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ORIGINAL ARTICLE

#### **Retrospective Study**

# Intrahepatic distribution of hepatitis B virus antigens in patients with and without hepatocellular carcinoma

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# Abstract

**AIM:** To study the intrahepatic expression of hepatitis B surface antigen (HBsAg) and hepatitis B core antigen (HBcAg) in chronic hepatitis B patients with and without hepatocellular carcinoma.

**METHODS:** A total of 33 chronic hepatitis B patients (mean age of 40.3  $\pm$  2.5 years), comprising of 14 HBeAg positive and 19 HBeAg negative patients; and 13 patients with hepatitis B virus related hepatocellular carcinoma (mean age of 49.6  $\pm$  4.7 years), were included in our study. Immunohistochemical staining for HBcAg and HBsAg was done using standard streptavidin-biotin-immunoperoxidase technique on paraffin-embedded liver biopsies. The HBcAg



and HBsAg staining distributions and patterns were described according to a modified classification system.

**RESULTS:** Compared to the HBeAg negative patients, the HBeAg positive patients were younger, had higher mean HBV DNA and alanine transaminases levels. All the HBeAg positive patients had intrahepatic HBcAg staining; predominantly with "diffuse" distribution (79%) and "mixed cytoplasmic/nuclear" pattern (79%). In comparison, only 5% of the HBeAq-negative patients had intrahepatic HBcAg staining. However, the intrahepatic HBsAg staining has wider distribution among the HBeAg negative patients, namely; majority of the HBeAg negative cases had "patchy" HBsAg distribution compared to "rare" distribution among the HBeAg positive cases. All but one patient with HCC were HBeAg negative with either undetectable HBV DNA or very low level of viremia. Intrahepatic HBcAg and HBsAg were seen in 13 (100%) and 10 (77%) of the HCC patients respectively. Interestingly, among the 9 HCC patients on anti-viral therapy with suppressed HBV DNA, HBcAg and HBsAg were detected in tumor tissues but not the adjacent liver in 4 (44%) and 1 (11%) patient respectively.

**CONCLUSION:** Isolated intrahepatic HBcAg and HBsAg can be present in tumors of patients with suppressed HBV DNA on antiviral therapy; that may predispose them to cancer development.

**Key words:** Hepatitis B virus; Chronic hepatitis B; Hepatocellular carcinoma; Hepatitis B core antigen; Hepatitis B surface antigen

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**Core tip:** This study described the distributions and patterns of intrahepatic hepatitis B core antigen and hepatitis B surface antigen (HBsAg) in patients with chronic hepatitis B using a novel, modified classification system. The HBeAg negative patients were found to have intense HBsAg in liver tissues despite their lower serum hepatitis B virus (HBV) DNA levels. For those with hepatocellular carcinoma (HCC), we observed high rate of HBV antigen detection in tumor tissues, but not in adjacent non-tumor livers, especially among those with optimal serum HBV DNA suppression. These data support that HCC can be derived from clonal expansion of infected hepatocytes with high carcinogenic potentials and selective resistance to antiviral agents.

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# INTRODUCTION

Hepatitis B virus (HBV) infection has a worldwide distribution affecting more than 240 million individuals<sup>[1,2]</sup>. It accounts near to 1 million deaths annually through complications such as liver failure and hepatocellular carcinoma (HCC)<sup>[3,4]</sup>. Clinical outcomes of HBV depend on the interaction between viral and host-specific factors.

Hepatitis B viral proteins include a structural nucleocapsid core protein (HBcAg), an envelope protein (HBsAg), and a soluble nucleocapsid protein (HBeAg)<sup>[5]</sup>. Serum HBeAg is a surrogate marker for active hepatitis B virus replication in hepatocytes<sup>[6]</sup>. HBcAg is an intracellular antigen that is detectable either in the nucleus or cytoplasm of HBV-infected hepatocytes<sup>[7]</sup>. There are reports that high nuclear expression of HBcAg reflects high level of circulating HBV DNA and normal alanine transaminases (ALT) level as seen in the "immune tolerance phase" of hepatitis B<sup>[8-10]</sup>. In contrast, the cytoplasmic expression reflects a lower level of circulating HBV DNA, but increased hepatocellular damage with higher ALT level during the "immune clearance phase"<sup>[11-13]</sup>. A study also suggested that the membranous expression of HBsAg relates closely to active viral replication<sup>[13]</sup>. However, immunohistochemical expression of HBcAg, HBsAg and its relationship to virological and histological activities has been inconsistent and limited, especially in HBeAg negative chronic hepatitis B and HBV related HCC<sup>[16,17]</sup>. The lack of a standardized system to describe the intrahepatic HBV antigen distribution may contribute to the inconsistent findings<sup>[18]</sup>.

