. A recent 2-year prospective study included 27 RT patients, 18 (67%) were treatment naïve and 9 (33%) had been previously treated with LAM but had no resistant mutations. ETV cleared HBV DNA in 70%, 74%, 96% and 100% of patients after 12, 24, 52 and 104 wk respectively^[41]. There was no change of creatinine clearance, and no episodes of lactic acidosis or muscle damage during treatment. There were higher rates of undetectable HBV DNA levels in ETV treated than LAM treated patients (32%, 37%, 63% and 63% at 12, 24, 52 and 104 wk, respectively; $P < 0.005$)^[41]. In an analysis excluding 9 patients from the ETV group who were also LAM experienced, the remaining 18 ETV naïve subjects exhibited a better virological response at 52 and 104 wk than 19 previously treated with LAM ($P < 0.05$)^[41].

Other studies reported results with ETV in cohorts including both naïve and LAM-R patients, unfortunately with limited number of patient^[42-45]. Experience regarding the use of ETV in RT recipients who had developed LAM- or ADV-resistance had been examined in a small study with 10 solid organ transplant recipients (8 kidney allograft recipients)^[42]. Treatment with ETV resulted in an appreciable drop in HBV DNA levels and a 50% HBV undetectability in both HBeAg positive and HBeAg negative patients after 16.5 mo of treatment without significant changes in glomerular filtration rate^[42]. In our small experience we reported ETV use in 11 patients with several chronic renal diseases: 1 with stage 4 CKD, 7 in dialysis, and 3 RT recipients^[43,44]. HBV DNA was cleared in 54.5% ($n = 6$); 77.7% of HBeAg-positive patients (7/9) seroconverted to antiHBe positive; and only one patient (9.1%) showed antiHBs seroconversion. There were no significant changes in renal or hematological biochemical parameters^[43,44]. In the most recent report, twenty-one RT patients (10 treatment naïve, 11 with LAM resistance) were treated with ETV for 34.7 ± 22.9 mo (range 6-75 mo)^[45]. The cumulative rate of HBV DNA undetectability at 12, 24, and 36 mo was 60%, 100%, and 100% for treatment naïve group, and 27%, 45%, and 45% for LAM-R group, respectively. Genotypic resistance to ETV emerged after 20.0 ± 3.5 mo with increase in ALT and HBV DNA in two patients with LAM-R, but was not observed in the treatment-naïve group. There were no significant changes in glomerular filtration rate^[45]. Also, ETV was used in RT patients who developed hepatic flares due to the appearance of LAM-R^[46]. Four patients were treated with ADV and two with ETV. After 18 mo, HBV DNA was $< 10^5$ copies/mL in 4 subjects and $< 10^2$ copies/mL in 1 subject. There were no remarkable adverse events and no changes in renal function^[46]. ETV appears as one of the best options for NUC naïve RT patients; it is less effective in LAM-R and better options are available.

Tenofovir was the last NUC to be approved for HBV monoinfection and is the other first line option together with ETV^[9,10,15]. It has a high antiviral potency, a high genetic barrier for resistance and a good safety profile^[9,10,15], but there is some concern about its potential nephrotoxicity^[47]. There is little data in the renal transplant setting: only one study reports the results of three RT treated patients together with 3 liver, and 1 heart transplant recipients^[48]. HBV DNA viral became significantly decreased and 3 patients cleared HBV DNA at the end of the study period. There were no adverse events related to tenofovir treatment. No episodes of acute rejection were reported under therapy. There were no statistically significant changes in renal function represented by stable creatinine levels, estimated creatinine clearance, serum phosphorus level, or daily microalbuminuria level^[48]. TDF appears

as one of the best options for both NUC naïve and LAM-R RT patients; treatment results have to be extrapolated from the general population since there is little experience in RT.

TIMING OF INITIATION OF TREATMENT: PREEMPTIVE OR PROPHYLACTIC THERAPY

Patients with chronic renal disease go through different phases: varying stages of renal failure, ESRD, hemodialysis (HD)/peritoneal dialysis (PD), and transplantation. Once transplanted could suffer various kidney disease and finally lose the graft and return to dialysis. HBV infection will go with the patient along the road. The timing of HBV treatment initiation may vary depending on the stage of renal disease.

Patients undergoing HD or PD who are not RT candidates can start NUC therapy if HBV DNA levels are ≥ 2000 IU/mL regardless of ALT levels, especially if they have moderate fibrosis in the liver biopsy (METAVR score $F \geq 2$) or estimated by a non-invasive methods^[11].

All HBsAg-positive RT recipients are considered candidates for NUC treatment. RT candidates with HBV DNA levels > 2000 IU/mL must initiate treatment at HBV diagnosis, those with HBV DNA ≤ 2000 IU/mL should start therapy at least 2 wk before RT. NUC therapy has to be continued indefinitely as long as the patients are under any immunosuppressive treatment^[9,10,11,15]. It should be remembered that compensated cirrhotic patients are not candidates for RT, and cirrhotic patients with decompensated disease should be evaluated for combined liver-kidney transplantation^[11,49].

As previously mentioned, RT candidates who are inactive carriers have an increased reactivation risk after transplantation. In this subgroup of HBsAg positive patients treatment can be used as prophylactic (HBV DNA undetectable, no hepatocellular injury), preemptive (HBV DNA ≤ 2000 IU/mL, no hepatocellular injury), and salvage therapy after reactivation (HBV DNA > 2000 IU/mL, with hepatocellular injury). Even if the prophylactic/preemptive initiation is the generally accepted treatment, the data comparing these treatments are few. The disappearance of viral load is a prerequisite for a HBV positive patient on hemodialysis to be enrolled in the RT list. Therapy with ETV, TDF or LAM on adjusted doses for renal function is included in the current guidelines for prophylaxis of HBV positive RT candidates. The optimal NUC regimen has not been proposed yet, so prophylaxis may start before or at the time of RT and continue thereafter^[9,10,15,50]. ETV should be the first line option for avoidance of short term resistance and ADV nephrotoxicity, while TDF had better be applied in case of LAM-R^[9,10,50].

Lamivudine is the most extensively drug used in prophylactic/preemptive therapy in RT patients. In a small study, LAM given as either prophylactic or preemptive treatment was proven superior to salvage therapy when liver dysfunction is evident^[51]. None of the HBsAg positive patients receiving prophylactic or pre-emptive therapy developed reactivation, while 50% of the patients not been treated suffered reactivation^[51,52]. These results were confirmed by others, but there is some controversy about the clinical impact of prophylactic/preemptive therapy vs salvage therapy^[51-55]. One study showed that there was no differences in survival between HBsAg positive RT patients treated preemptively with LAM and HBsAg negative controls. HBsAg positive patients transplanted without preemptive therapy had an increased mortality rate [relative risk of death, 9.7 ($P < 0.001$); relative risk of liver-related mortality, 68.0 ($P < 0.0001$)^[53]. Twenty five RT candidates received pre-transplantation prophylactic/preemptive NUC therapy, 22 (88%) were treated with LAM and 3 (12%) with ETV^[54]. When compared with a historical control group NUC treated patients has a significant improvement in 10 year graft (82% vs 34%) and patient (91% vs 57%) survivals. There was no liver-related death in NUC treated patients. In contrast, in untreated controls patient death (68%) was the most frequent cause of graft failure, which was mostly caused by liver diseases. Prophylactic and preemptive therapy resulted in the same graft and patient survival, but patients receiving preemptive therapy had a higher HBV reactivation incidence. NUC treatment was independently associated with better patient survival ($P = 0.005$)^[54]. On the contrary, a retrospective analysis using LAM in the majority of patients found no benefit of prophylactic/preemptive treatment^[55]. Ninety four RT candidates were evaluated, 56 received antiviral prophylaxis (Group 1), 51 with LAM and 5 with ETV, and 38 did not (Group 2). In group 2 20 patients experienced HBV reactivation: 16 received LAM, 2 received ETV and 2 received no antiviral treatment. Using the Cox-regression model, prophylactic treatment did not improve patient survival (OR = 1.29, 95%CI: 0.37-4.49, $P = 0.693$), graft survival (OR = 1.25, 95%CI: 0.45-3.46, $P = 0.666$) or reduce the risk of hepatic decompensation (OR = 2.01, 0.35-11.57, $P = 0.434$)^[55]. LAM-R occurred in 21 LAM-treated Group 1 and 4 LAM-treated Group 2 patients ($P = 0.243$), with mean times of resistance after RT of 82 and 132 mo, respectively ($P = 0.001$)^[55].

A recent retrospective study compared both treatment strategies^[17]. It included 58 HBsAg positive RT recipients: 24 in the prophylactic group (all patients used LAM) and the 34 in the preemptive group (32 patients used LAM and 2 patients used ETV). The graft/patient survival rates for HBsAg positive were the same as those of hepatitis-free recipients ($P = 0.18$). In the prophylactic group,

there were fewer hepatic dysfunctions (12.5% vs 30%, $P = 0.12$), viral breakthroughs (16% vs 32%, $P = 0.17$) and elevated alanine aminotransferase concentrations (37% vs 52%, $P = 0.24$), however these differences were not statistically significant. In the prophylactic group, one patient was switched to ETV and then to TDF due to partial response finally achieving complete virological response. In the preemptive group, LAM was withdrawn and changed to TDF in 3 patients and to ADV in another one achieving an adequate virologic/biochemical response. These NUCs were almost as safe as LAM, as there were no significant differences among proteinuria and estimated glomerular filtration rate^[17].

Results from these studies support the clinical guidelines recommendations: prophylactic or preemptive therapy with NUCs provides comparable graft/patient survival with hepatitis-free RT recipients and may be better in preventing hepatic dysfunction than salvage therapy. Given its high risk for developing resistant mutations, LAM is no longer a first option, and ETV should be the first one. TDF can be an effective and safe treatment for LAM-R in RT recipients (Figure 1)^[9,11,15,21,56].

TREATMENT IMPACT ON LONG TERM EVOLUTION

In the last years, several cohort studies had demonstrated that HBV infection is associated with higher patient mortality and risk of graft failure in RT patients^[15,40,50,57]. These results had been validated in two meta-analysis^[58,59]. The first meta-analysis was published in 2005 and included 6050 patients from six observational cohort retrospective studies. Pooled results showed that HBsAg positive status was a significant predictor for death (RR = 2.49, 95%CI: 1.64-3.78) and for graft loss after RT, when compared to seronegative patients (RR = 1.44, 95%CI: 1.02-2.04) (homogeneity test, $P < 0.0001$)^[58]. These results have been updated in 2014: ten observational studies involving 82690 unique RT recipients were included. In this study, HBsAg positive status was associated with an increase risk for all-cause mortality (adjusted RR = 2.214, 95%CI: 1.56-3.137, $P < 0.0001$) and for all-cause graft failure (aRR = 1.44, 95%CI: 1.26-1.63, $P < 0.0001$)^[59]. Both meta-analyses of observational studies concluded that untreated RT HBsAg positive patients have an reduced patient and graft survival.

As previously mentioned oral NUC therapy safely and effectively can suppress HBV replication in RT recipients. Several studies had shown that this antiviral effect may impact on long term graft and patients outcome^[31,54,60-62]. In 63 LAM treated HBsAg positive RT recipients 10-year survival rate was 81% and such results were nearly comparable to HBsAg

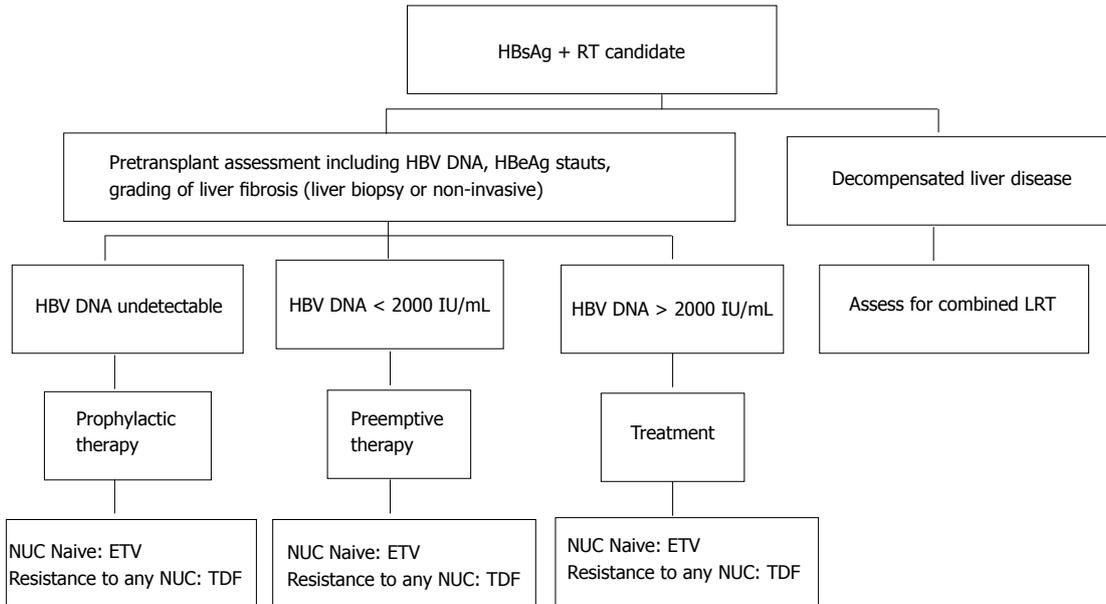


Figure 1 Treatment algorithm for management of renal transplant candidates with chronic hepatitis B virus infection. RT: Renal transplantation; LRT: Liver renal transplantation; NUC: Nucleos(t)ide analogue; ETV: Entecavir; TDF: Tenofovir disoproxil fumarate.

negative patients^[31]. Initiating treatment with LAM is associated with a 62% chance of developing drug resistance after 4 years of continuous therapy. Six months after beginning rescue therapy with ADV or ETV, HBV DNA decrease three-log in 75% of patients without significant adverse effects. When compared with untreated patients, those treated with NUCs showed a significant improvement in survival after 20 years of follow up (34% vs 83% respectively, $P < 0.006$). Even though NUC therapy reduced overall mortality by reducing liver related disease ($P < 0.036$), 40% of death in chronic HBV infected patients are still related to liver complications and 22.2% of them developed in patients being treated^[31].

Forty two RT patients were treated for long term with different NUCs regimens: at the end of follow up 18 patients were receiving monotherapy (9 LAM, 2 ADV, and ETV) and 24 combination therapy (11 LAM + ADV, 2 LAM + TDF, 4 ETV + ADV, 6 ETV + TDF and 1 TDF + emtricitabine, FTC)^[60]. At the end of the study 100% (18/18) of patients under monotherapy and 87.5% (21/24) of patients under combination therapy cleared HBV DNA. The 3 patients (12.5%) in the combination therapy group with detectable HBV DNA, had HBV DNA levels below 300 IU/mL. At the end of follow up, 92.8% of the entire cohort had cleared HBV DNA. Patient survival was 97.6% at 10 years, 95.2% at 15 years, and 90.4% at 20 years after renal transplantation, while graft survival was 100% at 5 years, 97.6% at 10 years, 95.2% at 15 years, and 88.1% at 20 years after renal transplantation. At the end of the study, 8 patients (19.04%) died and 1 received a liver transplantation due to end stage liver disease. Four

deaths were liver related: 4 patients (3 cirrhotics and 1 with only mild fibrosis at baseline) developed an hepatocellular carcinoma (HCC) despite complete virological response; three of the other patients died of non liver cancer and the remaining from stroke^[60]. During the study, 11.9% (6/42) of patients returned to dialysis due to chronic rejection leading to graft failure. The NUC dose was modified according to creatinine clearance in 45.2% (19/42) of the patients^[60].

Thirty RT patients underwent long term NUC therapy; at the end of follow up 25 were still alive and being treated and 24 were HBV DNA non detectable. Eight patients were receiving LAM monotherapy, 1 on ETV, 1 on TDF, 6 on LAM + ADV combination and 9 LAM + TDF^[61]. Five patients died from no liver related disease and 9 returned to dialysis after graft failure due to chronic allograft nephropathy. In this cohort, 10 year patient survival was 92% and 10 year graft survival was 86%. There were no renal adverse effects related to ADV/TDF therapy^[61].

Twenty five patients received pre-RT treatment with LAM (22 patients) and ETV (3 patients): 18 were HBV DNA undetectable (prophylactic group) and 8 were HBV DNA positive (preemptive group), and were compared to a historical control group^[54]. Unadjusted 10-year graft survival rates in the treatment cohort vs historical control cohort were 81.8% and 34.3%, respectively ($P = 0.003$). Graft lost occurred more frequently in the historical control than in the treated cohort (70.3% vs 4.3%, respectively); this was mainly related to patient death. Treated patients showed a better 10-year patient survival than the control group: 90.0% vs 57.4%; $P = 0.013$. Pre-transplantation NUC

Table 2 Definition of response to nucleos(t)ide analogue antiviral therapy of chronic hepatitis B

Category of response	
Biochemical (BR)	Decrease in serum ALT to within the normal range
Virologic (VR)	Decrease in serum HBV DNA to undetectable levels by PCR assays, and loss of HBeAg in patients who were initially HBeAg positive
Primary non-response	Decrease in serum HBV DNA by 2 log ₁₀ IU/mL after at least 24 wk of therapy
Virologic relapse	Increase in serum HBV DNA of 1 log ₁₀ IU/mL after discontinuation of treatment in at least two determinations more than 4 wk apart
Histologic (HR)	Decrease in histology activity index by at least 2 points and no worsening of fibrosis score compared to pre-treatment liver biopsy
Complete (CR)	Fulfill criteria of biochemical and virological response and loss of HBsAg
Time of assessment	
On-therapy	During therapy
Maintained	Persist throughout the course of treatment
End-of-treatment	At the end of a defined course of therapy
Off-therapy	After discontinuation of therapy
Sustained (SR-6)	6 mo after discontinuation of therapy
Sustained (SR-12)	12 mo after discontinuation of therapy

HBV: Hepatitis B; HBsAg: Hepatitis B surface antigen.

treatment was an independent factor for the improved patient survival [odds ratio (OR) = 0.052; $P = 0.005$]. Liver-related disease was the main cause of death in the historical control cohort (84.6% of the cases); sepsis was the second most frequent cause (15.4% of the cases)^[54]. Overall, graft (100% vs 71.4%) and patient survivals (100% vs 85.7%) were similar in the 2 treated cohorts ($P = 0.601$)^[54].

Only one study showed some conflicting results on NUC therapy impact in patients' survival^[62]. The study included 94 HBV-positive and 282 age/sex-matched HBV negative RT patients: 56 patients received an antiviral agent for prophylaxis (LAM 51, ETV 5), and other 18 for HBV reactivation. Although the patient survival rate was lower for HBV positive than HBV negative RTRs (89% vs 94% at 5 years, 78% vs 88% at 10 years, $P = 0.031$), there was no difference in graft survival between the two groups (86% vs 92% at 5 years, 73% vs 81% at 10 years, $P = 0.113$). In multivariate analysis, HBsAg positive status was a significant risk factor for death (OR = 2.19, 95%CI: 1.14-4.20, $P = 0.019$), but not significant for graft loss (OR = 1.64, 95%CI: 0.94-2.86, $P = 0.079$)^[62]. HBeAg and HBV DNA Pretransplant status was not available for all the patients. Of the 26 HBeAg-positive patients, 14 were receiving antiviral prophylaxis at transplantation: 8 showed reactivation while 6/12 of the untreated developed reactivation. All survived with stable liver chemistry, except for one dying from an HCC. Of 57 HBeAg-negative patients, 35 were started on antiviral prophylaxis at transplantation: 14 showed reactivation while 14/22 of the untreated developed reactivation. Among them, 12 died,

whereas the remaining 45 survived without hepatic dysfunction^[62]. Even though treated patients showed a reduced survival, it appears to be better than the survival reported in untreated patients.

NUC therapy in HBsAg positive RT patients is associated with a higher long term patient and graft survival rate. Studies have some limitations since most of the used LAM, which is not the best treatment option. More potent NUCs may add some benefit over LAM, but this still has to be demonstrated. Salvage therapy with TDF or ADV is safe and effective in patients developing LAM-R. Despite this clear benefit, all HBV infected patients must be closely follow up and HCC screening must be performed every six months, since the risk of HCC development may not entirely disappears even in the presence of virological response^[60,63].

DURATION OF THERAPY AND EVALUATION OF RESPONSE

In RT patients it is unclear what is the optimal treatment extent that assures long term viral suppression, preserving adequate liver function with the minimal risk of viral resistance development^[15]. Current guidelines clearly define how to monitor on treatment response, what are the therapeutic endpoints and when it is possible to stop treatment (Table 2)^[9,10]. In the case of NUC therapy, there are some terms regarding resistance that have also been defined. These is particularly important in this population, since some patients had initiated treatment long time ago with old NUCs such as LAM (Table 3)^[9,10].

Table 3 Definition of terms relating to antiviral resistance to nucleos(t)ide analogue treatment

Term	Definition
Virologic breakthrough	Increase in serum HBV DNA by > 1 log ₁₀ (10-fold) above nadir after achieving virologic response, during continued treatment
Viral rebound	Increase in serum HBV DNA to > 20000 IU/mL or above pretreatment level after achieving virologic response, during continued treatment
Biochemical breakthrough	Increase in ALT above upper limit of normal after achieving normalization, during continued treatment
Genotypic resistance	Detection of mutations that have been shown in "in vitro" studies to confer resistance to the NA that is being administered
Phenotypic resistance	<i>In vitro</i> confirmation that the mutation detected decreases susceptibility (as demonstrated by increase in inhibitory concentrations) to the NUC administered

NUC: Nucleos(t)ide analogue.

The duration of treatment depends on HBeAg status. HBeAg positive patients should be treated until HBV DNA and HBeAg are cleared and antiHBe seroconversion develops. Additional treatment, also known as "consolidation therapy", is needed for at least 6 to 12 mo after antiHBe seroconversion to prevent virological relapse. It is recommended to closely monitoring for relapse after treatment withdrawal. Relapse, even in patients achieving adequate virological response is a possibility, but their rates tend to be low^[64]. HBeAg negative patients should treated until HBsAg clearance is achieve^[9,10]. These recommendations might be applied to treatment in RT recipients to ensure treatment success, but outcomes after NUCs withdrawal in RT immunosuppressed patients is unknown.

A small recent study evaluated the long term results in HBV positive RT patients after NUC treatment discontinuation^[65]. Fourteen patients treated with LAM (11 patients), ADV (1 patient), ETV (1 patient), and LdT (1 patient) were included in this study. Patients were allowed to discontinue treatment if they have all of the following: (1) no clinical and histologic evidence of cirrhosis; (2) normal liver biochemistry; (3) negative for both HBV DNA and HBeAg; (4) no viral resistance; (5) antiviral therapy > 9 mo; (6) maintenance dosage of immunosuppressant for > 3 mo; and (7) no history of acute rejection during recent 6 mo^[65]. All patients were followed at 3 to 6 mo interval for liver biochemistry, viral serology, and HBV DNA level after treatment discontinuation. In 6 (42.9%) of 14 patients who meet the pre-specified criteria treatment was discontinued. In 4 of them (66.7%) it was successfully discontinued and HBV DNA was still undetectable for a median 60.5 mo (range, 47-82 mo). In the other 2 patients HBV reactivated, but HBV DNA was again cleared after immediately resuming NUC therapy^[65]. On the contrary, in LAM treatment discontinuation in 19 RT recipients after 2 years of treatment without adequate virological response, relapse rate was high (75%)^[66]. Even though evidence is scarce, it seems that in certain RT patients, after complete viral suppression

and sufficient duration, antiviral therapy can be successfully and safely withdrawn.

IMPACT OF NUCLEOS(T)IDE ANALOGUES ON RENAL FUNCTION

Nucleos(t)ide analogues are primarily eliminated without changes in the urine following ingestion, and appropriate dose modifications are proposed for patients with impaired renal function (eGFR < 50 mL/min) (Table 1). Treatment guidelines recommend that all patients initiating NUC treatment should be tested for serum creatinine levels and estimated creatinine clearance before therapy; and baseline renal risk should be assessed for all of them^[9,10]. High baseline renal risk includes one or more of the following clinical situations: decompensated cirrhosis, creatinine clearance < 60 mL/min, poorly controlled hypertension, proteinuria, uncontrolled diabetes, active glomerulonephritis, concomitant nephrotoxic medications and solid organ transplantation. In consequence, RT recipients may have many of these basal renal risk factors.

In clinical trials outside renal transplant setting, minimal decline in renal function have been showed with all NUCs, except for LdT which appears to improve renal function^[67,68]. Impact of LdT on renal function was analyzed from a database including all patients treated in the GLOBE Study (2 years), in the long term extension study CN04E1 (4 to 6 years) and in patients with decompensated cirrhosis (2 years)^[69]. Renal function improved in LdT treated patients in GLOBE trial (+8.5% increase in mean eGFR,) and it was sustained for 4 to 6 years. Improvement in renal function in LdT treated patients was also observed in those at increased risk for renal impairment: patients with baseline eGFRs of 60-89 mL/min per 1.73 m² (+17.2%), > 50 years (+11.4%), and with advanced liver fibrosis or cirrhosis (+7.2% for patients with Ishak fibrosis score 5-6). In patients with the highest renal risk such as decompensated cirrhotics, eGFR was also improved with LdT (+2.0%). In patients who received 2 years of LAM in GLOBE/015 studies and

rolled over to extension study to receive LdT for 2 additional years, eGFR also improved after treatment switch (+8.9%)^[69]. Although this data may suggest that LdT may be renal protective, it is not clear whether this protective effect is specific to this NUC. This potential benefit, particularly relevant in the RT population, does not overcome the high risk of treatment resistance and neuromuscular adverse events. As previously mentioned, this beneficial safety profile does not support the use of LdT as a first-line NUC in hepatitis B treatment^[69].

Nucleotide, specially ADV, appear to be more nephrotoxic than nucleoside analogues^[70-74]. In a real-life setting study, 145 patients ADV treated patients were compared with 145 untreated patients regarding its impact on renal function^[71]. During follow-up, 30% of ADV treated patients show a mild decrease in renal function (10%-20% reduction in eGFR from baseline) compared with 16% in the untreated group, 15% vs 6% showed a moderate decrease (20%-30%), and 7% vs 1% showed a severe decrease (> 30%) respectively ($P > 0.0001$). In the ADV group 6.9% of the patients discontinued treatment ($P > 0.004$). In a multivariate analysis ADV treatment significantly predicts renal dysfunction [hazard ratio (HR) = 3.94, $P = 0.03$]. In the same analysis, age > 50 years (HR = 3.49, $p = 0.087$), baseline mild renal dysfunction (HR = 4.49, $P = 0.073$), and hypertension and/or diabetes mellitus (HR = 2.36, $P = 0.074$) were not significant predictors^[71]. In a retrospective study, 687 patients receiving ADV monotherapy (18.2%) or in combination with LAM (81.8%) for 1 year or more were enrolled to evaluate the incidence and risk factors of renal impairment^[72]. Renal dysfunction was defined as mild (20%-30% reduction in eGFR), moderate (30%-50%), or severe (more than 50%). Patients were treated for a median of 27 mo, 10.5% ($n = 72$) developed renal dysfunction being mild in 77.8% of patients, moderate in 20.8% of patients, and severe in only 1 patient. The cumulative incidence of renal dysfunction at 1, 3, and 5 years was 2.6%, 14.8%, and 34.7%, respectively. ADV dose was modified in 7 patients and it was discontinued in 3 patients; after these changes, eGFR remained stable^[72]. In 271 ADV treated patients, after 6 years of treatment GFR ≤ 60 mL/min incidence was 38.3% and after 5 years, serum creatinine increased ≥ 0.5 mg/dL in 21.48%. Switching ADV to other NUC or reducing its dose was associated with reversal of renal dysfunction in almost all patients; there were no differences between the two approaches ($P = 0.737$)^[73]. On the contrary, a study including 46 HBeAg negative LAM-R patients treated with ADV add on for up to 90 mo found no impact on renal function when compared with a matched control group of untreated inactive HBV carriers^[74].

The number of patients treated with ADV in the

RT setting is smaller than in the general population. In this subgroup, ADV treatment may also impact on renal function. A significant decrease of estimated GFR and an increase in serum creatinine from 1.42 (± 0.39) to 1.6 (± 0.36) mg/dL, ($P = 0.02$) was found in 11 patients treated for 2 or more years^[37]. It was also associated with an increase in proteinuria, changes in renal tubular parameters and changes in phospho-calcic metabolism^[37]. Another study including also 11 LAM-R patients did not show significant changes in median creatinine clearance (CLcr), in serum phosphorus or in urinary protein level from baseline to the last available visit. However, after a median treatment time of 11 mo (range: 9-42), 54% ($n = 6$) of patients reduce ADV dose due to renal dysfunction. Renal function remained stable ($n = 5$) or improved ($n = 1$) 22 mo (range: 6-34) after dose modification^[38].

Fourteen patients were treated with long term ADV (5 monotherapy, and 9 ADV + LAM combination therapy). Eight patients (57.2%) developed impaired renal function; it was mild (5%-20% reduction in the eGFR compared to baseline values) in 4, moderate (20%-30%) in 2, and severe (> 30%) in the 2 remaining patients. Acute graft rejection was diagnosed by kidney biopsy in 2 of these patients. Calcineurin inhibitors nephrotoxicity was presumed in 2 of these patients and their doses were accordingly adjusted. ADV dose was reduced in 3 patients due to severe renal dysfunction (eGFR 30-50 mL/min) and it was discontinued in 1 patient (eGFR < 20 mL/min) without impact on virological response^[39]. Renal dysfunction in long-term ADV treated patients appears relatively frequent, but serious nephrotoxicity is unusual. Renal dysfunction can be safely managed by dose reduction or switching to another NUC without impact on virological response.

In TDF treated patients, also a nucleotide analogue, renal dysfunction is less frequently seen than with ADV. The majority of previously nephrotoxic events reported, which were similar to those reported under ADV treatment, were in HIV infected patients^[68]. There is recently presented data about TDF impact on renal function in HBV mono-infected patients^[75-80]. A study evaluated the pooled results from three global randomized clinical trials including 426 TDF treated patients for 144 wk. In this study 0.5% (2/426) of patients developed a creatinine increase ≥ 0.5 mg/dL from pre-treatment values and none showed an eGFR decrease < 50 mL/min, showing a minimal impact of TDF on renal function even in high risk patients such as cirrhotics or diabetics^[75]. Moreover, when 74 patients with mild renal dysfunction (CrCl 50-80 mL/min) were compared with 206 with normal renal function (CrCl ≥ 80 mL/min), none of them showed signs of renal impairment defined as a creatinine increase ≥ 0.5 mg/dL after 96 wk of therapy^[76]. Among 441 patients from the Vireal cohort, 114 with baseline impaired

renal function were classified as stage 2 (GFR 60-89 mL/min), stage 3 (GFR 30-59 mL/min), stage 4 (GFR 15-29 mL/min) and stage 5 (GFR < 15 mL/min or dialysis) and included in the study. When compared from baseline, after 48 wk of treatment, TDF did not significantly modified GFR in patients with stage 2 (76 mL/min vs 77 mL/min), 3 (50 mL/min vs 49 mL/min), or 4 (23 mL/min vs 23 mL/min) renal failure^[77]. Two RT recipients were included and had a stable GFR under therapy. Nine patients needed to adjust TDF dose. At the end of the study, 67% had a stable renal failure stage, 22% had an improvement and 11% had a decreased in it^[77]. In a retrospective study, 195 refractory patients were treated with TDF monotherapy for 30 ± 16 (6-90) mo were compared with 89 asymptomatic HBsAg carriers^[78]. After 48 mo, TDF treated patients in showed a significantly greater reduction in eGFR when compared to untreated patients [-16 ± 36 (-48 - +23) and -9.6 ± 36 (-21 - +22) mL/min, respectively, $P = 0.03$]. TDF dose was reduced in only 1 patient after 15 mo of treatment due to a 0.38 mg/dL increase in creatinine levels^[78]. In 26 LAM-R patients treated with TDF, there were no significant variations in phosphatemia and GFR from baseline after one year of treatment^[79]. Even if there is no impairment on renal function, TDF may have some potential effects on the proximal tubule. In 61 TDF treated patients for a mean time of 29 mo, there were no significant change in mean GFR in the overall population but 58% of patients showed an impairment in GFR (median 8.1%, range 0.01% to 20.5%) and two patients developed an GFR to < 60 mL/min^[80]. At least one sign of proximal tubular damage appeared in 26 (42%) individuals: glucosuria without diabetes mellitus, increased alpha1-microglobulinuria/creatinine ratio, hypophosphatemia, reduced tubular resorption of phosphate rate and reduced tubular maximum reabsorption rate^[80]. The effects of TDF on renal function were evaluated in 321 naïve patients treated for 4 years in clinical practice^[81]. In this large European cohort, there were no modifications in creatinine and phosphorus serum levels and eGFR was reduced from 84 to 80 mL/min. At year 4, patients with eGFR < 50 increased from 2% to 3% and those with eGFR < 60 mL/min increased from 7% to 11%. At the same time point, hypophosphatemia (serum phosphate < 2.3 mg/dL) increased from 2% to 5.1 %, while 1% of the patients had phosphate levels < 2.0 along the study period. TDF dose was reduced in 17 patients due to reduction in eGFR and in 2 due to hypophosphatemia. Seven patients had to withdraw treatment and were switched to ETV. Overall, some renal adverse effect was reported in 26 patients (7%)^[81].

