

# Quantification of Serum Hepatitis B Virus DNA

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## Abstract

Hepatitis B virus (HBV) infection is an important cause of acute and chronic viral hepatitis. Worldwide there are more than 300 million chronic carriers of HBV, with 100 million carriers in China. In addition to serology markers and liver enzymes, the detection and quantification of HBV DNA in serum appear to be the most reliable methods for monitoring HBV infection and assessing responses to antiviral treatment. This review will discuss the methodology, sensitivity, technical aspects, turn-around time and the clinical applications of the different HBV DNA assays.

*Key words: HBV DNA. Quantitation PCR. Real-time PCR.*

## Introduction

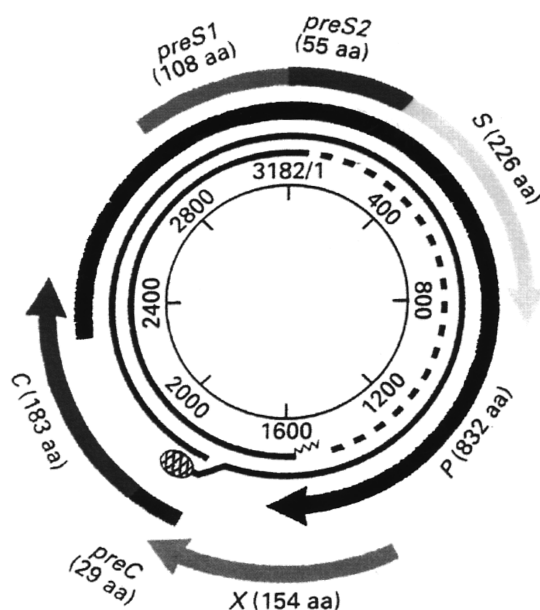
Although the institution of vaccination programmes has decreased the incidence of new HBV infections in many areas, HBV infection remains a significant global health problem with the number of chronic HBV carriers at more than 300 million worldwide. The HBV carrier rate in Hong Kong is about 9 %. Chronic HBV infection may result in an asymptomatic carrier state, reactive fulminant hepatitis, cirrhosis, or hepatocellular carcinoma.

The structure of HBV consists of a partly double-stranded circular DNA virus of the class Hepadnavirus<sup>1,2</sup>. The complete infectious virion (Dane particle) consists of a 42 nm spherical envelope carrying the HBsAg, a 27 nm inner shell of nucleocapsid with the hepatitis B core antigen (HBcAg), and a processing derivative hepatitis B e antigen (HBeAg). The inner nucleocapsid contains the viral genome and its tightly associated DNA polymerase. The partly double-stranded circular DNA genome is composed of approximately 3200 base pairs (Figure 1)<sup>3</sup>. The long DNA strand (L-) is the coding strand carrying four primary translational open reading frames: S, for the surface, pre-S1 and pre-S2 genes; C, for the core and pre-C genes; P, for the polymerase and reverse transcriptase genes; and X, for the X gene that activates viral and cellular promoters. The short DNA strand (S+) is of variable length capable of elongation at its 3' end by the activation of specific DNA polymerase to form a double-stranded molecule with the L- strand<sup>4</sup>.

The conventional diagnosis of the status of viral replication after HBV infection is based on the detection of serological markers HBsAg, anti-HBs, anti-HBc, IgM anti-HBc, HBeAg, and anti-HBe, which can indicate the stage of infection, and thus the degree of infectivity, as well as the individual's immune status<sup>5,6</sup>. However, the reliability of these serological tests can be influenced by the host's immune responsiveness. In addition, testing for HBeAg can not detect the presence or absence of viraemia in infections by pre-core or core promoter HBV mutants<sup>7-9</sup>. Rarely is HBV DNA detected in individuals sero-negative for HBsAg<sup>10-12</sup>. Whereas the measurement of liver enzymes reflects hepatocyte lysis, the detection of HBV DNA in serum provides direct evidence of viraemia. Recently, it has also been suggested that the determination of the virus load may provide clinically useful information during the course of infection. Moreover, monitoring of the changes in viral level during therapy of infected patients with interferon alpha and/or nucleoside analogues is of particular importance for the response in terms of virus elimination, viral reactivation, or development of drug resistant viruses<sup>13-17</sup>.

