Prevention of hepatitis B virus-related hepatocellular carcinoma

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How to cite this article: Lin CL, Kao JH. Prevention of hepatitis B virus-related hepatocellular carcinoma. Hepatoma Res 2021;7:9. http://dx.doi.org/10.20517/2394-5079.2020.125

Received: 15 Oct 2020  First Decision: 16 Nov 2020  Revised: 24 Nov 2020  Accepted: 30 Nov 2020  Published: 7 Jan 2021

Abstract

Hepatocellular carcinoma (HCC), especially hepatitis B virus (HBV)-related, remains a major cause of cancer-related mortality worldwide. Unless there is early detection with curative treatment, the 5-year survival rate of advanced HCC is less than 15%. The preventive strategies for HBV-related HCC are thus urgently needed to reduce the global burden of this disastrous cancer. Primary prevention involves the avoidance of viral infection through hepatitis B vaccination and interruption of viral transmission from patients with chronic HBV infection. Universal neonatal hepatitis B vaccination program has successfully reduced the prevalence of HBV carriage rate as well as HCC incidence in vaccinated cohorts. However, HBV elimination is still difficult to achieve. Regarding secondary prevention, long-term treatment with nucleos(t)ide analogues has been proven to reduce the risk of HBV-related HCC. Individual risk stratification and a periodic HCC surveillance in these patients could facilitate early HCC diagnosis. Finally, tertiary prevention can also be achieved by life-long treatment with NAs to reduce the risk of HCC recurrence after curative treatment of primary HCC. Challenges ahead include the fact that HBV is not yet curable by current antiviral agents. Combination therapy with direct anti-HBV agents and host-targeting immunomodulatory agents is under active development. In addition, HCC risk cannot be eliminated even in patients with HBsAg seroclearance or functional cure. Therefore, HCC surveillance is strongly recommended for every patient with chronic HBV infection.

Keywords: Anti-HBV therapy, hepatitis B vaccination, hepatitis B virus, hepatocellular carcinoma

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INTRODUCTION

Hepatocellular carcinoma (HCC) is one of the most prevalent solid tumors and leading causes of cancer-related death worldwide. In 2018, the estimated annual incidence and mortality increased to 841,000 cases and 782,000 deaths, respectively[1]. The highest HCC incidence rates are found in East and Southeast Asia and Northern Africa as well, but the incidence rates of HCC also increased in North America and Europe over the past few decades[2]. Although early detection by HCC surveillance provides the opportunity of curative treatment for early-stage HCC, the worldwide 5-year survival rate of HCC has been unsatisfactory over the past two decades[3]. In the last 10 years, emerging molecular target therapy and immunotherapy has constituted a major breakthrough for advanced HCC. However, overall survival has improved marginally[4]. The major reasons for the poor prognosis of HCC include the heterogeneous nature of HCC affecting treatment outcome and the majority of HCC occurring in countries where medical resources are limited[5]. Therefore, we urgently need effective measures to prevent the occurrence of HCC in the populations at risk. The current exploration of HCC risk factors provides us a good opportunity to develop preventive strategies. It is known that the etiologies of HCC include virological factors (such as hepatitis virus infection), chemical factors (such as aflatoxin B1 and alcohol) and genetic as well as metabolic factors (such as hereditary hemochromatosis, metabolic associated fatty liver disease (MAFLD) or non-alcoholic steatohepatitis (NASH))[6,7]. The contribution of different risk factors to HCC varies widely by country and region. Among them, chronic hepatitis B virus (HBV) infection is one of the most important risk factors. The global burden of HCC in 2015 revealed that the contribution of HBV to HCC mortality was geographically related. From highest (up to 40%-60%) in most areas of Asia and Sub-Saharan Africa to the less common cause of HCC death, approximately 9% in North America[8].

Since 1981, the development of hepatitis B vaccines has successfully prevented the spread of HBV across the world. In addition, the development of anti-HBV agents has led to a delay or block of the progression of chronic hepatitis B (CHB). As a consequence, prevention of HBV-related HCC in this setting should be expected by early 2050s. In this review, the mechanisms of HBV-induced HCC and individual prevention of HBV-related HCC will be summarized and discussed.