In comparison to nucleotide analogues, nucleoside analogues, such as ETV and LdT, show not significant renal toxicity^[67,68]. Studies have been performed

comparing ETV and TDF nephrotoxicity^[82-86]. After 2 years of treatment, there was no significant modifications in eGFR in 74 ETV and 50 TDF ± LAM treated patients^[82]. In the ETV group 2.7% showed a reduction $\geq 40\%$ in eGFR vs 3.92% in the TDF ± LMV group ($P = NS$). When compared with an untreated control group, in ETV treated patients eGFR was reduced by -7.6 mL/min (95%CI: -15.8+0.6, $P = 0.07$) and by -8.7 mL/min (95%CI: -18.3+1.0, $P = 0.08$) in TDF ± LMV treated patients. In untreated controls, eGFR remained stable or even improved by +7.4 mL/min (95%CI: 0.78-14.1, $P = 0.03$)^[82]. In another real-life cohort of 212 patients were treated with TDF and 79 with ETV and its impact on renal function was evaluated^[83]. No significant differences were found in urea, creatinine and phosphorus levels and in eGFR after 12 mo of TDF treatment. Also in the same group, there was no difference in the proportion of patients with eGFR < 60 mL/min when compared with baseline levels. In ETV treated patients, there was a significantly reduction in serum phosphorus (0.96 vs 1.06, $P = 0.016$), increased in creatinine levels (1.0 vs 0.89, $P < 0.05$) and reduction in eGFR (80 vs 89, $P < 0.05$) after 12 mo of treatment. In ETV treated patients, 3.8% of patients had a 25% increase in creatinine levels while 0.47% of TDF treated patients had a 25% decrease in eGFR after 12 mo of treatment^[83]. In a community-based retrospective cohort study, 80 patients treated with TDF monotherapy or in combination with other NUCs were matched with 80 ETV treated patients and incidences of serum creatinine increments and eGFR decrease were evaluated^[84,85]. More patients in the ETV group had creatinine increments ≥ 0.5 mg/dL (3 vs 11; $P = 0.025$), whereas more patients treated with TDF had eGFR reductions of < 60 mL/min (15 vs 6; $P = 0.022$) and at least 1 dose modification (13 vs 4; $P = 0.021$). In a multivariate analysis, previous organ transplantation (aOR, 6.740; 95%CI: 1.799-28.250; $P = 0.005$) and pre-treatment renal failure (aOR, 10.960; 95%CI: 2.419-48.850; $P = 0.002$) were significantly associated with increases in serum creatinine levels^[84]. Renal function was evaluated in 197 HBV mono-infected patients from two outpatient clinics and who were classified according to the received treatment: LAM ($n = 36$), ADV ($n = 32$), ETV ($n = 32$), TDF ($n = 37$), and untreated HBsAg-positive patients ($n = 60$)^[86]. The CKD-EPI equation was used to calculate eGFR and the individual change in eGFR over time was modeled with linear mixed effects models. Patients with previous renal dysfunction, diabetes mellitus, or arterial hypertension were excluded from the analysis. The yearly predicted median individual changes in eGFR according to this model were: -2.05 mL/min in untreated patients, and -0.92 mL/min, -1.02 mL/min, -1.00 mL/min, and -0.92 mL/min in LAM, ADV, ETV and TDF treated patients,

respectively. A decrease of eGFR > 20 mL/min from baseline developed in 3.3% of untreated patients, and in 5.5%, 0%, 6.25%, and 2.7% in LAM, ADV, ETV and TDF treated patients, respectively. Renal insufficiency stage 3 (eGFR of < 60 mL/min) was uncommon and not different between all patient groups^[86].

First line NUCs, ETV and TDF, appears to have little impact on renal function in the general population when compared with untreated controls and with the others NUCs. Markers of renal function indicated that TDF treated patients, suspected to be more nephrotoxic, have similar risks of developing changes in renal function than ETV treated patients. Although there is some evidence showing some degree of renal dysfunction in ETV treated patients, its clinical significance remains unclear and it may represent a physiological decrease in renal function in this group and/or reflect the potential limitations of standard biochemical tests of renal function in patients with liver disease^[83]. Baseline renal risk factors may play a role in the nephrotoxic effects of NUCs. Data on RT patients is limited but these results can be extrapolated to this population, taking into account that these RT recipients can be considered within the high renal risk group. Therefore, it is recommended in all HBV treated patients to measure serum creatinine levels and estimated creatinine clearance, and in ADV or TDF treated patients it is also recommended to measure serum phosphate levels, especially in patients at high renal risk. In patients at low renal risk these tests can be performed every 3 mo during the first year and every 6 mo thereafter, in case of no renal adverse events. In patients at high renal risk these tests can be performed every month for the first 3 mo, every 3 mo until the end of the first year and every 6 mo thereafter, in case of no renal adverse events. Closer renal monitoring is required in patients who develop reductions in creatinine clearance < 60 mL/min or reductions in serum phosphate levels < 2 mg/dL^[9,10,15,87].

CONCLUSION

Current guidelines clearly define who needs treatment, when to start, what is the first line therapy, how to monitor treatment response, when to stop, and how patients must be controlled for its safety. There is some data showing a favorable safety and efficacy profile of NUC treatment in the renal transplant setting. ETV, an agent without signs of major nephrotoxicity, appears to be the best option for NUC naïve patients and TDF is still the preferred agent in patients with resistance to LAM or any other NUC. Renal transplant recipients under antiHBV treatment should be closely monitored for its efficacy against HBV and for its safety, especially regarding its impact on renal function. Studies including a large

number of patients with long term treatment and follow up are still needed to better demonstrate the safety and efficacy of newer NUCs in this population.

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Recent advances in dietary supplementation, in treating non-alcoholic fatty liver disease

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cirrhosis, and failure or hepatocellular carcinoma. Since NAFLD is positively associated with the development of obesity, insulin resistance, and ultimately type 2 diabetes mellitus, it is often regarded as the hepatic manifestation of the metabolic syndrome. No pharmacologic treatment has yet been proven for this disease. For most patients with presumed or confirmed NAFLD, the only proven strategy is to offer lifestyle advice that can lead to sustained weight loss. Since insulin resistance, oxidative stress, inflammation, and necro-apoptosis are involved in NAFLD pathogenesis, it seems that every potential therapeutic agent should target one or some of these pathologic events. There are many well known anti-oxidants, anti-inflammatory, and insulin sensitizer dietary supplements which have shown beneficial effects on NAFLD improvement in animal and human studies. The purpose of this review is to explore the existing evidences on dietary supplements considered to have hepatoprotective properties, and to present some proposed mechanisms by which they may protect against NAFLD.

Key words: Nonalcoholic fatty liver disease; Dietary supplementation; Treatment

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Core tip: This review explores the existing evidences on dietary supplements considered to have anti-oxidant, anti-inflammatory, and/or insulin sensitizer properties, and their role in management of nonalcoholic fatty liver disease while addressing some of their proposed mechanism of action.

Abstract

Nonalcoholic fatty liver disease (NAFLD) is currently known as the most common liver problem, characterized by excessive lipid accumulation in hepatocytes, which may progress to other liver diseases such as nonalcoholic steatohepatitis, hepatic tissue fibrosis, liver

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INTRODUCTION

Non-alcoholic fatty liver disease (NAFLD) encompasses a range of conditions caused by fatty infiltration of the hepatocytes without significant amounts of alcohol use, that can be originated from multiple factors^[1]. NAFLD begins with simple hepatocyte steatosis, and progresses to nonalcoholic steatohepatitis (NASH), fibrosis of the hepatocytes, and liver cirrhosis, which can further progress to hepatocellular carcinoma (HCC)^[2]. Patients with NAFLD are usually asymptomatic and are diagnosed accidentally through routine checkup exams. Currently liver biopsies are considered gold standard for the diagnosis and staging of NASH, since there are no specific symptoms to differentiate between this disease and other liver disorders. Magnetic resonance spectroscopy (H¹-MRS) and Fibroscan are noninvasive modalities for diagnosis and staging, assessing a larger section of the liver in comparison to liver biopsy^[3,4]. Other clinical diagnostic indices such as increased serum levels of aspartate aminotransferase (AST) and alanine aminotransferase (ALT), as well as evidence of liver steatosis in ultrasonography are also routinely used^[5]. Developing effective therapies with minimal side-effects against NAFLD is critical for controlling the progression of this disease to end-stage liver disorders^[6].

EPIDEMIOLOGY

NAFLD is the most common diagnosis in subjects with altered aminotransferases in the Western world^[7], where one third of the population is affected^[8]. In Asia, recent reports showed a similar prevalence of NAFLD^[9,10]. About 20%-25% of adults with NASH have been reported to develop liver cirrhosis^[11]. About 30% to 40% of patients who develop cirrhosis secondary to NAFLD, will die of liver-related problems^[12]. The prevalence of NAFLD is different among men and women, and it increases with age, occurring in less than 20% of individuals younger than 20 years of age, and in more than 40% of those over the age of 60^[13]. NAFLD has also been identified in the pediatric population, prevailing at 2.6%, although it is estimated that its prevalence will increase to 22.5%-58.5% in obese children^[14]. Parallel to the rising prevalence of conditions such as obesity and type 2 diabetes mellitus (T2DM), the rate and prevalence of NAFLD is also increasing^[15]. NAFLD affects 40%-75% of patients with T2DM, 33%-76% of obese and 90% of morbidly obese people^[16].

PATHOGENESIS

The pathogenesis of NAFLD is complicated and while its exact mechanism remains largely unknown, different genetic factors and/or environmental elements seem to influence it^[12,17]. The "two-hit

hypothesis" of NASH, originally explained by Day and James suggests that lipid deposition in the liver (first hit) is followed by a series of other, oxidative and hepatotoxic processes (second hit), caused by a mechanism currently not known^[18]. Several factors such as genetics, epigenetic mechanisms, as well as environmental elements, appear to promote hepatocyte fat deposition and insulin resistance, both of which further lead to the secondary pathologic events^[19], such as oxidative stress, lipid peroxidation, increased inflammatory responses, hepatic fibrosis and apoptosis^[20]. Other triggers such as lipotoxicity, endotoxemia, and adipocytokines or other inflammatory signals released from fat-infiltrated hepatocytes and adipose tissue, may promote oxidative stress in the liver, inducing the progression of NAFLD to NASH^[21,22].

MANAGEMENT OF NAFLD

Currently, the only proven strategy for NAFLD management is lifestyle modification techniques such as weight loss through diet and exercise. Since obesity strongly influences the development of NAFLD, weight loss is again the main objective in NAFLD management, and the first-line therapy. All NAFLD patients are encouraged to follow a low caloric diet, increase their physical activity, and stop smoking (if applicable)^[11,23,24]. Moreover, a wide range of drugs and supplements, including antioxidants, anti-inflammations, insulin sensitizers, and lipid lowering agents, have been evaluated in patients and experimental models of NAFLD, however none of them have shown long term efficacy^[25,26].

In the recent years, however, the beneficial effects of dietary supplements on NAFLD progression have received increasing attention since these substances have several advantages such as being widely available, while having low or minimal side effects^[27]. In the present review, we mainly focus on the recent advances of dietary supplements in NAFLD amelioration.

RECENT FINDINGS FROM DIETARY SUPPLEMENTATION IN THE TREATMENT OF NAFLD

Antioxidants agents

Vitamin E and vitamin C: Since oxidative stress is one of the factors involved in the pathogenesis of NAFLD, it was thought that antioxidant agents could be beneficial in its treatment. Thus, many clinical trials have evaluated the effects of vitamin E and/or vitamin C, as the main dietary sources of antioxidants to treat NAFLD. Nobili *et al.*^[28] have shown that vitamin E supplementation does not provide a greater benefit for NAFLD treatment, than diet and physical exercise^[28]. Akcam *et al.*^[29] have reported that

metformin is more efficacious in reducing metabolic parameters such as insulin resistance, fasting insulin and lipid levels, than dietary advice and vitamin E use in obese patients with NAFLD. A clinical trial using atorvastatin and vitamins E + C vs placebo, showed improved hepatic steatosis on computed tomography scans. It was not however detected whether this improvement was due to the combination treatment or a single compound alone^[30]. The TONIC randomized controlled trial showed that neither vitamin E nor metformin are superior to placebo in sustaining a reduction in ALT levels of pediatric NAFLD patients^[31]. A recent review article concluded that vitamin E is only recommended in adults with NASH who do not have diabetes or cirrhosis, or an aggressive histology^[32]. In a meta-analysis, adjuvant vitamin E was not shown to have a significant effect on normalizing serum ALT levels. Using higher doses of vitamin E, a longer duration of therapy or adding vitamin C did not alter the effect of these antioxidants on the measured outcomes either^[33]. There seems to be lacking evidence on the long-term effects of vitamin E use on histological improvements of NAFLD patients, which calls for larger, well-designed randomized controlled trials (RCTs) with histological endpoints, to really determine the efficacy of its use.

Resveratrol: Resveratrol (3,5,4-trihydroxystilbene) is a natural phenol produced by certain plants and found in the skin of red grapes. Resveratrol has been widely accepted as a chemopreventive agent that exerts other positive health effects as well because of its ability to take part in many biological activities. Resveratrol is thought to have antioxidative, anti-inflammatory, anti-cancer, anti-obesity, anti-diabetic, and anti-aging properties. Its positive effects on animal NAFLD models have been shown in several studies. In different studies, Resveratrol decreased NAFLD severity in animal models in the following ways: through TNF-alpha inhibition and antioxidant activities^[34], through the activation of AMPK^[35,36], by induction of skeletal muscle SIRT1 and SIRT4 expression^[37], by increasing the number of mitochondria, and specially, by increasing hepatic uncoupling protein 2 expression^[38], decreasing hepatic LDL receptor and SR-BI mRNA and protein expressions^[39], and the reduction of nuclear factor-kappaB (NF-kappaB) activity^[40].

Clinical trials evaluating the effects of Resveratrol supplementation on NAFLD characteristics are scarce. A recent study, administering Resveratrol vs placebo for eight weeks, not only failed to show any significantly improvements in any NAFLD features in the Resveratrol group, it also showed an increase in hepatic stress, based on increased liver enzyme levels^[41]. A different trial however, did find a significant improvement in NAFLD characteristics after 12 wk of supplementation with 500 mg Resveratrol^[42]. It appears that the dose and duration

of Resveratrol administration is important in its efficacy. Future clinical studies with different dosages and durations are needed to clarify the true impact of Resveratrol treatment in NAFLD/NASH patients^[43].

Anthocyanin: Anthocyanins (ACNs) are water-soluble bioactive compounds of the polyphenol class that are present in many plant based products. It has been reported that ACNs decrease hepatic lipid accumulation and may counteract oxidative stress and hepatic inflammation in animal studies, but their benefits in patients with NAFLD has not yet been well elucidated^[44]. There is only one study evaluating the effects of ACN on NAFLD patients; Suda *et al*^[45] have reported that supplementation with 400 mg of acylated ACNs could reduce levels of liver enzymes, in particular gamma-glutamyl transferases in patients with NAFLD. This clinical trial had many limitations; liver damage was not directly assessed, fatty liver was not confirmed by direct imaging, and the effect of acylated ACNs was not compared to that of a control food or to the lack of intervention^[45]. More research studies are therefore required to evaluate the effects of ACNs supplementation on NAFLD features.

Green tea extract: It has been shown that the main important green tea polyphenol, epigallocatechin-3-gallate (EGCG), has a positive therapeutic effect on obesity, features of metabolic syndrome, and liver steatosis in mice^[46]. In experimental models of NAFLD, EGCG supplementation significantly decreased weight gain, total and visceral body fat, insulin resistance, liver steatosis, serum cholesterol, and monocyte chemoattractant protein concentrations^[46].

Both *in vitro* and *in vivo* experiments have revealed that green tea and EGCG could prevent steatosis by reducing dietary absorption of lipids and carbohydrates, and by the inhibition of adipose tissue breakdown, and *de novo* lipogenesis in both hepatic and adipose tissues, through the stimulation of β -oxidation and thermogenesis in the liver, and by improving insulin sensitivity. Furthermore, EGCG may inhibit the development of steatohepatitis from fatty liver disease, through its antioxidant and anti-inflammatory characteristics^[47]. Currently, there are no randomized, controlled trials in humans, evaluating the effects of green tea on NAFLD. These studies are needed to provide enough evidence that green tea can effectively prevent the development and/or progression of NAFLD^[48].

Coffee: Both epidemiological and animal studies have shown that drinking coffee on a regular basis can decrease the risk of T2DM development^[49-51]. A recent case-control study comparing coffee vs non-coffee drinkers showed that fatty liver occurred less frequently in coffee drinkers, and that

drinking coffee was inversely associated with the degree of liver brightness, as well as obesity and insulin resistance^[52]. Among NASH patients, coffee consumption has been shown to be significantly associated with a reduced risk of fibrosis^[53].

More research is needed to determine the protective properties of caffeine against NAFLD. Coffee contains certain phytochemicals with potential antioxidant properties, which may be protective against cardiovascular and liver diseases, and malignancies. The anti-oxidative, anti-inflammatory, and anti-fibrotic properties of coffee might explain its hepatoprotective effects in NAFLD^[54,55].

Garlic: Garlic-derived S-allylmercaptocysteine (SAMC) has a therapeutic role in diabetes and non-alcoholic fatty liver disease due to its properties in the regulation of lipogenesis and glucose metabolism^[56]. Results of two studies show that SAMC decreases the liver injury caused by NAFLD, while decreasing fat build-up, and collagen formation. This may occur because SAMC takes part in different activities at the molecular level that affect NAFLD, by for example decreasing lipogenesis and restoring lipolysis markers. The expression of pro-fibrogenic factors is also reduced by SAMC, as well as oxidative stress in the liver, by means of cytochrome P450 2E1-dependent pathway inhibition. SAMC may partially prevent NAFLD-induced inflammation as well, by reducing pro-inflammatory mediators, chemokines and suppressor of cytokine signaling. The protective effects of SAMC are also partly shown through its ability to restore the altered phosphorylation status of FFAs-dependent MAP kinase pathways, and to diminish the activity of nuclear transcription factors such as NF-kappaB and AP-1, while reducing apoptosis and enhancing autophagy during NAFLD development^[57,58].

In addition, garlic essential oil (GEO) and its major organosulfur component diallyl disulfide (DADS), also have therapeutic effects on the development of NAFLD. They exert anti-obesity and anti-hyperlipidemic effects by reducing weight gain, adipose tissue weight, and serum lipid parameters. They significantly decrease the release of pro-inflammatory cytokines in the serum, while at the same time elevating in the hepatic antioxidant capacity by inhibiting cytochrome P450 2E1 expression during NAFLD development. The anti-NAFLD effects of GEO and DADS are mediated through the down-regulation of sterol regulatory element binding protein-1c, acetyl-CoA carboxylase, fatty acid synthase, and 3-hydroxy-3-methylglutaryl-coenzyme^[59]. Clinical trials are needed to confirm these experimental studies.

Ginger: Several mechanisms have been proposed by which ginger may prevent NAFLD or slow its progression to other liver diseases, such as incr-

easing insulin sensitivity, inducing the activation of peroxisome proliferator-activated receptor gamma, which in turn induces adiponectin and down-regulates pro-inflammatory cytokines, changing the balance between adiponectin and tumor necrosis factor-alpha in favor of adiponectin, promoting considerable antioxidant effects and antidyslipidemic properties, and reducing hepatic triglyceride content which can prevent steatosis. These mechanisms indicate that ginger possesses interesting potentials for serving as a natural supplement for the prevention and treatment of NAFLD^[60]. It might suppress fructose-stimulated overexpression of carbohydrate response element-binding protein (ChREBP) at the mRNA and protein levels in hepatocytes, which results in down regulation of the ChREBP-targeted lipogenic genes responsible for fatty acid biosynthesis, while expression of neither peroxisome proliferator-activated receptor- (PPAR-) alpha and its downstream genes, nor PPAR-gamma and sterol regulatory element-binding protein 1c is altered^[61]. Randomized clinical trials are needed to confirm these effects in patients with NAFLD.

Anti-inflammatory agents

Polyunsaturated fatty acids and monounsaturated fatty acids supplementation: Polyunsaturated fatty acids (PUFAs), especially n-3 PUFAs, are used to promote weight loss, and to reduce hepatic triglyceride accumulation, while improving insulin sensitivity and reducing steatosis, and hepatic damage in patients with NAFLD^[62-64]. They are also thought to exert anti-inflammatory effects^[65]. N-3 fatty acids affect lipid metabolism by mediating genomic pathways and regulating the transcription of genes involved in lipid metabolism^[66]. They improve insulin sensitivity by decreasing hepatic TNF α expression, repress fatty acid synthesis by negatively controlling sterol regulatory element binding protein-1c (SREBP-1c), and enhance fatty acid oxidation by positively controlling peroxisome proliferator-activated receptor- α (PPAR α)^[67,68]. Several studies support the protective effects of n-3 PUFAs in NAFLD. Among these, Capanni *et al*^[69] investigated the effects of n-3 PUFA supplementation (1 g/d for 12 mo) in 56 NAFLD patients. Their results indicated that n-3 PUFAs improved NAFLD characteristics such as ALT, AST, GGT, triglyceride and fasting glucose concentrations. Another clinical trial conducted in 23 patients with NASH found the same results^[70]. A recent systematic review reported a beneficial effect of omega-3 supplementation on hepatic fat content and AST, although the effect size was relatively small^[71]. The optimal dose and duration of this therapy need to be addressed in larger clinical trials in the future. Data on omega-6 fatty acids are very limited and mainly restricted to animal models.

Moreover, dietary monounsaturated fatty

acids (MUFAs) may prevent the development of NAFLD by reducing the oxidization of low-density lipoprotein (LDL), serum concentrations of LDL and total cholesterol (TC) and triacylglycerols, while decreasing body fat accumulation and postprandial adiponectin expression. It is shown that the replacing dietary carbohydrate and saturated fat consumption with MUFAs, reduces the blood pressure and glucose concentrations, and increases serum high-density lipoprotein (HDL) levels^[72]. The probable mechanisms for the beneficial effects of MUFA on liver fat content may be related to their roles in the regulation of insulin sensitizing gene expression^[7], and in the reduction of inflammation^[73], as well as to their inhibitory effects on nuclear factor- κ B (NF- κ B)^[74]. In a study, MUFA decreased the expression of hepatic lipogenesis and gluconeogenesis genes and SREBP in fatty rats^[75]. Further investigations are warranted to ascertain the role of MUFA on NAFLD.

Vitamin D: Evidence supporting the immunoregulatory roles of vitamin D continues to increase. Recent studies have indicated that deficiencies in vitamin D can result in insulin resistance, metabolic syndrome, and NAFLD^[76]. In one study, rats who were fed a western diet along with vitamin D depletion had significantly more steatosis, lobular inflammation, and NAFLD activity scores in comparison to animals with sufficient vitamin D intakes^[77]. In humans, vitamin D deficiency has been correlated with a more severe NAFLD activity score and hepatic fibrosis^[78], perhaps owing to the greater oxidative stress resulting from vitamin D deficiency^[79]. Hepatic expression of vitamin D receptors, CYP2R1 and CYP 27A1, negatively correlates with the severity of steatosis, inflammation, and NAFLD scores in patients with this disease^[80]. A recent study found a significant association between NAFLD and low serum vitamin D levels^[81]; this relationship remained significant even after adjustments were made for the presence of other metabolic syndrome features. Evidence from liver biopsies have shown that serum vitamin D levels are significantly related with the stage of hepatic fibrosis^[82]. Clinical trials have not yet been published to evaluate the effect of vitamin D supplementation on NAFLD characteristics.

Probiotics, prebiotics and symbiotic: It is known that the liver is susceptible to the exposure of intestine-derived bacterial products because of a close anatomic and functional connection between the intestinal lumen and the liver through the portal system^[83,84]. The gut-liver axis is an important pathway in NAFLD development, which is associated with small intestinal bacterial overgrowth and increased intestinal permeability^[85,86]. The contribution of microflora in NAFLD progression is mainly based on increased oxidative stress in the liver, which is caused by the increased ethanol and

lipopolysaccharide production in the intestine, further causing the release of inflammatory cytokines^[87,88].

Probiotics are live microorganisms that are beneficial to human health when ingested^[88]. The therapeutic effects of probiotics have been demonstrated in several animal models of NAFLD^[86,89-91]; however clinical trials are scarce^[92-95]. In a recent double blind, placebo controlled, clinical trial, we found that 28 wk of synbiotic supplementation can significantly decrease liver enzymes, inflammatory cytokines, NF- κ B activity, and fibrosis scores so that this supplementation in addition to lifestyle modification was significantly superior to lifestyle modification alone; whether these effects will sustain with longer treatment durations remains to be determined^[89].

Insulin sensitizers and lipid lowering agents

Cinnamon: Cinnamon might play a potential role in the reduction of post-prandial intestinal glucose absorption through the inhibition of pancreatic enzymes such as α -amylase and α -glucosidase, and by stimulating cellular glucose uptake by membrane translocation of glucose transporter-4, which stimulates insulin release, glucose metabolism, glycogen synthesis, and inhibits gluconeogenesis. These actions may ameliorate fasting blood glucose, LDL, and hemoglobin A1c, and might increase HDL cholesterol and insulin concentrations^[96].

Since one of the most important therapeutic strategies for NAFLD is modulating insulin resistance and oxidative stress, we thought that cinnamon could have beneficial effects on NAFLD features too. Thus, we investigated this hypothesis in a double-blind, placebo-controlled trial, and found that 12 wk of cinnamon supplementation significantly decreases HOMA (Homeostatic Model Assessment) index, FBS (fasting blood glucose), total cholesterol, triglyceride, liver enzymes, and high-sensitivity C-reactive protein in patients with NAFLD, however we did not find any significant changes in serum HDL levels^[97]. Further clinical trials of longer durations are recommended to elucidate the exact effects of cinnamon on NAFLD characteristics.

Curcumin: It has shown that curcumin can reduce serum lipid levels, and liver steatosis. Furthermore, it may prevent fatty liver progression to steatohepatitis due to its potent antioxidant and anti-inflammatory activities^[98,99]. Curcumin can reduce the expression of lipogenic genes in the liver and inflammatory responses of adipose tissue^[100], while enhancing the antioxidant defense system, attenuating mitochondrial dysfunction and inhibiting apoptosis^[101,102]. We did not find any clinical trial evaluating the effect of curcumin in patients with NAFLD.

Quercetin: Quercetin, a plant-derived bioflavonoid, has been reported to provide an improved health

status to its consumers, particularly with regard to obesity and diabetes^[103]. Studies have demonstrated that quercetin can modestly reduce weight and regulate the expression of genes related to *in vitro* adipogenesis^[103,104]. Quercetin reduces inflammatory cytokine levels and improves lipid peroxidation and insulin resistance in animal models of NAFLD, and its beneficial effects are dose dependent^[103,105]. There is no clinical trial evaluating its effects on patients with NAFLD.

Carnitin: Carnitine is an essential component of mitochondrial beta oxidation. It takes part in the transportation of long-chain fatty acids into the mitochondria. Abnormalities in the mitochondria have been found to play an important role in NAFLD and NASH development. There are two published clinical trials evaluating the effects of carnitin supplementation on NAFLD characteristics. Lim *et al.*^[106] showed that 3 mo of carnitine supplementation improved NAFLD features by improving serum liver function tests and mitochondrial DNA copies^[106]. Malaguarnera *et al.*^[107] showed that the addition of an L-carnitine supplement to an individual's diet for 24 wk, reduced TNF- α and CRP, and improved liver function, plasma glucose levels, lipid profile, HOMA-IR, and histological manifestations of NASH^[107]; how long these effects will sustain was not evaluated.

CONCLUSION

Since there is no proven pharmacologic treatment for NAFLD, it is critically important to find dietary approaches to the prevention, attenuation, or reversal of hepatic steatosis, and its progression to steatohepatitis. As insulin resistance, oxidative stress, and inflammation are involved in pathogenesis of NAFLD, it seems that dietary supplements that can modulate these pathologies could be useful in the treatment of NAFLD. These supplements have shown beneficial effects in animal models of NAFLD, however clinical trials are scarce. Further clinical trials are needed to support the use of supplements, either as preventative or therapeutic agents that effectively prevent the development and/or worsening of liver steatosis in patients with NAFLD.

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Management of patients with hepatitis C infection and renal disease

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Abstract

Hepatitis C virus (HCV) infection in patients with end-

stage renal disease (ESRD) is associated with more rapid liver disease progression and reduced renal graft and patients' survival following kidney transplantation. Evaluations and management of HCV in patients with renal disease are challenging. The pharmacokinetics of interferons (IFN), ribavirin (RBV) and some direct acting antiviral (DAA), such as sofosbuvir, are altered in patients with ESRD. With dose adjustment and careful monitoring, treatment of HCV in patients with ESRD can be associated with sustained virological response (SVR) rates nearly comparable to that of patients with normal renal function. DAA-based regimens, especially the IFN-free and RBV-free regimens, are theoretically preferred for patients with ESRD and KT in order to increase SVR rates and to reduce treatment side effects. However, based on the data for pharmacokinetics, dosing safety and efficacy of DAA for patients with severe renal impairment are lacking. This review will be focused on the evaluations, available pharmacologic data, and management of HCV in patients with severe renal impairment, patients who underwent KT, and those who suffered from HCV-related renal disease, according to the available treatment options, including DAA.

Key words: Hepatitis C; Renal disease; Chronic kidney disease; Dialysis; Interferon; Direct acting antivirals; Cryoglobulinemia

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Core tip: Hepatitis C virus (HCV) infection in patients with end-stage renal disease (ESRD) is associated with more rapid liver disease progression and reduced graft and patients' survival following kidney transplantation. The pharmacokinetics of interferons (IFN), ribavirin (RBV) and sofosbuvir are altered in patients with ESRD. With dose adjustment and careful monitoring, treatment of HCV can be safely utilized and successful in most patients with ESRD. direct acting antiviral (DAA)-based regimens, especially IFN-/RBV-free regimens, are preferred for patients with ESRD and kidney

transplantation (KT). However, due to inadequate data on clinical safety and efficacy, DAA-based therapies are not currently recommended in patients with severe renal disease. This review will be focused on evaluations and management of HCV in ESRD, KT recipients and HCV-related renal disease, according to the available treatment options including DAA.

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INTRODUCTION

Chronic hepatitis C virus (HCV) infection is a leading cause of chronic liver disease and hepatocellular carcinoma worldwide which over is a worldwide health problem in that it has a global prevalence rate of approximately 3% and affects over 170 million individuals^[1]. In clinical practice, it is common to see HCV patients with pre-existing renal disease. Thus, the prevalence of HCV infection is apparently increased in patients with end-stage renal disease (ESRD) on chronic replacement therapy, especially hemodialysis. Importantly, the liver-related morbidity and mortality of HCV appear to be higher in patients with ESRD than in the general population^[2-4]. For patients undergoing kidney transplantation (KT), HCV infection is associated with an increased rate of liver fibrosis, and the possibility of negatively affecting the renal graft and patients' survival^[2-4].

Management of HCV in patients with renal disease presents unique challenges. The pharmacokinetics of interferons (IFN) and ribavirin (RBV) are altered in patients with renal disease, particularly with ESRD^[2-4]. With dose adjustment and careful monitoring, treatment with pegylated interferons (PEG-IFN) with or without RBV can eradicate HCV infection in 40%-50% of ESRD patients infected with genotype 1/4 and about 80% of ESRD patients infected with genotype 2 or 3, with an incidence of discontinuation of up to 33%^[2-4]. All HCV-positive KT candidates should be assessed to receive antiviral treatment prior to transplantation due to the increased risk of progression of liver disease with immunosuppressive therapy and the inability to receive IFN therapy after KT. Theoretically, the use of direct acting antiviral (DAA)-based regimens, especially the IFN-free and RBV-free regimens, is preferred for patients with ESRD and KT in order to increase sustained virological response (SVR) rates and to reduce treatment side effects. However, the data on pharmacokinetics, dosing safety and efficacy of DAA for patients with severe renal impairment are lacking. In addition, the availability of DAA is currently limited in many

countries, especially in the developing world, mainly due to socio-economic reasons. Therefore, this review will be focused on the evaluation and management of HCV in patients with severe renal impairment, patients who underwent KT, and those who suffered from HCV-related renal disease, according to the available treatment options.

NATURAL HISTORY OF HCV IN PATIENTS WITH RENAL DISEASE

End-stage renal disease

The natural history of HCV in patients with ESRD is relatively uncertain^[3]. Nevertheless, several studies have demonstrated that ESRD on dialysis is associated with an increased risk for all-cause and liver-related mortality^[3,5,6]. Cardiovascular disease remains, however, the major cause of death in dialysis patients irrespective of HCV status^[1]. Death from cirrhosis and hepatocellular carcinoma was notably higher among HCV-positive ESRD patients^[5]. A meta-analysis on survival in dialysis patients (7 studies; $n = 11589$) showed an estimated relative risk for death in anti-HCV positive patients of 1.34 (95%CI: 1.13-1.59) with liver-related complications contributing to poorer outcomes^[5]. Moreover, HCV infection can adversely affect the quality of life in this population^[7].

KT

The impact of immunosuppression on HCV disease progression following renal transplantation is unclear. Serum HCV-RNA levels typically increase. Most data suggested that HCV-infected patients have worsening of hepatic necroinflammation and accelerated hepatic fibrosis following KT^[8-10], though some studies reported that liver histology may remain stable or even improve^[11,12]. Not only affecting the liver, several studies demonstrated that patients with HCV infection have a poorer patient and graft survival after KT compared to those without^[3,13,14]. The exact reason for reduced renal graft survival in HCV patients is unknown, but it may partly explain by de novo immune complex chronic glomerulonephritis in the allograft induced by chronic HCV infection^[15,16]. Nevertheless, undergoing KT evidently conferred a long-term survival advantages, particularly on the cardiovascular death, over HCV patients with ESRD on maintenance dialysis on the waitlist, although there was a higher risk for infection-related death during the first 6 mo after KT^[11]. As cirrhosis is an key predictor of poor survival after KT, assessment of the stage of liver fibrosis in all HCV-positive KT candidates is recommended^[17]. For patients with established cirrhosis and portal hypertension who failed (or are not suitable for) HCV treatment, isolated KT may be inappropriate in this settings and a combined liver and KT should be considered^[18].

HCV infection acquired during or after RT showed a severe and rapidly progressive course, which is significantly different from HCV patients without transplantation^[19]. In addition, fibrosing cholestatic hepatitis (FCH) can occur in HCV-infected patients following KT^[20]. It typically develops during the period of maximal immunosuppression (1-4 mo after KT) and is associated with progressive cholestatic, mild elevation of serum alanine aminotransferase (ALT), and high HCV viremia level^[20-22]. FCH is associated with very high morbidity and mortality rates. IFN-based treatment is often ineffective and is associated with a risk of graft rejection^[20-22].

EVALUATIONS OF HCV IN PATIENTS WITH RENAL DISEASE

Serum aminotransferases

It is known that serum ALT levels in patients with ESRD are lower than in the general population, so it should not be used to screen for liver diseases^[23-25]. This is possibly due to suppression of ALT synthesis in hepatocytes, defective release of ALT into the blood stream, or accelerated clearance in patients with chronic renal insufficiency^[24,26]. The lower cut-off ALT level (≥ 27 U/L) was proposed for patients with ESRD to increase sensitivity (to 50%) and specificity (to 100%) for detecting HCV viremia^[23]. In addition, there is a weak correlation between ALT levels and liver disease activity in patients with ESRD, especially those on dialysis^[25].

Viral markers

Anti-HCV assay by enzyme immunoassay (EIA) technique is the most commonly used screening tool for HCV infection due to its simplicity, availability and low cost. The second generation EIA (EIA-2) assay was frequently associated with false negative results in patients with ESRD on dialysis, with a reported rate of 2.6%-7%^[27,28]. The third generation EIA (EIA-3) testing provided excellent accuracy, with 0.26% false-negative rate, and is the preferred screening tool in this setting^[3,25,29]. PCR-based molecular diagnostics are required to confirm viremia, viral load, and genotype to guide management decisions. Notably, HCV-RNA level is transiently decreased during hemodialysis and gradually returns to baseline level within 48 h^[3]. This may be explained by several mechanisms, such as interference with PCR technique by heparin used during dialysis, adsorption of HCV onto the dialysis membrane, destruction of HCV particles by the hydraulic pressure, escape of HCV into the dialysate, or increased plasma IFN levels during the dialysis^[3,25]. Therefore, it is recommended to determine HCV-RNA level before hemodialysis to avoid the possibility of underestimation^[3].