## DETECTION OF HBV DNA IN SERUM

Direct nucleic acid hybridization or polymerase chain reaction (PCR) assays are commonly used in the detection and quantification of serum HBV DNA. Examples of the former include dot blot hybridization and solution hybridization. Solution hybridization assays can utilize radioactivity (e.g. Genostics HBV



**Figure 1.** Diagrammatic representation of the hepatitis B virus genome. The viral DNA is partially double-stranded. The long strand of fixed length encodes seven proteins from four overlapping reading frames (surface S, core C, polymerase P, and the X gene X), and three up-stream regions (pre-C, pre-S1, and pre-S2). [aa: amino acids] (Reproduced from reference 3, Lee WM: *N Engl J Med* 1997; 337(24): 1733-45)

DNA, Abbott Laboratories), antibody capture (e.g. Hybrid Capture HBV DNA Tests, Digene Corp.), branched DNA (bDNA) signal detection systems (e.g. Quantiplex HBV DNA, Chiron Corp.), or photoactive cross-linking (NAXCOR) techniques. In general, assays based on PCR methodology (e.g. Amplicor HBV Monitor Test, Roche Diagnostics) are capable of detecting and quantifying lower levels of HBV DNA. This review will discuss the methodology, the technical aspects, and the clinical applications of the different HBV DNA assays.

### Detection of HBV DNA by Direct Nucleic Acid Hybridization

#### *Dot blot hybridization*

Dot blot hybridization (DB) has been used to detect HBV DNA in serum since the early 1980's<sup>18,19</sup>. The size and intensity of the hybridized dot on the autoradiogram semi-quantitatively indicates the amount of HBV DNA. The process has a sensitivity of approximately  $2.85 \times 10^6$  copies/ml or 10 pg/ml, and requires an assay time of 4-7 days.  $^{32}\text{P}$  radio-labeled probe was originally used for hybridization. With the use of non-radioactive probes such as alkaline phosphatase, digoxigenin, FITC and biotin, the assay time has been shortened to 2-3 days<sup>20,21</sup>. Imaging densitometry can be applied for more accurate and

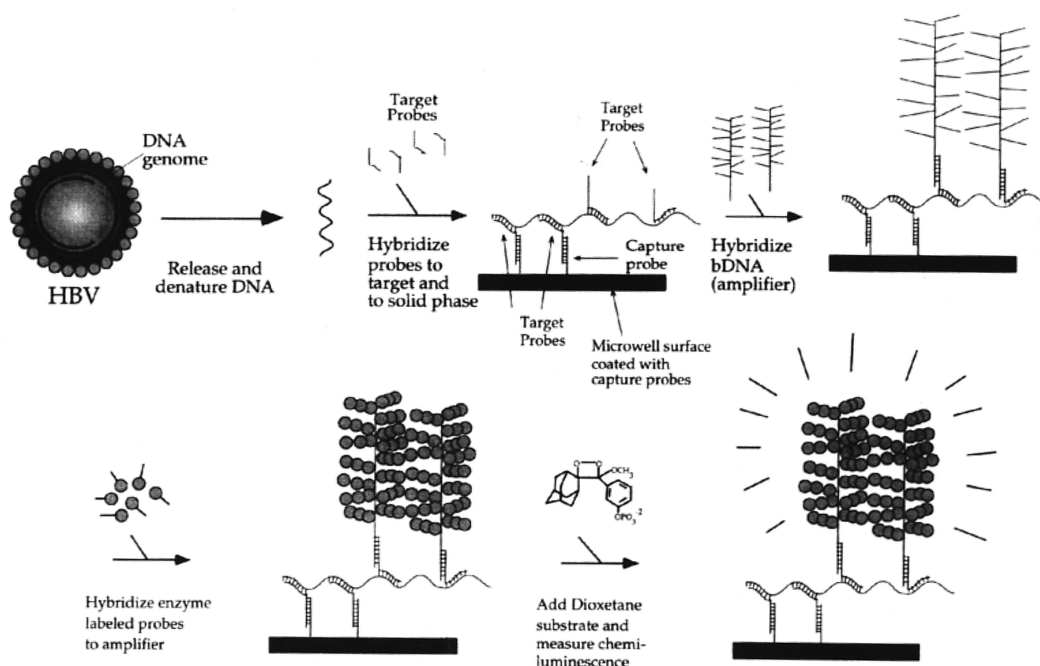
objective measurement of results<sup>22,23</sup>. However, DB assays have variable sensitivity and low reproducibility. Nevertheless, it can handle a large number of samples concurrently, and is relatively inexpensive.