In this study, we developed a modified classification system that allows consistency in the description of both the distributions and patterns of intrahepatic HBcAg and HBsAg. We applied the classification system to examine the HBcAg and HBsAg distributions in tissues from patients with HBeAg positive (+ve) and HBeAg negative (-ve) chronic hepatitis B (CHB). Since patients with optimal HBV DNA suppression on antiviral therapy are still at risk for HCC, we also examined the patterns of HBcAg and HBsAg in tumor and adjacent non-tumor tissues among HCC patients with and without antiviral treatment.

# MATERIALS AND METHODS

#### Patients

Liver biopsies from a total of 33 patients with chronic hepatitis B (CHB) and 13 patients with HBV-related hepatocellular carcinoma were included in the study. Fifty chronic hepatitis B cases were initially identified from our recent pathology database. Patients with HCV, HDV and HIV co-infection and inadequate biopsy size (< 2 cm) were subsequently excluded. Of the 33 HBsAg positive patients, 14 were HBeAg positive and 19 were HBeAg negative. None of the

	HBcAg	HBsAg
Distribution		
Diffuse	Contiguous without skip zones	Contiguous without skip zones
	Either in clusters or single cells	Either in clusters or single cells
Patchy	< 50% area with skip zones	< 50% area with skip zones
	Either in clusters or single cells	Either in clusters or single cells
Rare	Occasional area of entire field	Occasional area of entire field
	Either in clusters	Either in clusters
	or single cells	or single cells
Pattern	-Nuclear	-Membranous
	-Cytoplasmic	-Cytoplasmic

#### Table 1 Classification system for intrahepatic hepatitis B core antigen and hepatitis B surface antigen

HBcAg: Hepatitis B core antigen; HBsAg: Hepatitis B surface antigen.

CHB patients received antiviral or immunosuppressive therapy prior to liver biopsy. The 13 HCC patients were selected from our tumor clinic database. Patients with HCV, HDV, and HIV coinfection were excluded. Nine of the 13 (69%) HCC patients were on antiviral therapy prior to the development of liver cancer. Our liver pathologists reviewed histological scoring and invasiveness of HCC on all these 46 cases consistently. All patients' demographic and clinical characteristics were obtained from our electronic medical records at Beth Israel Deaconess Medical Center (BIDMC), Boston, Massachusetts, United States.

#### Liver pathology and immunohistochemical assays

The histological diagnosis was established using hematoxylin and eosin (HE) staining and masson trichrome stains of formalin fixed paraffin-embedded liver tissue. Hepatic fibrosis was graded by our liver histopathologists according to the Metavir system on a 5-point scale between 0 and 4.

HBeAg, anti-HBe, anti-HCV were tested by enzyme immunoassay and HBV DNA quantification was performed using the Cobas Ampliprep/Cobas Taqman HBV at the Molecular lab. Analysis of sections of paraffinembedded liver biopsies was performed using a previously reported procedure<sup>[7]</sup>.

Using standard streptavidin-biotin-immunoperoxidase technique, immunohistochemical staining for HBcAg and HBsAg was done<sup>[19]</sup> in our laboratory at BIDMC. Briefly, it involves 4 major steps: deparaffinization of tissue, antigen retrieval, tissue permeabilization and immunostaining. The distribution and patterns of intrahepatic HBcAg and HBsAg was uniformly recorded. We developed a clinically relevant, modified classification system to analyze the HBsAg and HBeAg patterns as shown in Table 1 and Figure 1.

# RESULTS

# Characteristics of HBeAg-positive and HBeAg-negative chronic hepatitis B patients

HBeAg positive patients were younger than the HBeAg

negative subjects. The gender distributions were not different between the groups with Asian males accounting for the majority of patients in both groups. Mean HBV DNA and ALT levels were higher in the HBeAg positive than the HBeAg negative patients. However, the histological characteristics were not different between the two groups. Over 90% of the patients in each group had mild to moderate grade 1-2 inflammation. Advanced Stage 3-4 hepatic fibrosis was present in 29% and 10% of the HBeAg positive and negative patients respectively (Table 2).