Assessment for liver fibrosis

Liver biopsy is the gold standard for assessing the degree of fibrosis in HCV patients. However, its use is limited by invasive nature, poor patient acceptance, and bleeding risk, especially in uremic patients. The use of non-invasive fibrotic markers has been evaluated in HCV patients with ESRD. The aspartate transaminase: platelet ratio index (APRI) can reliably predict liver fibrosis in HCV patients with ESRD, especially to exclude patients with significant fibrosis^[30-32]. In Schiavon *et al.*^[31] study (203 ESRD HCV-infected subjects), APRI < 0.40 accurately identified patients with fibrosis stage 0 or 1 with negative predictive value of 93%; APRI ≥ 0.95 can confirm significant fibrosis (\geq fibrosis stage 2) with positive predictive value of 66%^[31]. If biopsy indication was restricted to APRI scores in the intermediate range, about 50% of liver biopsies could be avoided^[31]. Transient elastography shows superior diagnostic accuracy to APRI in HCV patients with ESRD^[3,32]. The suggested optimized cut-off values were 5.3 kPa, 8.3 kPa, and 9.2 kPa, for fibrosis stage of $\geq F2$, $\geq F3$, and $F4$, respectively (sensitivity 93%-100% and specificity 88%-99%)^[32]. Further, a small study ($n = 22$) revealed a good correlation between transient elastography and fibrosis stage on histology in HCV-positive KT recipients^[33].

EPIDEMIOLOGY AND SCREENING OF HCV IN PATIENTS WITH ESRD ON DIALYSIS

The prevalence of HCV infection in patients with ESRD patients varies among geographical areas and dialysis centers, but it is obviously higher than that of the general population^[2]. Risk factors for acquiring HCV infection during dialysis include: the number of transfusions, duration of dialysis, number of procedures for dialysis access, type of dialysis; hemodialysis (HD) $>$ peritoneal dialysis (PD), prevalence of HCV infection and lack of compliance with universal precautions in the dialysis unit^[34,35].

In developed countries, the prevalence and incidence of HCV has been declining in the past decades^[2,36-38]. In the United States national surveillance ($n = 164845$), the prevalence of anti-HCV positivity has dropped from 10.4% in 1985 to 7.8% in 2002 (ranged from 5.5%-9.8%)^[37]. Similarly, the European multicenter survey reported that the prevalence of anti-HCV positivity has dropped steadily from 13.5% in 1991 to 6.8% in 2000 in the Belgian cohort ($n = 1710$); prevalence also decreased ($P < 0.05$) in France (42% to 30%), Sweden (16% to 9%) and Italy (28% to 16%), tended to decrease in United Kingdom (7% to 3%, $P = 0.058$) and Hungary (26% to 15%, $P = 0.057$), but

did not change (NS) in Germany (7% to 6%), Spain (5% to 12%) and Poland (42% to 44%)^[36]. Despite the elimination of post transfusion HCV infection, the incidence of HCV infection among patients on chronic dialysis treatment, with seroconversion rates ranging between 0.2%-15% per year of dialysis, continues to be a cause of concern^[2,36-38]. The data on epidemiology of HCV among patients with ESRD on dialysis in developing countries are less abundant and more heterogeneous, but the overall prevalence and incidence rates seem to be higher than developed countries^[2]. The prevalence of anti-HCV positivity in single center surveys from Brazil (2005), Turkey (2005), Tunisia (2006), Iran (2005), Saudi Arabia (2004), Morocco (2005) and Egypt (2000) were 8%, 19%, 20%, 25%, 43%, 76%, and 80%, respectively^[2]. Whereas in the Asia-Pacific dialysis registry (173788 HD; 27802 PD), HCV seroprevalences range between 0.7%-18.1% across different countries and were generally higher in HD vs PD populations (7.9% \pm 5.5% vs 3.0% \pm 2.0%, $P = 0.01$)^[38]. Thus, the annual incidence of HCV infections range from 0% in Thai PD patients to 18.1% in Indian HD patients with the rates were generally higher in HD patients than in PD patients (RR 0.33, 95%CI: 0.13-0.75)^[38].

Although prospective trials have shown a reduction in HCV transmission within dialysis units by complete isolation of HCV patients, but this practice has not been universally accepted^[2,39]. The Centers for Disease Control and Prevention of the US (CDC) and the Kidney Disease: Improving Global Outcomes (KDIGO) practice guidelines^[40] do not recommend dedicated machines, patients isolation, or a ban on reuse in HCV patients on hemodialysis^[40,41]. However, strict adherence to "universal precautions", careful attention to hygiene, and sterilization of dialysis machines is emphasized^[40,41]. Further, the CDC recommends that all HD patients should be screened for anti-HCV at baseline, and then subsequently tested semiannually^[41].

PHARMACOLOGIC ISSUES OF ANTIVIRAL AGENTS IN PATIENTS WITH RENAL DISEASE

Interferons and pegylated interferons

Interferon- α is a glycoprotein, produced by immune cells in response to foreign antigens, such as viruses, bacteria, parasites or tumor cells. The elimination half-life of IFN following subcutaneous injections is approximately 2-4 h, then it is filtered through the glomeruli and during proximal tubular reabsorption undergo lysosomal proteolytic degradation^[25,42]. Kidney is the main site of degradation of IFN molecule, while liver plays only a minor role. Due to a short elimination half-life of IFN following subcutaneous injections (2-4 h), sustained plasma levels

are not maintained when dosed 3 times weekly, which is believed in part to explain the suboptimal response rates^[25,42]. Accumulation of IFN occurs in patients with renal dysfunction, especially in ESRD^[25,42,43]. Although this may result in higher and more sustained plasma levels of IFN, which is preferable for the anti-viral activity against HCV, it may lead to serious adverse events in such patients as well^[25,42,43].

Combining a polyethylene glycol (PEG) polymer to IFN successfully created a molecule with a longer half-life, improved pharmacokinetic profile, and more importantly, a superior clinical response when dosed once weekly^[44,45]. PEG-IFN α -2a, a branched-PEG (40 kD) attached to an IFN α -2a molecule, is absorbed slowly (absorption half-life approximately 50 h), has a restricted volume of distribution (2-12 L), and a long elimination half-life (half-life approximately 77 h; peak through ratio 1.5-2)^[44,45]. PEG-IFN α -2b, a linear PEG molecule (12 kD) attached to IFN α -2b, is absorbed rapidly (absorption half-life approximately 4.6 h), has a large volume of distribution (0.9 L/kg) and a shorter elimination half-life (half-life approximately 40 h; peak through ratio > 10)^[44,45]. PEG-IFN α -2a is metabolized in the liver and kidneys while PEG-IFN α -2b is metabolized exclusively by the kidneys^[25,44,45]. The pharmacokinetics of PEG-IFN α -2a is less affected by renal failure, and less dose modifications are necessary in the setting of renal impairment as compared to PEG-IFN α -2b (Table 1)^[44,45]. In patients with severe renal impairment [creatinine clearance (CrCl) < 30 mL/min], the maximum plasma concentration and the area under the curve (AUC) of PEG-IFN α -2b are increased by approximately 90% and half-life is increased by approximately 40%^[46]. Similar pharmacokinetic profiles have been also observed for PEG-IFN α -2a^[25,45,46]. In addition, hemodialysis has only a small effect on IFN and PEG-IFN clearance^[25,45,46].

Ribavirin

After oral absorption, RBV is rapidly absorbed and distributed with a bioavailability of approximately 50%. It has extensive volume of distribution and the steady state is reached in 7-11 wk after multiple dosing and with a terminal half-life of 12 d^[47-49]. The route of RBV elimination is mainly by the kidney. Thus, body weight is also highly correlated with RBV clearance^[47-49]. Notably, the optimal dosing strategy of RBV must be calculated according to the renal function, as measured by CrCl^[47-49]. The AUC for RBV is increased by 2 folds in patients with CrCl 30-60 mL/min and by 3 folds in patients with CrCl 10-30 mL/min respectively, when compared to those with CrCl > 90 mL/min^[50]. Notably, the most pronounced side effect of RBV in patients with renal disease is hemolytic anemia. The mechanism of RBV-induced anemia is unclear, but evidently involves an extensive accumulation of active RBV metabolites

Table 1 Dosage modification for patients with renal impairment

Creatinine clearance	Pegylated-interferon alfa-2a	Pegylated-interferon alfa-2b	Ribavirin
30-50 mL/min	180 µg/wk	1.125 µg/kg per week (25% reduction)	Alternating doses; 200 mg and 400 mg every other day
< 30 mL/min	135 µg/wk (25%-45% reduction)	0.75 µg/kg per week (50% reduction)	200 mg/d
Hemodialysis	135 µg/wk (25%-45% reduction)	0.75 µg/kg per week (50% reduction)	200 mg/d

Table 2 Pharmacokinetic and metabolic parameters of selected direct acting antivirals

Drugs	Metabolism/excretion route	Interaction with CYP3A	Comments
NS3/4A protease inhibitors Boceprevir	Hepatic (CYP3A, aldoketoreductase)	Moderate CYP3A inhibitor	Significant DDI with other CYP3A substrate drugs
Telaprevir	Hepatic (CYP3A)	Strong CYP3A inhibitor	Significant DDI with other CYP3A and P-gp substrate drugs
Simeprevir	Hepatic (CYP3A)	Mild CYP1A2 and CYP3A inhibitor	Unconjugated hyperbilirubinemia commonly seen
Faldaprevir	Hepatic (CYP3A)	Moderate CYP3A inhibitor; weak CYP2C9 inhibitor	Inhibition of UGT1A1 results in unconjugated hyperbilirubinemia
ABT-450/ ritonavir NS5A replication complex inhibitors	Hepatic (CYP3A)	Strong CYP3A inhibitor (by ritonavir)	Unconjugated hyperbilirubinemia
Daclatasvir	Hepatic (CYP3A)	No/minimal	
Ledipasvir	Feces (major); hepatic and renal (minor)	No/minimal	
NS5B nucleos(t)ide polymerase inhibitors Sofosbuvir	Renal	No/minimal	Dose reduction if moderate to severe renal impairment
NS5B non-nucleoside polymerase inhibitors ABT-333	Hepatic (CYP2C8 60%, CYP3A4 30% and CYP2D6 10%)	No/minimal	

Adapted from Tischer *et al*^[55]. CYP3A4: Cytochrome P450 3A4.

and subsequent oxidative stress within red blood cells (RBC)^[51]. Ribavirin is actively transported into RBC and accumulates within RBC with the concentration greatly exceeding what is observed in plasma (up to 60-fold)^[47,51,52]. Once inside RBC, RBV is phosphorylated to RBV-triphosphate, then it is eliminated slowly from RBC^[47,51,52]. An increase in RBV-triphosphate concentrations in RBC enhances oxidative stress with subsequent RBC membrane damage, leading to premature extravascular destruction of RBC by reticuloendothelial system^[51,52].

With careful monitoring, reduced dose PEG-IFN plus markedly reduced dose RBV (170-400 mg/d) has been safely utilized in patients with renal impairment, as well as those patients on dialysis^[47,49]. However, the use of RBV in HCV patients on dialysis has not been well studied and it should be used with extreme caution and close follow. Further, minimal amount of RBV is removed by dialysis so that there is a potential risk of drug accumulation^[25]. Despite this, there are some data to support the use of RBV in dialysis patients by starting empirically with a low dose of 200 mg/d, then adjusting doses according to changes in hemoglobin^[47,49,53]. Notably, erythropoiesis-stimulating agents may be used

to counteract anemia and help maintain optimal RBV dose in patients with renal disease^[47,49]. Until recently, the use of RBV in persons with CrCl < 50 mL/min was contraindicated due to a markedly reduced ribavirin elimination rate, leading to severe hemolytic anemia. However, the US Food and Drug Administration (US-FDA) has approved a labeling change of RBV for patients with severe renal impairment with recommended dosage modifications since 2011^[54] (Table 1). It should be noted that breaking RBV tablet into half is not advised due to the potential environmental contamination of this serious teratogen. Given its extensive Vd and long half-life, RBV can be finely adjusted by alternating daily dosage^[47,49].

Direct acting antivirals

Most Direct acting antiviral (DAA) are metabolized primarily *via* the liver and dose adjustment is not necessary for patients with renal impairment^[55,56] (Table 2). However, it should be noted that these agents have not been adequately evaluated for the treatment of HCV in patients with renal impairment in clinical trials in terms of safety and efficacy. *In vitro* studies indicate that boceprevir (BOC) extensively

Timing	Available strategies
Moderate renal impairment (CrCl 30-60 mL/min)	Regular/reduced dose PEG-IFN plus RBV (approximately 200-400 mg/d) DAA-based therapy-should be utilized No dose reduction for BOC/TVR/SMV/SOF IFN-free/RBV-free regimens may be preferred
Severe renal impairment (CrCl < 30 mL/min), ESRD (CrCl < 15 mL/min), On dialysis	Reduced dose PEG-IFN plus RBV (approximately 200 mg/d) IFN/PEG-IFN monotherapy may be effective DAA-based therapy-limited clinical data No dose reduction for BOC/TVR/SMV SOF is not recommended IFN-free/RBV-free regimens are preferred Anti-HCV should be tested semiannually during dialysis Patients with established cirrhosis and PHT may be contraindicated for KT, especially if they are viremic
After KT	IFN-based therapy-generally contraindicated (↑rejection) DAA-based therapy-limited clinical data IFN-free/RBV-free regimens are preferred SOF/SMV have no/minimal DDI with IMS agents

Figure 1 Management of hepatitis C virus in patients with renal disease and kidney transplantation. Empiric dose changes should be done in conjunction with therapeutic drug monitoring. BOC: Boceprevir; TVR: Telaprevir; SMV: Simeprevir; SFV: Sofosbuvir; CYP3A4: Cytochrome P450 3A4; mTOR: Mammalian target of rapamycin; MMF: Mycophenolate mofetil.

undergoes metabolism through the aldo-keto reductase-mediated pathway and, to a lesser extent, oxidative metabolism mediated by CYP3A4 in the liver^[54]. Telaprevir (TVR) is primarily metabolized in the liver involving hydrolysis, oxidation, and reduction^[54]. No dose adjustment is required for BOC and TVR in patients with any degree of renal impairment^[54]. Simeprevir (SMV) is metabolized by liver CYP3A4 and kidney clearance plays an insignificant role (< 1%)^[57]. A pharmacokinetic study in volunteers demonstrated that SMV exposure after 7 d of 150 mg/d dosing was 62% higher in patients with severe renal impairment compared with matched healthy volunteers, but safety and tolerability were considered generally favorable^[57]. Daclastavir is also metabolized primarily by the liver, and the need for dose adjustment in patients with impaired renal function is unknown^[55,56]. Unlike the others, sofosbuvir (SOF) is excreted *via* the kidneys. A single 400-mg dose of SOF resulted in 56%, 90%, and 456% higher levels of the major systemic metabolite, GS-331007, among persons with mild, moderate, and severe renal dysfunction, respectively, compared with individuals with normal renal function^[58].

MANAGEMENT OF HCV IN PATIENTS WITH ESRD

Interferon or pegylated interferon monotherapy

There have been at least 4 meta-analyses demonstrating that IFN monotherapy is effective for HCV patients with ESRD, with the overall SVR rates of 33%-41% and the withdrawal rates of 17%-30%^[59-62]. The discontinuation of treatment were mainly due to

flu-like symptoms, gastrointestinal and hematological adverse events, which is more frequent than in patients with normal renal function^[59-62]. Notably, the use of PEG-IFN does not seem to provide an additional benefit in terms of SVR compared to conventional IFN monotherapy in patients on dialysis^[61,62]. This finding may largely explain by changes in the pharmacokinetics of IFN toward increasing half-life and AUC (more similar to that of PEG-IFN) in the setting of ESRD. In a recent meta-analysis of 25 studies (included 459 patients treated with IFN 38 patients treated with PEG-IFN and 49 patients treated with PEG-IFN/RBV), the overall SVR rate was 41% (95%CI: 33-49) for IFN and 37% (95%CI: 9-77) for PEG-IFN^[61]. Treatment discontinuation rates were 26% (95%CI: 20%-34%) for IFN and 28% (95%CI: 12%-53%) for PEG-IFN. The SVR rate tended to be higher when 3 million units or higher of IFN 3 times weekly were used^[61] (Figure 1).

Interferon or pegylated interferon plus ribavirin

In patients with normal renal function, the addition of RBV to IFN or PEG-IFN is generally required in order to achieve an optimal SVR, primarily by decreasing relapse rates^[47]. Several small studies have demonstrated that the use low-dose RBV (from 200 mg/wk to 400 mg/d) in combination with IFN-based therapy was feasible for treating HCV patients with ESRD^[3,50]. Despite the widespread use of high-dose erythropoietin, falling hemoglobin levels were commonly observed^[3,50]. Overall, SVR and treatment-related withdrawal rates after 24-48 wk of combination therapy ranged between 17%-90% and 0%-70%, respectively^[3,50]. Rendina *et al*^[63] reported a case series evaluating PEG-IFN 135 µg/

wk plus RBV 200 mg/d for 24 (non-G1) or 48 wk (G1) in HCV patients on dialysis^[63]. In this study, 35 patients received treatment (35 served as untreated controls), and 30 patients completed treatment (drop-out rate 14%)^[63]. Overall, 34 of 35 (97%) treated patients, including those with treatment discontinuation due to side effects, achieved SVR^[63]. Recently, an open-label, randomized trial in Taiwan randomized 205 HCV patients on hemodialysis to PEG-IFN 135 µg/wk plus RBV 200 mg/d ($n = 103$) or PEG-IFN 135 µg/wk alone ($n = 102$) for 48 wk^[53]. Compared with monotherapy, combination therapy had a greater SVR rate (64% vs 33%, $P < 0.001$)^[53]. More patients receiving combination therapy required a higher dosage and longer duration of epoetin-beta and had more hemoglobin levels < 8.5 g/dL than those receiving monotherapy (72% vs 6%, $P < 0.001$)^[53]. The adverse event-related withdrawal rates were similar between the two groups^[53]. The recent AASLD/IDSA guidance recommended patients with renal impairment/ESRD/HD, dosing of PEG-IFN and RBV to follow updated FDA recommendations or package insert recommendations based on calculated glomerular filtration rate^[64].

Boceprevir- or telaprevir-based triple therapy

The first wave DAA, boceprevir (BOC) and telaprevir (TVR), in combination with PEG-IFN/RBV has been the standard of care of HCV genotype 1 in many countries since 2011, with an improvement in SVR up to 65%-75% in treatment naïve patients and 60%-65% in previous relapsers/non-responders^[1]. Notably, patients with CrCl < 50 mL/min or those with ESRD were excluded from their registration trials^[1]. Despite the lack of required dose adjustment, there have been very few small case series evaluating the safety and efficacy of these protease inhibitors in patients with ESRD on dialysis^[65-67]. Therefore, the routine use of BOC/TVR-based triple therapy routine use in patients with severe renal impairment cannot be recommended^[54,64]. It is possible that treatment-related side effects, particularly anemia, will be increased in patients with renal disease. Thus, the potential drug-drug interactions between the BOC/TVR and concomitant medications that commonly used in patients with ESRD (*e.g.*, antihypertensive and lipid lowering agents) may further complicate the therapy.

Sofosbuvir- and simeprevir-based therapy

SOF and SMV have recently been approved for the treatment of HCV in United States and many countries in Europe. In settings where available, SOF- and SMV-containing regimens are preferred for over BOC/TVR-based triple therapy due to superior efficacy, more convenience dosing, and less drug-drug interactions^[64]. Ideally, patients with ESRD should receive an IFN-free, and if possible RBV-free

treatment regimen. Based on the available data, the recent AASLD/IDSA guidance advised that no dose reduction is needed when using SOF in patients with HCV infection with mild to moderate renal impairment (CrCl ≥ 30 mL/min). However, SOF is not recommended in patients with severe renal impairment and ESRD (CrCl < 30 mL/min) or those who require HD until more data available^[64]. For SMV, no dosage adjustment is required for patients with any degree of renal impairment. SMV has not been adequately studied in patients with ESRD, including those requiring HD^[64].

MANAGEMENT OF HCV IN KIDNEY TRANSPLANT RECIPIENTS

Interferon- or pegylated interferon-based therapy

Outcomes of IFN-based treatment after KT are somewhat disappointing. In a meta-analysis of 12 clinical trials (102 RT recipients with HCV), treatment with IFN with or without RBV is associated with the overall SVR and treatment-related withdrawal rates of 18% and 35%, respectively^[68]. More recent meta-analysis ($n = 140$) reported that the overall SVR rate, drop-out rate and graft rejection rate was 26.6%, 21.1% and 4%, respectively^[69]. Thus, PEG-IFN may be a more effective approach for treating HCV post-KT than standard IFN-based treatment (SVR: 40.6% vs 20.9%)^[69]. These suboptimal SVR rates may largely explain by the interruption of treatment by side effects and the limited efficacy of IFN in immunosuppressed patients. In addition, the immunostimulatory effects of IFN, including increased expression of cytokines and HLA antigens, and enhanced function of cytotoxic T cells and natural killer cells, can lead to an increased risk of acute allograft rejection in KT recipients^[4]. The early studies reported that graft dysfunction occurred in 15%-100% of HCV-positive KT recipients treated with IFN, with up to 20% resultant permanent allograft failure^[4,70-73]. Although more recent studies revealed that PEG-IFN/RBV treatment may be feasible for KT recipients and the development of graft rejection was in fact relatively rare (0%-5%), but the SVR rates are still relatively low (38%-50%)^[74-76]. Taken together, IFN-based therapy should only be initiated in KT recipients under specific clinical situations, such as FCH or severe de novo glomerulonephritis, where DAA are unavailable and when the risk of not treating HCV infection outweighs the risk of graft loss (Figure 1).

Boceprevir- or telaprevir-based triple therapy

Both BOC and TVR are substrates for and inhibitors of the CYP3A4 and the drug transporter P-glycoprotein, so that they are prone to interact with other medications involving this enzyme^[55,56,77]. Several studies have demonstrated that co-administration

of BOC and TVR significantly increased the dose exposure of cyclosporine, tacrolimus and mammalian target of rapamycin (mTOR) inhibitors by several folds^[55,56,77]. Lessons from LT suggested that the drug-drug interaction issues with BOC- and TVR-based triple therapy may be manageable with preemptive dose reduction and close monitoring of immunosuppressive drug levels^[55,56,77]. However, the use of BOC/TVR-based triple therapy may not be suitable for KT setting due to the possible rejection issue with IFN.

Sofosbuvir- and simeprevir-based therapy

KT recipients with an indication for HCV treatment should receive IFN-free regimen. SOF does not undergo metabolism *via* hepatic CYP3A, limiting the likelihood of drug-drug interactions with immunosuppressive agents that are inducers or inhibitors of this enzyme^[55,56]. However, SOF is a substrate for P-gp and should not be coadministered with potent P-gp inducers such as rifampin or St John's wort. SMV is a mild CYP3A inhibitor and weak drug-drug interaction may be observed when co-administration with immunosuppressive agents. For example, the AUC of tacrolimus decreased by 17% and that of cyclosporine increased by 19% with SMV co-administration^[78]. However, the package insert for SMV advises that no dose adjustments are needed during co-administration with cyclosporine or tacrolimus. Monitoring of sirolimus concentrations may be advisable when given with SMV.

HEPATITIS C VIRUS-RELATED RENAL DISEASE

Epidemiology

Considering HCV is both a hepatotropic and lymphotropic virus, in addition to causing a liver disease, it can be associated with a number of lymphoproliferative and immunological disorders of various organ systems, including the kidney^[79] (Figure 2). HCV-related renal disease can affect both glomerular and tubular component, which can be presented with a wide array of clinical manifestations, ranging from asymptomatic proteinuria, overt glomerular disease, to ESRD. Notably, the most common and well-established renal involvement of HCV is mixed cryoglobulinemia (MC)-associated glomerulonephritis^[16,79-81]. Whereas other conditions such as membranoproliferative glomerulonephritis (MPGN) and membranous nephropathy, focal segmental glomerulosclerosis, fibrillary glomerulonephritis, immunotactoid glomerulopathy, IgA nephropathy, mesangial proliferative glomerular nephritis, renal thrombotic microangiopathy, vasculitic renal involvement and interstitial nephritis are less commonly described^[16,80]. These pathologic findings are not specific for HCV and the diagnosis of HCV-related renal disease has to be made cautiously with exclusion of the

other secondary causes. In most circumstances, renal involvement itself does not affect overall survival in HCV infected patients but it could modify natural course of disease significantly^[82].

The prevalence of HCV-related renal disease varies among reports and also seems to be geographical heterogeneity. For example, the prevalence of MC is clearly more prevalent in Southern Europe (more than 50% of HCV-infected individuals) than in Northern Europe, North America, and Asia^[79,81]. One large epidemiologic study among US veterans showed that cryoglobulinemia and MPGN, not membranous glomerulopathy, are more prevalent in HCV-infected patients ($n = 34204$) compared to non-HCV controls ($n = 136816$): 0.57% vs 0.05%; $P < 0.001$, 0.36% vs 0.05%; $P < 0.001$ and 0.33% vs 0.19%; $P = 0.86$, respectively^[83]. In addition, HCV infection is independently associated with proteinuria^[84,85]. Besides, many epidemiological studies have shown an association between HCV and diabetes mellitus, especially type 2^[86]. The processes seem to involve direct viral effects, insulin resistance, proinflammatory cytokines, chemokines, and other immune-mediated mechanisms^[86]. Therefore, proteinuria and renal disease in such patients may be secondary from diabetic nephropathy, not directly related to HCV itself.

Diagnosis of mixed cryoglobulinemia

Cryoglobulinemia is a chronic systemic disease characterized by the presence of serum immunoglobulins that reversibly precipitate at low temperature^[87]. Classically, cryoglobulemia is divided into 3 types. Type I composed of a monoclonal immunoglobulin associated mainly with overt lymphoproliferative disorders and is classically presented with vascular occlusion or hyperviscosity syndrome^[80,88]. Type II and III, so-called MC, consisted of polyclonal IgG and monoclonal/polyclonal IgM with rheumatoid factor. These two types are associated with infectious, immunological and neoplastic disorders (also can be idiopathic), and often present with vasculitis syndrome, *e.g.*, peripheral neuropathy, skin ulcers as well as glomerulonephritis^[80,88]. The strong association between HCV and MC type II and III has been supported by several epidemiological studies^[16,79-81,87]. HCV appears to have an important etiologic role in MC, as the evidence of HCV infection can be found in 76%-95% of patients with MC^[89,90]. On the other hand, low level of cryoglobulinemia is commonly found in unselected HCV patients with prevalence 19%-54%^[91,92]. Most of these patients are asymptomatic and about 5%-20% have an overt clinical of MC^[91,92].

Diagnosis of MC is based on clinicopathological and laboratory work-up including cryoglobulin testing, quantitative serum protein and globulins, complement levels, virologic markers, and urine analysis. The

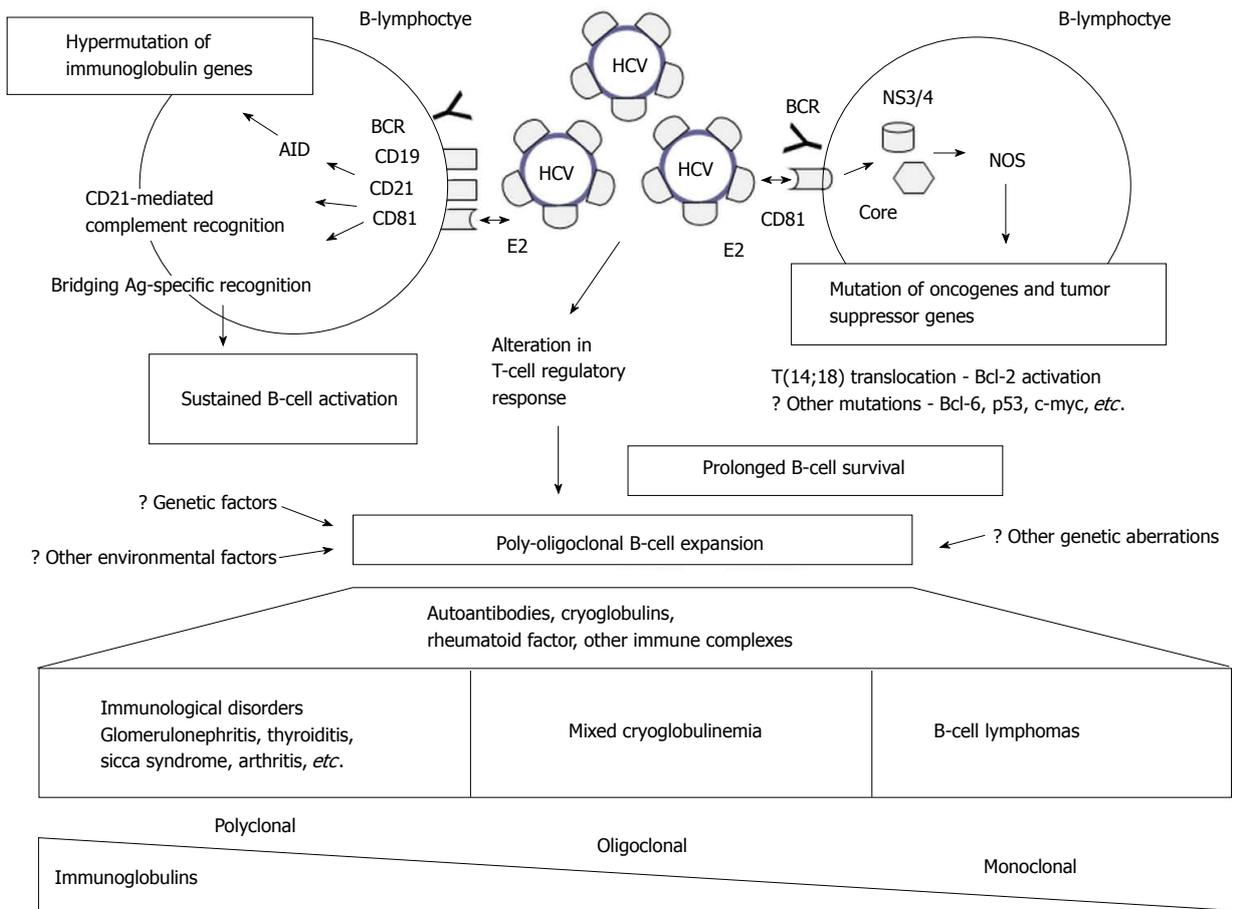


Figure 2 Proposed pathogenesis of hepatitis C virus-related immunogenic and lymphoproliferative disorders. HCV: Hepatitis C virus; NOS: Nitric oxide synthase; ROS: Reactive oxygen species; AID: Activation-induced deaminase.

clinical syndrome of MC caused by deposition of circulating immune complexes in small to medium-sized blood vessels in multiple organs, inducing systemic vasculitis. The main clinical manifestations of MC are palpable purpura, arthralgia, myalgia, peripheral neuropathy and hypocomplementemia^[88,90]. Cryoglobulins can precipitate *in vitro* at temperatures of less than 37 °C (typically at 4 °C^[80]) and re-dissolve after rewarming^[93].

Renal involvement is presented in up to one third of patients and represents a strong negative prognostic factor, even if their course may vary^[90,94]. Nephropathy is observed in 20% at the diagnosis of MC, and in 35%-60% during follow up, in which the majority occurs within a few years^[90,95]. Clinically, MC-associated glomerulonephritis may range from asymptomatic abnormal urinalysis (microscopic hematuria, or sub-nephrotic range proteinuria with normal, or mildly impaired, renal function), overt nephritis (20%-25%) and nephrotic syndrome (20%), with variable progression to end-stage renal disease in 10%-33% of patients^[90,95]. The typical renal histopathologic pattern is type I MPGN, which can be differentiated from idiopathic MPGN by the presence of capillary thrombi, composed of

precipitated cryoglobulins, and diffuse IgM deposition in the capillary loops^[80,90,96,97].

Treatment of mixed cryoglobulinemia-related glomerulonephritis

Several small studies reported a beneficial effect of PEG-IFN/RBV in patients with HCV-related MC with 62%-78% SVR rates^[87,98]. PEG-IFN/RBV is generally well-tolerated (treatment-related side effects 22%-54%) in cryoglobulinemic patients and the dosage should be adjusted according to renal function^[86,97]. Notably, the chance to achieve SVR is not affected by the presence of cryoglobulinemia^[87,98]. Importantly, significant improvement of clinical MC syndrome and immunologic parameters is typically observed in patients who attained SVR. Cryoglobulin level often declines or even disappears after successful treatment. Though, MC-related vasculitis has been reported to persist or to relapse after achieving in a small proportion of patients^[94,99]. A meta-analysis of controlled clinical trials suggested that IFN-based therapies were more effective than immunosuppressive agents in lowering proteinuria of patients with HCV-related cryoglobulinemic glomerulonephritis (OR = 3.86; 95%CI: 1.44-10.33)^[100].

Recent studies in HCV-related MC with or without renal involvement demonstrated that triple therapy with TVR, BOC or SOF plus PEG-IFN/RBV are safe and effective in cryoglobulinemic patients^[100-102]. However, such therapeutic regimens should be administered cautiously considering the high rate of side effects (up to 35% discontinuation rates)^[101,103]. More clinical trials, especially with IFN-free regimens, are eagerly awaited.

Corticosteroids and cyclophosphamides have shown to effectively induce clinical remission in patients with severe MC. However, their effects are not sustainable and they can be associated with significant side effects, liver toxicity, and subsequent increase in HCV viremia^[86,99]. Rituximab, a chimeric monoclonal antibody specifically directed against CD20 antigen, has been proven to be safe and effective in the treatment of MC with or without HCV^[87,98,104]. A randomized controlled trial comparing between PEG-IFN/RBV with or without rituximab in treatment-naïve MC patients demonstrated that a combination of PEG-IFN/RBV plus rituximab is well tolerated and more effective than PEG-IFN/RBV alone^[105]. Thus, its effect may last for more than 3 years^[105]. Therefore, a combination of rituximab and antiviral therapy is recommended to treat in HCV-related MC with progressive kidney disease^[40,87,106]. In addition, the removal of circulating cryoglobulins by therapeutic plasmapheresis combined with immunosuppressive agents, such as pulse corticosteroids, may be considered as an adjunctive therapy for severe exacerbation of vasculitis, especially rapidly progressive glomerulonephritis^[40,87,106,107].