#### *Solution hybridization*

Another method of detection uses solution hybridization with a radio-labeled probe. An example of this is the Genostics HBV DNA assay (Abbott Laboratories, North Chicago, IL, USA). Double-stranded HBV DNA in serum is denatured into single-stranded DNA and then allowed to hybridize with  $^{125}\text{I}$ -labeled single-stranded HBV DNA probes in solution for 18 hours at  $65^\circ\text{C}$ .  $^{125}\text{I}$ -labeled hybridization products are eluted with a sepharose column and radioactivity is measured in a scintillation counter<sup>24</sup>. Although the assay is simple to perform, the use of only one standard has resulted in sub-optimal reliability and reproducibility of results<sup>25,26</sup>.

#### *Signal Amplification by bDNA*

The Quantiplex HBV DNA assay (bDNA) (Chiron Corp., Emeryville, CA, USA) is a sandwich nucleic acid hybridization assay that utilizes branched DNA (bDNA) for signal amplification (Figure 2)<sup>27</sup>. HBV DNA in serum is first released and denatured by the



**Figure 2.** Schematic representation of the branched DNA (bDNA) assay for HBV DNA quantitation. (Reproduced from reference 27, Hendricks DA, et al: Am J Clin Pathol 1995; 104: 537-46)

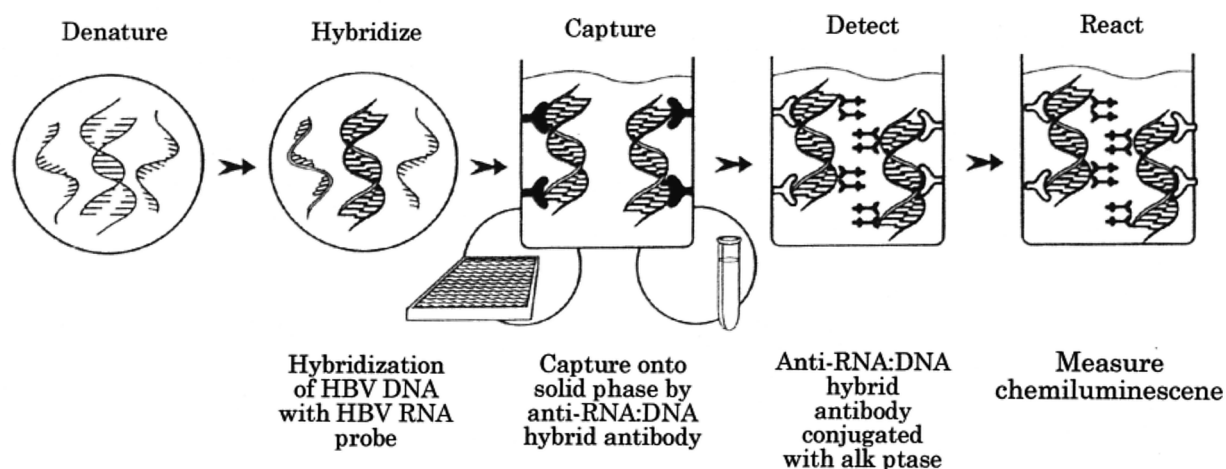
addition of a lysing agent and heating. It then hybridizes with two different sets of probes complementary to the conserved regions of the surface gene and core gene respectively. One set of probes also binds to microplate wells while the free ends of the other set of probes bind to the synthetic bDNA. Each bDNA molecule has multiple repeat sites for the binding of alkaline phosphatase labeled probes. The amount of HBV DNA captured is indicated by chemiluminescence after the addition of dioxetane substrate. This assay takes approximately 24 hours to perform and the detection range is  $7 \times 10^5 - 5 \times 10^9$  copies/ml (2.5 – 17,700 pg/ml). Apart from the increased sensitivity, the bDNA assay has demonstrated good reproducibility. Intra and inter-assay coefficients of variation have been found to be less than 15%<sup>27-29</sup>.