#### Comparison of HBcAg/HBsAg immune-staining characteristics in HBeAg-positive and HBeAg-negative CHB

Intrahepatic staining of HBcAg was seen in all 14 HBeAg positive patients with predominantly 'diffuse' distribution (79%) and "mixed cytoplasmic/nuclear" pattern (79%). Among the 19 HBeAg-negative patients, only one (5%) had positive intrahepatic HBcAg. It was 'rare' with mixed cytoplasmic/nuclear pattern. Intrahepatic HBsAg was seen in all the patients irrespective of their HBeAg status and was predominantly cytoplasmic (> 70%) in both groups. HBeAg negative patients actually had stronger intrahepatic HBsAg distributions. Majority (85%) of the HBeAg negative cases had either diffuse or patchy HBsAg distribution compared to 50% among the HBeAg positive cases (Table 3).

# Clinical features of patients with hepatocellular carcinoma

All 13 patients (9 males and 4 females) were Asian patients with age range between 25 and 69 years. All but one patient were HBeAg negative. Eight of Twelve (67%) HBeAg negative HCC patients had HBcAg in their adjacent non-tumor tissues. This is in distinct contrast to the HBeAg negative CHB patients without cancer, where only 1/19 (5%) had positive HBcAg. However, HBsAg in the adjacent non-tumor tissues was less prevalent among patients with HCC (67%) compared to the HBeAg negative CHB patients without



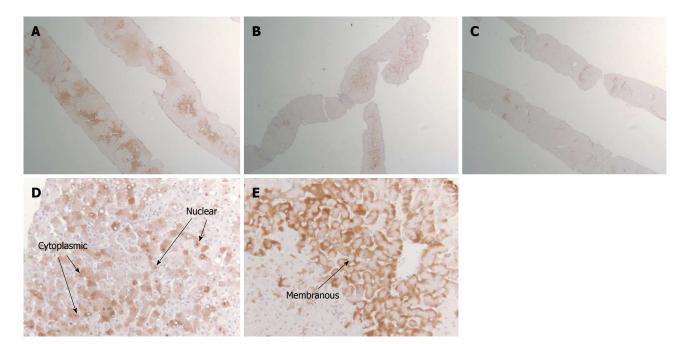


Figure 1 Images of pathology slides illustrating the classification system for intrahepatic hepatitis B surface antigen and hepatitis B core antigen. A: Diffuse distribution of HBsAg (× 2); B: Patchy distribution of HBcAg (× 2); C: Rare distribution of HBsAg (× 2); D: Cytoplasmic/nuclear pattern of HBcAg (× 20); E: Membranous pattern of HBsAg (× 20).

Table 2	Clinical,	virological,	and	histopatho	logical
characterist	ics of patie	ents with chroi	nic he	patitis B <i>n</i> (	%)

Variables	HBeAg-positive $(n = 14)$	HBeAg-negative $(n = 19)$
Mean age in years	38.5	42
Race		
Asian	13 (93)	10 (53)
White	1 (7)	1 (5)
African/African American	0 (0)	7 (37)
Hispanic	0 (0)	1 (5)
Male: female, n	11:3	13:6
HBV DNA (IU/mL), mean	28 million	7.3 million
ALT (U/L), mean	124	33
Inflammatory grades		
0	0 (0)	1 (5)
1-2	13 (93)	18 (95)
3-4	1 (7)	0 (0)
Fibrosis stage		
0	3 (21)	10 (53)
1-2	7 (50)	7 (37)
3-4	4 (29)	2 (10)

CHB: Chronic hepatitis B; ALT: Alanine aminotransferase.

#### cancer (100%) (Table 4).

Patients with HCC were further classified into two groups based on whether they were on antiviral therapy prior to the diagnosis of HCC. All four HCC patients who were not on anti-viral therapy had invasive carcinoma and replicative serum HBV DNA. In contrast, 8 of 9 (89%) patients on anti-viral therapy had undetectable serum HBV DNA, and only 3 developed invasive carcinoma.

