Ultrastructural Evidences of Hepatitis B Infection in Human Liver Biopsies Disclose Complex Assembly and Morphogenesis Pathways for Hepatitis B Virus

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Abstract: Despite of recent advances on acknowledge of hepatitis B virus (HBV) structure and biology, little is known about the morphogenesis and release of virions. In this study, the ultrastructural analysis of HBV components in liver biopsies from chronically HBV-infected patients disclosed complex assembly and morphogenesis pathways for HBV. HBV core (HBcAg) and surface (HBsAg) antigens were specifically identified in all liver biopsies from HBV-infected patients. HBcAg containing core-like particles 24-28 nm in diameter were observed both in nucleus and cytoplasm of hepatocytes. Dane’s-like particles were detected either budding to or into the lumen of ER close to HBsAg containing tubular structures. Besides, Dane’s-like particles were detected in different vesicular compartments resembling multivesicular endosomes. Other kind of enveloped HBV-like particles similar to the previously described cobra-shaped and horn-shaped particles were also observed in hepatocytes. Some of these particles were connected to the vesicle membrane through a stalk 20-22 nm in diameter. On the other hand, spherical subviral particles (SVP) were frequently observed in cytoplasmic vesicles. Moreover, a minor proportion of enveloped HBV-like particles budding through the plasma membrane to the extracellular space and bile canaliculi were detected. Interestingly, Dane’s-like particles in the bile canaliculi and entering into ductular-like cells were shown. Strikingly, Dane’s-like particles close to tubular structures and specific immunolabeling for HBcAg both in cytoplasm and nuclei of stellate-like cells were detected. Remarkably, HBV-like particles appeared entering hepatocytes from large cytoplasmic processes of fibrocytes raising the interesting possibility of a cell to cell passage of HBV through direct transcellular migration.

Key words: Hepatitis B virus, HBV morphogenesis, Dane’s particles, electron microscopy

INTRODUCTION

Hepatitis B viruses (HBVs; hepadnaviruses), are small, enveloped viruses and a causative agent of acute and chronic viral hepatitis. They replicate by transcription of RNA pregenomes and reverse transcription to produce the DNA genomes that are found in the virus particles secreted from infected hepatocytes. Hepatitis B virus (HBV) virions exhibit unique electron-microscopical features. The complete virus, called the Dane’s particle, is 42-47 nm in diameter. The nucleocapsid, or core particle, which is composed of single core protein species (HBcAg), contains a double-stranded DNA genome and the covalently attached viral polymerase and is surrounded by a membrane shell with two or three viral envelope...
proteins embedded\[4\]. The protein envelope present on HBV surface consists of three protein types: Small (S), Medium (M) and Large (L) proteins. The S protein (HBsAg) represents the main component of viral envelope and of subviral particles (SVP)\[4,5\]. SVP are seen in the form of spherical or elongated tubular structures of various lengths and 20-22 nm in diameter\[4,5\]. At the beginning, HBsAg is synthesized as an integral transmembrane Endoplasmic Reticulum (ER) protein, budding to the lumen in the form of particles, which undergo secretion\[13\].

Dane’s particles have been described in liver biopsies from HBV-infected patients\[6-10\]. They were similar to those visualized in human sera\[11,11\]. SVP have also been observed in great excess over virions. Moreover, other ultrastructural forms of HBV named tadpole- or cobra-shaped HBV with a head measuring 42 nm and a tail with 22 nm in diameter and horn-shaped HBV with a head and two branches of different length have also been reported\[11,11\].

Despite of recent advances on acknowledge of HBV structure and biology, little is known about the morphogenesis and release of virions\[14\]. Among factors complicating study of these processes, HBV replicates at a low rate and, therefore, the amount of HBV antigens in the liver is small, rendering its detection difficult\[4,5\]. Thus, it has been shown that the direct observation of HBV budding by electron microscopy is very difficult to achieve\[16\]. However, electron microscopy studies might be powerful tools to elucidate some of the mechanisms involved in HBV replication and virion morphogenesis as well as HBV pathogenesis\[17\]. To address these issues and to better understand virus-host interactions in HBV replication, an ultrastructural analysis of HBV components in liver biopsies from chronically HBV-infected patients was examined in this study.