#### Ultraviolet (UV) cross-linking hybridization

The use of UV results in a highly specific and rapid hybridization process of approximately 30 minutes. An example is the NAXCOR HBV DNA assay (NAXCOR, Menlo Park, California, USA), which can measure HBV DNA concentrations of  $5 \times 10^5 - 5.7 \times 10^8$  copies/ml (2 – 2000 pg/ml) in 6 hours<sup>30,31</sup>.

#### RNA: DNA hybrid capture (HC)

Very specific RNA:DNA hybrids are formed between the HBV DNA in serum and RNA probes with specificity to HBV ad and ay strains. An example of this is the Digene HC HBV DNA test (Digene Diagnostics, Silver Springs, MD, USA). Using an anti-RNA:DNA antibody, the hybrids are captured onto the surface of microplate wells. A second set of alkaline phosphatase conjugated anti-RNA:DNA antibodies then binds to these immuno-captured hybrids. The chemiluminescence after the addition of substrate is measured on a luminometer (Figure 3). The first-generation Digene HC HBV DNA test has a detection range of  $1.4 \times 10^6 - 5.7 \times 10^8$  copies/ml (5 - 2000 pg/ml) and an assay time of 5 hours. The second-generation assay (Digene HC II HBV DNA test) shows improved sensitivity and detection range ( $1.4 \times 10^5 - 1.7 \times 10^9$  copies/ml, or 0.5 – 6000 pg/ml), and a shorter turn around time of 3.5 hours. More recently, the HC II assay also offers an ultra sensitive format which can quantify HBV DNA concentrations down to 5000 copies/ml (17 fg/ml), made possible by an initial concentration step of high speed centrifugation for 2 hours. Apart from the relatively simple technical requirements, the HC II assay also has low variability and high specificity<sup>32-34</sup>.



**Figure 3.** Schematic representation of the RNA:DNA hybrid capture assay. (Reproduced with permission from Digene Diagnostics, Silver Springs, MD, USA)

### Detection of HBV DNA by Polymerase Chain Reaction (PCR)

The most sensitive method of HBV DNA detection involves HBV DNA extraction, target sequence amplification by PCR, and detection of the PCR product<sup>35-38</sup>. There have been modifications aiming to improve specificity and reduce contamination. The extraction and denaturation of HBV DNA from serum samples can be achieved by phenol chloroform precipitation<sup>39</sup>, alkaline treatment<sup>36</sup>, microwave treatment<sup>40,41</sup> or with commercial DNA isolation kit. The number of PCR cycles and the primers which target different regions of the surface can vary with different assays. The amplified DNA product is usually detected by ethidium bromide staining following gel electrophoresis, and the sensitivity is around  $3 \times 10^4$  copies/ml (0.1 pg/ml). In theory, improved sensitivity of detection down to 100 copies/ml (0.3 fg/ml) should be achievable when the amplified DNA product is measured by dot blot assay (PCR-DB)<sup>42,43</sup>, by DNA enzyme immunoassay (PCR-DEIA)<sup>44</sup>, or by nested PCR (nPCR)<sup>45</sup>. However, PCR based methods in general have significant inter-assay and inter-laboratory variability<sup>46</sup>. Qualitative HBV DNA detection by PCR is useful when serological markers yield non-conclusive results. Although a negative result may indicate the absence of HBV in serum, the sensitivity of this assay as well as the possibility of the virus resident in infected cells should be considered. On the other hand, PCR methodology has also been incorporated into quantitative HBV DNA assays.