Detection of HBV Antigens in HCC patients on anti-viral therapy: Despite on antiviral therapy, Table 3 Immunochemical characteristics of hepatitis B core antigen/hepatitis B surface antigen in hepatitis B envelop antigen-positive and hepatitis B envelop antigen-negative patients n (%)

Variables	HBeAg-positive $(n = 14)$		HBeAg-negative $(n = 19)$	
Distribution,	HBcAg	HBsAg	HBcAg	HBsAg
Diffuse	11 (79)	5 (36)	0 (0)	6 (32)
Patchy	1 (7)	2 (14)	0 (0)	10 (53)
Rare	2 (14)	7 (50)	1 (5)	3 (15)
Pattern				
Cytoplasmic (+)	1 (7)	-	0 (0)	-
Nuclear (+)	2 (14)	-	0 (0)	-
Mixed: Cyto/nuclear	11 (79)	-	1 (5)	-
Cytoplasmic (+)	-	11 (79)	-	14 (74)
Membranous (+)	-	1 (7)	-	0 (0)
Mixed: Cyto/memb	-	2 (14)	-	5 (26)

HBcAg: Hepatitis B core antigen; HBsAg: Hepatitis B surface antigen; HBeAg: Hepatitis B envelop antigen.

intrahepatic HBcAg was detected in all nine patients in the treatment group. The majority had diffuse distribution. Figure 2 illustrated diffuse HBcAg in both tumor and adjacent tissue of a patient. Interestingly, 4 (44%) patients with suppressed serum HBV DNA were found to have diffuse distribution of HBcAg only in the tumor but not in the adjacent liver (representative case shown in Figure 3). Intrahepatic HBsAg was detected in seven of nine (78%) patients. The majority had patchy distribution. Only one of the nine (11%) patients had HBsAg detected in the tumor tissue alone (Table 4).

**Detection of HBV Antigens in HCC patients not on anti-viral therapy:** Intrahepatic staining of HBcAg



Table 4 Features of hepatocellular carcinoma patients based

on the antiviral therapy status		
Variables	HCC group on anti-viral therapy (n = 9)	HCC group on no therapy (n = 4)
Mean age ± SD, yr	$53.2 \pm 8.5$	$46.5\pm20.5$
Male: female	7:2	2:2
HBeAg negative	8/9	4/4
Undetectable HBV DNA	8/9	0/4
Cirrhosis	4/9	0/4
Invasive HCC	3/9	4/4
Tumor:		
Well differentiation	6/9	1/4
Moderate differentiation	3/9	3/4
Intrahepatic HBcAg staining		
(+) Only in tumor tissues	4/9	1/4
(+) Only in non-tumor tissues	3/9	1/4
(+) Both in tumor and non-tumor	2/9	2/4
tissues		
Intrahepatic HBsAg staining		
(+) Only in tumor tissues	1/9	0/4
(+) Only in non-tumor tissues	3/9	2/4
(+) Both in tumor and non-tumor	3/9	1/4
tissues		

SD: Standard deviation; HCC: Hepatocellular carcinoma; HBsAg: Hepatitis B surface antigen; HBcAg: Hepatitis B core antigen; HBeAg: Hepatitis B envelop antigen.

was seen all four patients with only one having patchy HBcAg present in the tumor tissue alone. Intrahepatic HBsAg was detected in three of four patients with none presenting only in the tumor tissue alone. Intrahepatic staining of both HBcAg and HBsAg in tumor tissues was seen only in one patient (Table 4).

#### DISCUSSION

In this study, we sought to examine the intrahepatic HBcAg and HBsAg distribution and patterns using a standardized, modified classification system in patients with chronic hepatitis B and in those with HBV-related hepatocellular carcinoma.

HBeAg that is considered to be a marker of HBV replication and infectivity; is usually associated with high levels of serum HBV DNA<sup>[5,20]</sup>. In our study, we found HBeAg positive patients to be younger, had higher serum levels of HBV DNA and ALT when compared to the HBeAg negative patients. These observations are in concordance with previous reports<sup>[21-23]</sup>.

The presence of intrahepatic HBcAg in 100% of our HBeAg positive subjects with high level of viremia confirm the previous studies that the presence of HBcAg in hepatocytes serves as as a marker of viral replication<sup>[24,25]</sup>. The prevalence of intrahepatic HBcAg expression in HBeAg negative patients has been inconsistent and was reported between 0% and 90%<sup>[26-31]</sup>. In our current study, we detected HBcAg in only 5% of the HBeAg negative patients. Even though intrahepatic HBcAg expression was infrequent in

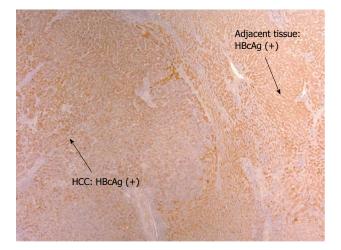


Figure 2 Diffuse hepatitis B core antigen distribution of mixed cytoplasmic, nuclear pattern in both tumor and adjacent non-tumor tissue (x 10). HCC: Hepatocellular carcinoma; HBcAg: Hepatitis B core antigen.