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Micro RNAs in the development of non-alcoholic fatty liver disease

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the underlying pathogenesis of the disease. Research efforts are ongoing to identify biological targets and signaling pathways that mediate NAFLD. Emerging evidence has implicated a role for micro RNAs (miRNAs), short single-stranded molecules that regulate gene expression either transcriptionally, through targeting of promoter regions, or post-transcriptionally, by blocking translation or promoting cleavage of specific target mRNAs. Several miRNAs have been associated with NAFLD, although our understanding of the biology underlying their role is still emerging. The goal of this review is to present an overview of the current state of knowledge of miRNAs involved in the development of NAFLD across a range of *in vitro* and *in vivo* models, including miRNAs that contribute to pathological mechanisms related to fatty liver in humans. Much less is known about the specific targets of miRNAs in cells, nor the molecular mechanisms involved in the development and progression NAFLD and related outcomes. More recently, the identification and validation of miRNA signatures in serum may facilitate the development of improved methods for diagnosis and clinical monitoring of disease progression.

Key words: MiRNA; Nonalcoholic fatty liver disease; Cell culture; Mouse; Human

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Core tip: Available data on miRNAs in nonalcoholic fatty liver disease (NAFLD) are largely derived from various cell culture and animal models. Reflecting an emerging field, little cross-model concordance is present and few human data are available for comparison with cell culture and animal model results. Although the generation of human data may be limited by the availability of tissue samples, recent reports of circulating miRNAs from NAFLD patients hold promise for significant progress for diagnosis and clinical

monitoring of disease progression.

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INTRODUCTION

Nonalcoholic fatty liver disease (NAFLD) represents a spectrum of conditions resulting from excessive accumulation of fat in hepatocytes, a condition known as steatosis. NAFLD can be categorized into nonalcoholic fatty liver representing simple steatosis and non-alcoholic steatohepatitis (NASH) with coincident hepatocyte injury, liver inflammation, and fibrosis. NAFLD is the major cause of chronic liver disease (CLD), which is associated with substantial morbidity and mortality in developed countries^[1]. In tandem with the rising rates of obesity and type 2 diabetes mellitus (T2D), the prevalence of NAFLD is also increasing, and is expected to double by 2030 in the United States. NAFLD/NASH has thus become a leading cause of cryptogenic cirrhosis^[2] and is currently the 3rd leading clinical indication for liver transplantation in the United States and expected to soon become the primary indication^[3]. At present liver biopsy is the only method to accurately assess the severity of liver fibrosis^[1] or predict progression of NAFLD to clinically severe forms^[4-6], which limits early diagnosis of NAFLD patients who are at high risk for development of liver-related morbidity and mortality. Given the substantial public health burden of NAFLD, novel therapeutic targets are also urgently needed to facilitate the development of improved pharmacological therapies for the treatment and prevention of the disease.

Although genetic predisposition, environmental exposures, and lifestyle factors contribute to the development of NAFLD, little is known about the underlying pathogenesis of the disease. Ongoing research efforts to identify biological targets and signaling pathways that mediate NAFLD are expected to provide the insight necessary to begin to distinguish among different clinical forms of the disease.

Emerging evidence has implicated a role for epigenetic factors, including micro RNAs (miRNAs) in the development of NAFLD. MiRNAs are endogenous, single-stranded RNAs (21-25 nucleotides in length) that regulate gene expression either post-transcriptionally, by blocking translation or promoting cleavage of specific target mRNAs, or transcriptionally, through targeting of promoter regions^[7]. Thus, they do not code for proteins, but

instead serve to regulate the expression of certain genes. More than 2500 miRNAs may be encoded in the human genome, residing in intergenic regions, introns, and within exons^[8]. MiRNAs go through a complex processing pathway following transcription by RNA polymerase II (RNA Pol II), which has been described in detail elsewhere^[9]. MiRNAs can be found both in cells and circulating in the blood, and have the potential to be taken up at sites distant from the cell of origin; as such they may serve as biomarkers of disease processes. They may also act in an endocrine or paracrine fashion to regulate expression at multiple sites. It is not yet known whether miRNAs have specific cell surface receptors or whether they target cells that express their target mRNAs.

MiRNAs have been found to regulate processes relevant to the development and progression of NAFLD; however, our understanding of the biology underlying these processes is presently in its infancy. The goal of this review is to present an overview of the current state of knowledge of miRNAs involved in the development of NAFLD across a range of appropriate *in vitro* and *in vivo* models, with a special focus on miRNAs that contribute to pathological mechanisms related to fatty liver in humans.

In vitro studies

A number of studies have been conducted on miRNAs using cell culture models of NAFLD, primarily those derived from hepatocyte cell lines (Table 1). In one study, immortalized human liver-derived L02 cells, cultured with high levels of free fatty acids (HFFA-treated) to serve as a model for hepatic steatosis, were analyzed using miRNA micro-array^[10]. A total of 17 and 15 miRNAs were up- or downregulated, respectively, in these HFFA-treated L02 cells. Of these, miR-10b was the most up-regulated miRNA, and HFFA-cultured L02 cells transfected with anti-miR-10b showed significantly decreased lipid content and the triglyceride level (*i.e.*, steatosis). Peroxisome proliferator-activated receptor- α (*PPARA*) was identified as a potential target for miR-10b, and expression of both transcript and protein levels of *PPARA* were reduced in steatotic L02 cells. Overexpression of miR-10b in HFFA-cultured L02 cells, led to decreased *PPARA* protein levels, while miR-10b knockdown increased *PPARA*, indicating that this miRNA may regulate the development of hepatic steatosis through mechanisms involving the *PPARA* pathway. Further investigation involving both animal and human studies will be necessary to confirm this relationship.

In an independent study to identify miRNAs involved in the formation of lipid droplets, the human hepatocellular carcinoma-derived cell line Huh7 was transiently transfected with a library of 327 miRNAs^[11]. The Huh7 cell line spontaneously accumulates lipid droplets in culture and is lipogenic, making it an

Table 1 Summary of *in vitro* studies of miRNAs relevant to fatty liver

Model	miRNAs	Biological effect	Validated targets	Ref.
L02	miRNA-10b	Inc expr, decrease steatosis	PPARA	[10]
Huh7	miR-181d	Inc expr, decrease steatosis	Not determined	[11]
Huh7 and Hep3B	miR-122	Dec exp, decrease steatosis	SOCS3	[14]
HepG2	miR-613	Incr exp, increase steatosis	LXR α	[15]
HepG2 and primary human hepatocytes	miR-107	Incr expression, increase steatosis	FASN	[16]

appropriate *in vitro* model for steatosis^[12]. Following primary and secondary screening, eleven miRNAs were identified that either increased or decreased intracellular lipid content. Of these, miR-181d showed the strongest influence on steatosis, decreasing lipid droplet formation by approximately 60%. Huh7 cells were also used, along with Hep3B human hepatocellular carcinoma cells, to link expression of miR-122 to decreased fatty acid and cholesterol levels^[13]. Expression of the suppressor of cytokine signaling 3 (SOCS3) gene is also regulated by miR-122^[14]. SOCS3 protein increases expression of sterol regulatory element-binding protein 1 (SREBP1), a transcription factor that regulates cholesterol and lipid metabolism. Silencing of miR-122 in Huh7 cells corresponded with reduced SOCS3 expression, which in turn decreased SREBP1 levels, while restoration of SREBP1 expression when miR-122 levels were depleted through RNA silencing could be achieved by over-expression of SOCS3.

In addition to L02 and Huh7 cells, HepG2 cells have also been used as a model in which to study the role of miRNAs in hepatic lipid metabolism. In these cells, over-expression of miR-613 reduced expression of the nuclear receptor liver X receptor α (LXRA) and several of its target genes including acetyl-CoA carboxylase, sterol-regulatory element binding protein 1c, and fatty acid synthase, and led to the formation of lipid droplets^[15]. miR-613 was shown to bind to the 3'-untranslated region of the LXRA mRNA. Similarly, miR-107 was shown to bind to the 3' UTR of the fatty acid synthase gene (FASN), reducing its expression and causing malonyl CoA and lipid accumulation in both HepG2 cells and primary hepatocytes^[16].

As evident from the studies described, the inherent advantage of cultured cell systems for the study of miRNA physiology in NAFLD is in their ease and economy of manipulating levels of specific molecules. However, NAFLD occurs in the context of multiple cell types that constitute the liver and with molecular interactions with distant cell types and organs; therefore studies involving *in vivo* models have been important in enhancing our understanding of the role miRNAs play in the development of the disease.

***In vivo* studies**

A number of studies of miRNAs have been conducted

using animal models of fatty liver, primarily in mice and rats^[17], in whom NAFLD is typically induced with a high fat diet. A summary of the main findings from these studies is shown in Table 2. In one study, microarray analysis of 350 miRNAs in liver samples of sprague-dawley rats with diet-induced NASH showed downregulation of miR-122, miR-451, and miR-27a and upregulation of miR-429, miR-200a, and miR-200b compared to animals fed a standard diet^[18]. In a similar microarray-based study of diet-induced NASH in sprague-dawley rats, the authors observed that upregulation of miR-146a, miR-210, miR-29c, miR-103, miR-20b-5p, miR-106b, miR-212, miR-31, miR-10a, miR-203, miR-27b, miR-199a, miR-107, let-7b, and downregulation of miR-33, miR-145, miR-196b, miR-93, let-7d, miR-19 could differentiate between steatohepatitis and steatosis^[7]. No common mRNA targets were found for the 14 upregulated miRNAs, but 12 common targets were found for the six downregulated miRNAs including stearoyl-coenzyme A desaturase 1 (Scd1). Hepatic expression of miR-15b was also shown to be upregulated in liver RNA of Sprague Dawley rats fed a high fat diet for 16 wk^[19]. Surprisingly, no miRNAs were replicated across these rat studies, possibly due to differences in dietary composition and regimen, as well as phenotypic endpoint.

MiRNA microarray analysis of liver RNA from both C57BL/6J and DBA/2J inbred strains of mice fed a lipogenic methyl-deficient diet to induce a form of fatty liver injury similar to human NASH identified significant upregulation in the expression of miR-34a, miR-155, and miR-200b, and downregulation of miR-29c^[17]. A strain-specific effect was seen, with more significant changes occurring in DBA/2J mice.

In livers of C57BL/6J mice fed a high fat diet for eight weeks, miR-467b expression was significantly decreased, corresponding to an increase in hepatic lipoprotein lipase (LPL) expression^[20]. The authors utilized bioinformatics sequence analysis to identify LPL as a direct target of miR-467b and confirmed the miRNA-mRNA interaction *in vitro*. Interestingly, the interaction between miR-467b and its target gene was associated with insulin resistance, which strongly increases the risk of NAFLD. In a separate study, apoE(-/-) mice treated with intra-peritoneal injection of an miR-467b mimic or agomirna (synthetic chemically modified RNA duplexes) led to reduced lipid accumulation and inflammatory cytokine secretion by macrophages *via* downregulation of LPL expression,

Table 2 MiRNAs associated with nonalcoholic fatty liver disease

miRNA	Species	Model	Ref.
Downregulation of miR-122, miR-451, and miR-27a and upregulation of miR-429, miR-200a, and miR-200b	Sprague-dawley rats	High fat diet induced NASH <i>vs</i> standard diet	[18]
Upregulation of miR-146a, miR-210, miR-29c, miR-103, miR-20b.5p, miR-106b, miR-212, miR-31, miR-10a, miR-203, miR-27b, miR-199a, miR-107, let-7b, and downregulation of miR-33, miR-145, miR-196b, miR-93, let-7d, miR-19 miR-15b	Sprague-dawley rats	High fat diet induced NASH <i>vs</i> steatosis	[7]
	Sprague-dawley rats	High fat diet induced NASH <i>vs</i> standard diet	[19]
Decreased expression miR-29c, and increased expression miR-34a, miR-155, and miR-200b	C57BL/6J and DBA/2J mice	methyl-deficient diet induced NASH	[17]
Decreased miR-467b	C57BL/6J mice	High fat diet <i>vs</i> standard diet	[20]
Increased miR-103 and miR-107	C57BL/6J mice	High fat diet <i>vs</i> standard diet	[22]
Decreased miR-21	C57BL/6J mice	High fat diet <i>vs</i> standard diet	[23]
miR-122 decreased in all strains, while expression of miR-34a, miR-200b, and miR-181a	A/J, C57BL/6J, C3H/HeJ, 129S/SvImJ, CAST/EiJ, PWK/PhJ, and WSB/EiJ strains	choline- and folate-deficient diet	[24]
miR-122 (-/-)	129SvJ	Steatosis on a Normal Diet	[25]
miR-155 (-/-)	C57BL/6	High fat diet <i>vs</i> standard diet	[26]
Up regulation of miR-122, miR-24, miR-195a, miR-106b, miR-15b, miR-802, miR-185, miR-214, miR-378, and let-7c; downregulation of miR-224, miR-126, miR-7a, miR-128, miR-455, miR-452, miR-135b, miR-145, miR-18a, and miR-196a	<i>ob/ob</i>	Standard diet	[27]
Increased miR-16, miR-122, miR-126 decreased miR-27b	Gankyrin transgenic zebrafish	Standard diet	[29]

NASH: Non-alcoholic steatohepatitis.

leading to protection from atherosclerosis in these animals^[21]. Together, these findings suggest that the miR-467b-LPL interaction may play an important role in lipid accumulation, which may exert diverse effects on the development of both hepatic steatosis and atherosclerotic vascular disease.

In LDL receptor knockout (LDLR^{-/-}) mice derived from a C57BL/6J background and fed a high-fat diet for 10 wk, increased expression of miR-103 and miR-107 was abolished by daily dosing of a mixture of concentrated plant-derived polyphenol compounds^[22], although weight gain and liver steatosis were ameliorated. The expression of miR-122 was not altered by the high fat diet, but was decreased by dietary polyphenols. Further studies on polyphenol administration for 8 wk to C57BL/6J mice fed a high-fat diet demonstrated that lycopene also ameliorated hepatic steatosis and prevented down-regulation of miR-21^[23]. Expression of fatty acid-binding protein 7 (FABP7) was downregulated *via* interaction of miR-21 with the FABP7 3' UTR.

Administration of a choline- and folate-deficient diet for 12 wk to induce NAFLD-like liver injury in inbred male mice of the A/J, C57BL/6J, C3H/HeJ, 129S/SvImJ, CAST/EiJ, PWK/PhJ, and WSB/EiJ strains induced strain-related differences in levels of hepatic and plasma miRNAs^[24]. Hepatic expression of miR-122 decreased in all strains, while expression of miR-34a, miR-200b, and miR-181a increased and was correlated with histological severity of liver injury. Serum levels of miR-34a, miR-122, miR-181a, miR-192, miR-200b, and miR-221 were correlated with histological severity across all strains, providing evidence that release of miRNAs may serve as

biomarkers of liver injury.

Knockout studies have also been conducted targeting specific miRNAs, although few have been reported thus far. A recent study by Hsu *et al.*^[25], in which mice with either germline or liver-specific knockdown of miR-122 were generated, showed that deficient animals developed steatosis, steatohepatitis, fibrosis, and spontaneous tumors that were histologically similar to hepatocellular carcinoma. These findings support a critical role for miR-122 in mediating the progression of steatosis to more clinically severe phenotypes and the subsequent development of cancer.

An independent study also using knockout mice investigated the role of a second miRNA, miR-155, in the development of hepatic steatosis^[26]. In that study, miR-155^{-/-} mice fed a high fat diet for six months developed significantly more hepatic steatosis, which was associated with increased liver weight and lipid levels, than C57BL/6 wild-type controls. Hepatic expression of genes involved in glucose regulation, fatty acid uptake, and lipid metabolism were also elevated in the miR-155^{-/-} mice. Among the differentially expressed genes, the authors identified and validated only one, Nr1h3 (LXR α) as a direct target of miR-155. Together these data indicate that miR-155 plays a protective role in liver lipid metabolism and that downregulation of miR-155 expression may contribute to the development of hepatic steatosis.

In addition to microarray and knockdown studies of specific miRNAs, approaches using next generation sequence analysis have also been used to quantify miRNA expression in NAFLD. Sequencing of liver

non-coding RNA in *ob/ob* and control mice identified 37 differentially expressed hepatic miRNAs^[27]. Although miR-122 showed the greatest alteration in expression between the two groups, miR-24, miR-195a, miR-106b, miR-15b, miR-802, miR-185, miR-214, miR-378, and let-7c were also significantly upregulated. In contrast, levels of miR-224, miR-126, miR-7a, miR-128, miR-455, miR-452, miR-135b, miR-145, miR-18a, and miR-196a were significantly downregulated. To determine whether overexpression of miR-126 or inhibition of miR-24 played a mechanistic role, AML-12 liver cells were treated with free fatty acids. Up-regulation of hepatic miR-126 using a miR-126 mimic or down-regulation of hepatic miR-24 using antagomiR-24 was correlated with decreased fat accumulation, suggesting that both may potentially mediate liver steatosis. Additional studies will be necessary to address this possibility.

In addition to mice and rats, zebrafish are presently becoming recognized as suitable models for lipid-related diseases, including hepatic steatosis^[28]. In this model, transgenic over-expression of gankyrin, a small ankyrin-repeat protein that plays a role in cellular proliferation, led to increased lipid content in > 90% of viable adult fish. Overexpression of gankyrin led to the development of hepatic steatosis and was associated with increased levels of miR-16, miR-122, and miR-126, and decreased miR-27b^[29]. This study provides evidence supporting a link between gankyrin and miRNAs in modulating the development of hepatic steatosis in zebrafish; however, the role of this network in humans is not yet known.

HUMAN STUDIES

A large number of studies on the role of miRNAs in viral hepatitis and hepatocellular carcinoma in humans have been reported^[30], which is in contrast to the relatively limited investigations of these molecules in modulating the pathogenesis of NAFLD and liver-related outcomes. Of the studies that have been published, most have been performed using very small sample sizes, thereby limiting the impact of any conclusions drawn from the results. For example, one study recently profiled liver miRNA in 15 individuals with NASH and 15 individuals with normal liver histology^[31]. Out of the 474 miRNAs represented on the array, six showed differential expression between the two groups. The authors confirmed overexpression of miR-34a and miR-146 b and underexpression of miR-122 using RT-PCR, although miRNA levels were not associated with NASH severity. Examination of miR-122 target genes, including SREBP-1c, FAS, and HMG-CoA reductase, showed significant increases in mRNA and protein levels in individuals with NASH, which is consistent

with *in vitro* findings in HepG2 cells following miR-122 silencing. Similarly, an inverse correlation between miR-122 and levels of SOCS3 and SREBP1 was observed in human liver samples^[13]. In obese individuals, levels of miR-34a were approximately 2-fold higher in mild NASH, increasing to more than 3-fold higher in severe NASH relative to steatosis^[32]. In comparisons of steatotic and severe NASH liver samples, levels of miR-122, miR-143, and miR-451 were decreased.

In addition to studies using liver tissue, miRNA levels in adipose tissue have also been investigated. A total of 664 miRNAs were profiled in visceral adipose tissue obtained from 12 extremely obese bariatric surgery patients with biopsy-proven NASH and 12 with without NASH^[33]. Expression of miR-132, miR-150, miR-433, miR-28-3p, miR-511, miR-517a, and miR-671-3p were all significantly decreased in individuals with NASH. In addition, expression of miR-197 and miR-99 were also decreased in NASH peri-sinusoidal fibrosis compared to non-fibrosis. *IL6* was identified as a target gene for all seven miRNAs, and the authors found that serum IL6 levels were inversely correlated with levels of these candidates. A study with a similar sample size was also conducted using visceral adipose tissue obtained from patients undergoing bariatric surgery^[34]. In that study, *Drosha*, *DGCR8*, and *Dicer1*, all of which represent key components of miRNA processing, and seven pri-miRNAs including pri-miR-125b-2, pri-miR-16-2, pri-miR-26a-1, pri-miR-26a-2, pri-miR-7-1, pri-miR-7-2, and pri-miR-7-3 were assayed. Of these, levels of *Dicer1*, *Drosha*, *DGCR8*, and pri-miR-7-1 were significantly increased in NASH patients compared to normal controls. These results indicate that even in the context of severe obesity, specific miRNAs may serve to differentiate liver function, although the small sample sizes of these studies limit the generalizability of the results.

In addition to expression in tissue, miRNAs can also circulate freely in blood or be packaged within microvesicles that provide a high level of protection from degradation. Some studies report a correlation of miRNA levels in tissue and biofluids. A summary of findings from studies of circulating miRNAs in NAFLD is shown in Table 3. For example, levels of miR-122 in serum and liver were significantly correlated ($R = 0.461$; $P = 0.005$) in patients with NAFLD^[35], suggesting that miR-122 released from hepatic cells enters the bloodstream. Serum levels of miR-122 were lower in individuals with mild steatosis, compared to those with severe steatosis, but higher in patients with mild fibrosis compared to those with severe fibrosis. This result is in agreement with those of previous studies, reporting decreased levels of hepatic miR-122 at advanced stages of fibrosis in patients with liver disease^[36]. The reason for the

Table 3 Changes in circulating miRNAs associated with nonalcoholic fatty liver disease in humans

Population	Study design	Source of miRNA	miRNAs	Effect	Ref.
NAFLD	Mild <i>vs</i> Severe steatosis; Severe <i>vs</i> Mild fibrosis	Serum	miR-122	Decrease	[35]
NAFLD	NAFLD <i>vs</i> normal	Serum	miR-122, miR-34a, miR-16, and miR-21	Increased	[36]
NAFLD	NAFLD <i>vs</i> normal	Serum	miR-15b	Increased	[19]
NAFLD	NAFLD <i>vs</i> normal	Serum	miR-122, miR-192, miR-19a, miR-19b, miR-125b, and miR-375	Increased	[37]

NAFLD: Nonalcoholic fatty liver disease.

discrepancy in miR-122 levels in NAFLD stage may represent the loss of hepatocytes in worsening liver injury. Because hepatocytes are the primary source of miR-122 and since worsening of liver fibrosis results in the replacement of hepatocytes with extracellular matrix, hepatic miR-122 levels may be expected to decrease with severe fibrosis. These results indicate that levels of miR-122 may have significant prognostic value for patients with NAFLD.

Similarly, Cermelli *et al.*^[36] investigated serum levels of four miRNAs commonly dysregulated in liver fibrosis: miR-122, miR-34a, miR-16, and miR-21. In a study sample comprised of 34 individuals with NAFLD and 19 healthy controls, serum levels of miR-122, miR-34a, and miR-16 were significantly higher in NAFLD patients. Levels of miR-21 showed no difference between the two groups. Interestingly, levels of miR-122 and miR-34a were positively correlated with disease severity from simple steatosis to steatohepatitis, supporting the potential value of these two miRNAs to serve as noninvasive biomarkers for progressive NAFLD.

Zhang *et al.*^[19] also recently examined miR-15b as a potential biomarker for NAFLD in 69 individuals with fatty liver and 42 healthy controls. Levels of miR-15b were higher in the NAFLD patients compared to the control group. However, because there were significant differences in BMI, blood glucose, triglyceride levels, total cholesterol, and ALT between the two groups, these findings must be interpreted with caution. Additional studies will be necessary to demonstrate the appropriateness of miR-15b as a biomarker for fatty liver disease.

A case hyphen control, multi-phased study that analyzed 84 miRNAs in serum of patients with biopsy-proven NAFLD and healthy controls identified a greater than 2-fold up-regulation of miR-122, miR-192, miR-19a, miR-19b, miR-125b, and miR-375 with steatosis or more advanced NAFLD^[37]. Only miR-122 was associated with severe *vs* no or mild fibrosis. Interestingly, miR-122 was 10-fold and miR-192 2-fold down-regulated in the liver. In situ hybridization of miR-122 in liver showed that the miRNA staining was concentrated at the hepatocyte membrane, not more broadly distributed throughout the cytoplasm, consistent with preparation for export

to the circulation.

CROSS STUDY COMPARISONS

About a dozen miRNAs have been analyzed in two or more model systems (Table 4). miR-122, the most studied, was also the most discrepant across systems. miR-122, the most abundantly expressed miRNA in hepatocytes has been associated with a variety of liver diseases^[38], suggesting complex regulation. Indeed, each miRNA may play a role in regulating the expression of tens to hundreds of genes, which may be regulated by multiple miRNAs. This complex regulatory landscape may thus have species specificity, accounting for differences within or across model systems. Mechanistic differences in the protocols for inducing NAFLD may also be reflected in differential miRNA expression. For example, high fat diet induced NAFLD has substantial differences with a choline- and folate-deficient diet that are very different than dyslipidemic mouse knockout models. Indeed, evidence suggests strain background also has substantial effect on miRNA levels, further implicating complex regulation.

The discrepancy between a decrease observed for human liver and an increase in circulating miR-122 is more difficult to reconcile, although little is known about the metabolism of miRNAs in circulation. In addition, serum levels will reflect the total body levels of the miRNA, thus contributions from other tissues may compensate for decreased levels in the liver. The relative severity of NAFLD may also be related to the levels in the serum, particularly if there is significant ballooning degeneration and hepatocyte cell death with subsequent release of miRNAs.

Despite the discrepant results, the change in expression with NAFLD of 10 of the 12 miRNAs were found to be in the same direction, although seven were only studied in rats and mice. However, three zebrafish miRNAs were concordant with those found in either mice or rats. The general agreement in results between two rodent species, as well as between rodents and zebrafish, suggests that the biology may be conserved. Cross-model concordance between evolutionarily distant species

Table 4 Cross-study comparison of miRNA changes with nonalcoholic fatty liver disease

	Cell lines	Mice	Rats	Zebrafish	Human liver	Human serum
MiR-122		INC/NC/DEC	DEC	INC	DEC	INC
miR-34a	INC/DEC	INC	INC		INC	
miRNA-107		INC	INC			
miR-29c		INC	INC			
miR-200b		INC	INC			
miR-103		INC	INC			
mir-106b		INC	INC			
miR-15b		INC	INC			
miR-126		DEC		DEC		
MiR-145		DEC	DEC			
miR-451			DEC		DEC	
miR-27b			INC	INC		

also suggests functional relevancy. Unfortunately, few human data are available for comparison with animal model results. The need for additional human data is largely limited by the availability of tissue samples. Future studies based on large cohorts of well annotated and high quality biological samples and clinical data are needed.

CONCLUSION

miRNAs may act either independently or interactively with environmental exposures and lifestyle factors to affect susceptibility to hepatic fat accumulation. Thus, the link between miRNA and progression of fatty liver disease represents a key area of focus for research endeavors in the development of novel therapies targeting control and prevention of NASH. Given the substantial public health burden of NAFLD, which is increasing at alarming rates due to the rising prevalence of obesity, novel therapeutic targets are urgently needed to facilitate the development of improved pharmacological therapies for the treatment and prevention of the disease.

In NAFLD, a small number of miRNAs, most notably miR-122, have emerged as potential participants in the regulation of biological processes relevant to the disease. However, much less is known of the specific targets of candidate miRNAs, and how, in fact, they affect disease development and progression and variability in liver-related outcomes in affected individuals. Studies aimed at delineating specific miRNA/mRNA networks will enhance our understanding of the complex pathogenesis of NAFLD and enable exploitation of relevant miRNAs as novel targets for therapeutic interventions. Notably, identification and validation of circulating miRNA signatures may facilitate the development of improved methods for diagnosis and clinical monitoring of disease progression. At present, current findings, combined with the rapidly expanding field of miRNA research, are expected to yield new insights into the complex pathogenesis of NAFLD and may eventually

lead to the identification of novel, noninvasive biomarkers for the disease.

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Image-guided therapies in the treatment of hepatocellular carcinoma: A multidisciplinary perspective

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thermal ablation, transarterial chemoembolization, and radioembolization whilst stereotactic body radiation therapy also uses imaging to target the radiation. Both survival rates and cure rates have improved markedly since the introduction of these techniques. This review article describes the image guided techniques used for the treatment of HCC.

Key words: Ablation; Chemoembolization; Radiation; Hepatocellular carcinoma

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Core tip: This review article provides an updated description of the image guided therapies for hepatocellular carcinoma including stereotactic radiation, set in the context of a multidisciplinary approach.

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INTRODUCTION

Hepatocellular carcinoma (HCC) is the fifth most common cancer worldwide and the third most common cause of cancer related deaths^[1]. Treatment depends on the stage of the tumor, performance status, and liver function, as well as on the multidisciplinary capabilities of the managing team of hepatologists, gastroenterologists, surgeons, radiologists and oncologists. Curative resection, liver transplantation, ablative therapies, trans-arterial chemoembolization (TACE), radioembolization and systemic therapy all

Abstract

A multidisciplinary approach to the treatment of patients with unresectable hepatocellular carcinoma (HCC) has led to improvements in screening, detection, and treatments. Interventional techniques include

lie within the range of treatments available to this team^[2,3].

In recent years surveillance strategies for patients with viral hepatitis or with cirrhosis have improved, leading to earlier diagnosis in many patients. These patients have a chance of gaining a curative response to treatment^[4,5]. In contrast, delay in treatment leads to worse survival^[6]. Resection remains the first option for patients who are suitable for surgery, as defined by the Barcelona Cancer of the Liver Clinic (BCLC) staging system. However, several different image guided minimally invasive therapies have emerged and evolved to improve the treatment of HCC at an early stage. These complement therapies provided by surgical and radiation/oncology services. Selection of treatment pathways is determined by a multidisciplinary approach^[7-9] and is most commonly based on the BCLC staging system^[10]. For early and intermediate stage hepatocellular carcinoma (HCC) (stages A and B) locoregional treatments including ablative therapies and TACE are used. Radioembolization is used for intermediate and advanced stage HCC who are poor candidates for TACE, and who have portal vascular invasion^[11]. It is also used with limited evidence base for the downstaging of tumors so that more curative treatments can be employed^[12,13].

For patients who have either failed locoregional therapies or who present with more advanced HCC, Sorafenib induces a clinically relevant improvement in time to progression and in survival.

ABLATIVE THERAPIES

Liver transplantation and surgical resection remain the primary options for curative treatment in appropriate patients. The Milan criteria^[14] provide strict guidelines for transplantation eligibility, whilst surgical resection is suitable only for patients with single nodules and Child Pugh class A liver function^[15]. The limitations on these treatment options offer up a substantial number of patients who can benefit from locoregional therapies. Radiofrequency ablation has become the most accepted treatment for patients with very early and early stage (BCLC 0 and A) disease who are not eligible for surgery^[16,17]. In three independent meta-analyses^[17-19] which include five randomized controlled trials, better local control and increased survival has been demonstrated in comparison with percutaneous ethanol ablation. When compared with surgical resection, there is conflicting evidence. In randomized controlled trials Huang *et al.*^[16] indicate improved results for surgery over RFA who were followed up for 5 years while Feng *et al.*^[20] showed that although there was a greater risk of local recurrence with RFA, there was no significant difference in overall survival. Similar conflicts are demonstrated in meta-analyses. Liu *et al.*^[21] found equivalent survival rates despite higher rates of

local recurrence with RFA, whilst Zhou *et al.*^[22] found better survival rates in surgical patients for tumors measuring greater than 3 cm, and equivalent rates in smaller tumors. Survival ranges from 78%-94% at 1 year and 58%-96% at 3 years^[17-22].

RFA employs low-voltage alternating current to provide sufficient heat to kill cells^[21-24]. The probes are inserted under ultrasound or computed tomography (CT) guidance. The procedure is performed under moderate sedation or general anesthesia and patients can be discharged on the same day or the following day. Complication rates are lower than those of surgery and include abscess formation, tumor seeding along the electrode track, burns from the grounding pads, bile duct injury and thermal injury to adjacent organs^[25]. The procedure is also less expensive than surgery.

Cryoablation is similar in terms of technical approach to RFA, but creates tissue injury from low temperatures of -20 °C to -60 °C^[26]. More than one needle is usually required. The procedure can be applied with lower rates of complication than RFA when close to the gall bladder^[27] or bowel loops, and is less painful when employed for lesions which are contiguous with the diaphragm^[28]. The procedure can be performed under moderate sedation^[29]. It is possible to follow the ablative effect on CT by visualization of the ice ball^[23,28-30]. One and 3 year survival rates are demonstrated at 81.4% and 60.3%, similar to those of RFA^[30]. A single meta-analysis shows an advantage for RFA over cryotherapy in terms of recurrence rate^[28]. There is no study comparing the survival rates.

Microwave therapy also works by heating the local tissues. It achieves a larger ablation zone in a shorter period of time than RFA^[31]. Early studies have shown that there is a larger rate of local recurrence with microwave than with RFA, but there are no large studies or randomized studies to support this.

With the advent of RFA, percutaneous ethanol injection has decreased in popularity. The procedure is low cost, but requires several sessions of treatment. It is performed with a fine needle under ultrasound guidance. Tumor recurrence rates and survival rates are inferior in comparison with those of RFA^[32].

CHEMOEMBOLIZATION

HCC is preferentially supplied by the hepatic arterial inflow, in contrast to the normal liver parenchyma which is largely supplied by the portal vein. The TACE procedure exploits these blood supply dynamics. Techniques vary according to resources and expense, but the principal is that an intra-arterial catheter is placed in the vessel(s) supplying the tumor(s) and high concentrations of a chemotherapeutic agent is delivered along with an embolic agent to achieve the dual purposes of targeted chemotherapy and reduction in arterial supply to the tumor.

TACE has been performed since 1980. Chemotherapeutic drugs, most commonly doxorubicin, cisplatin and mitomycin, are delivered locally along with an embolic agent, normally lipiodol, an oil emulsifying agent, thereby avoiding systemic toxicity. Other embolic agents used are gelatin sponge and PVA particles.

Drug eluting beads (DEB-TACE), although not yet the standard, are becoming increasingly popular largely due to the decreased side effect profile in comparison with the standard TACE cocktail of drugs. DEB-TACE delivers small beads which have been soaked for several hours, normally in doxorubicin. The loaded beads occlude the feeding vessels of HCC, while the anticancer drug is released gradually, creating tumor necrosis and increasing chemotherapeutic concentrations locally. Bead size varies from 75 micron to 700 micron, the choice of size being dependent on tumor size and the desired level of concentration within the treated volume. Improved results are achieved when chemoembolization is performed selectively to segmental or subsegmental arteries feeding the tumor(s)^[33].

TACE is recommended as the standard of care for intermediate stage HCC without vascular invasion or distant metastases. Although there has been some heterogeneity in the results of several randomized controlled trials, TACE has been shown to achieve at least a partial response in 15% to 62% of patients, and improves survival from 16 mo to 20 mo^[34-40]. The variability in results is likely explained by the fact that intermediate stage HCC covers a broad spectrum of disease burden, that there is variability in the chemotherapeutic agents and embolization materials administered to patients, and that the procedure is performed on both Childs A and Childs B liver disease populations. DEB-TACE has been shown to achieve improved outcomes in patients with Child-Pugh B, bi-lobe disease and recurrent disease^[41].

There remains debate about the optimal degree of arterial embolization to achieve tumor ischemia^[42,43]. There is some evidence which indicates that complete tumor ischemia may stimulate angiogenesis, resulting in an increased susceptibility to tumor growth rather than suppression. It is therefore suggested that arterial patency be maintained, not only to prevent this angiogenic effect, but also so that patients can receive repeated treatments^[44].