### Quantitative Polymerase Chain Reaction (qPCR)

Competitive PCR is used in these assays for HBV DNA quantification. An example is the Amplicor HBV Monitor Test (Roche Diagnostics, Branchburg, NJ, USA), in which common biotinylated primers allow co-amplification of HBV DNA and a known amount of an internal standard. The primers amplify a 104 base-pair (bp) sequence in the conserved pre-C/C gene region of the HBV genome, and the synthetic internal standard is double-stranded DNA with a 21-mer internal sequence different from the amplified HBV sequence. The PCR products are quantified by hybridization in microplate wells with a dinitrophenyl-labeled HBV specific probe and another dinitrophenyl-labeled probe with specificity for the internal standard, and colorimetric immunoenzymatic detection. Sensitivity of the Amplicor HBV assay is 1000 copies/ml (3.5 fg/ml) and the assay time is 6 hours. An automated assay with the COBAS analyzer can detect HBV DNA at 200 copies/ml (0.7 fg/ml) in 4 hours<sup>47,48</sup>.

### Real-time PCR

Recent advances in PCR instrumentation and fluorometric detection of PCR products have enabled the detection of PCR product during each amplification cycle using real-time PCR instrumentation. This is possible through the use of specially designed fluorescent probes such as the TaqMan probe, hybridization probes, and molecular beacons. Examples of such systems are the ABI PRISM 7700 Sequence Detection System (PE

Applied Biosystems, Foster City, Calif., USA), the Light Cycler (Roche Diagnostics, Mannheim, Germany) and the iCycler (Bio-Rad Laboratories). Continuous fluorescence measurement and quantification of the amplified products is generated using laser technology or other light detecting systems. Recent studies have shown that these systems can quantify HBV DNA within 5 hours from 200 to  $2 \times 10^9$  copies/ml with high reproducibility and specificity<sup>49-50</sup>, thus covering the detection range of earlier qPCR assays and the hybridization assays. Besides a wide range of linearity, amplification and signal detection take place in a closed tube system which does not require post-handling of PCR product, thereby reducing the risk of contamination. Furthermore, this methodology may allow for simultaneous detection of more than one virus<sup>50</sup>.

## CLINICAL APPLICATIONS OF HBV DNA ASSAYS

The PCR method is a sensitive assay to detect low level viraemia, but the clinical usefulness of qualitative PCR may be limited to rare situations when serological markers yield non-conclusive results. Quantification of viraemia is important in the selection of patients for antiviral therapy and to monitor them during treatment. Prior to the availability of quantitative assays for HBV, HBeAg was often used as a surrogate marker of the replicative status. The indirect nature and the presence of pre-core or core promoter mutants present obvious disadvantages of this marker. Furthermore, while disappearance of both HBV DNA and HBeAg from serum correlate with clinical and biochemical improvements<sup>51</sup>, HBV DNA is lost earlier than HBeAg and is a more direct index of response to therapy<sup>52</sup>. Although DB hybridization assays are less costly and are able to handle a large set of samples, they do not have sufficient sensitivity and accuracy to corroborate low-grade replication, which is especially important in patients under treatment. The improved sensitivity and reproducibility of nucleic acid hybridization tests with bDNA, UV cross-linking, or hybrid capture confer clinical usefulness, especially in serial monitoring of HBV DNA levels, e.g. during anti-viral treatment or to look for increased viral replication after immunosuppression<sup>53</sup>. Nevertheless, cost, technical complexity, and assay turn-around time remain valid concerns. Based on these considerations, the second generation hybrid capture (HCII) assay appears superior to other non-PCR

based assays because of its technical simplicity, rapidity (5.5 hours), and sensitivity (5000 copies/ml, or 17 fg/ml). The cost, technical and hardware requirements of quantitative PCR assays are inherent restrictions on their general clinical use, while the linearity and widened detection range make them valuable research tools.

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