FACTORS ASSOCIATED WITH HBV-RELATED HCC

Thanks to the advances in the molecular biology of HBV, the pathogenesis of HBV-related HCC is largely clarified [Figure 1]. The pathogenesis of HCC in patients with CHB can be divided according to HBV-related direct and indirect mechanisms. Direct oncogenic effects of HBV include HBV-host genome integration and HBV-encoded oncogenic protein. The integration of HBV DNA into the host genome induces both host chromosomal instability and insertion mutagenesis of HCC-related genes[9]. This integration in the host genome most frequently happens in the telomerase reverse transcriptase region and myeloid/lymphoid or mixed-lineage leukemia 4 gene. Dysregulation of telomerase and histone methyltransferase expression subsequently increase the risk of hepatocarcinogenesis[10]. Truncated HBV pre-S/S mutation induces surface protein synthesis imbalance and may increase endoplasmic reticulum stress and decrease tumor suppressor gene expression, and thus, it may play a primary role in oncogenesis[11]. Among HBV-encoded proteins, HBV X protein has the most oncogenic effect in hepatocarcinogenesis. HBV X protein is a multifunctional regulator, and it is involved in several cancer-related molecular mechanisms, including alteration of signal pathways (e.g., inhibition of P53 and activation of the Jak1/STAT pathway), DNA repair (e.g., alteration of p53/ERCC3), apoptosis (inhibition of P53), mitochondrial function (e.g., activation of calcium-dependent kinase pathway, epigenetic modification (e.g., epigenetic silencing of ASPP gene), and expression of noncoding RNA (e.g., activation of the NF-κB pathway)[12,13]. Through these mechanisms, HBV X protein directly contributes to HCC development.

Hepatitis B viral load is the strongest risk factor associated with an increased HCC risk, independent of HBeAg status and cirrhosis. Chronic inflammation caused by long-term exposure to high levels of HBV
DNA will increase the risk of subsequent HCC. Clinically, several cross-sectional and longitudinal cohort studies in Asia revealed a positive correlation between serum HBV DNA level and HCC development\textsuperscript{[14-20]}. One of the most important longitudinal studies tracking HBV DNA and risk of HCC is the REVEAL-HBV study. REVEAL-HBV study providing convincing evidence of a dose-response relationship between HBV DNA levels and risk of HCC development in HBeAg-negative carriers. In the natural course, patients with persistently high HBV viremia had the greatest risk of HCC development\textsuperscript{[14]}. After a long-time evolution, at least 10 HBV genotypes (A to J) with distinct geographic and ethnic distributions have been identified\textsuperscript{[21]}. Briefly, genotypes B and C are highly prevalent in the Asia-Pacific region. Genotypes A and D are widespread in Africa, Europe and North America. The risk of HCC is significantly different between HBV genotypes. Several studies from Asian countries have revealed that patients with genotype C infection have a greater risk of developing HCC than those with genotype B infection\textsuperscript{[20-23]}. In a prospective study, Yu et al.\textsuperscript{[20]} found that genotype C patients had significantly higher serum levels of HBV DNA than did genotype B patients. Furthermore, genotype C with high viral load had a substantial increase in HCC risk than those with genotype B with low viral load. Thus, through chronic inflammation caused by high HBV viral load and its interaction with host immunity, an increased risk of HCC can be predicted in HBV genotype C patients. In the regions where genotypes A and D are prevalent, the comparison of HCC risk between patients with genotype A and D infection remains controversial. A meta-analysis including more than 14,500 patients revealed that the risk of HCC was similar between genotypes A and D\textsuperscript{[24]}. On the contrary, in a cohort study of Alaska native persons with HBV genotype A, B, C, D and F revealed that individuals with genotype A, C and F had significantly higher HCC risk than those with genotype B and D\textsuperscript{[25]}. These studies supported the notion that HBV genotypes show a strong correlation with HCC risk. Beside HBV genotype, mutations in specific regions are also related to HCC development. Core promoter mutations, especially basal core promoter mutations, are confirmed to be...
associated with an increased risk of HCC\textsuperscript{[21,23,26]}. Pre-S deletion mutations may block hepatocyte DNA repair and lead to genomic instability\textsuperscript{[27]}. Taken together, these naturally occurring mutations may contribute to hepatocarcinogenesis in patients with chronic HBV infection.