DEB-TACE causes fewer side effects than conventional TACE. Side effects associated with both DEB-TACE and conventional TACE include nausea, vomiting and right upper quadrant pain (post embolization syndrome), cardiac toxicity related to the doxorubicin, bone marrow aplasia, hepatic abscess and cholecystitis^[36,38,45]. Two recent randomized controlled trials have shown improved side effect profiles^[46,47]. One trial showed equivalent survival rates^[42,46] whilst the

other showed longer time to progression for DEB-TACE in comparison with conventional TACE^[47]. A single meta-analysis demonstrated equivalent tumor response rates^[48].

RADIOEMBOLIZATION

Radioembolization for primary hepatic cancer with Yttrium-90 (90Y) was first described in 1965 who used isotope embedded 50 μm ceramic microspheres to embolize hepatic cancer *via* a surgically placed catheter based in the hepatic artery^[49]. Today, the technique has evolved away from an open surgical approach to a minimally invasive fluoroscopically guided microcatheter based technique using either 90Y embedded non-biodegradable glass microspheres measuring $25 \pm 10 \mu\text{m}$ (Theraspheres, Nordion Incorporated, Ottawa, Ontario, Canada) or 90Y embedded non-biodegradable glass resin based microspheres measuring 29-35 μm (SIR-Spheres, Sirtex Medical Incorporated, Lake Forest, Illinois). Radioembolization, similar to TACE, exploits the preferential arterial blood supply of an HCC by delivering radiotherapy in an embolic agent directly to the tumor bed while preserving the blood flow to the normal liver parenchyma, which is supplied primarily by the portal vein. Unlike TACE, which uses 75-700 μm beads to occlude medium to large sized arteries leading to tumor ischemia, 90Y radioembolization uses these smaller beads to act as a microembolic agent to deposit radiotherapy directly within the tumor *via* an intratumoral vessel. Once deposited at the target lesion, 90Y delivers tumoricidal doses of a pure high energy beta emitter (937 KeV) with a short tissue penetration (mean 2.5 mm and maximum 11 mm) and short half-life of 2.67 d. The short tissue penetration and half-life of 90Y make it an ideal radioisotope for intra-arterial radiotherapy as there is minimal dose deposited in the adjacent liver parenchyma and the patient can immediately be safely discharged home without fear of radiation being delivered to others.

Radioembolization with 90Y has generally been reserved for patients who have intermediate/advanced BCLC stage hepatocellular carcinoma and who are not candidates for TACE due to portal vein invasion^[10,50,51]. Sorafenib is generally considered the treatment of choice for advanced HCC^[52]. However, Sorafenib is often not well tolerated^[53], and 90Y radioembolization is a suitable alternative for patients with advanced HCC given the equivalent median overall survival of 13.2 mo in the radioembolization group vs 14.4 mo in the Sorafenib group^[54]. 90Y radioembolization has also been proposed as an alternative treatment option to prevent progression of disease in eligible transplant patients and to downstage patients in order to become eligible transplant recipients based on the Milan criteria^[12-14].

According to the Radioembolization Brachytherapy

Oncology Consortium, it is recommended that patients undergo preembolization planning and treatment simulation with intrarterial injection of technetium-99m labeled macroaggregated albumin (^{99m}Tc-MAA) and CT to rule out > 30 Gy radiation exposure to the lung from hepatopulmonary shunting and to measure liver volumes^[51]. There is currently no consensus on the recommended radiation dose to deliver to effectively treat HCC. In patients with advanced stage inoperable HCC, however, Lau *et al.*^[55] did demonstrate a median survival benefit of 55.9 wk vs 26.2 wk in patients receiving > 120 Gy vs < 120 Gy, respectively. A randomized controlled trial is underway examining the efficacy of radioembolization when compared with chemoembolization (Seinstra)^[56].

COMBINATION TACE AND RFA

Ablative techniques demonstrate diminished efficacy when tumor diameter is greater than 3 cm^[57,58]. This failure to achieve complete tumor necrosis is largely attributed to the "heat sink" effect: cooling by blood flow resulting in a reduction in temperature adjacent to vessels within or adjacent to the ablation zone^[59].

Adjuvant locoregional therapies have been employed to achieve higher rates of efficacy in the treatment of larger tumors (3-5 cm). The most common of these is chemoembolization. The embolic effect of the lipiodol or beads decreases the "heat sink" effect caused by local vessels, whilst the addition of the chemotherapeutic drug improves overall tumor kill efficacy^[60-62]. A single randomized controlled trial has shown decreased rates of tumor progression in the combination group in comparison with the RFA only group^[63], although no significant difference in survival was demonstrated.

SURGICALLY ASSISTED RFA

Radiofrequency ablation was widely adapted in the 1990's as a method to treat lesions deemed unresectable at the time of open hepatectomy. As technology improved, RFA moved from the operating room to the IR suite where percutaneous ablations could be performed without the morbidity of a laparotomy. While percutaneous image guided ablative therapies are a useful tool in the armamentarium for the loco-regional treatment of liver lesions, there are some limitations. These include difficulty in localizing lesions, potential for injury to extra hepatic structures and decreased efficacy in close proximity to liver vasculature. Many of these limitations can be addressed by performing surgically assisted RFA using a laparoscopic approach^[64].

Because percutaneous ablation relies on the ability to localize lesions with ultrasound, obese patients with thick abdominal walls can provide a challenge, particularly with lesions in the dome.

While this has in large part been abrogated by use of CT and magnetic resonance imaging (MRI) guidance, some tumors are difficult to localize on cross sectional imaging. The ability to perform ultrasound directly on the liver surface allows for more accurate tumor localization and may result in more efficacious tumor treatment when compared to percutaneous ablation^[65,66]. The most widely used technique for laparoscopic assisted RFA is with insufflation of the abdomen after induction of general anesthesia. A laparoscopic ultrasound probe is then introduced and used to guide a percutaneously placed RFA needle. Additional laparoscopic ports can be introduced to manipulate the liver as well as other extra-hepatic structures. The ability to manipulate the peri-hepatic environment can protect structures such as the colon, stomach, small bowel and diaphragm from transmitted heat. It also allows for potential removal of the gallbladder prior to RFA, preventing injury and heat sink. Other techniques to protect peri-hepatic structures include instillation of artificial ascites which can absorb heat without transmission to surrounding viscera^[67].

An additional benefit of surgically assisted RFA is the ability to occlude hepatic vascular inflow, which in theory reduces the heat sink from major vessels. With minimal mobilization of the liver, a temporary ligature can be placed around the porta hepatis and tightened immediately prior to application of energy. In-vivo animal studies have indicated an increase in tumor necrosis around blood vessels, although human data is lacking^[68,69].

TACE + RADIOTHERAPY FOR HCC

Locoregional relapse remains an important issue for HCC. In early stage HCC stereotactic body radiotherapy (SBRT) has been used in conjunction with TACE in an effort to improve cure rates^[70,71]. In the locally advanced setting, three dimensional conventional radiation therapy (3DCRT) has shown promising results following TACE in promoting tumor necrosis and reducing local relapse.

SBRT entails the delivery of highly conformal, high dose, ablative radiotherapy to a liver lesion in a short period of time (typically over 1-2 wk). Selection criteria for liver SBRT is similar to that of TACE: Childs A liver function, 1-3 lesions, more than 700 cc uninvolved liver, tumors less than 5 cm, and well controlled extrahepatic disease^[72]. Prior to SBRT patients undergo a 4 dimensional CT planning scan to delineate the target lesion and its real time movement across several phases of respiration. Ultrasound guided insertion of tumor fiducial markers is often useful for image guided radiotherapy where the markers act as a surrogate for tracking the lesion's location for radiation delivery. Liver SBRT prescriptions can vary from 50 Gy/5 fractions to

60 Gy/3 fractions, in contrast to 3DCRT where the conventional daily dose of radiation is 1.8 to 2.0 Gy/fraction and the total dose is 45-50 Gy in 25 fractions. Care is taken to avoid excess dose to adjacent bowel and the remaining liver during SBRT given the potential for severe complications with high doses.

Data is also emerging that SBRT may be an effective salvage strategy for patients who experience local failure post TACE. Patients with Childs A disease and tumors measuring less than 10 cm who have undergone partial or incomplete TACE may have 2 year local control rates as high as 94.6% when salvaged with SBRT^[73-75]. High grade toxicity resulting in duodenal or gastric perforation is rare (approximately 5%) if dose constraints are respected^[73]. However, the presence of tumor vascular thrombosis is a risk factor for severe and even mortal toxicity^[76,77]. Combination TACE + SBRT appears to be a potentially promising treatment for early stage HCC and likely merits a multi-institutional phase III study as the existing literature consists of single institution retrospective data or small phase I / II trials^[78].

In locally advanced HCC, 3DCRT or chemoradiation post TACE or partial TACE may confer better outcomes than Sorafenib. One study compared 67 patients with BCLC stage C disease who received TACE + 3DCRT with a cohort that was given Sorafenib as first line treatment. While this study did not examine local control, the median survival of the TACE + RT group was 14.1 mo compared to 3.1 mo in the Sorafenib group^[79,80]. Combining TACE and conventional radiation treatments for locally advanced HCC may also be an effective treatment in patients with extensive portal vein thrombosis^[74,81-85]. One year progression free rates in patients who receive TACE + 3DCRT for unresectable HCC can be as high as 70% compared to TACE alone (40%)^[85]. Patients who have failed 1-2 TACE treatments and who received subsequent 3DCRT have been reported to have as high as 68% response rate post radiotherapy with 70% achieving stable disease at 1 year^[86,87]. As prognosis for locally advanced HCC remains poor, the use of local therapies in conjunction with chemotherapy is also being explored. Clinical trials of concurrent chemoradiation and TACE in advanced disease have shown promise in improving local control and progression free survival^[88,89].

IMAGING AFTER IMAGE-GUIDED THERAPIES

With the variety of image-guided liver directed therapies available, it is important to know the different expected post-therapy appearances and be able to differentiate these from abnormal imaging findings. Contrast-enhanced CT or MRI are the preferred imaging modalities for post

therapy surveillance. Post therapy imaging should be performed at scheduled intervals, although a standard interval has not been established. At our institution, we perform contrast enhanced CT or MRI at 6 wk and then at 3, 6, 9 and 12 mo intervals.

ABLATION

An ablation zone encompasses the tumor with a variable margin, and is therefore usually larger than the tumor on initial imaging. Unenhanced CT and MRI images are obligatory as the ablation zone may be hyperattenuating or have intrinsic hyperintensity on pre-contrast T1-weighted images due to coagulative necrosis and hemorrhage making evaluation for arterial enhancement more difficult. Subtraction MRI is particularly useful when there is T1 hyperintensity on unenhanced images. In a completely treated lesion, contrast enhanced images demonstrate a non-enhancing well-defined ablation zone.

Familiarity with normal periablation changes is also important. Transient hyperemia and edema can be present around the ablation zone due to thermal injury to the surrounding parenchyma, manifesting as a concentric thin rim of enhancement on arterial and sometimes portal venous phase contrast enhanced CT and MRI and a hyperintense rim on T2-weighted images. Peripheral geographic arterial enhancement can also be seen post ablation, often related to injury to the portal vein branches and subsequent increase in perfusion from the hepatic artery. These changes usually resolve within several months^[90,91]. Residual disease in contrast demonstrates an area of irregular or thick, peripheral arterial enhancement.

MRI is particularly helpful for residual disease evaluation, as this demonstrates focal hyperintensity on T2-weighted images and often increased signal on diffusion-weighted images. Recurrent disease has similar imaging characteristics to residual disease, but can occur within or adjacent to the ablation zone. New disease occurs in other areas of the liver or in extra-hepatic locations. In some cases, it may be difficult to differentiate expected a post ablation peripheral rim of enhancement from residual or recurrent tumor. In these cases, closer follow-up imaging may be necessary. Risk factors for residual or recurrent disease include large tumor size, aggressive histology, difficult location, and heat sink effect, specifically in radiofrequency ablation^[92].

Transient bile duct dilatation peripheral to the ablation zone is often seen. Leakage of bile from injured ducts can result in a biloma, which appear as a non-enhancing fluid collection.

If there is injury to larger vessels, parenchymal or intraperitoneal hemorrhage can occur and be detected on CT or MRI. If both the portal and hepatic arteries are injured, hepatic infarction can occur, which

appears as non-enhancing parenchyma peripheral to the ablation zone. Other vascular complications, such as arteriovenous fistula or pseudoaneurysm can also be identified on arterial phase imaging.

Hepatic abscess is an additional complication which can be seen after ablation. This usually presents a few weeks after the procedure and demonstrates peripheral enhancement and development of gas within or adjacent to the ablation zone. Injury to adjacent structures is an additional complication to be aware of after ablation: for example, adjacent bowel or the diaphragm^[92].

TACE, CONVENTIONAL TACE, AND RADIOEMBOLIZATION

Imaging following TACE and transarterial radioembolization is similar to ablation with a few additional caveats. The treated lesion again should demonstrate lack of enhancement, but also may demonstrate a peripheral rim of enhancement, geographic arterial enhancement or both. After transarterial radioembolization, there may be heterogeneous parenchymal enhancement in a perivascular distribution due to radiation effect. This can mimic tumor and may need shorter term follow-up. In patients treated with lipiodol MRI has been shown to be superior to CT given the ability to perform diffuse weighted images and image subtraction^[93]. Residual or recurrent disease appears as nodular arterially enhancing tumor, often in the periphery, similar to ablation.

SBRT

Treatment response assessment for SBRT is evolving. As with other image guided therapies, tumor response after SBRT is recognized as non-enhancement of tumor. However there are other unique imaging characteristics. After SBRT, recurrence can occur within the planned target volume, suggesting that an inadequate dose was used, or can occur along the margin of the high dose region, suggesting incomplete coverage of the tumor margin. This marginal recurrence may be due to patient respiratory motion. Additionally, focal peritumoral enhancement may be seen on any phase of imaging, likely representing radiation induced changes and inflammation of the surrounding normal liver parenchyma. These areas of enhancement can persist for months and should not be confused with recurrent tumor. Additionally, it can take time for the initial tumor enhancement to disappear, and therefore continued follow-up is necessary. Sanuki *et al.*^[94] demonstrated in 38 patients a median time of 5.9 mo to reach complete treatment response with a range of 1.2 to 34.2 mo.

RESPONSE ASSESSMENT

Different criteria have been developed to evaluate tumor treatment response. Conventional size measurement, such as Response Evaluation Criteria in Solid Tumors (RECIST), is predominantly useful for evaluation of cytotoxic systemic agents but does not work well for evaluation after locoregional therapy, as tumor necrosis is the goal and may not always manifest as a decrease in lesion size. The European Association for the Study of the Liver (EASL) measures the arterially enhancing area in two dimensions, while a modified RECIST classification uses a single largest diameter of arterially enhancing tumor. The modified RECIST criteria has been recommended as the preferred criteria for tumor response by the EASL and European Organisation for Research and Treatment of Cancer^[95].

CONCLUSION

Most patients presenting with HCC are ineligible for surgical curative treatment. Advances in locoregional therapy, both catheter based and ablative, have led to improvements both in cure rates and in survival. A multidisciplinary approach is optimal for the planning of treatment given that there are treatment contributions from gastroenterology, surgery, interventional radiology, and oncology.

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Surgical management of hepatocellular carcinoma

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Abstract

Hepatocellular carcinoma (HCC) is the second most common cause of death from cancer worldwide. Standard potentially curative treatments are either resection or transplantation. The aim of this paper is to provide an overview of the surgical management of HCC, as well as highlight current issues in hepatic resection and transplantation. In summary, due to the relationship between HCC and chronic liver disease, the management of HCC depends both on tumour-related and hepatic function-related considerations. As such, HCC is currently managed largely through non-surgical means as the criteria, in relation to the above considerations, for surgical management is still largely

restrictive. For early stage tumours, both resection and transplantation offer fairly good survival outcomes (5 years overall survival of around 50%). Selection therefore would depend on the level of hepatic function derangement, organ availability and local expertise. Patients with intermediate stage cancers have limited options, with resection being the only potential for cure. Otherwise, locoregional therapy with transarterial chemoembolization or radiofrequency ablation are viable options. Current issues in resection and transplantation are also briefly discussed such as laparoscopic resection, ablation vs resection, anatomical vs non-anatomical resection, transplantation vs resection, living donor liver transplantation and salvage liver transplantation.

Key words: Hepatocellular carcinoma; Liver surgery; Liver resection; Liver transplantation; Laparoscopic liver surgery

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Core tip: Surgical management through either resection or transplantation are the only potentially curative treatment for hepatocellular carcinoma. The decision for the management strategy depends on tumour factors, hepatic functional reserve, organ availability, wait time as well as local expertise and resources.

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INTRODUCTION

Hepatocellular carcinoma (HCC) is a prevalent cancer. It is estimated, by the World Health Organisation, to affect 782000 people and caused 746000 deaths worldwide in 2012. In fact, it is the second most

common cause of death from cancer worldwide^[1]. Its incidence is the highest in East and South East Asia, which is related to the prevalence of chronic hepatitis B in these regions. Standard potentially curative treatments for this cancer are either resection or transplantation, although radiofrequency ablation is considered curative therapy in some cases^[2]. Because of its relationship with liver cirrhosis, the assessment of hepatic function is an important consideration in determining management. It is now well established that liver transplantation is the treatment of choice for early stage HCC in patients with decompensated cirrhosis^[3-5].

The aim of this paper is to review the surgical management of hepatocellular carcinoma, as well as highlight current issues in this area.

SURGICAL RESECTION

Only 10%-37% of patients with HCC are amenable to liver resection at the time of diagnosis^[6-8]. An Australian study of 235 patients demonstrated that only 17% and 16% of HCC patients were treated with liver resection or transplantation respectively^[9]. In fact, best supportive treatment was the most common management strategy employed for this cohort. An important aspect of diagnosis in HCC is that there are risk factors which are known to increase the risk of its development. Both the AASLD and EASL-EORTC guidelines recommend 6 monthly surveillance using abdominal ultrasonography in high risk patients, although the definition of which patients are considered high risk varied^[10,11]. Interestingly, a systematic review of the benefits and harms of HCC screening in patients with chronic liver disease found only poor quality evidence to support the benefits of screening. Of the two randomised controlled trials reviewed, only one demonstrated a survival benefit (ultrasound screening) and one found no difference in all-cause mortality (alpha-feto protein screening)^[12].

Improvements in surgical technique and perioperative care have led to a fall in the morbidity and mortality of liver resection over the last two decades. Currently, in high volume centres, the mortality of liver resection is expected to be less than 4%^[13-16]. There is also evidence to suggest that there is a volume-outcome relationship in hepatic resection surgery^[17]. Whilst the 5 years overall survival rates of around 50% is now achievable, the recurrence rate remains high which remains an important cause of late deaths^[18,19].

The decision for liver resection depends on the assessment of tumour factors, hepatic function and remnant size. With regards to tumour factors, one of the most commonly used staging system in Western countries is the Barcelona Clinic liver Cancer (BCLC) system which classifies patients into early,

intermediate, advanced and terminal stages^[20]. This system is utilised by both the American Association for the Study of Liver Disease and the European Association for the Study of Liver guidelines on the management of HCC^[21,22]. According to this system, the surgical treatment of HCC is limited to the early stage cancers, that is, those which satisfy the Milan Criteria (a single HCC \leq 5 cm in diameter or up to 3 HCCs \leq 3 cm in diameter^[5]) and have good hepatic function (Child-Pugh class A in the absence of portal hypertension) and performance status. However, such a criteria for resection is considered restrictive, for size and number of tumours is not a contraindication for resection provided there is adequate hepatic reserve and that the tumour is resectable. Certainly, long term disease-free survival is possible in these patients^[23]. Tumour size and number are not the most important factors influencing survival^[9,19]. In fact, for those patients with HCCs who do not satisfy the Milan Criteria, the only hope of cure is through hepatic resection. Ng *et al.*^[24] demonstrated that large or multinodular HCC could be safely resected, with a five-year overall survival of 39% and disease-free survival of 26% being achievable. In general, tumours which are extensively multifocal and bilateral, involve the main portal vein or inferior vena cava are considered contraindicated for surgery.

Hepatic function can be classified using a variety of measures. The simplest and most commonly used is the Child-Pugh Score^[25]. Resection is really only considered in patients with Child A cirrhosis and early Child B cirrhosis. In the former, up to 50% resection may be considered, whilst in the latter, up to 25% resection may be performed. On the other hand, in patients with entirely normal hepatic function with no history of cirrhosis could tolerate the resection of up to 75% of liver parenchyma^[26]. In Asian countries, the use of ICG clearance at 15 min is also prevalent, with a cut-off of greater than 20% precluding major liver resection^[27,28]. Model of End-Stage Liver Disease (MELD) score is an alternative score used to classify patients into risk groups. A MELD score of $<$ 9 is associated with minimal perioperative mortality^[29,30]. In addition to hepatic function, the other aspect which precludes hepatic resection is significant portal hypertension. This can be objectively measured using a transhepatic caval approach (hepatic vein pressure gradient). This is a measure of the pressure difference between the wedged hepatic venous pressure (an estimation of portal venous pressure) and the free hepatic venous pressure (inferior vena caval pressure). A pressure gradient of greater than 10 mmHg is associated with poorer outcomes post resection^[31]. Other indicators of clinically relevant portal hypertension include splenomegaly, oesophageal varices and thrombocytopaenia.

Given the relevance of liver function on the permitted resection size, the size of the liver remnant is important. This can be measured using CT volumetry^[32]. If adequate future liver remnant is not achievable, then portal vein embolization (PVE) should be considered. The aim of PVE is to induce compensatory hypertrophy in the non-embolised side. Generally, this is performed by the percutaneous transhepatic approach. A recent meta-analysis has demonstrated that PVE is safe and effective in inducing liver hypertrophy and preventing liver failure^[33]. It has also been shown to increase resectability^[34,35].

It should be noted that the recurrence rate after hepatic resection is high. In a systematic review and meta-analysis of resection vs transplantation, the 5 year disease-free survival rate of resection varied from 18%-51% compared to 54%-84% for transplanted patients^[18]. In patients with intermediate and advanced stage HCC (multiple tumours or macrovascular invasion), 5 year disease free survival range from 0%-31%^[36]. Follow-up for recurrence is therefore mandatory and recurrence should be managed using a multimodal approach including re-resection, TACE and ablative therapy.

CURRENT ISSUES IN HEPATIC RESECTION

Laparoscopic liver resection

With the advent of minimally invasive surgery, there is increasing uptake of the laparoscopic techniques for liver resection. Initially, the experience of laparoscopic liver resection was restricted to benign pathologies, and peripheral lesions/left lateral sectionectomy, although now major resections are being conducted laparoscopically^[37]. There have been several systematic reviews with meta-analyses on this topic. The most recent and the largest, a meta-analysis of 32 studies by Rao *et al.*^[37], found that laparoscopic hepatic resection was associated with significantly lower blood transfusion requirements, blood loss and length of stay but longer operating time. The overall complication rate was significantly lower (OR = 0.35, $P < 0.001$) in the laparoscopic group. Whilst overall survival was not different between the two groups, the rate of positive resection margins were found to be lower in the laparoscopic group. Note however, that the vast majority of studies were retrospective studies, with no randomised controlled trials, and therefore there may be significant selection bias. Unsurprisingly, these findings echo those of an earlier meta-analysis of 26 studies^[38]. However, in relation to oncological outcomes, this meta-analysis analysed HCC outcomes separately to other malignant diseases and found that there was a significant trend for improved overall survival (OR: 1.5 - 1.0-2.2; $P = 0.049$) in the

laparoscopic group. Another meta-analysis restricted only to studies evaluating laparoscopic resection for patients with HCC has demonstrated similar findings - lesser blood loss and blood transfusion requirements, lesser overall morbidity, cirrhotic decompensation and shorter length of stay^[39]. However, no differences in oncological outcomes (margins and survival) were found. Whilst the above studies point to potential advantages of performing laparoscopic hepatic resection, the major weakness of this systematic review is that the majority of studies included only patients who underwent minor hepatic resections. Their findings therefore may not be applicable to major laparoscopic hepatic resections. The efficacy of major liver resections is still under evaluation although early reports would suggest that they are comparable to the open procedure in terms of short and long term outcomes. For instance, Martin *et al.*^[40] compared 90 laparoscopic hepatectomies (left or right) to case-matched open hepatectomies and found lesser blood loss, lesser use of Pringle manoeuvre, lesser operative time, and lesser incidence of any type of complication. At present there exist only a few case series on robotic major hepatic resections-so while it is possible, the limited experience makes any conclusion about its comparative efficacy and risks difficult to make at the present time^[41].

Whilst the above results are encouraging, these should be interpreted with caution as there is likely significant publication and selection biases in the above studies. Also, laparoscopic liver resection has a learning curve, both for the surgeon and the institution^[42]. Results in centres where such expertise is available may not be generalizable to other centres. For instance, Viganò *et al.*^[43] demonstrated over the course of a 12 year period, operative time, pedicle clamping, blood loss, morbidity and hospital stay all decreased^[43]. They estimated the learning curve for minor laparoscopic liver resection is 60 based upon cumulative sum analysis on conversion rates. The same group reviewed the major hepatectomies performed laparoscopically at 6 experienced centres around the world and found similar improvements in operative time, conversion rate, blood loss and pedicle clamping^[42]. A larger study of 365 patients over a 14 year period estimated the learning curve for laparoscopic liver surgery to be in the order of 30-40 cases^[44].

Ablation vs resection

The alternative to resection in early HCC (satisfying Milan criteria) is ablation, either percutaneous ethanol injection (PEI), radiofrequency (RFA) or microwave ablation. In theory, ablation could treat a tumour of up to 5 cm in diameter - a size which correlates with the Milan criteria. Indeed, the European Association for the Study of the Liver

(EASL) and the European Organisation for Research and Treatment of Cancer (EORTC) guidelines suggest that ablation with either radiofrequency ablation or percutaneous ethanol injection is recommended as standard of care for patients with BCLC stage 0 or A who are unsuitable for surgery^[11]. A recent meta-analysis analysing outcomes from three randomised controlled trials and 25 non randomised studies has suggested little difference in survival or recurrence early after intervention but at 5 years, significantly different survival began to be observed. This appeared to be more pronounced in larger tumours than smaller ones and more in the non-randomised studies than the randomised trials. Consistently however, the complication rate and length of stay favoured RFA^[45]. Another recent systematic review which pooled the findings of 6 randomised controlled trials and 4 cohort studies comparing RFA/PEI to resection came to the same conclusion - that early outcomes (survival and recurrence) were equivalent (at one-year) but the differences became more pronounced with longer duration of follow-up. This seemed to apply even for small cancers (tumour size ≤ 3 cm). Complication rates however significantly favoured ablative therapy^[46]. Therefore one can conclude that in appropriately selected patients, surgical resection is the preferred management even for small cancers although ablative treatment had the advantage of lower morbidity.

Anatomical vs non-anatomical resection

The debate surrounding anatomical vs non-anatomical resection remain a controversial one. In theory, hepatocellular carcinomas recurrence is strongly related to microvascular tumour emboli, therefore, resection of the vascular territory of the tumour makes oncological sense^[47]. On the other hand, HCC often occurs in cirrhotic livers and the preservation of hepatic parenchyma to prevent postoperative liver failure suggests a non-anatomical approach. Indeed, numerous studies have been performed to elucidate the benefits of either approach, but none in a randomised fashion^[48]. Two meta-analysis of these non-randomised studies favoured anatomical resection although only one found statistically significant difference between the two groups in terms of both overall and disease-free survival^[49,50]. To further complicate one's understanding of this debate, a meta-regression performed by Cucchetti *et al*^[48] demonstrated that much of the heterogeneity of overall and disease-free survival results arose from the presence or absence of cirrhosis as a covariate. That is, non-anatomical resection had poorer outcomes because of the higher prevalence of cirrhosis in that group^[48]. Hence, only a randomised trial whereby the baseline characteristics are randomised can we make any final conclusions regarding this ongoing debate. An

interesting addition to the debate comes from the development of preoperative 3D simulation, which facilitates subsegmental and segmental anatomical resection, potentially allowing anatomical resection to be performed in patients who have limited hepatic reserve and allowing for a quality indicator of success or otherwise of anatomical resection^[51].

LIVER TRANSPLANTATION

The first successful liver transplantation in humans was performed in 1967^[52]. The attraction of using liver resection to manage HCC is that not only is the HCC treated with maximal resection margins, the underlying liver disease (and hence, premalignant field change) is also treated. In a landmark paper by Mazzaferro *et al*^[5], the Milan criteria were established in 1996. Mazzaferro *et al*^[5] described that patients operated within this criteria had excellent outcomes post liver transplantation which were comparable to those of patients operated on for non-cancer indications. The overall and recurrence-free survival rates at 4 years were 85% and 92% respectively^[5]. This criteria was subsequently expanded by the University of California San Francisco (UCSF) group who nonetheless demonstrated equivalent excellent outcomes - 5 year survival of 72%^[53]. These results have been validated in the Australian/New Zealand cohort by Chen *et al*^[54], who demonstrated a 5 year survival rate of 74% and 73% in those satisfying the Milan and UCSF criteria respectively^[54]. Those outside Milan or UCSF criteria were found to have significantly poorer outcomes. On the other hand, the use of UCSF criteria as preoperative selection criteria was found by a French group to have resulted in a 5 year survival of less than 50% despite a short waiting time^[55].

Other complications of transplantation include rejection as well as complications from immunosuppression may limit the long term survival of transplant recipients^[56]. The major limitation of liver transplantation in the treatment of HCC is the limited availability of donor livers. These patients with HCC also compete with non-cancer patients for transplants. As a result, strict listing criteria are used, such as mentioned above, to limit transplantation to those patients whose outcomes are comparable to those who do not have HCC. In the context of donor shortage, it is often accepted that transplanted patients should have 5 year survival rates of at least 50%. Unfortunately, whilst outcomes of transplantation are good, the potentially significant period on the waiting list may lead to dropout due to disease progression. This could be as high as 25%-38% in 12 mo, although it is highly variable^[55,57,58]. In fact, studies included in a recent systematic review reported median time to transplantation varying from 30 to 231 d^[18].

CURRENT ISSUES IN LIVER TRANSPLANTATION

Liver resection vs liver transplantation

The question of whether liver transplantation or liver resection is more efficacious in the treatment of HCC depend very much on the clinical scenario. For instance, for the patient with early HCC with inadequate hepatic reserve for resection, transplantation may be the only potentially curative option. On the other hand, a patient with a large HCC but with preserved hepatic function would only have resection as the curative option. The controversial case is of patients with early HCC with well compensated cirrhosis. In this case, not only are the outcomes of resection and transplantation important, the availability of donor livers, and therefore the dropout rate is highly relevant. Analysis of survival on an intention-to-treat basis would be more reflective of the relative efficacy of each treatment strategy in the real world. There have been some recent meta-analyses conducted to evaluate the question of relative efficacy. Rahman *et al*^[18] in 2012 looked at nine studies comparing liver resection and transplantation for early stage HCC^[18]. The key finding was that all these studies were retrospective and only a few reported intention-to-treat survival data for the transplantation group. Five-year overall survival ranged from 40%-70% for resection and 52%-81% for transplantation. Pooling of data from studies which conducted intention-to-treat survival analysis demonstrated no significant difference in survival between the two treatment strategies at 5 years. Another meta-analysis on an intention-to-treat basis also found no significant difference between the outcomes of the two groups^[59]. The other key finding highlighted by these two systematic reviews is the lack of prospective/randomised or even simply well-matched studies in the literature. Certainly, as yet there are no randomised controlled trials to guide treatment^[60].

Living donor liver transplantation

With the limitation of the availability of cadaveric liver transplants, there is increasing interest in the use of living donor liver transplant (LDLT). Clearly this requires the donation of a liver graft from a donor - a procedure not without its risks. The risk of mortality is estimated to be 0.1% for donor left hepatectomy and 0.5% for donor right hepatectomy; with a morbidity rate of up to 20%^[61,62]. This risk to the donor, without direct beneficial effects to the person also brings about an ethical dilemma to transplant surgeons and physicians alike - "first do no harm". However, the advantage of LDLT is that as the liver is obtained outside the usual donor pool, this strategy expands the number of organs available for transplantation. As a result, the criteria

for liver transplantation can often be extended beyond the usual Milan or UCSF criteria. The Kyoto criteria (≤ 10 tumours, less than 5 cm, PIVKA-II ≤ 400) and "up-to-seven" criteria are examples of extended transplant criteria which have been used in the context of living donor transplants^[63,64]. Using a decision analytical model taking into account the risk of dropout while waiting (4% per month), the expected survival of the recipient (70% at five years) and the risk for the donor (0.3% to 0.5% mortality), Sarasin *et al*^[65] demonstrated that patients with HCC waiting more than seven months for a deceased donor liver would benefit from LDLT. Early reports have suggested a higher recurrence rate in LDLT as compared to Deceased donor liver transplantation (DDLT), however, this is hypothesised to be due to the "fast-tracking" of LDLT which therefore allowed transplantation of patients with more aggressive HCC^[61]. Two recent meta-analysis of LDLT vs DDLT were reported. These found overall survival rates to be similar between the two groups. One of these, by Grant *et al*^[66] found LDLTs to be associated with decreased disease-free survival rates. On the other hand, Liang *et al*^[67] performed a subgroup analysis of patients within the Milan criteria and found similar survival outcomes between the two groups.

Bridging therapy and salvage liver transplantation

Out of the need to minimise dropout during waiting, strategies such as bridging therapy or resection with salvage transplantation has been developed. Bridging therapies such as RFA or TACE are frequently used. Whilst bridging therapy does seem to be useful in decreasing dropout rate whilst awaiting transplant, its role in improving survival after transplantation has not been established^[68]. An alternative strategy to primary transplantation is primary resection followed by salvage transplantation. The advantage to this is to minimise the need for organs and to use resection as the ultimate bridging therapy to prevent progression whilst waiting for transplantation. A recent systematic review of 16 studies found that of those 7 studies which reported salvage transplant rates, the median rate of salvage transplantation was 41% after a median time to recurrence of 21 mo^[69]. Whilst a meta-analysis was not performed, they found a median 5 year survival to be 67%. Interestingly, half of studies reported a mortality rate of higher than 5% and two studies reported mortality rates of greater than 10%^[70,71]. A meta-analysis by Zhu *et al*^[72] analysed 14 studies, of which 10 overlapped with Chan's systematic review. Zhu *et al*^[72] found that compared to primary liver transplantation, salvage liver transplantation was associated with longer operative time, greater blood loss but failed to find a significant difference in postoperative mortality. With regards to long term survival, primary liver transplant was found to have

better five-year disease free survival but not overall survival^[72]. These results would suggest that salvage liver transplant is a viable strategy in appropriately resourced transplant centres.