MATERIALS AND METHODS

Patients and samples: Patients with chronic HBV infection hospitalized for hepatitis in the Institute of Gastroenterology, Havana, Cuba were recruited after informed consent in writing was obtained. Liver needle biopsies samples were taken at the time of routine diagnostic biopsy from all patients. Seven chronically HBV-infected patients (3 females and 4 males, aged 20-40 years) were selected based upon they were serologically positive to HBV enzyme immunoassays (Tecnosuma International, Havana, Cuba). They also were histologically confirmed as bearing chronic hepatitis and had abnormal serum alanine aminotransferase levels for at least six month before the biopsy was performed. None were seropositive for markers of hepatitis C virus, hepatitis A virus and human immunodeficiency virus by enzyme immunoassays (Tecnosuma International, Havana, Cuba). In addition, liver needle biopsies samples were taken from two HBV-uninfected healthy donor livers for transplantation purpose and from hepatitis C-infected patients as negative controls.

Antibodies: The following mouse monoclonal antibodies (mAbs) were used: anti-HBcAg CBSS-HepB.1 mAb specific for HBcAg and anti-HBsAg CBSS-HepB.2 mAb specific for HBsAg\[17\].

Transmission electron microscopy: The hepatic tissue samples were fixed for 1 h at 4°C in 1% (v/v) glutaraldehyde and 4% (v/v) paraformaldehyde, rinsed in 0.1 M sodium cacodylate (pH 7.4), post-fixed for 1 h at 4°C in 1% OsO4 and dehydrated in increasing concentrations of ethanol. The embedding was done as previously described\[18\]. Samples were examined with a JEOL/JEM 2000 EX transmission electron microscope (JEOL, Japan).

Immunoelectron microscopy (IEM): Samples of hepatic tissue were fixed with 4% (v/v) paraformaldehyde containing 0.2% (v/v) glutaraldehyde in 0.1 M phosphate buffer (pH 7.3) at 4°C for 3 h and washed with 0.1 M phosphate buffer pH 7.3. Fixed cells were dehydrated as described above, embedded in Lowicryl and polymerized by exposure to ultraviolet light at Room Temperature (RT) for 72 h. Ultrathin sections of liver biopsies were incubated with either CBSS-HepB.1 or CBSS-HepB.2 mAbs in phosphate buffer, for 45 min at RT. The sections were rinsed three times for 30 min at RT with 0.1% bovine serum albumin in phosphate-buffered saline pH 7.3 (BSA-PBS) and incubated for 1 h at RT with gold-labeled (15 nm) anti-mouse IgG (Amersham, UK) diluted 1:100 in BSA-PBS. As control the primary antibody was substituted by normal mouse serum. All sections were stained and analyzed with a transmission electron microscope as mentioned above.

Immunofluorescence staining: Samples were immediately fixed with 4% paraformaldehyde in PBS at 4°C and then mounted on gelatine-coated glass slides and stored for 2 days at -200°C. Later on, mounted samples were hydrated for 10 min in PBS and incubated with 0.2% Triton X in PBS during 10 min. To block non-specific antibody reaction, best results were obtained by incubating the sections with 0.2% BSA (free of IgG) (Sigma Chemical Co. St. Louis, USA) for 30 min. Samples were then rinsed with 0.1 M Tris-buffered saline pH 7.4 (TBS) and then incubated with primary antibodies diluted 1:200 in TBS containing 3% (v/v) normal goat serum, 3% (v/v) sucrose and 0.5% (v/v) Triton X in a humid chamber at 4°C for 72 h. Samples were rinsed three times for 30 min in TBS, and then incubated with fluorescein-isothiocyanate-conjugated (FITC-conjugated) donkey anti-mouse IgG (Amersham, USA) diluted 1:50 in TBS containing 3% (v/v) normal goat serum, 3% (v/v) sucrose and 0.5% (v/v) Triton X for 30 min at RT. Samples were then rinsed in TBS three times for 30 min, mounted in 90% glycerol and observed with a Zeiss Axioskop 2 (Carl Zeiss, Germany) fluorescence microscope equipped with a UV filter. Results were photographed with a Leica DFC 420 camera (Leica, Wetzlar, Germany) and analyzed with the software Leica Application Suite V3.4 (Leica, Wetzlar, Germany).
Mo, USA), for 10 min at RT. After two washes in PBS-T, samples were incubated overnight at 4°C with either CBSS-HepB.1 or CBSS-HepB.2 mAbs (dilutions 1:20 in PBS-T). Incubations were followed by washes with PBS-T. The second incubations were accomplished with FITC-conjugated anti-mouse IgG (dilutions 1:60 in PBS-T, Vector laboratories, Inc., Burlingame, CA., USA) for 1 h at RT. After three washes with PBS-T the sections from all samples were counterstained with propidium iodide (dilution 1:1000, Vector laboratories, Inc. Burlingame CA., USA), followed by extensive washing in PBS-T. Immunostained samples were coverslipped in Vectashield mounting medium (Vector Laboratories, Inc. Burlingame, CA., USA), sealed with nail polish and viewed on a confocal laser scanning microscope. Negative controls were performed by substituting the primary antibodies with normal mouse serum.