The indirect mechanism of HBV-related hepatocarcinogenesis is through the process of chronic necroinflammation, regeneration and fibrosis, associated with HBV or concomitant HBV and other causes of inflammation, such as MAFLD or nonalcoholic fatty liver disease (NAFLD). Although several studies have reported that the prevalence of NAFLD is lower in HBV carriers than general population\textsuperscript{[22]}, the concurrent NAFLD increases disease progression and the risk of HCC in CHB\textsuperscript{[26,28]}. In particular, NAFLD was also associated with HCC development in CHB patients with effective anti-HBV therapy\textsuperscript{[30]}. Regarding non-viral factors, a systematic review revealed that smoking as an environmental factor augmented HCC development with an odds ratio of 1.90 (95%CI: 1.68-2.14; \( P < 0.00001 \))\textsuperscript{[32]}. In a recent study from Taiwan, Wang \textit{et al.}\textsuperscript{[33]} further found that the plasma levels of interferon-g and natural killer cell levels in HBV carriers who smoked were significantly lower than in those who did not. Through the reduction of antiviral immunity, HBV carriers who smoked had persistently higher viral load and serum ALT levels compared with nonsmokers over the long-term follow-up period.

Finally, host predisposing genetic variations also contribute to HCC risk in HBV carriers. The application of genome-wide association study has identified several susceptibility loci or genetic variants for HBV-related HCC. Recent uncovered host genetic variants associated with HBV-related HCC included HLA class II region, \textit{KIF1B} gene, \textit{STAT4} gene and \textit{GRIK1} gene\textsuperscript{[34]}. Genomic studies also elucidated a number of somatic mutated genes affecting signal pathways associated with HBV-related HCC. For example, \textit{CTNNB1} mutation in the Wnt/b-catenin pathway was associated with hepatic cell proliferation and differentiation. Mutations of \textit{TP53} in the cell cycle pathway was associated with genomic stability\textsuperscript{[10]}. However, functional assessment is warranted to confirm the causal relationship with HBV-related HCC.

**PREVENTIVE STRATEGIES FOR HBV-RELATED HCC**

In the past few years, the incidence and mortality of HBV-related HCC has continuously increased\textsuperscript{[1]}. Thus, efforts in the development of prevention measures is urgently needed to decrease the global burden of HBV-related HCC. From the literature, the three levels of preventive strategies, including primary, secondary, and tertiary prevention will be effective for the prevention of HBV-related HCC [Figure 2].

**Primary prevention**

The goals of primary prevention of HBV-related HCC include the avoidance of HBV infection among susceptible populations and the interruption of mother-to-child transmission (MTCT) of HBV.

\textit{HBV vaccination}

Enhancement of host immunity by vaccination remains the most effective way for the prevention of HBV infection\textsuperscript{[35]}. Since the development of the first hepatitis B vaccine in 1981, universal infant HBV vaccination has been introduced in 187 countries\textsuperscript{[36]} with 75% of the global coverage rate among infants\textsuperscript{[37]}. After hepatitis B vaccine was implemented, the prevalence of chronic HBV infection was markedly reduced in most countries. For example, Taiwan is a hyperendemic area for HBV infection, where the prevalence of HBsAg was found to be more than 12% in the general population\textsuperscript{[38]}. After the implementation of the hepatitis B vaccination program in 1984\textsuperscript{[39]}, the prevalence of chronic HBV infection rapidly changed from 10.9% in children who were born before 1984 to 0.5% in children who were born after 1984\textsuperscript{[40]}. This dramatic change also appeared in countries where universal infant HBV vaccination was adopted. The global prevalence of HBV in children decreased from 4.7% before vaccine adoption to 1.3% in 2015\textsuperscript{[41]}. Surprisingly, the incidence of HCC in children also decreased after implementation of universal infant
HBV vaccination. In Taiwan, the incidence of HCC in children was 0.23 per 100,000 person-years in the vaccinated children, compared with 0.92 per 100,000 person-years in the unvaccinated cohorts\textsuperscript{[42]}. According to the population-based cancer registry in Taiwan, the annual incidence of HCC increased slightly, 1.3%, in adults during 2003 to 2011. In contrast, the annual incidence of HCC decreased 16.6% in children during the same period and decreased to zero in 2011\textsuperscript{[43]}. Similarly, HCC incidence significantly decreased 84% in China\textsuperscript{[44]}, 70% in Korea\textsuperscript{[45]} and down to zero in Alaska\textsuperscript{[46]} after implementation of a national hepatitis B vaccination program. Ample evidence from Taiwan and other countries revealed that the hepatitis B vaccine significantly reduces the incidence of HCC in children, teenagers and young adults. Therefore, hepatitis B vaccine has been recognized as the first anti-cancer vac the history of medical development\textsuperscript{[47]}.