CONCLUSION

HCC is currently largely managed through non-surgical means as the tumour-related and hepatic function considerations for surgical management is still largely restrictive. For those who have tumours eligible for surgical therapy from the tumour point of view (early stage tumours), those with good hepatic function and significant functional liver remnant would be candidates for either resection or transplantation depending on local resources. Those with poor hepatic function may be placed on the liver transplant list, with or without bridging therapy. Patients with intermediate stage cancers have limited options, with resection being the only potential for cure. Otherwise, regional therapy with TACE or RFA are viable options. With further development of surgical techniques, including salvage liver transplantation, the indications for surgical management of HCC may continue to expand. With this, the outcomes of HCC may further improve.

Surgical therapy is the only curative hope for patients with HCC. The selection of patients for transplantation and resection will depend on local resources, but both have potentially good outcomes in appropriately selected patients.

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Occult hepatitis B virus co-infection in human immunodeficiency virus-positive patients: A review of prevalence, diagnosis and clinical significance

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(HIV) and hepatitis B virus (HBV) co-infection is high as they share similar mechanisms of transmission. The development and widespread use of highly sensitive tests for HBV diagnosis has demonstrated that a significant proportion of apparently healthy individuals with evidence of exposure to HBV continue to carry fully functional HBV DNA in their hepatocytes, a situation that predisposes them to the development of progressive liver disease and hepatocellular carcinoma. The presence of co-infections frequently influences the natural evolution of each of the participating infections present by either facilitating their virulence or competing for resources. Furthermore, the drugs used to treat these infections may also contribute to changes in the natural course of these infections, making the analysis of the impact of co-infection more difficult. The majority of studies has examined the impact of HIV on overt chronic hepatitis B, finding that co-infection carries an increased risk of progressive liver disease and the development of hepatocellular carcinoma. Although the effect of HIV on the natural history of occult hepatitis B infection (OBI) has not been fully assessed, all available data suggest a persisting risk of repeated flares of hepatitis and progressive liver disease. We describe studies regarding the diagnosis, prevalence and clinical significance of OBI in HIV-positive patients in this short review. Discrepancies in worldwide prevalence show the urgent need for the standardization of diagnostic criteria, as established by the Taormina statements. Ideally, standardized protocols for testing should be employed to enable the comparison of data from different groups. Additional studies are needed to define the differences in risk for OBI without HIV and in HIV-HBV co-infected patients with or without overt disease.

Key words: Hepatitis B virus; Occult hepatitis B; Human immunodeficiency virus; Prevalence; Diagnosis; Clinical significance

Abstract

The prevalence of human immunodeficiency virus

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Core tip: The prevalence of human immunodeficiency virus (HIV) and hepatitis B virus (HBV) co-infection is high. However, as HBV infection may be occult, its diagnosis requires the routine use of highly sensitive tests. Although viral load or replication in these patients is low, they still have an increased risk of viral reactivation, chronic liver disease and hepatocellular carcinoma development. The majority of our knowledge on occult hepatitis B infection is derived from studies performed in patients with mono-infection or with HIV co-infection. This review summarizes the latest contributions in the field, clearly revealing that more studies are needed to evaluate the full impact of HIV in patients with occult HBV disease.

Maldonado-Rodriguez A, Cevallos AM, Rojas-Montes O, Enriquez-Navarro K, Alvarez-Muñoz MT, Lira R. Occult hepatitis B virus co-infection in human immunodeficiency virus-positive patients: A review of prevalence, diagnosis and clinical significance. *World J Hepatol* 2015; 7(2): 253-260 Available from: URL: <http://www.wjgnet.com/1948-5182/full/v7/i2/253.htm> DOI: <http://dx.doi.org/10.4254/wjh.v7.i2.253>

INTRODUCTION

The presence of co-infections frequently influences the natural evolution of each of the participating infections present, either by facilitating their virulence or competing for resources^[1]. Furthermore, the drugs used to treat these infections may also contribute to changes in the natural history of these infections^[2], complicating the analysis of the impact of co-infection.

For overt chronic hepatitis B virus (HBV) infection, it is known that the virus does not substantially alter the progression of human immunodeficiency virus (HIV) disease, nor does it influence HIV suppression or CD4 cell responses following the initiation of antiretroviral treatment^[3-5]. In contrast, it is clear that HIV infection has a negative effect on HBV disease, for both acute and chronic infections. In cases of acute hepatitis, progression to chronic infection is approximately 4 times more frequent in patients with HIV than in those without HIV infection (20% vs 5% and most likely higher if the CD4 count is low)^[5]. For chronic hepatitis, HIV infection results in a faster progression of fibrosis, a faster development of cirrhosis and hepatocellular carcinoma, and a lower rate of spontaneous hepatitis B e antigen (HBeAg) or hepatitis B surface antigen (HBsAg) seroconversion^[5]. There is also a greater risk of HBV reactivation in inactive carriers.

Occult HBV infection (OBI) occurs when viral DNA persists in the liver (with detectable or undetectable HBV DNA in serum, with or without HBV antibodies)

in individuals testing negative for HBsAg^[6]. By convention, OBI has been defined as the persistence of isolated anti-HBc in patients who may or may not have detectable serum HBV DNA. However, OBI may also occur in patients without anti-HBc antibodies. Thus, in a workshop of the European Association for the Study of Liver held in Taormina (Italy) in 2008, experts joined together to re-define and standardize diagnostic criteria, which resulted in a series of new criteria known as the Taormina statements^[7]. It is expected that the universal use of these criteria will allow the comparison and proper evaluation of studies by different research groups.

In patients with HBV infection, the absence of HBsAg can occur in two main scenarios: firstly, in the early phases of acute infection prior to the development of antibodies and the detection of HBsAg in serum; secondly, during chronic HBV infection following the decline of HBsAg to an undetectable level, which is sometimes associated with the appearance of anti-HBs^[8]. In chronic occult infections, viral covalently closed circular DNA (cccDNA) persists as a stable chromatinized episome in the nucleus of infected cells. This viral genome remains competent for replication and able to synthesize minute amounts of antigens, which are undetectable by the available technical approaches but sufficient to maintain an HBV-specific T cell response^[6]. Fragments of the HBV genome may integrate into the host hepatic cell genome but this integration does not have a role in the replicative cycle of HBV and should not be strictly considered as occult infection^[6]. In addition, chronic occult infection may be associated with the presence of one or more specific antibodies in serum; therefore, individuals with occult infections are conventionally divided into seropositive [anti-hepatitis B core antigen (anti-HBcAg) and/or anti-hepatitis B surface antigen (anti-HBsAg) positive] and seronegative (anti-HBc and anti-HBs negative) groups^[6,7]. More than 20% of occult-infected individuals are negative for all HBV serum markers^[9]. The development of occult infections is mainly determined by host factors, such as the immune response, and the status of the infection can be modified by the presence of co-infections, such as with hepatitis C virus (HCV) or HIV, or the administration of drugs, including immune suppressors and/or antiretrovirals^[10]. Multiple viral variants have been identified in the liver of occult HBV-infected patients and it is believed that viral factors are not major determinants for the development of occult infections^[6,11].

We describe the diagnosis, prevalence and clinical significance of OBI in HIV-positive patients in this short review.

OBI DIAGNOSIS

The presence or absence of full infective virions is not assessed in clinical practice. Surrogate markers

for the detection of circulating virus are either the presence of HBV DNA or HBV proteins (HBsAg, HBeAg and HBcAg). Typically, the presence of anti-HBs antibodies and the absence of HBsAg suggests a resolved infection; however, the persistence of only anti-HBc may be associated with OBI^[12]. Overt HBV infection (acute or chronic) is defined as the presence of circulating HBsAg; according to the Taormina statements, OBI is defined as the absence of circulating HBsAg and the presence of HBV DNA^[7]. However, the sensitivity and specificity of diagnosis strongly depend on the sensitivity of the assays used; indeed, for many years, HBV DNA tests were not very sensitive. Thus, the presence of anti-HBc was used as a surrogate marker and according to the Taormina statements, it still can be used in regions where modern molecular assays are not available^[13,14]. The gold standard for OBI diagnosis is the demonstration of the presence of HBV DNA in the liver. To rule out the possibility that amplified fragments correspond to partial regions of the viral genome integrated into the host genome, several regions of the genome must be identified to suggest that full-length cccDNA is present. It is accepted that the diagnosis is frequently underestimated as liver biopsies are only rarely available and therefore diagnosis is usually based on blood samples. There is no evidence to date that HIV infection modifies the sensitivity or specificity of these tests.

Markers for screening OBI

HBsAg: It is of crucial importance to define the best methodology to test HBsAg to prevent false positive results, which is dependent on the HBsAg assay sensitivity. Several problems of this aspect are associated with the virus, the host or the test kits employed in practice. The quantification of HBsAg should be performed by comparing the sample with a standard curve generated with the second International Standard for HBsAg (World Health Organization code number 00/588, document WHO/BS/03.1987); 1 international unit is equivalent to 5.6 Abbott ng, 1.9 French ng and 0.43 PEI units^[8]. Unfortunately, some HBV variants have mutations in the *HBs* gene and the encoded proteins are not detected with conventional commercial kits. Accordingly, alternative methods should be tested to detect common mutants^[15].

HBV DNA: It has been established that HBV-DNA is the only reliable diagnostic marker for OBI. The experts meeting at Taormina recommended that new generation assays with detection limits of less than 10 copies of HBV DNA per reaction should be employed^[7]. The estimated viral load in OBI is usually below 200 IU/mL.

New generation assays for DNA detection include nested-polymerase chain reaction (PCR), real-time PCR and transcription-based mediated amplification.

Indeed, advances in the development of these DNA detection technologies has allowed a decrease in the lower detection limit (< 5 IU/mL of HBV DNA), which is particularly important in OBI because DNA levels vary at -5-10 IU/mL (range < 10 to 425 copies/mL)^[8].

HBV genome regions for diagnosis: According to the Taormina statements^[7], the primers used must be specific for different HBV genomic regions and be complementary to highly conserved nucleotide sequences. The *S* and *X* genes are the regions most commonly amplified by PCR for diagnosis; it has been found that the *X* gene is the most sensitive for the liver, whereas the *S* gene is better for serum samples^[9]. To avoid the problems of cross-contamination, appropriate controls in each PCR assay as well as amplicon sequencing are recommended.

PREVALENCE

The prevalence of infections in open populations is frequently estimated using data obtained from serological testing performed using blood donor samples; however, HBV tests can detect either overt infections or previous exposure to the virus in blood donors. Therefore, the available data reflect only the prevalence of overt HBV infections. OBI is only screened in specific scenarios, such as in areas of high endemicity of HBV, intravenous drug users, organ transplant patients, patients on maintenance hemodialysis or patients with HIV and/or HCV. For this review, we analyzed 34 papers that examined the prevalence of OBI in HIV patients during the period 2003-2014 (Table 1). The range of reported prevalence in these studies varied from 0.63% to 88.4%, which is similar to the prevalence reported in a previous review (0%-89.5%)^[16]. This extremely wide range of prevalence reflects the diverse nature of published studies. Some of the differences are explained by the individual prevalence of HIV and HBV in the different populations studied. Although there are reports from central and south America^[17-22], the majority of the studies are from regions of Africa, India and the Far East, regions where the prevalence of both HIV and HBV is high^[23-27]. Differences also arise from the type of high-risk group to which the co-infection patients studied belong (*e.g.*, hemodialysis patients, homosexuals, intravenous drug users). Another source of variability depends on differences in the sensitivity of the diagnostic test used, with more recent studies using more sensitive HBV DNA detection assays than earlier studies^[28]. Clearly, prospective, longitudinal studies in well-defined populations are needed to fully evaluate the prevalence and impact of HIV in the natural history of OBI. These studies should contain detailed clinical and demographic data, including age, sex, risk group liver function tests and, whenever possible, the evaluation

Table 1 Reported prevalence of occult hepatitis B virus infection in HIV-infected subjects *n* (%)

Ref.	Country	OBI prevalence			
		Overall	Anti-HBsAg-/anti-HBcAg+	Anti-HBsAg+/anti-HBcAg+	Anti-HBsAg-/anti-HBcAg-
Alvarez-Muñoz <i>et al</i> ^[17]	Mexico	24 (49.0)	5 (10.2)	8 (16.3)	11 (22.4)
Araujo <i>et al</i> ^[18]	Brazil	6 (14.0)	1 (2.3)	5 (10.2)	ND
Attia <i>et al</i> ^[45]	Africa	40 (21.3)	40 (21.3)	ND	ND
Azadmanesh <i>et al</i> ^[46]	Iran	3 (13.6)	2 (9.1)	ND	1 (4.5)
Bagaglio <i>et al</i> ^[47]	Italy	9 (31.0)	9 (31.0)	ND	ND
Bell <i>et al</i> ^[48]	Africa	45 (15.1)	16 (5.4)	17 (5.7)	12 (4.0)
Bloquel <i>et al</i> ^[38]	France	3 (0.8)	2 (0.5)	ND	1 (0.3)
Chadwick <i>et al</i> ^[49]	England	15 (4.5)	5 (1.5)	10 ¹ (3.0)	ND
Coffin <i>et al</i> ^[50]	Canada	19 (42.0)	ND	19 (42.2)	ND
Dapena <i>et al</i> ^[51]	Spain	6 (2.4)	2 (0.8)	4 (1.6)	ND
Filippini <i>et al</i> ^[13]	Italy	17 (20.0)	11 (12.8)	3 (3.5)	3 (3.5)
Firnhaber <i>et al</i> ^[23]	Africa	38 (88.4)	38 (88.4)	ND	ND
Gupta <i>et al</i> ^[30]	India	24 (45.3)	13 (24.5)	11 (20.8)	ND
Hakeem <i>et al</i> ^[52]	Scotland	2 (2.8)	2 (2.9)	ND	ND
Jardim <i>et al</i> ^[19]	Brazil	8 (5.0)	2 (1.3)	6 (3.8)	ND
Khamduang <i>et al</i> ^[35]	Thailand	47 (23.5)	47 (23.5)	ND	ND
Liang <i>et al</i> ^[53]	Taiwan	3 (2.3)	3 (2.3)	ND	ND
Lo Re <i>et al</i> ^[54]	United States	17 (10.0)	10 (5.6)	7 (3.9)	ND
Loustaud-Ratti <i>et al</i> ^[55]	France	31 (44.3)	20 (28.6)	11 (15.7)	ND
Morsica <i>et al</i> ^[56]	Italy	27 (15.4)	9 (5.1)	18 (10.3)	ND
Mphahlele <i>et al</i> ^[57]	Africa	31 (18.6) ²	5 (3.0)	26 (15.6)	ND
N'Dri-Yoman <i>et al</i> ^[24]	Africa	51 (10.0)	51 (11.8)	ND	ND
Neau <i>et al</i> ^[58]	France	1 (0.6)	1 (0.6)	ND	ND
Nebbia <i>et al</i> ^[59]	England	48 (14.0)	48 (14.0)	ND	ND
Opaleye <i>et al</i> ^[25]	Nigeria	21 (11.2)	8 (4.3)	9 (4.8)	2 (1.1)
Panigrahi <i>et al</i> ^[26]	India	12 (10.7)	9 (8.0)	3 (2.7)	ND
Santos <i>et al</i> ^[20]	Brazil	16 (15.8) ²	4 (4.0)	12 (11.9)	ND
Sen <i>et al</i> ^[27]	India	1 (5.6) ²	1 (5.6)	ND	ND
Shire <i>et al</i> ^[60]	United States	4 (10.5)	4 (10.5)	ND	ND
Shire <i>et al</i> ^[61]	United States	12 (30.2)	3 (7.0)	5 ¹ (11.6)	5 (11.6)
Sucupira <i>et al</i> ^[21]	Brazil	6 (18.8) ²	3 (9.4)	3 (9.4)	ND
Torres Barranda <i>et al</i> ^[22]	Mexico	7 (18.4)	1 (2.6)	1 (2.6)	5 (13.2)
Tramuto <i>et al</i> ^[62]	Italy	24 (5.9)	8 (2.0)	7 ¹ (1.7)	9 (2.2)
Tsui <i>et al</i> ^[63]	United States	8 (2.0)	8 (2.0)	ND	ND

¹In some studies the anti-HBsAg positive group was also included; ²Prevalence calculated using the reported data; anti-HBsAg+, antibodies against hepatitis B surface antigen positive; anti-HBcAg+ antibodies against hepatitis B core antigen positive. Prevalence (%) were included for each group of patients studied according the HBV serological markers (Anti-HBsAg-/anti-HBcAg+, Anti-HBsAg+/anti-HBcAg+, Anti-HBsAg-/anti-HBcAg-). ND: Not determined because this group was not included in the study; OBI: Occult hepatitis B infection; anti-HBsAg: Anti-hepatitis B surface antigen; anti-HBcAg: Anti-hepatitis B core antigen.

of liver damage by biopsy. Ideally, standardized protocols for testing should be employed so that studies from different groups can be compared.

In general, OBI prevalence appears to be higher among patients at high risk for HBV infection and with liver disease than among individuals at low risk of infection and without liver disease^[9,29-31]. As mentioned above, patients with OBI may be negative for all HBV serum markers and there is also evidence that levels of viremia are correlated with levels of anti-HBc and/or anti-HBs^[8]. Unfortunately, the majority of studies have been performed in seropositive patients (usually isolated anti-HBc), whereas seronegative patients who are more difficult to identify have not been fully assessed. Furthermore, the available data suggest seronegative patients have a different clinical evolution and should therefore be evaluated separately. Another factor that is common in HIV patients and that is known to affect

the evolution of HBV infection is HCV co-infection; the presence of HCV should always be screened and those with OBI/HCV should be analyzed separately.

CLINICAL SIGNIFICANCE

The significance of chronic OBI relates to the risk of transmission, reactivation, progression to chronic liver disease and development of hepatocellular cancer^[6,8,32,33]. The rate of transmission is directly proportional to the number of viable virions in blood. Unfortunately, the true level of viremia, *i.e.*, the number of infectious HBV particles within 1 mL of serum or plasma, is difficult to measure in a bioassay^[34]. Therefore, surrogate markers of HBV infectivity are being used to measure the risk of transmission. The best available marker for the presence and number of infectious HBV particles is the number of HBV DNA molecules. However, it is

important to note that the detection of HBV DNA in serum does not always correspond to infectivity or to the number of HBV progeny viruses released from hepatocytes. Indeed, in the majority of viral infections, the number of physical virus particles is much larger than the number of fully infectious virions^[34]. There also appears to be a correlation between levels of HBV DNA and serological status among patients with OBI: HBV DNA levels are lowest in seronegative patients, intermediate in anti-HBc negative and anti-HBs positive patients, and highest in subjects who are anti-HBc-positive but anti-HBs negative^[8]. This last group is more likely to be infectious. The possibility of transmission is crucial for non-immunocompromised individuals who are potential donors of blood or other organs, for health care providers who can infect their patients, and for pregnant women *via* vertical transmission. Although known HIV patients are not candidates for organ donation, there remains a risk of HIV/HBV transmission either through sexual encounters, needle sharing or other risky behaviors. The potential risk of vertical transmission in pregnant HIV/OBI women is clear but the prevalence rate has not been fully evaluated. In a recent study of 1682 HIV-infected pregnant women in Thailand who were fully evaluated for HBV infection, 216 were HBsAg negative and anti-HBc positive (14%). It was also possible to assess the levels of HBV DNA in 200 of these women with OBI; all 200 women had HBV DNA < 1000 IU/mL, with 153 showing HBV DNA below the limit of detection (15 IU/mL), 44 with an HBV DNA level between 15-100 IU/mL, and 3 showing HBV DNA between 101 and 1000 IU/mL. However, none of these women transmitted the disease to their infants^[35]. Based on the available information, a group of experts in the United States has provided guidelines to manage pregnant women with HIV-HBV co-infection; the full guidelines have been published at the AidsInfo site of the National Institute of Health (<http://aidsinfo.nih.gov/guidelines/html/3/perinatal-guidelines/159/hiv-hepatitis-b-virus-coinfection>). It is recommended that all pregnant women should have full screens for both HBV and HCV; these tests would identify most cases of OBI in HIV patients. They also strongly recommend that the management of HIV/HBV co-infection in pregnancy should be performed with the advice of an expert in HIV and HBV, with close monitoring of the viral activities in pregnant woman. In their review, the experts conclude that women who screen negative for HBV (*i.e.*, HBsAg-negative, anti-HBc-negative and anti-HBs-negative) should be vaccinated for HBV, stating that the presence of isolated anti-HBc can represent a false-positive case or can indicate a previous exposure and a posterior loss of anti-HBs or most likely an "occult" HBV infection. The possibility of OBI needs to be confirmed by HBV DNA detection. These experts also

recognize that the clinical significance of isolated anti-HBc is still unknown. For that, they recommend, with the panel of their peers on opportunistic infections in HIV-infected adults and adolescents, to test for HBV DNA in HIV+/anti-HBc+ patients before HBV vaccination and treatment initiation or prophylaxis to avoid the risk of a paradoxical exacerbation of HBV infection and the incidence of immune reconstitution inflammatory syndrome (http://aidsinfo.nih.gov/contentfiles/lvguidelines/adult_oi.pdf). The risk of vertical HBV transmission in mothers with OBI is most likely small due to the low levels of HBV viremia. Clearly, the decision to treat a pregnant woman with OBI should be made by an expert.

The natural history of chronic hepatitis B is highlighted by spontaneous flares of the disease. The reasons for HBV reactivation are not clear but most likely can be explained by subtle modifications in virus replicative fitness due to host immunological control, such as that which occurs with herpes virus^[36]. The flares are potentially important clinically because they can have severe or even fatal consequences. Most frequently, the reactivation of HBV replication occurs in patients with overt chronic HBV infection (HBsAg-positive) who receive cytotoxic or immunosuppressive therapy^[10,33]. In contrast, HBV reactivation occurs more rarely in patients with OBI. Both in overt and occult infection, the risk of HBV reactivation is high, particularly in patients with hematological malignancies, in those receiving hematopoietic stem cell transplantation and in those treated with either anti-CD20 (rituximab, which destroys B-cells) or anti-CD52 (alemtuzumab, which targets mature lymphocytes) monoclonal antibodies (reviewed in^[10]). In these cases, HBV reactivation is associated with a mortality rate of approximately 20% due to hepatic failure or to progression of the underlying disease owing to the discontinuation of treatment.

There are multiple anecdotal reports of the reactivation of OBI in HIV-patients, with the majority of cases being identified in seropositive patients. Many factors, similar to those demonstrated for overt infection, have been implicated in the recurrence of HBV replication in HIV/OBI-co-infected patients, including the interruption of HAART^[37-39], recovery of immune responses after HIV-treatment^[14], development of resistance to lamivudine therapy^[13], and appearance of HBV immune-escape^[40].

The overall prevalence of OBI reactivation in HIV patients and the frequency of the specific triggers of reactivation need to be assessed in prospective longitudinal studies^[14]. In an effort to evaluate the full impact of HIV-infection in OBI, a multicenter prospective study on 115 consecutive anti-HIV+, HBsAg-negative, treatment-naïve patients was performed^[13]. Of the 86 patients with at least 6 mo of follow-up, 13 were HBV DNA positive on

admission and four in the subsequent testing. The HBV DNA+ frequency for the anti-HBs-negative/anti-HBc-positive group was 36% and 21% for the anti-HBs-positive/anti-HBc-positive group; the lowest frequency was reported in the anti-HBs-negative/anti-HBc-negative group (9%). Episodes of reactivation were detected in 32% of the patients and were more common in patients with detectable HBV DNA than in those without it (65% vs 25%). These preliminary data await confirmation in larger studies^[10].

HBV infection causes liver inflammation and hepatocellular carcinoma. The lifetime risk of developing HBV-related cirrhosis or hepatocellular carcinoma has been estimated to be between 15% and 40% for males who acquire infection during early life^[36]. There is evidence to suggest that the risk of progression of liver disease from inflammation, to fibrosis, to cirrhosis and to cancer is directly related to the level of transcription of HBV DNA^[41-43]. However, the prevalence and rate of progression to cirrhosis are also related to differences in the clinical and serological features of the disease, such as repeated flares, age and HBV genotype^[36]. There is an increased risk of hepatocellular carcinoma even in patients with inactive HBV infection^[44]. It is widely accepted that the risk of progression is particularly high in patients with OBI/HCV co-infection^[8]. However, the impact of HIV infection in the oncogenicity of OBI HIV patients has not been studied and requires specific evaluation^[10].

CONCLUSION

The real impact of co-infection can only be fully established when the presence of OBI is routinely assessed in all patients with HIV. As routine liver biopsy of all HIV patients is not possible, at least all HIV patients should be screened initially with a complete panel of HBV tests. As both the reactivation of hepatitis (with or without the emergence of drug resistance) and an increase in the risk of transmission are associated with higher levels of HBV DNA, the monitoring of OBI should always include regular highly sensitive measurements of circulating HBV DNA. More studies are needed to define when to repeat the full set of HBV tests and when a liver biopsy is indicated in these patients.

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Role of surgical resection for hepatocellular carcinoma based on Japanese clinical guidelines for hepatocellular carcinoma

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in hepatectomy and perioperative management of hepatocellular carcinoma.

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Core tip: In the Algorithm for Diagnosis and Treatment in the Japanese Evidence-Based Clinical Practice Guidelines for Hepatocellular Carcinoma, the treatment strategy is determined by three major factors: liver function and the number and size of tumors. The algorithm is quite simple, consisting of fewer components than the Barcelona-Clinic Liver Cancer staging system. In this article, we describe the roles of the treatment algorithm in hepatectomy and perioperative management of hepatocellular carcinoma.

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Abstract

In the Algorithm for Diagnosis and Treatment in the Japanese Evidence-Based Clinical Practice Guidelines for Hepatocellular Carcinoma, the treatment strategy is determined by three major factors: liver function and the number and size of tumors. The algorithm is quite simple, consisting of fewer components than the Barcelona-Clinic Liver Cancer staging system. In this article, we describe the roles of the treatment algorithm

INTRODUCTION

Hepatocellular carcinoma (HCC) is the sixth most common neoplasm worldwide and the third most frequent cause of cancer-related death. More than 0.7 million people were diagnosed with HCC in 2008, indicating an incidence of 16 per 0.1 million people^[1]. The distribution of HCC is regional, with approximately 80% of HCC cases found in Eastern Asia and central Africa. The risk factors in these

Table 1 Staging systems for hepatocellular carcinoma

Classification	Year	Background	Variables		
			Tumor status	Liver function	Health status
Okuda staging	1985	850 Japanese patients	50% liver involvement	Ascites, bilirubin, albumin	-
BCLC staging	1999	Selected papers	Size, number, vascular invasion, Okuda stage	Child-Pugh, bilirubin, porta hypertension	Performance status
CLIP score	2000	435 Italian patients	50% liver involvement, vascular invasion, AFP	Child-Pugh	-
CUPI	2002	926 Chinese patients	TNM, AFP	Bilirubin, albumin, alkaline phosphatase	Presence of symptoms
JIS score	2003	3334 Japanese patients	TNM (Japanese)	Child-Pugh	-
m-JIS score	2006	42269 Japanese patients	TNM (Japanese)	Liver damage	-
Tokyo score	2005	403 Japanese patients	Size, number	Bilirubin, albumin	-

BCLC: Barcelona Clinic Liver Cancer; CLIP: Cancer of the liver Italian Program; CUPI: Chinese University Prognostic Index; JIS: Japan Integrated Staging; m-JIS: Modified Japan Integrated Staging; AFP: Alpha-fetoprotein. Revise from ref. [22].

areas are hepatitis B and aflatoxin, but those in North America, Europe, and Japan are hepatitis C and alcohol.

The spread of the concept of evidence-based medicine (EBM) has provided an opportunity for development of treatment guidelines. In Western countries, the Barcelona Clinic Liver Cancer (BCLC) staging system was published as practice guidelines in 2005 and updated in 2011, and is recommended for use by the American Association for the Study of Liver Diseases (AASLD)^[2] and the European Association for the Study of the Liver (EASL). In Japan, the Clinical Practice Guidelines for Hepatocellular Carcinoma were published in 2005^[3,4] and then revised in 2009 and 2013 to add new information^[5]. The “treatment algorithm” listed in the guidelines has become well disseminated as a standard method for selection of optimal treatment based on liver function and tumor conditions^[6]. Here, we describe the roles of the treatment algorithm in hepatectomy for HCC and we discuss current knowledge on hepatectomy in Japan.

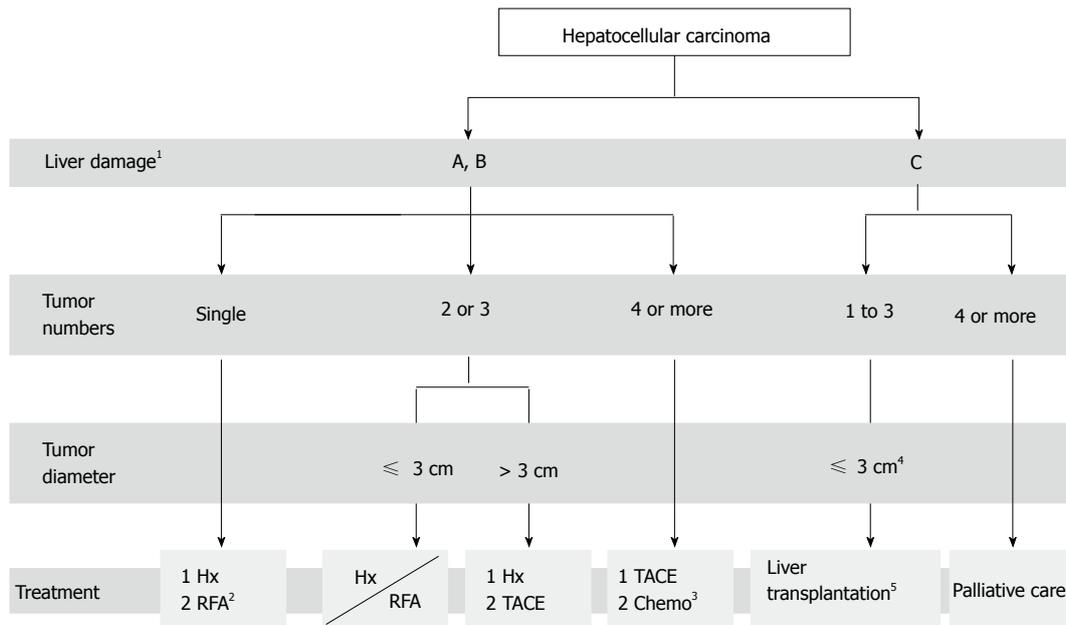
STAGING SYSTEM FOR HCC

Staging systems for liver cancer have three elements: (1) tumor stage (TNM system); (2) hepatic functional reserve; and (3) integrated stage, a combination of (1) and (2). The International Union Against Cancer (UICC) published the UICC TNM classification of malignant tumors in 1968 and added liver cancer to the TNM classification in 1987. Now, the seventh edition is used from 2009^[7]. The UICC-TNM classification is based on the staging system of the American Joint Committee on Cancer (AJCC) Cancer Staging Manual, with a database from multicenter research by the International Cooperative Study Group on Hepatocellular Carcinoma^[8]. The UICC-TNM classification is a simplified version of the AJCC Manual, and the 7th edition set the cutoff for tumor size as 5 cm. In 1983, the Liver Cancer Study Group of Japan published the “General Rules for Clinical

and Pathological Studies of Primary Liver Cancer” (henceforth referred to as the “General Rules”), which included the Japanese TNM classification^[9] and was prepared based on a database developed by the Liver Cancer Study Group. In the latest edition, the stages are classified using a cutoff tumor size of 2 cm, single/multiple lesions, and vascular invasion. In a comparison of these two staging systems in Japanese patients, Minagawa *et al.*^[10] found that both systems allowed clear stratification of patients into prognostic groups, but that the General Rules were more appropriate for stratifying patients with early-stage HCC^[10].

The Child-Pugh classification is most commonly used for evaluation of hepatic functional reserve^[11,12]. This classification has five parameters: serum bilirubin, serum albumin, ascites, hepatic encephalopathy, and prothrombin activity, which are used to assess liver function in three classes: A, B and C. The indocyanine green retention rate at 15 min (ICGR₁₅) is also used in Japan, eastern Asia, and some European countries as a more detailed index for assessment of hepatic functional reserve. ICGR₁₅ is useful for prediction of postoperative mortality^[13] and as a marker of liver function for determining the extent of hepatectomy^[14]. The General Rules also have a liver damage classification system that uses ICGR₁₅, as well as serum bilirubin, serum albumin, ascites, and prothrombin activity^[9,15]. The degree of liver damage has replaced the Child-Pugh classification to evaluate liver function in Japan. For serious liver failure patients, model for end stage liver disease (MELD) is used to indicate liver transplantation^[16].

Integrated Stage score for liver function and tumor stage, including OKUDA^[17], Cancer of the Liver Italian Program (CLIP)^[18], Chinese University Prognostic Index (CUPI)^[19], Japan Integrated Staging (JIS)^[20], modified-JIS^[15], and Tokyo^[21], is effective for prognostic assessment in HCC (Table 1). Kudo *et al.*^[20] proposed the JIS score, which unified TNM staging in the General Rules and the



For patients with the severity of liver damage class A accompanied by vascular invasion, hepatectomy, chemotherapy, transcatheter arterial embolization may be selected. For patients with Child-Pugh class A with extrahepatic metastasi, chemotherapy is an option.

¹When taking non-surgical treatment into consideration, using of Child-Pugh classification is possible; ²Selected when the tumor measuring is 3 cm or less; ³Oral or arterial infusion chemotherapy may be selected; ⁴A single tumor measuring 5 cm or less in diameter; ⁵Patients 65 years of age or younger.

Figure 1 Treatment algorithm for hepatocellular carcinoma. Revise from ref. [5]. Hx: Hepatectomy; RFA: Radiofrequency ablation; TACE: Transcatheter arterial chemo embolization; Chemo: Chemotherapy.

Child-Pugh classification^[20]. The JIS is superior to the CLIP system [a combination of the Child-Pugh classification, tumor morphology, α -fetoprotein (AFP), and portal vein tumor thrombosis] in terms of (1) clear stratification of scores; (2) prognostic predictive power in HCC with a score of 0; and (3) differentiation of scores in patients with a poor prognosis. Thus, the JIS score is useful for prediction of prognosis of patients, but is not appropriate for comparison of treatment modalities or selection of optimal treatment.

CLINICAL GUIDELINES FOR HCC

The BCLC staging system, which is recommended by AASLD and EASL, is used worldwide to plan treatment for HCC. In contrast, in Japan, the treatment algorithm in the Clinical Practice Guidelines for Hepatocellular Carcinoma is commonly used for selection of optimal treatment based on liver function and tumor conditions (Figure 1). BCLC system links stage stratification to a treatment strategy and recommends standard care for a given patient, whereas the Japanese guidelines are not directly associated with clinical tumor stage, such as the JIS score^[22]. The another major difference between the treatment algorithm used in Japan and the BCLC system is the indication of hepatectomy for HCC with ≤ 3 lesions and a diameter ≤ 3 cm on Child-Pugh A/B.