**Laser confocal scanning imaging and serial section collection:** Samples were viewed with a 60x (NA 1.4) objective on a Nikon microscope with attached laser confocal scanning system MRC 600 (BioRad, Watfod). Ten to twelve fields were imaged from each sample. Four to fifteen serial optical z-sections (0.2-0.5 μm thick) were collected from each observed field using the dual channel imaging with green exciter filter (554 nm) for propidium iodide excitation (rhodamine channel) and the blue exciter filter (494 nm) for the fluorescein channel. Each series of confocal optical sections was scanned through a total of 25 μm. The resulting optical sections were fully projected onto two-dimensional planes using the imaging processing system of the microscope (Camos package).

**RESULTS**

Reaction products suggestive of HBV antigens within hepatocytes were observed in liver biopsies from the seven HBV-infected patients (Fig. 1a,b). No signals were observed in the liver biopsy specimens from the control subjects (Fig. 1c). The specific stain reaction was blocked completely by incubation with normal mouse serum as primary antibodies (not shown). Hepatocytes containing such reaction products were distributed sporadically or focally in some lobuli. Samples immunolabeled with anti-HBsAg mAbs showed perinuclear and cytoplasmic reactions (Fig. 1b). On the other hand, HBCAg was immunolabeled in both the cytoplasm and nucleus (Fig. 1a).

In five out of seven liver biopsies showing positive reactions for HBV antigens, the presence of HBCAg within cell nuclei of infected hepatocytes in the form of ring-like structures (core-like particles) of 2-28 nm in diameter were demonstrated by IEM (Fig. 2a). Most of the nuclear core-like particles were likely empty inside. However, electron-dense core-like particles (probably containing viral genome) were observed in the cytoplasm of these samples (Fig. 2b). They form groups or are spread all over the cell.

**Fig. 1:** Immunofluorescence staining with either anti-HBcAg mAbs (a): Or anti-HBsAg mAbs (b): In liver biopsy specimens from HBV-infected patients. (c): Liver biopsy from a healthy subject, no immunostaining was observed using any of these mAbs. (Bar = 8 μm)

**Fig. 2:** Electron micrographs of core-like particles either in the nuclei or cytoplasm of hepatocytes. Fragment of either hepatocyte nuclei (a): Or cytoplasm (b): From a liver biopsy specimen of a chronically HBV-infected patient showing abundant core-like particles of 24-28 nm in diameter. Inset in a (Magnification): Immunolabeling of HBCAg in the nuclei of hepatocytes by IEM using anti-HBcAg mAbs. (Bar = 200 nm)
Fig. 3: Electron micrographs of Dane’s-like particles in liver biopsies from HBV-infected patients. (a): Part of a hepatic cell showing vesicles containing Dane’s-like Particles of 42-47 nm of diameter (black arrows) into larger cytoplasmic vesicle with a membrane invagination (white arrowhead). (b): Cytoplasmic vesicle resembling a multivesicular body containing Dane’s-like particles of 42-47 nm of diameter. (Bar = 200 nm)

In liver biopsies showing intranuclear core-like particles, enveloped HBV-like particles resembling Dane’s-like particles with diameters of 42-47 nm were observed into different cytoplasmic vesicles of hepatocytes (Fig. 3a, black arrows). Interestingly, some of these HBV-like particles containing vesicles were inside larger vesicles showing membrane invaginations (white arrowhead). Dane’s-like particles also appeared in vesicles resembling Multivesicular Bodies (MVB) (Fig. 3b).

In addition, the cytoplasm of hepatocytes from these liver biopsies contained tubular structures 20-25 nm in diameter (Fig. 4a, white arrows). Some cells were packed with these structures; in others they were only present in part of the cytoplasm. However, these structures were not detected in liver biopsies from either healthy subjects or from HCV-infected patients (not shown). Tubular structures were specifically immunolabeled with anti-HBsAg mAbs (Fig. 4b). No immunogold staining was observed in sections incubated with normal mouse immunoglobulins as primary antibodies (not shown). Besides, Dane’s-like particles with diameters ranging from 45-55 nm were localized close to the tubular structures (Fig. 4a, black arrows). They appeared budding into the lumen of ER. Moreover, spherical SVP with 20-22 nm in diameter were frequently observed inside cytoplasmic vesicles close to tubular structures (Fig. 4a, Lower Inset: black arrows).