**Antiviral prophylaxis in pregnant women**

Despite that effective HBV vaccination dramatically decreased the prevalence of HBV in children in the post-vaccine era\textsuperscript{[41]}, MTCT of HBV has not been completely blocked. In a prospective study which enrolled 303 mother-infant pairs, all of the mothers were HBV carriers, and the infants received a complete immunization program. There were 3.3% of children with chronic HBV infection. All of the children with chronic HBV infection were born to HBeAg-positive mothers with high viral load. In addition, the risk of
MTCT of HBV was positively correlated with maternal viral load. To completely block MTCT of HBV, anti-HBV therapy for HBsAg-positive mothers was suggested. Afterwards, two prospective studies from Taiwan and China enrolled HBeAg-positive pregnant women with high HBV DNA levels. Anti-HBV prophylaxis with tenofovir disoproxil fumarate (TDF) 300 mg daily was administrated in the third trimester of pregnancy. Compared to the mothers without anti-HBV prophylaxis, the infants born to mothers with TDF prophylaxis had significantly lower rates of HBV infection. Currently, antiviral prophylaxis in the third trimester of pregnancy in HBsAg-positive mothers with high viral load is recommended in international liver societies.

**Interruption of horizontal transmission**

Although universal HBV vaccination is the key to primary HBV prevention, it is equally important to prevent horizontal HBV transmission in adults. HBV vaccination for individuals at risk of contracting HBV infection, including healthcare workers, people who inject drugs, men who have sex with men, HIV carriers and sex workers, is effective in interrupting horizontal transmission. A new hepatitis B vaccine with simple schedule, only two doses over one month, can increase patient compliance and coverage in adults. In addition, appropriate information to promote safer sex and intention of receiving HBV vaccination also aids in the effort to prevent horizontal HBV transmission.

**Secondary prevention**

Secondary prevention for HBV-related HCC involves anti-viral treatment to delay or prevent liver disease progression and early development of HCC.

**Anti-HBV therapy**

Currently, standard or pegylated interferon-α (PEG-IFN-α) and NAs are widely approved for the anti-HBV treatment of CHB. According to inhibitory efficacy and resistance incidence, the first-line treatments include PEG-IFN-α, entecavir (ETV), tenofovir disoproxil fumarate (TDF) and tenofovir alafenamide. Several earlier randomized control studies and meta-analyses reported that interferon-α and PEG-IFN-α therapy reduced the risk of CHB progression and HCC incidence by 30%-40%. Regarding NAs, the prevention effect in HBV-related HCC in cirrhosis patients is more obvious. The cohort studies from Taiwan, Japan and Hong Kong demonstrated that CHB patients with ETV therapy had significantly lower cumulative HCC incidence, with HCC risk reduction by 45%-60%, especially in a cirrhosis setting. A recent study enrolled 1,088 patients with and without TDF treatment in Hong Kong. Compared to untreated patients, TDF treatment significantly reduced risks of HCC [adjusted hazard ratio (aHR) 0.46, P < 0.01]. Similarly, Nguyen et al. conducted a retrospective study in which 6,914 CHB patients with and without TDF treatment were recruited. During 8 years of follow-up, TDF treatment had a 77% reduction in the risk of HCC (aHR, 0.23; 95%CI: 0.56-0.92) in cirrhotic patients and a 73% reduction (aHR = 0.27; 95%CI: 0.07-0.98) in non-cirrhotic patients. According to the national healthcare database in Taiwan, Hsu et al. found that the HCC annual incidence decreased with an adjusted incidence rate ratio of 0.73 (95%CI: 0.66-0.80) per year in patients receiving ETV or TDF treatment.

Recently, several studies have compared the preventive effects for HBV-related HCC of first-line NAs. A Korean nationwide cohort study revealed that patients with TDF treatment had a significantly lower risk of HCC (HR = 0.61; 95%CI: 0.54-0.70) than patients with ETV treatment. In a meta-analysis that included seven articles with 3,698 patients, the incidence of HCC was significantly lower among patients with TDF treatment than those with ETV treatment (rate ratio: 0.66; 95%CI: 0.49-0.89; P = 0.008). However, three hospital-based studies in Taiwan and Korea demonstrated that the annual HCC incidence was not different between the patients receiving ETV and TDF treatment. The recent international study including 19 centers from 6 countries or regions also revealed that TDF and ETV had a similar effect in the prevention of HCC in patients with CHB. In the latest meta-analysis investigating 31 studies that enrolled 119,053
patients, the incidence of HCC was similar in patients with TDF or ETV treatment. Taking these lines of evidence together and unless we could have a large-scale head-to-head comparison study, the preventive effects for HBV-related HCC of first-line NAs remain unsettled.