The BCLC system recommends liver transplantation or radiofrequency ablation (RFA) for HCC with 2 or 3 nodules and a diameter ≤ 3 cm. In contrast, the treatment algorithm in Japan recommends hepatectomy for HCC with ≤ 3 lesions if liver function is good, regardless of the tumor size. The recommended treatment strategy also differs for HCC with portal hypertension (Table 2). The BCLC system states that liver transplantation or RFA, instead of hepatectomy, is indicated in such patients, but the treatment algorithm in Japan advises that aggressive hepatectomy based on ICGR₁₅ should be performed because the therapy must yield positive results^[23].

DIAGNOSIS OF CLASSICAL HCC AND TREATMENT FOR EARLY HCC

Classical HCC is diagnosed based on CT images with early arterial enhancement and delayed washout (EASL criteria)^[1,24]. Various guidelines have also adopted these criteria. Early HCC generally has stromal invasion in the portal region with remaining tumor^[25] and has a macroscopically small nodular type with indistinct margins. Diagnostic imaging identifies this type as an ischemic mass. Prolongation of survival time by liver resection for early HCC is not significant and is limited due to the lead time bias^[26]. This suggests that early HCC should be followed up

Table 2 Treatment options for hepatocellular carcinoma in barcelona clinic liver cancer system and Japanese guidelines

Tumor number	Tumor size (cm)	Child-Pugh class	Treatment	
			BCLC system	Japanese guidelines
Single	2	A, B	Resection	1 Resection 2 Ablation
	2.1-3	A, B	1 Resection 2 Transplantation or ablation	1 Resection 2 Ablation
	3.1-5	A, B	1 Resection 2 Transplantation	Resection
2 or 3 nodules	≤ 3	A, B C	Transplantation or ablation Palliative care	Resection or ablation Transplantation
	> 3	A, B	Chemoembolization	1 Resection 2 Chemoembolization
4 or more nodules		A, B	Chemoembolization	1 Chemoembolization 2 Chemotherapy
		C	Palliative care	Palliative care

Degree of Liver damage replaced Child-Pugh classification as liver function in Japan Revise from ref. [22].

without treatment based on the risk of a second primary cancer. This strategy is accepted according to the HCC management based on the consensus in the Japan Society of Hepatology^[27].

EVIDENCE FOR EFFICACY OF HEPATECTOMY

The indication of hepatectomy for HCC is determined by the balance between liver function and tumor conditions. Excessive liver resection to completely remove lesions based on overestimation of hepatic functional reserve may cause hepatic failure, whereas minimal resection that does not correspond to the degree of tumor progression and focuses only on safety may increase the risk of early recurrence of HCC. Therefore, it is important to select an optimal approach that is appropriate for the degree of tumor progression based on the indication for hepatectomy. The major methods for preoperative assessment of liver function are the galactose tolerance test, ^{99m}Tc-GSA liver scintigraphy, and the ICG loading test. Makuuchi's criteria are particularly useful for patients with chronic hepatitis or hepatic cirrhosis^[14]. These criteria are based on three factors: ascites, serum bilirubin, and ICGR₁₅. Patients who still have ascites after diuretic administration or those with a serum bilirubin level that is consistently > 2.0 mg/dL are not indicated for surgery. The patients with 1 < bilirubin level ≤ 2.0 mg/dL are indicated for limited liver resection. For eligible patients with serum bilirubin in the normal range of ≤ 1.0 mg/dL, the extent of resection is then determined based on ICGR₁₅ as resection of 2/3 of the total liver volume (TLV) (*e.g.*, right lobectomy) in patients with normal ICGR₁₅ of < 10%; 1/3 of the TLV (*e.g.*, left lobectomy) in those with ICGR₁₅ of 10%-19%; and 1/6 of the TLV (Couinaud segmentectomy) in those with ICGR₁₅ of 20%-29%. If ICGR₁₅ is ≥ 30%,

limited resection or enucleation should be applied. A surgical mortality rate of 0% has been reported in 1056 consecutive hepatic resections performed in accordance with these criteria^[28].

In portal venous invasion of HCC^[29], the area supplied by the portal vein branches should be systemically removed as much as possible within the acceptable range of liver function. A new procedure of systematic subsegmentectomy has been developed to overcome the potential incompatibility between cure of cancer and preservation of liver function^[30]. A study of survival after hepatectomy indicated a good prognosis in cases with a tumor diameter < 5 cm, a single lesion, capsule formation, no vascular invasion, serum albumin < 4.0 g/dL, and pathological TNM (pTNM) stage I or II. Of these parameters, pTNM stage is the most reliable prognostic factor^[31]. A study of recurrence-free survival also identified the significant prognostic factors as the tumor stage, tumor size, number of tumors, and capsule formation, and also found that vascular invasion was a poor indicator of long-term survival^[32]. Risk factors for early recurrence within 2 years postoperatively include non-anatomical resection, microscopic vascular invasion, and AFP ≥ 32 ng/mL^[33]. A retrospective study showed that the cumulative survival rate was significantly greater after anatomical resection compared to that after non-anatomical resection, which suggests that the surgical technique can influence prognosis^[34]. A future prospective study is required to clarify all of these findings.

Determination of the acceptable liver remnant volume after hepatectomy is an important task. In general, it is desirable to preserve the 20%-40% of the total liver volume (TLV) or the standard liver volume (SLV) in normal livers^[35-42]. The MD Anderson group proposed that the smallest acceptable liver remnant volume is ≥ 20% of the SLV in cases

without chronic underlying liver disease^[36], with the validity of this proposal supported by an analysis of 301 consecutive patients after extended right lobectomy^[43]. On the other hand, there was a mortality rate on postoperative day 60 of 4.7% in this literature cited. However, HCC often develops in livers with chronic hepatitis or hepatic cirrhosis, and major hepatectomy such as lobectomy may induce hepatic failure due to insufficient liver remnant volume. Portal vein embolization (PE) prevents hepatic failure since the portal vein branches in hepatectomy are blocked to induce compensatory hypertrophy in the remnant liver area^[44]. PE can be applied to cases with ICGR₁₅ < 10% and a ratio of nontumorous parenchymal volume of the resected liver to that of the whole liver (R2) ≥ 60%, and those with ICGR₁₅ ≥ 10% - < 20% and R2 of 40%-60%^[35]. Three-dimensional CT permits simple and accurate determination of the relative positions of major blood vessels and the tumor, resection ranges, and liver remnant volume^[45].

HEPATIC RESECTION

In liver surgery, hepatic parenchymal transection is associated with increased intraoperative blood loss, postoperative hemorrhage, and early complications such as bile leakage and surgical site infection (SSI). In addition to hemostasis, new devices have been developed to stop bleeding from the resection margin, which allows performance of safer and more secure hepatic resection. The Pringle maneuver, which blocks hepatic inflow once by manual compression of the hepatoduodenal ligament to minimize blood loss during hepatic resection, is also widely used. Several randomized clinical trials (RCTs) have shown that the Pringle maneuver reduces blood loss without affecting liver function^[46,47]. Hemihepatic vascular occlusion has also been applied when resection is limited to one lobe^[48,49].

Bleeding from the hepatic vein occurs most commonly during hepatic resection. Intraoperative hemorrhage is positively associated with central venous pressure (CVP) and several RCTs have shown that a decrease of CVP to ≤ 5 cm H₂O during hepatectomy reduces intraoperative blood loss and stabilizes hemodynamics^[50,51]. In contrast, infrahepatic inferior vena cava clamping with a low CVP has been shown not to reduce blood loss during hepatectomy^[52], and thus the effects of low CVP require further study.

Hepatic parenchymal transection is performed using methods such as clamp crushing^[53] and devices including the cavitron ultrasonic aspirator (CUSA)^[54], Tissue Link^[55], water jet scalpel^[56], harmonic scalpel^[57], floating ball^[58], and LigaSure. In clamp crushing, a Pean clamp is used to ligate and resect remaining blood vessels after the hepatic parenchyma is crushed using the clamp. In RCTs, there were

no differences in operating time, volume of blood loss, and incidence of postoperative complications between patients treated with clamp crushing and CUSA, but clamp crushing was superior in terms of complete appearance of landmark hepatic veins on the cut surface^[53]. However, volume of blood loss and incidence of postoperative complications have also been reported to be lower using CUSA compared with clamp crushing^[59]. A RCT comparing clamp crushing with Tissue Link found no differences in operating time, volume of blood loss and incidence of postoperative complications^[60]. Another RCT showed the superiority of the LigaSure Vessel Sealing System for liver resection compared to vascular ligation based on clamp crushing^[61], but a second RCT found no differences between these techniques^[62].

PERIOPERATIVE MANAGEMENT

Since 1990, hepatectomy for HCC has been performed with acceptable blood loss of approximately 500 ml at many high-volume medical centers^[28,63-67]. Allogenic blood transfusion in the perioperative period should be avoided when possible because it is likely to promote cancer recurrence and to induce hyperbilirubinemia and hepatic failure, and lower hematocrit is also desirable for microcirculation in the liver^[68]. Autologous blood transfusion avoids homogenous red blood cell transfusion and does not increase the frequency of cancer recurrence^[69]. The use of fresh frozen plasma (FFP) has been recommended for supplement of coagulation factors, maintenance of an effective plasma volume, and volume substitution^[70]. However, FFP transfusion has also been reported to have no effect on the post-hepatectomy course^[71] and to be unnecessary in Child-Pugh class A cases with intraoperative blood loss of < 1000 mL and serum albumin levels > 2.4 g/dL on postoperative day (POD) 2^[72].

PREVENTION OF COMPLICATIONS

Bile leakage is a complication that is specific to hepatectomy and may be intractable. A RCT of the efficacy of a bile leakage test on prevention of bile leakage from the liver resection margin showed no difference in the incidence of bile leakage between patients who did and did not receive the test^[73], whereas another RCT found that the test was able to prevent bile leakage and complications after hepatic resection^[74]. Thus, more cases are required to evaluate the utility of this test.

Other post-hepatectomy complications include hemorrhage and intra-abdominal abscess, and these conditions may be fatal if diagnosis is delayed. Intraperitoneal drain placement is required for monitoring and treatment of these complications, but the efficacy of elective hepatectomy with standardized drain placement has been questioned and the

Centers for Disease Control and Prevention (CDC) guidelines state that such routine drain placement is not necessary: "If drainage is necessary, use a closed suction drain. Place a drain through a separate incision distant from the operative incision. Remove the drain as soon as possible"^[75]. RCTs conducted in several countries on the need for drainage have also concluded that drain placement is not necessary^[76-81]. Due to differences in the healthcare environment and health insurance system, drain placement has not been completely withdrawn in Japan, but early removal of drains has been recommended^[82]. A RCT has also shown that subcutaneous drainage is not effective for prevention of SSI^[83].

Immunity is weak after hepatectomy and this may result in hepatic failure and disseminated intravascular coagulation (DIC). A RCT of the efficacy of steroid administration for improvement of liver function after hepatectomy compared the post-hepatectomy liver function in patients treated with and without 500 mg/body hydrocortisone before hepatectomy^[84]. Serum bilirubin levels were significantly lower in the steroid group on POD 2 compared with the non-steroid group and there were significant differences in serum bilirubin and prothrombin levels until POD 7, which shows the efficacy of steroid administration prior to hepatectomy. To unify the definition of post-hepatectomy liver failure (PHLF), in 2010 the International Study Group of Liver Surgery (ISGLS) proposed defining PHLF as an increased international normalized ratio (INR) and concomitant hyperbilirubinemia on or after POD 5^[85]. PHLF seems to be the more efficient indicator comprehensively compared to 50-50 criteria^[86] and MELD score because it is significantly associated with both of the incidence of post-hepatectomy complications and the post-hepatectomy mortality^[87]. As for 50-50 criteria, it was not significantly related to the incidence of post-hepatectomy complications. As for MELD score, it revealed less strong association of the odds ratio (2.06) to the post-hepatectomy mortality.

CONCLUSION

In this article, we have described evidence-based techniques for hepatectomy and perioperative management of HCC. Improved assessment of liver function and development of surgical devices are likely to contribute to safe and effective hepatectomy and a good prognosis for patients.

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Immune response to hepatitis B vaccine among patients on hemodialysis

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electronic systematic search of the published literature was carried and data on the immunological response to hepatitis B vaccination among hemodialysis patients was extracted from relevant studies. End stage renal disease patients on hemodialysis have a lower or an absolutely negative response to HBV vaccine. Several means have been tried to improve this response with some success, nevertheless none have been universally adopted. Genetic investigations are foreseen to make a break through concerning HBV vaccination.

Key words: Hemodialysis; Chronic kidney disease; Immune response; Vaccine; Adjuvant

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Core tip: This article discussed the history of immunological response to various types of Hepatitis B vaccines among patients on hemodialysis based on published findings of an array of studies up to this year. Moreover, it tackled the possible causes for such a response and possible future ways out of this dilemma.

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Abstract

Infection with hepatitis B virus (HBV) poses a major health threat worldwide, where the magnitude and overburden of chronic carrier state approaches 150 million chronic carriers. The prevalence of HBV is greater among dialyzed patients compared to the general population owing to their increased vulnerability to blood and its products, along with hazards posed by contaminated hemodialysis tools and devices. An

INTRODUCTION

Infection with hepatitis B Virus (HBV) poses a major health threat worldwide, where the magnitude and overburden of chronic carrier state approaches 150 million chronic carriers. The prevalence of HBV is greater among dialyzed patients compared to the general population, this could be attributed to the

fact they are needy for the blood and its products and thus more vulnerable, along with the jeopardy posed by contaminated hemodialysis tools and devices^[1]. Therefore HBV immunization is highly advised for patients suffering chronic kidney disease (CKD), whether pre-emptive or dialysis dependent, who are potential nominees for kidney transplant along with those on dialysis^[2]. In spite of this, surges of the aforementioned infection among patients on hemodialysis are persistently encountered even in advanced countries^[3-5].

The extent of renal failure has been described to determine the immunological riposte to hepatitis B immunization among this group of patients, where it has generally been reported to be suboptimal^[6-8]. The seroconversion rates and antibody titers among chronic renal disease patients has been shown to be less than the general population along with a shorter duration of seroprotection^[9]. The flawed effectiveness of HBV immunization among dialysis dependent patients can be justified by a group of determinants, noticeably the defective immunity owing to; azotemia, age^[10], sex^[11], body mass^[12], nourishment of such patients^[13], concomitant infection with HCV^[14] or HIV^[15], history of transfusion of blood or blood products^[16] and having the major histocompatibility complex also to be associated with this response^[17-19], along with failure to complete the full course of HBV vaccine^[20].

In order to enhance the riposte degree to hepatitis B immunization in end-stage kidney disease a group of strategies have been embraced and these are; building up the vaccine dose^[20], supplementary vaccine injections along with resorting to intradermal injections rather than the intramuscular in order to supplement the vaccination^[21]. Hepatitis B vaccination among pre-dialysis chronic kidney disease patients results in higher seroprotection rates when compared to those patients on dialysis^[22]. Adjuvants has been proposed to be of some help in raising the immunity, a good example for these adjuvants is levamisole, which is an antihelminthic with characteristics enabling it to stimulate suppressed T-cell action along with potentiation of B lymphocyte action^[23-25]. The target of this review is to discuss inadequate immunological riposte to hepatitis B immunization, its determinants and the possible solutions.

MANAGEMENT

Search strategy and data extraction

An electronic systematic search of the published literature was carried and data on the immunological riposte to hepatitis B vaccination among hemodialysis patients was extracted from studies of relevance. The databases were searched with the words "Hepatitis B immunization", "dialysis", "immunological riposte", "retarded riposte", "non-responders" and

"adjuvants" were used interchangeably in MEDLINE, Pubmed, MiPc library and Google.

Epidemiology of HBV among hemodialysis

Hepatitis B infection has been declining in the last two decades in artificial kidney facilities, a status that mirrors the outcomes of efforts made in providing efficient prophylaxis measures^[26]. Variable prevalence of HBV infection among dialysis patients were reported from the different continents ranging between (6%), and (1.2%)^[27-29]. In a large scale study including 8615 adult dialysis patients from different dialysis facilities in the Western world, hepatitis B prevalence rates ranged from 0% to 6.6%^[5]. A principal determinant hindering the transmission of such an infection in artificial kidney facilities was the preservation of universal infection control measures. CDC guidelines advice segregation of patients who are antigen-positive, dedicating an independent nursing group and it further more prohibits sharing medicaments in artificial kidney facilities^[30]. Undiminished vulnerability percentages to hepatitis B infection were particularly observed in renal facilities dealing with HB (S) Ag carriers. Such vulnerability could be controlled by the strict cohesion to the global precautions; however causal incidences can to an outbreak the whole facility^[30,31]. History of immunization against hepatitis B targeting end stage renal disease sufferers started by the utilization of live attenuated virus derived from plasma, although it was initially reported to mount enough immunity, however, it was found later not to have induced a sufficient immunity. Currently supplied vaccines possess an outstanding safeness and immunogenicity account, providing protection rates falling just below 100% of the immunized group^[32]. However, some population subgroups, including some people of normal health and immune-deficient individuals, riposte inadequately to immunization. Part of such sets, are chronic kidney disease patients, including pre-emptive and dialysis patients, whom are regarded to have vulnerability to contract HBV owing to transmission to those on dialysis through surrounding surfaces, expendables, or apparatuses during hemodialysis^[33,34]. Hepatitis B vaccination, when combined with application of the other precautions, ended up in a definite and appreciable decline in new infections among hemodialysis patients and kidney facility personnel in Western countries^[35,36]. Despite the fact that the frequency of HBV is absolutely squatty, a big proportion of vulnerable chronic kidney disease sufferers have to get the vaccine. Regulations meant with control of transmission of infections in renal facilities gives a feeling of protection to working personnel, nevertheless, chronic kidney disease sufferers' vaccination is yet regarded a subsidiary and pricy procedure, there for resulting in

a greater proportion of unimmunized chronic renal disease sufferers in some countries. Hepatitis B poses an intimidation for chronic renal sufferers on hemodialysis, regardless of precautions secured, let alone meanwhile they are subjected to hemodialysis in their local facilities, but again meanwhile accommodated by other units when considering superior chronic kidney disease sufferers' acclimation, vacation enjoyment. During such plots, the susceptible chronic kidney disease sufferers' acts as a probable HBV infection aim (undertaking dialysis in machines, units dedicated for infected patients) moreover they act as a probable harbor for the disease taking it back to their local facilities^[37]. It should be noted that HBV can also be transmitted to dialyzed chronic kidney disease sufferers (as the rest of the general population) by other means known to transmit the disease (sexual route *etc.*). Those who caught the hepatitis infection can spread the disease in their home unit in turn, ahead of hepatitis B infection detection, unless active steps take place to protect patients from such an infection in terms of a fruitful immunization plan^[38]. A great proportion of hemodialyzed chronic kidney disease sufferers acquiring HBV have a tendency to progress to chronic hepatic disease (unable to eliminate their virus). Such chronic kidney disease sufferers are rendered to have greater vulnerability on attempting kidney transplantation, further more they pose a potential harbor for the disease to both other chronic kidney disease sufferers and non-immune working personnel^[39-41]. The aforementioned fact makes a comprehensively fruitful immunization plan imperative for chronic kidney disease sufferers and staff safeguard against this lifelong, enduring and possibly killing infection^[42].

BOOSTING THE IMMUNOLOGICAL RESPONSE TO VACCINATION

Evolution of the vaccine

Comparative to numerous other infections, immunization, as a protective strategy, performs a crucial role in limitation of the HBV infection and its consequences^[9]. Hitherto, there are triumvirate derivations of HBV vaccines. Saul Krugman's observation about immunogenicity of HBsAg and the immunizing properties of anti-HBs antibody facing HBV was a real breakthrough that resulted in producing the early vaccine derivation^[26] incorporating an inactive HBsAg extracted out of the plasma of the HBV carrier persons. Merck and Pasteur institute simultaneously produced the early vaccine derivation making use of aforementioned observation. Then, Food and Drug Administration of the United States of America approved it during 1981^[26]. The second derivation of HBV vaccine was engineered using recombinant DNA technology utilizing the yeast

Saccharomyces cerevisiae resulting in the formulas; Engerix B along with Recombivax HB. Both vaccine formulas encompass HBsAg. Nevertheless, the third vaccines derivation mounts an appreciably greater protection if compared to HBsAg owing to the use of pre-S1 along with pre-S2 immune triggers; however, their availability is yet limited. Recombinant DNA technology has also been used to produce the third generation vaccines by mammalian cells^[26,43]. Different American, European and Asian states adopted The WHO recommendation 1991 concerning large-scale HBV immunization by year 1997. Therefore there is an appreciable drop in pervasiveness of the HBV infection^[43,44], and its complications including HCC and fulminating hepatitis^[44].

Adjuvants

Several methods have been suggested to potentiate the outcome of HBV immunization and its riposte among chronic kidney disease sufferers on hemodialysis. Use of adjuvants was suggested to potentiate the riposte to immunization. Examples for such adjuvants are; high thymopentin doses^[45]. Levamisole is another adjuvant that has been suggested to improve vaccination results among such patients. It is probable that it can also be efficient in boosting HBV vaccination riposte among HD dependent patients^[46], however in terms of taking a rather mature decision it is prudent to conduct further research in this field in order to now the pros and cons of such agents. Fabrizi *et al.*^[47] found in their meta-analysis that a better immune response is mounted when GM-CSF is added as an adjuvant to HBV vaccine. Polymethylmethacrylate is another adjuvant that has been proposed to improve the immunity post vaccination^[48]. HBV-AS04 encompassing the synergist 3-O-desacyl-40-monophosphoryl lipid A that is consistent with Engerix B customary, is a further adjuvant improving the immune response^[26]. A Recent research conducted by Saade *et al.*^[49] has found that Advax (a polysaccharide adjuvant) induces a potent humoral and cellular induced immunity with minimal reactions in the preclinical phase. Yet most of the studies in this context have some methods limitation such as lack of randomization^[50]. Thus a long term data about the sustainability of the effect of these adjuvants is to be verified.

Changing the route of administration and booster doses

Currently, HBV vaccinations, particularly the second derivation, are administered through the intramuscular (deltoid) route three times (on 0, 1 and 6 mo's period). Antibodies' to Anti-HBs titers above 10 IU/L are deemed effective. Considering revaccinating subjects or boosting a currently administered vaccine is required in situations where

the titers level falls beneath 10 IU/L such as time linked falls in titers, or those seen among high risk groups such as immunosuppressed, smokers, obese persons, kidney failure patients and those suffering hepatic disease^[9,43].

Currently, administration of HBV immunization through either the intradermal and intramuscular injection routes is under evaluation in chronic kidney disease sufferers who undergo HD. Short term follow up, reflected that the former route of HBV immunization can mount a more potent immune riposte if compared to the later route^[26]; nevertheless, such a believe has been refuted by long term follow up.

FUTURE PROSPECTS

Genetic investigations might help in the improvement of hepatitis B vaccines and there for it may lead to reduction in the proportion of vaccine failures^[51]. Increased interferon (IFN)-gamma production was shown to be associated with positive response to vaccines^[52,53]. As IL-18 is involved in IFN-gamma production^[54-57], it was used as adjuvant to DNA vaccines against HBV^[53,57,58]. Channarong *et al*^[57] has devised a recombinant plasmid bearing a gene encoding HBsAg combined to DNA segment encoding full-length murine IL-18. All immunized mice showed a remarkable serum anti-HBsAg IgG response following two intramuscular injections of the vaccines on comparison to the level of mice vaccinated with the vaccine devoid of the DNA segment encoding IL-18. Recently Hu *et al*^[59] found on animal experiments that adding calcineurin B subunit to Engerix mounted an higher hepatitis B antibodies both in dose and time dependent manner through promoting an inflammatory response where IFN- γ , IL-6, TNF- α are produced^[59]. It is probable that in the near future all people throughout the world will be vaccinated on the mandatory basis.

CONCLUSION

Chronic kidney disease patients on hemodialysis tend to have no or at most a lower response to HBV vaccine. Several means have been tried to improve this response with some success, nevertheless, absolutely none have been universally adopted. Genetic investigations are foreseen to make a breakthrough concerning HBV immunization.

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Retrospective Study

Post-transplantation hepatocellular carcinoma recurrence: Patterns and relation between vascularity and differentiation degree

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Ethics approval: This work has been carried out in accordance with the Declaration of Helsinki (2000) of the World Medical Association.

Informed consent: All patients gave their verbal consent to the use of their data for scientific purposes at the moment of their admission to the Transplantation Department.

Conflict-of-interest: All authors have no conflicts of interest to declare.

Data sharing: Technical appendix, statistical code, and dataset available from the corresponding author at anna_pecchi@yahoo.com.

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Abstract

AIM: To evaluate the relationship between hepatocellular carcinoma (HCC) vascularity and grade; to describe patterns and vascular/histopathological variations of post-transplantation recurrence.

METHODS: This retrospective study included 165 patients (143 men, 22 women; median age 56.8 years, range 28-70.4 years) transplanted for HCC who had a follow-up period longer than 2 mo. Pre-transplantation dynamic computed tomography or magnetic resonance examinations were retrospectively reviewed, classifying HCC imaging enhancement pattern into hypervascular and hypovascular based on presence of wash-in during arterial phase. All pathologic reports of the explanted livers were reviewed, collecting data about HCC differentiation degree. The association between imaging vascular pattern and pathological grade was estimated using the Fisher exact test. All follow-up clinical and imaging data were reviewed for evidence of recurrence. Recurrence rate was calculated and imaging features of recurrent tumor were collected, classifying early and late recurrences based on timing (< or ≥ 2 years after transplantation) and intrahepatic, extrahepatic and both intrahepatic and extrahepatic recurrences based on

location. All intrahepatic recurrences were classified as hypervascular or hypovascular and the differentiation degree was collected where available. The presence of variations in imaging enhancement pattern and pathological grade between the primary tumor and the intrahepatic recurrence was evaluated and the association between imaging and histopathological variations was estimated by using the χ^2 test.

RESULTS: Of the 163 patients with imaging evidence of viable tumor, 156 (95.7%) had hypervascular and 7 (4.3%) hypovascular HCC. Among the 125 patients with evidence of viable tumor in the explanted liver, 19 (15.2%) had grade 1, 56 (44.8%) grade 2, 40 (32%) grade 3 and 4 (3.2%) grade 4 HCC, while the differentiation degree was not assessable for 6 patients (4.8%). A significant association was found between imaging vascularity and pathological grade ($P = 0.035$). Post-transplantation recurrence rate was 14.55% (24/165). All recurrences occurred in patients who had a hypervascular primary tumor. Three patients (12.5%) experienced late recurrence; the location of the first recurrence was extrahepatic in 14 patients (58.3%), intrahepatic in 7 patients (29.2%) and both intrahepatic and extrahepatic in 3 patients (12.5%). Two patients had a variation in imaging characteristics between the primary HCC (hypervascular) and the intrahepatic recurrent HCC (hypovascular), while 1 patient had a variation of histopathological characteristics (from moderate to poor differentiation), however no association was found between imaging and histopathological variations.

CONCLUSION: A correlation was found between HCC grade and vascularity; some degree of variability may exist between the primary and the recurrence imaging/histopathological characteristics, apparently not correlated.

Key words: Contrast media; Hepatocellular carcinoma; Liver transplantation; Cell differentiation; Recurrence

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Core tip: During hepatocarcinogenesis, besides the differentiation loss, blood supply changes occur. Recently, a correlation between higher histopathological grades and hypervascular dynamic-imaging enhancement pattern has been demonstrated. Hepatocellular carcinoma recurrence after transplantation is relatively common, however the issue of possible variations in imaging and histopathology between the primary and the recurrent tumor, and particularly the relationship between enhancement and grade changes, has never been investigated. We demonstrated a correlation between vascularity and pathological grade in a large population of transplanted patients, and some degree of variability between the primary and the recurrent tumor vascularity was found, though not associated with histopathological changes.

Pecchi A, Besutti G, De Santis M, Del Giovane C, Nosseir S, Tarantino G, Di Benedetto F, Torricelli P. Post-transplantation hepatocellular carcinoma recurrence: Patterns and relation between vascularity and differentiation degree. *World J Hepatol* 2015; 7(2): 276-284 Available from: URL: <http://www.wjgnet.com/1948-5182/full/v7/i2/276.htm> DOI: <http://dx.doi.org/10.4254/wjh.v7.i2.276>

INTRODUCTION

Hepatocellular carcinoma (HCC) is the most frequent primary hepatic malignancy^[1], and it represents an important health issue due to its increasing incidence and poor survival^[2]. HCC usually arises in the background of chronic liver diseases, in particular cirrhosis is the substrate of HCC in 80%-90% of cases^[3]. Carcinogenesis in HCC is typically a multistep process, that comprehends low-grade and high grade dysplastic nodules possibly with neoplastic foci, early HCC, and overt HCC. Pathologically, HCC is graded based on differentiation degree into four degrees of cellular dysplasia and architectural tissue disarrangement (well-differentiated, moderately differentiated, poorly differentiated and undifferentiated, grade 1 to 4)^[4]. During the hepatocarcinogenesis multistage process, besides the size growth and the loss of differentiation of the nodule, blood supply changes progressively occur, so that the hepatocellular nodule becomes more and more dependent on newly formed arteries and in parallel less dependent on the portal contribution^[3].

Contrast-enhanced dynamic imaging, performed both with multidetector CT (MDCT) and magnetic resonance imaging (MRI), provides information about HCC vascularity. In particular, two different hemodynamic patterns have been featured: the typical, much more common, hypervascular pattern, and the less frequent hypovascular variant of HCC, which shows no arterial phase hypervascularity^[5]. Recently, some studies have compared contrast enhanced dynamic imaging findings with histopathological differentiation degree, showing a correlation between higher pathological grades and hypervascular enhancement pattern^[6-10], even though some investigators have reported a subsequent decrease in arterial blood supply in the late stage of HCC development (grade ≥ 3)^[7-9].

Liver transplantation (LT) is the preferred treatment for selected patients with HCC. However, even after the introduction of selection criteria such as Milan Criteria^[11], HCC recurrence rate after LT has been estimated to be 8%-17%^[12-15]. Many investigators have reported on the spectrum of imaging findings of HCC recurrence after LT, distinguishing different patterns on the basis of recurrence location^[12,13,16] or timing^[14]. In particular, the majority of tumor burden in recurrent HCC is typically in extrahepatic locations^[12,15,16] and the average time to recurrence

Table 1 Demographic and clinical characteristics of the study population

Demographic and clinical characteristics (n = 165)				
Sex	Male (%)			143 (86.7)
	Female (%)			22 (13.3)
Median age (range)				56.8 (28; 70.4)
HIV + (%)				15 (9.1)
Etiology of the underlying hepatic disease	Viral (%)	136 (82.4)	HBV-related	28 (17.0)
			HCV-related	92 (55.8)
			Mixed	16 (9.7)
	Not viral (%)	29 (17.6)	Alcoholic	16 (9.7)
			Cryptogenetic	6 (3.6)
			Other	7 (4.2)

ranges between 1 and 2 years after LT^[14,16]. The hemodynamic imaging characteristics of recurred HCC have been scarcely reported, however, similarly to the primary HCC, the majority of recurred HCC appear as hypervascular lesions.

To our knowledge, the issue of possible variations in imaging and histopathological characteristics between the primary and the recurrent HCC, particularly referring to the relationship between enhancement pattern changes and differentiation degree changes, has never been investigated. The preliminary objective of this study was to evaluate the relationship between HCC contrast-enhanced dynamic imaging pattern and pathological differentiation degree in a population of patients transplanted for HCC. Additional aims were to describe the patterns and imaging features of HCC recurrence after LT and to evaluate the variations in imaging and histopathological characteristics between the primary HCC and the intrahepatic recurrence, particularly elucidating whether differentiation degree variations may justify contrast-enhanced imaging pattern changes.

MATERIALS AND METHODS

Patients

Between October 2004 and November 2011, a total of 172 consecutive patients with known HCC underwent LT at our hospital. During this period, another patient, transplanted without known HCC, had pathologic diagnosis of incidental HCC in the liver explant. Of these 173 patients, 8 patients were excluded because of a short follow up period (≤ 2 mo), as a result of perioperative mortality. The remaining 165 patients (143 men, 22 women; median age 56.8 years, range 28-70.4 years), who had a follow-up period longer than 2 mo, were included. Fifteen of them were HIV-infected patients. Clinical data about the etiology of the underlying liver disease, were collected. Table 1 summarizes demographic and clinical characteristics of the included patients.

Pre-transplantation imaging

All available MDCT and MRI dynamic hepatic examinations were reviewed, selecting those with evidence

of viable tumor, and thus where vascularity was assessable. MDCT examinations were performed using a 64-slice CT scanner (Lightspeed VCT, GE Medical Systems, Milwaukee, Wisconsin, United States) with contrast enhancement and bolus-tracking technique to obtain a multiphase (arterial, portal and hepatic venous phases) examination after an unenhanced scan. Image reconstruction was obtained with a 2.5 mm slice thickness and a 2.5 mm interval. Dynamic MRI studies were conducted on a 1.5-T high field magnet (Philips Achieva, Philips Medical System, Best, The Netherlands) with a Phased Array coil. The protocol included axial T1- and T2-weighted sequences with and without fat suppression and axial dynamic three-dimensional T1-weighted fat-suppressed GRE sequences obtained before and after a bolus injection of gadopentetate dimeglumine (Gd-DOTA) in arterial, portal and hepatic venous phases.

Pre-transplantation imaging examinations were retrospectively reviewed by two experienced radiologists by consensus reading, both blinded to the results of the pathologic reports. From the last pre-LT examination with evidence of viable tumor, data about the HCC enhancement pattern were collected. In particular, based on the presence or absence of wash-in, which was defined as present when the lesion was hyperattenuating compared to the surrounding hepatic parenchyma during the arterial phase, lesions were classified into hypervascular and hypovascular.

Histopathology of the explanted liver

A pathologist experienced with liver pathologies reviewed all pathologic reports of the explanted livers, collecting data about the differentiation degree, scored according to the World Health Organization criteria^[4] into well-differentiated (Grade 1), moderately differentiated (Grade 2), poorly differentiated (Grade 3), and undifferentiated (Grade 4) types. When different degrees were reported in the same explanted liver, the prevailing grade (the one demonstrated by the larger number of nodules) was considered.

Recurrence analysis: Imaging and histopathology

All available postoperative dynamic imaging examin-

Table 2 Pre-Transplantation imaging enhancement pattern and histopathological differentiation degree

Pre-LT imaging and transplant pathology		
Pre-LT imaging		
Enhancement pattern (n = 163)	Hypervascular (%)	156 (95.7)
	Hypovascular (%)	7 (4.3)
Histopathology		
Differentiation degree (n = 125)	Grade 1 (%)	19 (15.2)
	Grade 2 (%)	56 (44.8)
	Grade 3 (%)	40 (32)
	Grade 4 (%)	4 (3.2)
	Not assessable (%)	6 (4.8)

ations (MDCT or MRI) were retrospectively reviewed for evidence of recurrent HCC. Proof of recurrence was made on the basis of biopsy or growth of new lesions with appropriate radiologic features, combined with rising AFP levels or with negative work-up for another primary malignancy. Imaging features of recurrent HCC were collected. Based on recurrence timing, recurrences were divided into early (< 2 years after LT) and late (≥ 2 years after LT). With respect to tumor location at the moment of the first recurrence, three different patterns were distinguished: intrahepatic recurrence (allograft itself), extrahepatic recurrence and both intrahepatic and extrahepatic recurrence.