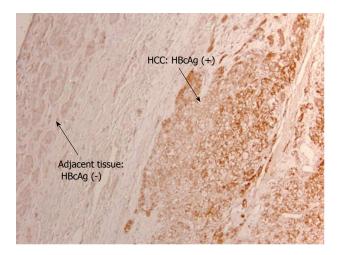


Figure 3 Diffuse hepatitis B core antigen distribution of mixed cytoplasmic, nuclear pattern in tumor tissue alone but not in adjacent non-tumor tissue (× 10). HCC: Hepatocellular carcinoma; HBcAg: Hepatitis B core antigen.

HBeAg negative CHB, its absence cannot exclude the presence of modest levels of HBV replication<sup>[18]</sup>.

The mixed cytoplasmic/nuclear pattern of HBcAg staining in the majority of our HBeAg-positive patients correlates positively with the diffuse HBcAg distribution and high HBV DNA levels. These results are consistent with previous reports<sup>[7,8,13]</sup>. Among HBeAg-negative patients, only one had positive but rare distribution intrahepatic HBcAg. These results are consistent with previous reports that higher level of viral replication (or serum HBV DNA) is associated with diffuse distribution and mixed cytoplasmic/nuclear pattern of HBcAg<sup>[7,8,11,1,13,18,24,30]</sup>.

In contrast to HBcAg, all patients had positive intrahepatic HBsAg regardless of their HBeAg status, and its distribution did not correlate with HBV DNA levels. In fact, HBeAg negative patients had more intense intrahepatic HBsAg distributions. These findings



are consistent with the hypothesis that the increased accumulation of intracellular HBsAg during inactive viral replication is secondary to the diminished synthesis of HBcAg and, therefore, reduced export of HBsAg<sup>[32,33]</sup>. Both HBeAg positive and negative patients had predominantly cytoplasmic HBsAg pattern regardless of their serum HBV DNA levels. In the literature, either cytoplasmic or membranous expression of HBsAg had been associated with active viral replication<sup>[7,28]</sup>. We could not comment on the isolated membranous pattern for the majority of our patients had either cytoplasmic or mixed cytoplasmic, membranous HBsAg pattern.

HBcAg expressions have been thought to induce liver necroinflammation through HBV specific and nonspecific T cells. Previous studies indicated that patients with predominantly nuclear expression of HBcAg had lower hepatocellular damage compared to those with predominantly cytoplasmic expression<sup>[21,22,34]</sup>. Studies from European and Asian population, however, reported no significant correlation between the patterns of HBcAg, HBsAg and hepatic injury<sup>[35-40]</sup>. Similarly, our study did not find a relationship between intrahepatic HBV antigen expression and histological activity in chronic hepatitis B. These inconsistent results may signify the complex interactions between the host and the hepatitis B virus.

Chronic hepatitis B is a major cause of hepatocellular carcinoma (HCC) globally. Its risk factors include advanced age, male sex, HBV genotype, longer duration of infection, higher viral load, elevated ALT, presence of cirrhosis and co-infections<sup>[41,42]</sup>. It is known that the cumulative risk of HCC persists after the clearance of HBV DNA, HBeAg and HBsAg in CHB<sup>[43-45]</sup>. In this study, HCC patients were older (mean age,  $49.6 \pm 4.7$  years) compared to those with chronic hepatitis B. The majority (92%) of our HCC patients were HBeAg negative, had undetectable HBV DNA on antiviral therapy and only 23% had underlying cirrhosis. These observations provided further evidence that HCC risk persists despite HBeAg seroclearance and optimal serum HBV DNA suppression. Detailed studies on the intrahepatic HBV antigens in HCC patients are limited. We detected intrahepatic HBcAg and HBsAg in 100% and 77% of our HCC patients respectively in spite of their undetectable serum HBV DNA and HBeAg negative status. These findings strongly suggest that HBV replication in HCC occurs more commonly than previously perceived by other studies<sup>[16,46-48]</sup>. Suzuki et al<sup>[48]</sup> reported 11.6/4.2% HBsAg/HBcAg in HCC and the highest reported was from Hsc et al and coworkers around 32/14.7% respectively.