The adherence to NA treatment also influences the protective effect of HCC prevention. In a long-term follow-up study of CHB patients receiving naïve ETV treatment, poor adherence significantly increased the risk of HCC development with a HR of 2.86 (95% CI: 1.76-4.64; \(P < 0.001\)) particularly among patients with cirrhosis.  

**Non-anti-HBV therapy**

Although hepatitis B viral suppression with long-term treatment with NAs reduced the incidences of HCC, HCC risk persists in HBV carriers, especially those with advanced fibrosis. Thus, effective non-anti-HBV therapy to prevent HCC is urgently needed.

The chemoprotective effects of aspirin against cancers have been addressed for decades, including gastrointestinal tract-related cancer and lung cancer. Recently, the Taiwan nationwide cohort study compared 2,123 patients with daily aspirin for 90 or more days and 8,492 patients without aspirin therapy, the investigators found that aspirin therapy significantly decreased HCC risk (HR: 0.71; 95%CI: 0.58-0.86; \(P < 0.001\)). In the Swedish nationwide cohort including chronic hepatitis B and C patients, the use of aspirin decreased the cumulative incidence of HCC (aHR: 0.69; 95%CI: 0.62-0.76). In addition, the protective effect was associated with duration of aspirin treatment.

Statins, 3-hydroxy-3-methylglutaryl CoA reductase inhibitors, have potential anticancer effects, such as anti-inflammation, antiproliferation, and anti-invasion, which are associated with carcinogenesis. In an earlier hospital-based study from Hong Kong, statin use reduced HCC risk by 32% in patients with chronic HBV infection. Recently, in the Swedish nationwide cohort including chronic hepatitis B and C patients, the use of lipophilic statins decreased the cumulative incidence of HCC (aHR: 0.41; 95%CI: 0.32-0.61). A recent meta-analysis, including 7 studies with HBV carriers revealed that statin therapy reduced the risk of HBV-related HCC with a relative risk of 0.52 (95%CI: 0.40-0.68; \(P < 0.001\)). Although the chemopreventive effects of these non-anti-HBV drugs are encouraging, further prospective studies or randomized trials are warranted to confirm these observations.

**Early detection of HCC**

Early detection of HCC in HBV patients is another important strategy for secondary prevention of HCC. To identify those at high risk of HCC development, risk stratification is needed in clinical practice. Currently, several predicting models have been developed to evaluate the risk of HBV-related HCC development in CHB patients, regardless of without or with anti-HBV therapy. The components of the models included clinical features (age, sex, platelet count, serum ALT and cirrhosis) and hepatitis B viral factors (HBV DNA, HBeAg, genotype and levels of HBsAg). For example, there was the REACH-B model with a 17-point risk score, which included age, male sex, ALT, HBeAg-positive and HBV DNA as variables. The estimated accuracy by area under receiver operating characteristic (AUROC) curve to predict risk of HCC was 0.796 (95%CI: 0.775-0.816) at 5 years and 0.769 (95%CI: 0.747-0.790) at 10 years. The REAL-B model included 8048 CHB patients who received oral anti-HBV treatment for prediction of the 5-year HCC risk. This model was developed on the basis of the simplified variables, including male sex, age, alcohol use, diabetes, baseline cirrhosis, platelet count, and alpha-fetoprotein (AFP). The estimated 5-year cumulative HCC incidence rates were 0.3%-1.38% for low risk (REAL-B score \(\leq 3\)), 2.29%-10.24% for moderate risk (REAL-B score 4-7), and 16.5%-90.37% for high risk (REAL-B score 8-13). The AUROC curves for HCC risk at 3, 5, and 10 years were all greater than 0.8. Based on the risk
### Table 1. Summary of current predicting models of hepatocellular carcinoma development in treatment-naïve chronic hepatitis B