Other kinds of enveloped HBV-like particles were detected close to the tubular structures. Dane’s-like particles with a long tail 20-25 nm in diameter and a length of 230-270 nm resembling the previously described cobra-shape particles were observed (Fig. 4a, arrowhead). In addition, Dane’s-like particles with two long tails 20-25 nm in diameter with a variable length of 300-400 nm resembling the previously described horn-shape particles were detected (Fig. 4a, Upper Inset: white arrowhead).
Fig. 5: Immunolabeling of HBcAg by IEM in stellate-like cells from liver biopsies of HBV-infected patients. (a): Liver biopsies from healthy subjects, no immunostaining was observed using anti-HBcAg mAbs and gold-labeled anti-mouse IgG. (b): Immunostaining with anti-HBcAg revealed HBcAg in cytoplasm and the nuclei (black arrows). GL: Lipid droplets. (Bar = 500 nm)

Fig. 6: Electron micrographs of liver biopsies from HBV-infected patients. (a, b): HBV-like particles protruding from large processes of fibrocytes and entering neighbour cells (white arrows). (c): HBV-like particles entering ductular-like cells (black arrows) (Bar = 300 nm in A and C, 500 nm in B)

Interestingly, tubular structures and Dane’s-like particles with diameters ranging from 45-55 nm budding into the lumen of ER were also observed in stellate-like cells (Fig. 4c, black arrows). Note the presence of Dane’s-like particles connected to the membrane through a stalk 20-22 nm in diameter (white arrowhead). Remarkably, HBCAg was shown to localize in stellate-like cells (Fig. 5b, arrows). Specific immunolabeling appeared distributed in the cytoplasm as well as in the nuclei. No immunogold staining was observed in liver sections from uninfected individuals (Fig. 5a) or in sections incubated with normal mouse immunoglobulins as primary antibodies (not shown).

Enveloped HBV-like particles were also localized in the extracellular space (Fig. 4d) and in bile canaliculi (not shown). Some of them appeared budding from the plasma membrane (white arrowhead). Most of these particles were clearly composed of an inner electron-dense core-like particle surrounded by an envelope. However, a proportion of empty particles were observed (white arrow). In addition, these particles were not detected in hepatocytes from normal controls (not shown).

It is interesting to note that Dane’s-like particles appeared protruding from large processes of fibrocytes and entering neighbour cells (Fig. 6a and b, white arrows). These cells were in close contact. Moreover, Dane’s-like particles entering ductular-like cells were seen (Fig. 6c, black arrows).

DISCUSSION

Both immunofluorescence and IEM studies specifically identified HBV antigens in all liver biopsies from HBV-infected patients. In agreement with previous studies, HBcAg was disclosed in the cytoplasm and cell nuclei of hepatocytes\cite{9,19}. On the other hand, HBsAg was noted in the cytoplasm as well as on tubular structures\cite{7,10}. It is interesting to note that the HBV genome has been detected in hepatocytes from these liver biopsies using an in situ hybridization assay\cite{20} followed by confocal microscopy (Falcon et al, unpublished results). Thus, data support the infection of hepatocytes by HBV in these samples.

Some ultrastructural lesions associated with chronic type B hepatitis are related to molecular structure of HBV\cite{7,21}. In the present work we were able to detect core and enveloped HBV-like particles in liver biopsy samples from HBV-infected individuals with chronic hepatitis. Core-like particles 24-28 nm in diameter were located both in nucleus and cytoplasm. Nuclear HBcAg possibly represents accumulation of empty nucleocapsids whereas cytoplasmic HBcAg may be considered as active virus replication\cite{19}. Enveloped HBV-like particles showed a hepadnavirus-like morphology (Dane’s particles) with ultrastructural
HBV virion budding and release currently are still not fully understood. Results from this study suggest complex assembly and morphogenesis pathways for HBV. Dane’s-like particles were detected either budding to or into the lumen of ER close to HBsAg containing tubular structures. This may represent a primary site for HBV virion assembly and budding[6,7]. On the other hand, Dane’s-like particles were observed into vesicles resembling MVB. Besides, vesicles containing Dane’s-like particles were detected inside larger vesicles heterogeneous in size showing membrane invaginations. It is possible that these vesicles derived from multivesicular endosomes (MVE)[22]. Thus, they may represent newly forming intermediates in either the degradation pathway or the HBV virion maturation. This is in agreement with a recent report showing assembly and budding of virion particles into a novel type of intracellular vesicles using duck HBV as a model system[23]. These vesicles derived from ER having unique identity and properties of ER, intermediate compartment, endosomes and MVB.