To our knowledge, this is the first study to describe the distribution or patterns of HBV-associated antigens in HCC patients based on treatment status. Perhaps the most significant observation in our study is the high rate of HBV antigen detection in tumor tissue alone, but not in adjacent non-tumor liver, among those with optimal HBV DNA suppression. We identified five of 9 (56%) of HCC patients with either HBcAg or HBsAg present only in tumor tissues despite prolonged nucleoside analogue treatment. These data may imply that these tumors were derived from clonal expansion of infected hepatocytes with high rate of proliferation, carcinogenic potentials and selective resistance to antiviral agents<sup>[49,50]</sup>. Our findings also support the possible mechanisms of HBV-induced carcinogenesis secondary to HBV DNA integration into host genome leading to potential activation of oncogenes and induction of genetic instability. For instance, protein HBx is reported to be involved in activation of numerous signaling pathways and cellular promoters and to host DNA mutations<sup>[51,52]</sup>.

This study, while providing new observations on the potential mechanisms of HCC development, has a number of limitations. It is a retrospective study with especially patients with HCC referred from other health centers. Hence, details about other potential confounding factors of tumor development such as alcoholism and other environmental exposure were unavailable for the analyses. Lastly, the enrolled subjects are mostly Asians, and hence our findings should be confirmed with other populations.

In conclusion, the presence of intrahepatic HBcAg is associated with HBeAg positive status. Hepatocyte expression of diffuse, mixed (cytoplasmic/nuclear) expression of HBcAg correlates with serum HBV DNA levels, but not with the histological activity. Intrahepatic HBsAg was detected in all chronic hepatitis B patients regardless of their HBeAg status or levels of viremia. While intrahepatic HBcAg is rarely detectable in HBeAg negative chronic HBV patients, majority of the HBeAg negative patients with hepatocellular carcinoma had positive HBcAg even with optimal viral suppression. The presence of either HBcAg or HBsAg within tumor tissues of HCC patients on treatment may indicate that these tumors were derived from clonal expansion of infected hepatocytes with high carcinogenic potentials and drug resistance.

#### COMMENTS

#### Background

Hepatitis B virus (HBV) infection is a globally prevalent liver disease that can progress to cirrhosis and hepatocellular carcinoma (HCC). The outcome of the chronic HBV infection depends on both viral and host-specific factors. The disease outcomes have been correlated to serum HBV DNA levels and HBeAg status. Intra-hepatic distributions and localization patterns of viral antigens detected by immunohistochemistry could potentially provide additional prognostic values. In this study, the authors developed a modified classification system to characterize the expressions of HBV core antigen (HBcAg) and HBV surface antigen (HBsAg) in liver tissues of patients with chronic hepatitis B with and without HCC.

#### **Research frontiers**

The authors observed a high rate of HBV antigen detection in tumor tissue alone, but not in adjacent non-tumor liver in HCC patients with optimal HBV DNA suppression. This raised the question of HBV DNA integration into host genome leading to potential activation of oncogenes, induction of genetic instability and antiviral drug resistance.

#### Innovations and breakthroughs

The authors developed a modified classification system to standardize the reporting of the intrahepatic distributions and patterns of HBcAg and HBsAg. The authors discovered distinct differences in the distributions of these antigens in HBeAg positive and HBeAg negative chronic hepatitis B patients. Interestingly, authors observed variable HBV antigen expressions in the tumor tissues. These results provided insights into the potential mechanisms of disease persistence and carcinogenesis.

#### Applications

These findings open up new research study to confirm that these liver cancers were derived from clonal expansion of infected hepatocytes with high rate of proliferation, carcinogenic potentials and selective resistance to antiviral agents.

#### Terminology

Immunohistochemical staining of liver tissues: it applies standard streptavidinbiotin-immunoperoxidase technique to identify the various proteins in the liver.

#### Peer-review

The author of this paper evaluated the distributions and patterns of intrahepatic HBcAg and HBsAg in patients with chronic hepatitis B using a novel, modified classification system. The HBcAg expression is associated with high levels of serum HBV DNA and HBeAg status. The extensive HBsAg distribution even in HBeAg negative patents with low level of viremia explains the persistence of hepatitis B. The distinct expressions of viral antigens in tumor tissues deserve further evaluations to understand the carcinogenesis of HBV.

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