| Developed area | Study design | Number of patients (derivation/validation) | Percentage of cirrhosis (derivation/validation) | Components of model | Accuracy for prediction of hepatocellular carcinoma (5-year AUROC) |
|----------------|-------------|-------------------------------------------|-----------------------------------------------|--------------------|---------------------------------------------------------------|
| GAG-HCC (87)   | Retrospective cohort | 820/0 | 15%/0 | Age, Male, BCP mutation, Cirrhosis, HBV DNA | 0.88              |
| CU-HCC (88)    | Retrospective cohort | 1005/424 | 0/18% | Age, Male, BCP mutation, Cirrhosis, HBV DNA | 0.76              |
| REACH-B (89)   | Retrospective cohort | 3584/1505 | 0/0 | Age, Male, ALT, HBeAg-positive, HBV DNA | 0.796             |
| REACH-B II (90) | Retrospective cohort | 2221/1113 | 0/0 | Age, Male, ALT, HBeAg-positive, HBV DNA | 0.89              |
| REACH-B IIa (91) | Retrospective cohort | 3584/314 | 32%/31% | Age, Male, ALT, HBeAg-positive, HBSag | 0.811-0.823       |
| REACH-B IIb (92) | Retrospective cohort | 3584/3114 | 13.7% | Age, Male, Cirrhosis, Alpha-fetoprotein, HBV DNA | 0.83              |
| LSM-HCC (93)   | Prospective-retrospective cohort | 1035/520 | 32%/31% | Age, Male, ALT, HBeAg-positive, HBSag | 0.88              |
| RWS-HCC (94)   | Prospective-retrospective cohort | 583/3353 | 13.7% | Age, Male, Cirrhosis, Alpha-fetoprotein, HBV DNA | 0.876             |
| DAS (95)       | Prospective-retrospective cohort | 971/507 | 0/0 | Age, Male, HBeAg-positive, HBV DNA | 0.76              |
| AGED (96)      | Prospective-retrospective cohort | 628/1663 | 0/0 | Age, Male, HBeAg-positive, HBV DNA | 0.76              |

ALT: alanine aminotransferase; AUROC: area under receiver operating characteristic; BCP: basal core promoter; LS: liver stiffness

### Table 2. Summary of current predicting models of hepatocellular carcinoma development in chronic hepatitis B with antiviral treatment

| Developed area | Study design | Number of patients (derivation/validation) | Percentage of cirrhosis (derivation/validation) | Components of model | Accuracy for prediction of hepatocellular carcinoma (5-year AUROC) |
|----------------|-------------|-------------------------------------------|-----------------------------------------------|--------------------|---------------------------------------------------------------|
| PAGE-B (97)    | Retrospective cohort | 1325/490 | 20%/48% | Age, Male, Platelet | 0.82              |
| mPAGE-B (98)   | Retrospective cohort | 2001/100 | 19.1%/20.1% | Age, Male, Platelet | 0.82              |
| HCC-RESCUE (99) | Retrospective cohort | 990/1071 | 39%/35% | Age, Male, Platelet, Albumin | 0.77              |
| APA-B (100)    | Retrospective cohort | 883/442 | 36%/36% | Age, Male, Platelet, Albumin, Cirrhosis | 0.827             |
| CAMO (101)     | Retrospective cohort | 23851/19321 | 27%/7% | Age, Male, Platelet, Albumin, Cirrhosis, Alpha-fetoprotein | 0.76              |
| AASL-HCC (102) | Retrospective cohort | 944/298 | 39%/39% | Age, Male, Platelet, Albumin, Cirrhosis | 0.802             |
| REAL-B (103)   | Retrospective cohort | 5365/2683 | 20.2%/22.1% | Age, Male, Platelet, Albumin, Cirrhosis, Alpha-fetoprotein | 0.80              |
| aMAP (104)     | Retrospective cohort | 3688/9400 | 19.3%/17.6% | Age, Male, Platelet, Albumin, Cirrhosis, Alpha-fetoprotein | 0.82              |

Harrell's C-index was used to access the performance of model discrimination. AUROC: area under receiver operating characteristic.
stratification through these clinical HBV-related HCC risk predicting models, HCC surveillance should be provided to HBV carriers with different risk levels. Among patients with a low HCC risk, a follow-up with regular interval is suggested. For patients with intermediate risk, a close follow-up is needed. For high-risk patients, in addition to close follow-up, anti-HBV treatment should also be considered.