The MVB pathway is responsible to sort cargo proteins destined for degradation, lysosomal functions, or exosomal release toward intraluminal vesicles[22]. MVB functions have been implicated in budding by different enveloped RNA viruses[24]. Most of these viruses access the MVB machinery through its virus-encoded late assembly domains and bud through the plasma membrane. In addition, recent communications have reported the involvement of MVB functions in HBV virion maturation and egress using virus-replicating liver cell lines[25,26]. HBV virions have been suggested to bud into internal MVB-related compartments and exit the cell by the exosomal pathway.

In this study, a minor proportion of enveloped HBV-like particles budding through the plasma membrane to the extracellular space were detected. Some of these particles consist of empty particles with no electron-dense nucleocapsid possibly representing either defective interfering particles[25] or SVP[23]. In addition, enveloped particles with an electron-dense nucleocapsid likely corresponding to virions were observed. Some authors have shown that HBV envelope proteins do not appear in the plasma membrane[5]. So, based on the molecular characterization of the hepadnaviral morphogenesis it is assumed that hepatitis B virions are formed by budding of the capsids through an intracellular membrane[4]. However, other reports had shown that HBV envelope proteins could be immunodetected in the plasma membrane of hepatocytes from liver biopsies of HBV-infected patients[6,19,28]. In addition, HBV-like particles budding from the plasma membrane of hepatocytes has been reported[10,29]. Thus, it seems that a proportion of HBV could bud through plasma membrane during chronic infection in humans. Taken together, it is possible to suggest that similar to enveloped RNA viruses that use components of the MVB sorting pathway for the formation of progeny particles that mostly bud at the plasma membrane of infected cells, HBV may utilize MVB functions for budding not only into internal MVB-related compartments but also through the plasma membrane.

On the other hand, large tubular structures (that were specifically immunostained with anti-HBsAg mAbs) were detected in liver biopsy samples from HBV-infected patients. These structures were similar to those previously described in liver biopsies from HBV-infected patients[6,7]. In addition, spherical SVP were observed inside vesicles. Previous ultrastructural studies have proposed that HBV virions and SVP exploit a common morphogenetic pathway[5]. Nevertheless, results from this work showed that Dane’s-like particles and SVP localize in different cellular compartments. This is in agreement with recent evidences showing that the viral and subviral HBV assembly pathways seemingly differ in their requirement for cell functions and trafficking routes[25,29]. While HBV virions budding required the involvement of MVB functions, the mechanisms of SVP production showed to be clearly distinct and likely independent of MVB functions[25,29].

Strikingly, Dane’s-like particles were also observed close to tubular structures in stellate-like cells. In addition, stellate-like cells were specifically immunolabeled with anti-HBcAg in cytoplasm and nuclei suggesting that HBV could replicate in this cell type. Stellate cells have a central role in the pathogenesis of chronic liver injury[30]. Our finding raises the possibility that HBV could directly modulate the stellate cells functions playing a role in HBV pathogenesis. Further studies are necessary to define this interesting topic both at cellular and molecular levels.

In addition, other kind of enveloped HBV-like particles were observed in this study. They resembled the previously described tadpole or cobra-shaped and horn-shaped particles[1,7,11-13]. A recent report suggested that these particles are the native replicative form of HBV[13]. However, in agreement with this study, previous reports had shown the presence of Dane’s particles connected to the vesicle membrane through either a short or a long stalk 20 nm in diameter[10,5]. Besides, it has been shown that interfering with some of the components of the machinery that catalyzes MVB vesicle budding inhibited HIV-1 assembly and budding, resulting in the accumulation of partially budded
virions that remain connected to the plasma membrane. Therefore, another interesting hypothesis would be that these particles arise due to MBV functions impairment leading to deficient fission events during viral budding.

Our results showed Dane’s-like particles in the bile canaliculi and entering into ductular-like cells suggesting the involvement of the biliary system in the replicative cycle of HVB and the pathogenesis of HVB-induced liver disease. These findings support previous observations showing HBV components in ductular cells as well as Dane’s particles in bile canaliculi and budding of core particles from hepatocytes into the bile canaliculi.

By using the DHBV model, it has been previously suggested that liver sinusoidal endothelial cells play a key role in the initial uptake of DHBV into the liver. The authors proposed that DHVB initially scavenged by LSEC were thereafter released to infect adjacent hepatocytes. In this study, HVB-like particles appeared entering hepatocytes from large cytoplasmic processes of fibrocytes. This raises the interesting possibility that a cell to cell passage of HBV through direct transcellular migration may occur in vivo. A more profound study in this field could further contribute to clarify the life cycle of HBV.

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