All follow-up dynamic imaging examinations were also reviewed to describe the enhancement pattern of all intrahepatic recurrences (those that occurred at the moment of the first recurrence and subsequent ones), in particular classifying lesions into hypervascular and hypovascular based on the presence of wash-in.

Intrahepatic recurrence histopathological differentiation degree was obtained by a review of the available pathological reports (in case of biopsy or resection of the recurred HCC).

Statistical analysis

The association between pretransplantation HCC enhancement pattern (hypervascular or hypovascular) and explanted liver HCC differentiation degree was evaluated by using the Fisher exact test. Recurrence rate was calculated. The presence of variations in imaging features (enhancement pattern) and histopathological characteristics (differentiation degree) between the primary and the intrahepatic recurred HCC was evaluated and the association between imaging and histopathological variations was estimated by using the χ^2 test. For all statistical analyses, a $P < 0.05$ was considered to indicate a statistically significant difference.

The statistical methods of this study were reviewed by Marta Di Nicola from Department of Experimental and Clinical Sciences, Laboratory of Biostatistics, University of Chieti, Italy.

RESULTS

Pre-transplantation imaging and histopathology of the explanted liver

Of the 165 patients who were included, 2 had no evidence of viable tumor in all dynamic imaging examinations performed within 6 mo to LT. Of the remaining 163 patients in whom evaluation of enhancement pattern was possible, 156 (95.7%) had evidence of hypervascular HCC (Table 2). Of these 163 patients, 125 patients had evidence of viable tumor in the explanted liver, while the remaining 38 patients had completely necrotic nodules as a result of pre-transplantation loco-regional therapies performed after imaging examinations. The distribution of the pathological differentiation degree of the primary HCC over the 125 patients who had viable tumor is summarized in Table 2. The differentiation degree was not assessable for 6 patients (4.8%). Different degrees were shown in the same explanted liver in 5 cases, and in such cases the prevailing grade was considered.

Both enhancement pattern and differentiation degree were available for 113 patients. Among them, a significant association was found between imaging enhancement pattern and histopathological differentiation degree ($P = 0.035$). As shown in Table 3, 50% (3/6) of the patients with a hypovascular HCC had a well-differentiated tumor, vs 14% (15/107) of those with a hypervascular HCC. An explicative case of a well-differentiated HCC which was characterized by an atypical enhancement pattern in the pre-LT dynamic MDCT examination is depicted in Figure 1.

Recurrence analysis

Of the 165 patients included, 24 (14.55%) had evidence of HCC recurrence after the LT. The 1-, 3-, 5-years cumulative disease-free survival rates according to the Kaplan-Meier method were 92.96%, 83.9% and 82.84%, respectively. The mean duration of recurrence-free survival was 40.15 mo.

Time to development of the recurrence ranged from 1.55 to 41.85 mo after LT, with a median value of 12.36 mo. Three patients (12.5%) experienced late recurrence (≥ 2 years after LT), with a rate of 1.7% (3/175). All recurrences occurred in patients who had a hypervascular primary HCC, while none of the patients with hypovascular primary HCC had a recurrence after LT.

With respect to recurrence location at the moment of the first recurrence, 14 patients (58.3%) showed extrahepatic recurrence, 7 patients (29.2%) had intrahepatic recurrence and 3 patients (12.5%) showed both intrahepatic and extrahepatic recurrence. Eight patients had more than one recurrence site at the moment of the first recurrence. Only one of the 3 patients with late recurrence had intrahepatic recurrence. Recurrence patterns and most common imaging features of recurrence are further described

Table 3 Distribution of dynamic imaging enhancement patterns according to histopathological differentiation degrees

Dynamic imaging enhancement pattern and histopathological differentiation degree (<i>n</i> = 113)					
Enhancement pattern	Grade 1	Grade 2	Grade 3	Grade 4	<i>P</i> value
Hypovascular	3	1	1	1	0.035
Hypervascular	15	52	37	3	

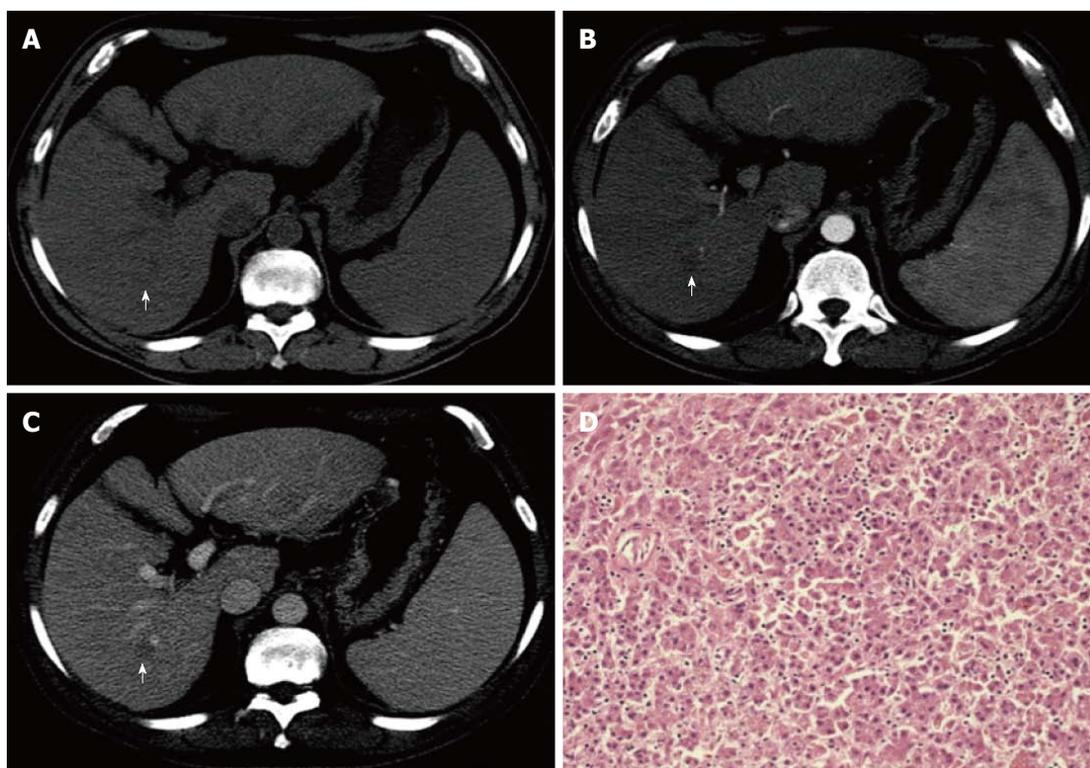


Figure 1 Pre-contrast (A), hepatic arterial (B) and hepatic portal venous (C) phases multidetector CT scans demonstrate a hypovascular hepatocellular carcinoma (arrows), with no hypervascularization during the arterial phase (wash-in). Pathologically (HE stain, $\times 100$) (D), it was classified as well-differentiated (Grade 1).

in Table 4 and some recurrence cases illustrated in Figure 2. When extrahepatic recurrences had a solid component which was large enough to allow vascularity evaluation, they showed a contrast enhancement similar to the primary hypervascular HCC, hence usually already evident in the arterial phase of the examination. Frequently they also presented with some necrotic intralesional component.

Among the 14 patients who didn't show intrahepatic recurrence at the moment of the first recurrence, 4 had evidence of hepatic recurrence in further follow-up imaging examinations. On a total of 14 patients who firstly or subsequently developed an intrahepatic recurrent HCC, 12 had hypervascular and 2 had hypovascular recurrent HCCs, classified based on the presence of wash-in. Evidence of wash out was shown by all intrahepatic recurrences. Nine of them underwent liver biopsy with histological diagnosis of HCC, 4 scored as moderately and 5 scored as poorly-differentiated. Two patients (14.3%) showed a variation in imaging characteristics

between the primary HCC (hypervascular) and the intrahepatic recurrent HCC (hypovascular). However, they didn't show any variation in histopathological characteristics (Figure 3). Only 1 patient, who had hypervascular enhancement pattern on pre-LT imaging as well as on recurrence imaging, had a variation of histopathological characteristics (from moderate to poor differentiation) (Figure 4). Therefore, no association was found between imaging and histopathological variations.

DISCUSSION

It is well known that during HCC development progressive changes in the vascular supply occur^[3]: as dysplastic nodules undergo malignant transformation, abnormal neoplastic arterial supply increases while portal supply decreases. Classic HCC is exclusively supplied by the hepatic artery and lacks a portal venous supply, leading, on dynamic imaging examinations, to the typical hypervascular pattern

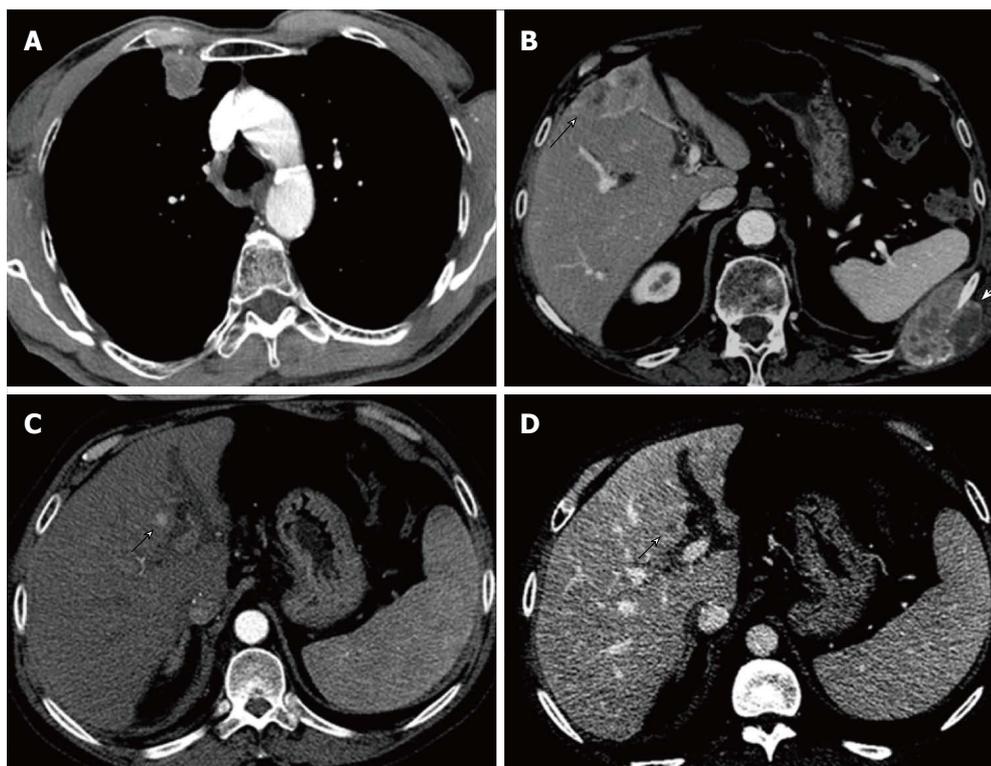


Figure 2 Representative multidetector CT images of different recurrence patterns based on the location at the moment of the first recurrence: (A) Extrahepatic recurrence presenting with a solitary pleural-based lung nodule; (B) Both intrahepatic and extrahepatic recurrence presenting with bone lesions and an intrahepatic lesion in segments IVb and V; (C and D) Intrahepatic recurrence presenting with a solitary hypervascular nodule in segment IV.

Table 4 Recurrence patterns based on the location at the moment of the first recurrence and most common imaging features of recurrence

Recurrence patterns (n = 24)		
Intrahepatic		7 (29.2%)
	Solitary nodule	3 (12.5%)
	Multifocal lesions	4 (16.7%)
Extrahepatic		14 (58.3%)
	Lung	9 (37.5%)
	Solitary nodule	4 (16.7%)
	Multiple nodules	5 (20.8%)
	Consolidation	/
	Bone	5 (20.8%)
	Osteolytic	5 (20.8%)
	Osteoblastic	/
	Lymph nodes	2 (8.3%)
	Brain	1 (4.2%)
	Spleen	1 (4.2%)
	Adrenal	1 (4.2%)
Intrahepatic and extrahepatic		3 (12.5%)

characterized by both wash-in in the arterial phase and wash-out in portal or delayed phases. Less frequently HCCs present as hypovascular lesions, enhancing less than the surrounding liver both on arterial and venous phase imaging, probably as a result of a dual blood supply, both arterial and portal^[17,18].

Some investigators have recently reported a correlation between histopathologic grade and HCC

blood supply in radiological and pathologic analyses. A tendency towards higher grades in tumors with hypervascular pattern was demonstrated^[6-10], however there is some evidence that in the late stage of HCC development, the arterial blood supply decreases again^[7-9].

In our population, comparatively with previous studies^[9], the majority of HCCs were moderately or poorly differentiated, almost all of them being characterized by a typical hypervascular pattern. On the contrary, the few atypical hypovascular HCCs were predominantly distributed in the well-differentiated group, resulting in 50% of patients with hypovascular HCC vs only 15% of those with typical HCC having a grade 1 tumor. Therefore, despite the fact that in our population a very small number of patients had evidence of atypical HCC (4.3%), a correlation was found between differentiation degree and enhancement pattern ($P = 0.035$).

Recently, some authors have hypothesized the utility of imaging pattern as a prognostic factor for tumor outcome after locoregional treatment^[19] or surgery^[20,21]. In our study, the pre-LT imaging examinations of all the 24 patients who experienced HCC recurrence after LT showed hypervascular tumors, and all of them were graded as moderately or poorly differentiated on histopathology ($n = 14$ grade 2, $n = 9$ grade 3, $n = 1$ not assessable). No patient with hypovascular HCC at pre-LT imaging developed

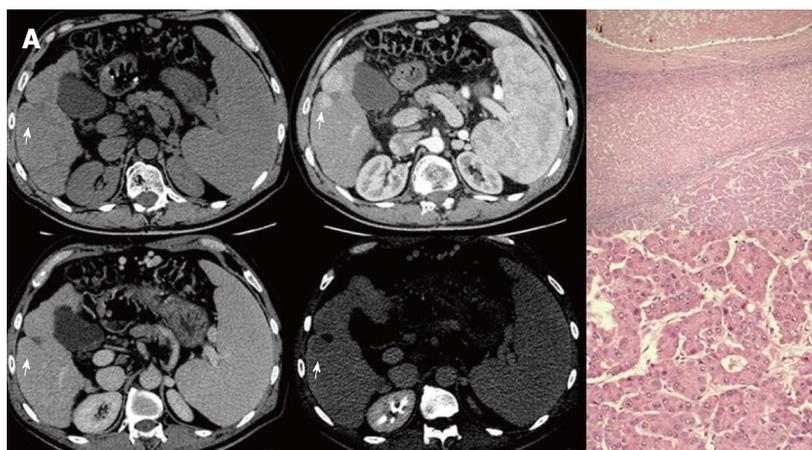


Figure 3 Multidetector CT scans demonstrating two hypervascular primary hepatocellular carcinoma nodules which were graded as moderately differentiated on histopathology (HE stain, magnification × 40 and × 200) (A); Multidetector CT scans of the same patient 2 years after LT demonstrating a hypovascular intrahepatic recurrent hepatocellular carcinoma, which was still graded as moderately differentiated on histopathology after resection (HE stain, magnification × 40 and × 200) (B).

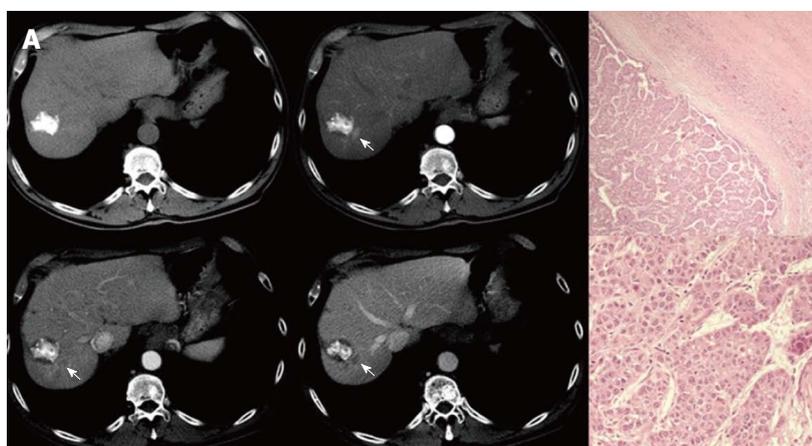
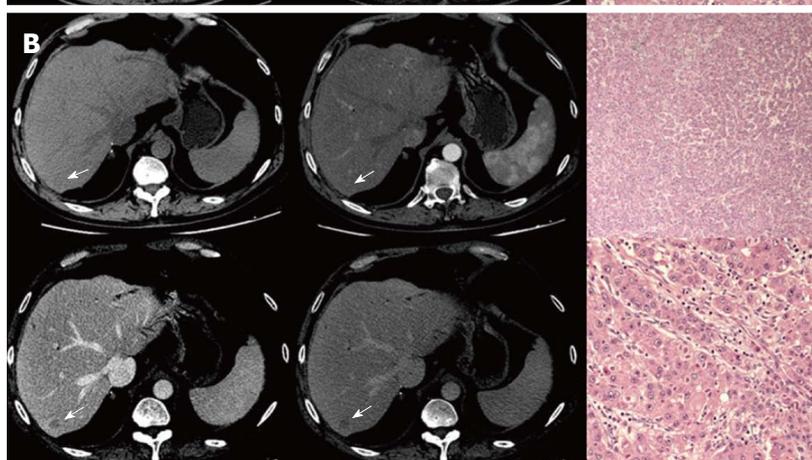
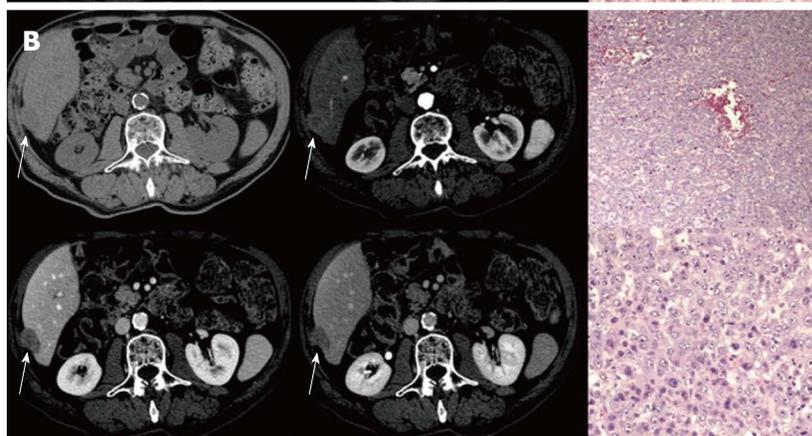


Figure 4 Multidetector CT scans demonstrating a hypervascular area representing viable hepatocellular carcinoma next to a previously lipidolized nodule, graded as moderately differentiated on histopathology (HE stain, magnification × 40 and × 200) (A); Multidetector CT scans of the same patients 1 year after LT demonstrating a hypervascular intrahepatic recurrent hepatocellular carcinoma, which was graded as poorly differentiated on histopathology after resection (HE stain, magnification × 40 and × 200) (B).



post-LT recurrence, not even the few of them with a high histopathological degree. These findings suggest, albeit the number of cases is limited, a lower tendency towards recurrence and a longer recurrence-free survival in patients with hypovascular rather than hypervascular HCCs, underlying the potential prognostic role of vascular supply along with the pathological grade.

In our population the tumor recurrence rate after transplantation was 14.55%, which is similar to the rates observed in other studies with comparable follow-up period and in which Milan selection criteria were adopted^[12,14,15]. Extrahepatic recurrence was the most common recurrence pattern with respect to tumor location, comparatively with other studies in which the focus was on the appearance of early recurrence^[12,15,16]. Consistently with previous studies^[14], the median time to recurrence was 12.36 mo and late recurrence was less common, representing only 12.5% of cases, with a rate of 1.7%.

Focusing on the variations in imaging and histopathological characteristics between the primary and the intrahepatic recurred HCC, among the 14 patients with intrahepatic recurrence, enhancement pattern changes (from hypervascular to hypovascular) were experienced by two patients and histopathological changes (from moderate to poor differentiation degree) were recorded in one patient, underlying that some degree of variability may exist between the primary and the recurred HCC. Even though the number of patients who had changes in imaging or pathological characteristics is small, it is to be noted that no correlation was found between enhancement and differentiation degree changes. In particular, the two patients who changed enhancement pattern did not show any variation in histopathology between the primary and the recurred HCC. On the other hand, in one patient a shift from moderately to poorly differentiated HCC was observed, while no change in the enhancement pattern was registered. This result agrees with the higher prevalence of hypervascular pattern among both grade 2 and 3 HCCs.

Some limitations of this study should be noted. First, it was designed retrospectively. Second, in the correlation analysis between imaging pattern and differentiation degree of the primary tumor, wash-out was not considered, while recently some studies have focused on hypervascular HCCs signal intensity during portal venous phase as an additional predictive factor for differentiation degree^[22]. Finally, this study lacked a genomic and immunophenotypical analysis of both the primary and the recurrent HCC.

In conclusion, this study confirms the recently explored correlation between HCC differentiation degree and dynamic-imaging enhancement pattern. Moreover, our preliminary results show that some degree of variability may exist between the primary and the recurred HCC imaging characteristics. If tumoral histopathological characteristics do not seem

to justify enhancement pattern variations, a possible explanation is perhaps to be found in changes which occur in liver parenchymal structure and vascularity following transplantation^[23]. However, more studies with larger patient groups are needed to better explore the presence of and the reasons for enhancement pattern changes between the primary and the recurred HCC.

COMMENTS

Background

A correlation between histopathological grade and vascularity of hepatocellular carcinoma (HCC) has been recently demonstrated. More frequently HCC is hypervascular, an enhancement pattern that usually corresponds to moderate or poor differentiation degree. Liver transplantation is the optimal treatment for HCC, however recurrence rate still ranges from 8% to 17%. HCC recurrence is usually similar to the primary tumor, but some variability in enhancement pattern and histopathological characteristics may be expected.

Research frontiers

The relation between HCC pathological grade and vascularity introduces the possible role of imaging enhancement pattern as a prognostic factor for tumor outcome after transplantation. Moreover, in the field of HCC recurrence after transplantation, the reasons for variability between the primary tumor and the recurrence characteristics are still to be understood.

Innovations and breakthroughs

The recently introduced correlation between HCC vascularity and differentiation degree has been discussed in other studies, however the authors aimed to better explore this relationship by applying it to the issue of HCC recurrence after transplantation.

Applications

The most relevant future application of the study is the possible use of imaging enhancement pattern as a prognostic factor for tumor recurrence after transplantation; moreover the study of recurrence variability may allow to better understand the factors involved in post-transplantation recurrence.

Terminology

HCC is the most common type of liver tumor, arising from hepatocytes and typically originating in patients with chronic liver diseases; liver transplantation is the optimal, potentially curative treatment for HCC; recurrence of hepatocellular carcinoma after transplantation reflects hematogenous spread, lymphatic spread and peritoneal seeding of neoplastic cells.

Peer review

A very interesting topic, with a somewhat novel approach that justifies the small number of patients included.

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Lesser celandine (pilewort) induced acute toxic liver injury: The first case report worldwide

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The causality assessment of several reports provided evidence for the existence of Greater Celandine hepatotoxicity. However, there hasn't been any case report published thus far, about lesser celandine induced liver injury. Here, we present a case of 36-year-old woman admitted to the hospital with acute hepatitis and jaundice on her sclera with no history of drug abuse or alcohol consumption. However, the patient had a recent history of lesser celandine extract consumption for hemorrhoids, for about 10 d, prior to the admission. Viral hepatitis, autoimmune hepatitis, and drug induced toxic hepatitis were ruled out by further imaging studies and laboratory analysis. Using the Council for International Organizations of Medical Sciences scale, the type of liver injury was assumed as hepatocellular and was scored as 7 which shows probable causality. Immediate discontinuation of lesser celandine extract resulted in rapid decrease of the elevated enzymes. Herbs have been reported to cause liver injury and therefore should be suspected in the case of acute hepatitis with an unknown etiology. This case is important to be the first to explain hepatotoxicity caused by lesser celandine. Physicians should consider lesser celandine as a causative agent for hepatotoxicity.

Key words: Celandine; Acute liver toxicity; Hepatitis; Pilewort; Herb

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Core tip: Herbs have been reported to cause liver injury and therefore should be suspected in the case of acute hepatitis with an unknown etiology. This case is the first to explain hepatotoxicity caused by lesser celandine. Physicians should consider lesser celandine as a causative agent for hepatotoxicity.

Abstract

Lesser celandine, also known as *Ranunculus ficaria*, is a herbaceous perennial plant that commonly utilizes piles and is taken either internally or used externally.

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INTRODUCTION

Numerous herbs can be hepatotoxic. This has been demonstrated by several case reports, case series and literature reviews^[1-4]. However, the potential hepatotoxicity of a variety of chemicals in any herb and case presentations lacking diagnostic exclusion made it difficult to have a clear clinical assessment^[5].

Greater Celandine (*Chelidonium majus L.*) has been used in traditional Chinese medicine as well as in the Western world for its numerous biological activities^[6,7]. Teschke *et al*^[8,9] reviewed several reports from Europe regarding Greater Celandine (*Chelidonium majus*) hepatotoxicity. The causality assessment of these reports provided evidence for the existence of Greater Celandine hepatotoxicity^[8,9].

Lesser celandine, also known as *Ranunculus ficaria*, is a herbaceous perennial plant. Lesser celandine is a herbal astringent that commonly utilizes piles and is taken either internally or used externally, and for this feature it is also known as pilewort^[10]. A review of literature revealed that there hasn't been any case report published thus far, about lesser celandine induced liver injury. Here, we present the first case of toxic hepatitis associated with lesser celandine consumption.

CASE REPORT

A 36-year-old woman was admitted to the hospital for acute hepatitis. Her past medical history and family history did not reveal any significant disease. She did not have any history of alcohol consumption or drug abuse. However, the patient had a recent history of lesser celandine extract consumption for hemorrhoids, for about 10 d, prior to the admission. A detailed anamnesis showed that the patient consumed lesser celandine as tea, one cup per day for 3 d. Physical examination revealed jaundice on her sclera. Laboratory abnormalities included alanine aminotransferase (ALT 1830 IU/L; normal range: 0 to 45 U/L), aspartate aminotransferase (AST 1520 IU/L; normal range: 0 to 45 U/L), alkaline phosphatase (ALP 225 IU/L; normal range: 30 to 120 U/L), and total bilirubin (3.4 mg/dL; normal range: 0.174 to 1.04 mg/dL). Anti-HBs IgG was positive, anti-HCV, HCV PCR, Anti-HAV IgM, and anti-HEV IgM were negative. There was no serologic evidence for recent infections with herpes simplex virus (HSV), Epstein-Barr virus (EBV), cytomegalovirus (CMV), or varicella zoster virus (VZV). Autoimmune marker analysis included anti-nuclear antibodies (ANA), anti-mitochondrial antibodies (AMA), anti-neutrophil

cytoplasmic antibodies (ANCA), smooth muscle antibody (SMA), liver kidney microsomal antibody type 1 (LKM-1) and all markers were found to be negative. Abdominal ultrasonography was normal. Liver biopsy wasn't performed, because the patient did not give an informed consent for the procedure. Lesser celandine extract was immediately discontinued. Using the Council for International Organizations of Medical Sciences scale (CIOMS), the type of liver injury of our case was assumed as hepatocellular (ALT > 5N and R ≥ 5) and was scored as "7: probable" (Table 1)^[11]. After discontinuation of lesser celandine, rapid recovery was observed in patient and liver enzyme levels returned to normal in 3 wk.

DISCUSSION

Hepatocellular, cholestatic, and mixed type (hepatocellular and cholestatic) injuries are the three types of herb induced liver injuries^[12]. In the decision of different injury types, ratio R (ALT/ALP activity at the time liver injury is suspected, activity is measured by multiples of the highest point of the normal values) is used. If ALT > 2N or R ≥ 5, hepatocellular injury is assumed; if ALP > 2N or R ≤ 2, injury is cholestatic; if ALT > 2N and ALP is increased, with R > 2 and R < 5, mixed injury is assumed^[11]. There are no diagnostic tests or specific criteria for herb induced hepatotoxicity. Careful history taking, insightful evaluation of laboratory findings and histopathology are essential for diagnosis. Liver biopsy might be helpful for the assessment of liver injury, however Teschke *et al*^[13] reported that it is not essential for the diagnosis. The best way to determine causal agent is re-challenging. But this, for obvious reasons, is not ethically acceptable.

Lesser celandine can easily be confused with greater celandine. A case report from Germany provides an evidence for that by presenting a case with toxic hepatitis caused by greater celandine, however mentioning the name of the herb as lesser celandine in the abstract^[14]. Herbs have been reported to cause liver injury and therefore should be suspected in the case of acute hepatitis with an unknown etiology. Our case suggests that physicians should consider lesser celandine as a causative agent for hepatotoxicity.

COMMENTS

Case characteristics

A 36-year-old woman was admitted to the hospital for fainting and a prior diagnosis of acute hepatitis from another hospital. She had a recent history of lesser celandine extract consumption for hemorrhoids, for about 10 d prior to the admission.

Clinical diagnosis

Physical examination revealed jaundice on her sclera.

Differential diagnosis

Viral hepatitis, autoimmune hepatitis and drug induced toxic hepatitis were

Table 1 Score of the presented patient according to Council for International Organizations of Medical Sciences Scale

Items for hepatocellular injury	Score	Result of the presented case ¹
1 Time to onset from the beginning of the drug/herb		
5-90 d (rechallenge: 1-15 d)	2	+
< 5 or > 90 d (rechallenge: > 15 d)	1	-
Alternative: Time to onset from cessation of the drug/herb		
≤ 15 d (except for slowly metabolized chemicals: > 15 d)	1	-
2 Course of ALT after cessation of the drug/herb		
Percentage difference between ALT peak and N		
Decrease ≥ 50% within 8 d	3	+
Decrease ≥ 50% within 30 d	2	-
No information or continued drug/herb use	0	-
Decrease ≥ 50% after the 30 th day	0	-
Decrease < 50% after the 30 th day or recurrent increase	-2	-
3 Risk factors		
Alcohol use (drinks/d: > 2 for women, > 3 for men)	1	-
Alcohol use (drinks/d: ≤ 2 for women, ≤ 3 for men)	0	+
Age ≥ 55 yr	1	-
Age < 55 yr	0	+
4 Concomitant drug(s) or herbs(s)		
None or no information	0	-
Concomitant drug or herb with incompatible time to onset	0	+
Concomitant drug or herb with compatible or suggestive time to onset	-1	-
Concomitant drug or herb known as hepatotoxin and with compatible or suggestive time to onset	-2	-
Concomitant drug or herb with evidence for its role in this case (positive rechallenge or validated test)	-3	-
5 Search for non drug/herb causes		"+" if negative
Group I (6 causes)		
Anti-HAV-IgM	+	-
HBsAg, anti-HBc-IgM, HBV-DNA	+	-
Anti-HCV, HCV-RNA	+	-
Hepatobiliary sonography/colour doppler sonography of liver vessels/endosonography/CT/MRC	+	-
Alcoholism (AST/ALT ≥ 2)	+	-
Acute recent hypotension history (particularly if underlying heart disease)	+	-
Group II (6 causes)		
Complications of underlying disease(s) such as sepsis, autoimmune hepatitis, chronic hepatitis B or C, primary biliary cirrhosis or sclerosing cholangitis, genetic liver diseases	+	-
Infection suggested by PCR and titer change for CMV (anti-CMV-IgM, anti-CMV-IgG)	+	-
EBV (anti-EBV-IgM, anti-EBV-IgG)	+	-
HEV (anti-HEV-IgM, anti-HEV-IgG)	+	-
HSV (anti-HSV-IgM, anti-HSV-IgG)	+	-
VZV (anti-VZV-IgM, anti-VZV-IgG)	+	-
Evaluation of group I and II		
All causes-groups I and II - reasonably ruled out	2	+
The 6 causes of group I ruled out	1	-
5 or 4 causes of group I ruled out	0	-
Less than 4 causes of group I ruled out	-2	-
Non drug or herb cause highly probable	-3	-
6 Previous information on hepatotoxicity of the drug/herb		
Reaction labelled in the product characteristics	2	-
Reaction published but unlabelled	1	-
Reaction unknown	0	+
7 Response to unintentional readministration		
Doubling of ALT with the drug/herb alone, provided ALT below 5N before reexposure	3	-
Doubling of ALT with the drug(s) and herb(s) already given at the time of first reaction	1	-
Increase of ALT but less than N in the same conditions as for the first administration	-2	-
Other situations	0	+
Total Score		7

¹The score of the patient for each 7 item for hepatocellular injury was indicated as "+" in the correspondent cell. HBV: Hepatitis B virus; HCV: Hepatitis C virus.

ruled out.

Laboratory diagnosis

Alanine aminotransferase (ALT 1830 IU/L; normal range: 0 to 45 U/L), aspartate aminotransferase (AST 1520 IU/L; normal range: 0 to 45 U/L), alkaline phosphatase (ALP 225 IU/L; normal range: 30 to 120 U/L), and total bilirubin (3.4 mg/dL; normal range: 0.174 to 1.04 mg/dL) were assessed. Anti-HBs IgG was

positive, anti-HCV, HCV PCR, Anti-HAV IgM, and anti-HEV IgM were negative. There was no serologic evidence for recent infections with herpes simplex virus, epstein-barr virus, cytomegalovirus, or varicella zoster virus. All autoimmune markers were negative.

Imaging diagnosis

Abdominal ultrasonography was normal.

Treatment

Lesser celandine extract was immediately discontinued.

Related reports

This is the first case of toxic hepatitis associated with lesser celandine consumption.

Term explanation

Greater Celandine (*Chelidonium majus* L.) is a perennial herb and is used in western phytotherapy and traditional Chinese medicine for its wide variety of biological activities. Lesser celandine, *Ranunculus ficaria* (syn. *Ficaria verna*, *F. ranunculoides* or *F. grandiflora*), also known as pilewort, is a herbaceous perennial plant.

Experiences and lessons

Herb induced liver injury is an important problem in clinical setting, because it can be an etiology of undiagnosed acute hepatitis. This case is important to be the first to explain hepatotoxicity caused by lesser celandine. Physicians should consider lesser celandine as a causative agent for hepatotoxicity.

Peer review

Lesser celandine (pilewort) induced acute toxic liver injury by Bulent Yilmez *et al* is a relatively good and interesting report.

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