Currently, periodic screening by using ultrasonography with or without AFP among populations at risk of HCC is the most effective strategy for early detection of HCC in several clinical practice guidelines. AFP is a long time and widely used biomarker for the diagnosis of HCC. However, the detection rates are only 30%-40% for the early-stage HCC. Protein induced by vitamin K absence II (PIVKA II) also known as Des-γ-carboxy-prothrombin was identified as a serum biomarker for HCC. In an early case-control study, PIVKA II had high diagnostic efficacy with a sensitivity of 61% and specificity of 70% for early stage HCC. In a meta-analysis with 14 studies, the diagnostic ability of PIVKA II combined with AFP was better than that of each biomarker used alone. The sensitivity and specificity were 91% and 83%, respectively. The pooled AUROC curve was 0.94 (95%CI: 0.91-0.95). In the future, the combination of different imaging modalities and non-invasive biomarkers are likely to improve the ability of early HCC detection.

Tertiary prevention

Anti-HBV therapy

Although the advance in the treatment of HCC provides opportunities for curative treatment, a high incidence of recurrence, up to 50% at 5 years after curative treatment, was observed in patients with underlying chronic HBV infection. Therefore, it is urgent to identify risk factors associated with HCC recurrence and develop tertiary prevention for HBV-related HCC after curative treatment. Ample evidence indicates that high viral load before curative treatment is associated with HCC recurrence post curative treatment. High viral load is known to result in severe hepatic inflammatory and fibrosis scores, which are related to HCC recurrence after curative treatment. Furthermore, curative treatment such as radiofrequency and surgical resection might be associated with HBV reactivation after these treatments. In a retrospective study, Li et al. analyzed a cohort of 2362 HBV-related HCC patients with surgical resection and demonstrated that high preoperative viral load was associated with high incidence of microvascular invasion. Of particular note, preoperative NA treatment significantly reduced the risk of microvascular invasion and recurrence of HCC. Therefore, all CHB patients with HCC who are planning surgical treatment should receive anti-HBV therapy to prevent ongoing hepatic inflammation and HCC recurrence.

Although standard or pegylated IFN-α are effective in the treatment of CHB, the effect of adjuvant IFN-α therapies on the prevention of HBV-related HCC recurrence after curative treatment remains unsatisfactory. In an earlier randomized controlled trial in Taiwan, the median recurrent free survival rates were similar in patients with and without adjuvant IFN-α therapy after curative treatment of primary HCC. Furthermore, two meta-analyses both revealed that adjuvant IFN-α therapy cannot decrease HCC recurrence after curative treatment. On the contrary, NAs are more effective than IFN-α in preventing HCC recurrence. In a randomized controlled study, NAs treatment significantly decreased HCC recurrence after surgical resection (HR = 0.48; 95%CI: 0.32-0.70). Subsequently, another randomized controlled trial enrolled 200 HCC patients with or without NA treatment after surgical resection. The patients with NA treatment had better recurrence-free survival and overall survival. NA therapy significantly decreased the incidence of late recurrence (HR = 0.34; 95%CI: 0.17-0.68; P = 0.002). Several large cohort studies in Asian countries also revealed that adjuvant NA treatment significantly reduced the risk of HCC recurrence after curative treatment of primary HCC. Recently, a meta-analysis including 21 studies confirmed that HCC patients receiving NA therapy after curative treatment had a lower cumulative incidence of HCC recurrence with relative risk of 0.76 (95%CI: 0.65-0.90; P = 0.001). Nowadays, NAs are recommended for the tertiary prevention of HBV-related HCC.
Non-anti-HBV therapy

Similar to secondary prevention, non-antiviral therapies also have a potential effect on tertiary prevention of HBV-related HCC recurrence after curative therapy. A hospital-based study, which included 430 patients with HBV-related HCC after curative therapy, revealed that aspirin reduced the risk for HCC recurrence with an aHR of 0.22-0.24 \cite{118}. A study from Korea revealed that liver transplant recipients with statin therapy had a significantly lower rate of HCC recurrence than those without statin therapy \cite{119}. Another Japanese study with 734 HCC patients after surgical resection also demonstrated that patients with statin treatment had significantly higher recurrent-free survival than those without statin treatment \cite{120}. The protective effect of aspirin or statin on the risk of HCC recurrence should be further validated through randomized clinical trials.

THE CHALLENGES IN FIGHTING HBV-RELATED HCC

HBV is not yet curable

Although IFN-\(\alpha\) and NAs are effective in the control of disease activity in CHB, covalently closed circular DNA (cccDNA) are persistently present in HBV patients and even in those who experience HBsAg seroclearance or functional cure \cite{121}. The challenges ahead include eliminating or at least silencing cccDNA, inhibition of virus replication and blockage of HBV intrahepatic dissemination \cite{122}. Several novel therapeutic approaches for HBV cure have been actively investigated. The direct anti-HBV agents (DAA) include entry inhibitor targeting sodium taurocholate cotransporting polypeptide, engineered site-specific nucleases targeting cccDNA and RNA interference targeting HBV RNA \cite{123}. The immunomodulatory host targeting agents (HTA) include innate immunity receptors agonist and adaptive immunity stimulation by therapeutic vaccine \cite{123-125}. In the foreseeable future, the most promising strategies towards achieving an HBV cure will be based on a combination of DAA and HTA \cite{126,127}.

HCC risk persists even with functional cure

It is not clear whether the risk of HCC can be completely eliminated after HBsAg seroclearance or seroconversion. In a long-term follow-up study, the estimated annual incidence of HCC was 0.31% and 0.16% in patients after spontaneous and NA-induced HBsAg seroclearance, respectively \cite{128}. Similarly, a retrospective study from Korea revealed that the estimated annual incidence of HCC was 0.29% in non-cirrhotic patients achieving HBsAg seroclearance. Notably, the estimated annual incidence of HCC was up to 2.85% in cirrhotic patients after HBsAg seroclearance \cite{129}. These observations were also confirmed by a systemic review showing that the incidence of HCC was quite low but present in patients with HBsAg seroclearance \cite{130}. In addition, the increased prevalence of comorbidities such as MAFLD or NAFLD, NASH and metabolic syndrome in HBV patients may be associated with HCC risk, regardless of the status of hepatitis B. In a large cohort study of Taiwanese HBV carriers, the presence of metabolic risk factors was associated with an increased risk of HCC in patients with low serum level of HBV DNA (< 2000 IU/mL) \cite{131}. Although effective anti-HBV therapy has led to a small proportion of patients with seroclearance or seroconversion of HBsAg, they are still at a small risk of developing HCC. Therefore, a periodic surveillance for HCC is still recommended even in patients after HBsAg seroclearance or seroconversion. This is particularly important for cirrhotic patients.

CONCLUSION

Since the discovery of HBsAg by Dr. Baruch Blumberg in the early 1960s, tremendous efforts have been made to reduce the complications, morbidity and mortality of CHB. After three decades of global promotion, universal hepatitis B vaccination has effectively interrupted MTCT of HBV. Elimination of HBV in neonates and infants is achievable \cite{35}. Despite the successful prevention of HBV spreading by hepatitis B vaccination, it is expected that HBV-induced cirrhosis and HCC will increase in people who have been chronically infected with HBV. Early anti-HBV treatment to prevent liver disease progression is the preferred strategy.
Ample evidence has shown that anti-HBV therapy can reduce the risk of HCC development in patients with chronic HBV infection. However, anti-HBV therapy does not completely eliminate the risk of HCC development. Therefore, HCC surveillance is recommended in every HBV patient. Through long-term profound viral suppression by anti-HBV therapy and early detection of HCC by optimal surveillance, secondary prevention of HBV-related HCC can be reached successfully. Finally, the benefit of adjuvant anti-HBV therapy in the reduction of HCC recurrence after curative treatment makes tertiary prevention of HBV-related HCC possible. The combination of individual risk-stratification approach and three levels of preventive strategies for HBV-related HCC will substantially decrease the global burden of HBV-related HCC and improve the long-term prognosis of people living with HBV as well.

DECLARATIONS

Authors’ contributions
Helped in acquisition of data, planned and conducted this project, and wrote this manuscript: Lin CL
Helped in acquisition of data, planned this project, supervised this study, obtained funding, and wrote this paper: Kao JH

Availability of data and materials
Not applicable.

Financial support and sponsorship
This work was supported by grants from the National Taiwan University Hospital, the Ministry of Health and Welfare, and the Ministry of Science and Technology, Executive Yuan, Taiwan.

Conflicts of interest
All authors declared that there are no conflicts of interest.

Ethical approval and consent to participate
Not applicable.

Consent for publication
Not applicable.

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