# STUDY IN BETWEEN SERUM HBsAg, HBV DNA & BLOOD IMMUNE CELLS IN VIRAL HEPATITIS-B INFECTION OF NORTH INDIANS

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## ABSTRACT

The severity of viral hepatitis B (HBV) infection is attributed to inappropriate functioning of cellmediated immunity. Besides the importance of measuring serum HBV DNA and HBV surface antigen (HBsAg) as markers of viral replication and exposure, respectively, studies regarding their influence on immune cell status in chronic HBV infection are still scarce. Because such studies of chronic HBV patients have not been reported for India, we attempted to evaluate the relationship between serum concentrations of HBsAg, HBV DNA, and percentage of immune cells in peripheral blood of North Indian subjects with chronic HBV infection. Thirty-one HbsAg-positive subjects were evaluated for serum HB-e antigen (HBeAg), anti-HBe, and alanine transferase status by standard enzyme-linked immune -sorbent assay (ELISA) and biochemical procedures. Serum HBV DNA level was determined by real-time TaqMan® polymerase chain reaction assay. Serum HBsAg level was measured by a thirdgeneration sandwich ELISA kit. Peripheral immune cell profiling was done by multi-fluorometric flow cytometry analysis, for which 21 healthy subjects were included as controls. The majority (93.5%) of the study subjects were HBeAg-negative and anti-HBeAg-positive. Mean viral load, HBsAg, and alanine transferase levels were 4.20  $\pm$  1.96 log copies/mL, 5.98  $\pm$  4.62 log IU/mL, and 74.5  $\pm$  110 IU/mL, respectively. In comparison with controls, total T cell and cytotoxic T cell populations were significantly (P,0.05) reduced in HBV-infected subjects, while the status of B cells, natural killer cells, T helper cells, and ratio of T helper to cytotoxic cells remained unaltered. It is concluded that a suppression of the peripheral cytotoxic T cell population in chronic HBeAg-negative chronic HBV infection is influenced by increased viral load. Serum HBsAg concentration appeared independent of serum HBV DNA level and immune cell status. Non elevation of natural killer cell and T helper cell numbers in subjects harboring lower to moderate HBV loads is further indicative of non- induction of innate as well as a coordinated adaptive immune response favoring severity of the viral disease.

Key Words: Hepatitis B Virus, HBsAg Quantitation, Viral Load, Peripheral Lymphocytes, Chronic HBV Infection

# INTRODUCTION

The complexity of the pathogenesis of hepatitis B virus (HBV) has not been clarified properly as yet. The host's response through cell-mediated immunity rather than a direct cytopathic effect of the virus is said to be responsible for the damage to infected hepatocytes (Mukherjee, 2009). Peripheral blood mononuclear cells containing an aggregate of immune-competent cells, including subsets of T lymphocytes, natural killer cells, and lymphokine-activated killer T cells, are thought to play an important role in the persistence or control of HBV infection (Wang, 2003). Moreover, a strong association between cytotoxic T cells, liver cell damage and HBV clearance has also been demonstrated. Intra hepatic compartmentalization of HBV-specific CD8 cells, with a progressive decline in human leukocyte antigen (HLA)-DR activation markers on these cells, has been reported in resolving acute HBV infection (Sprengers, 2006). Apart from the antiviral response on the host's part, measurement of serum HBV DNA, being a marker of viral persistence and replication, is the common tool used to monitor treatment response in HBV infection. Besides the variable treatment outcome in patients with chronic HBV treated with nucleoside/nucleotide analogs and interferon's, a high response rate in patients having a low baseline level of HBV DNA has been demonstrated (Rijckborst, 2012). The prime determinant of the likelihood of liver injury and its intensification to cirrhosis has therefore been attributed to the interplay of the host

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immune response and the replication ability of the virus. Recent studies of HBV pathogenesis in animal models demonstrated that enhanced recruitment of virus-specific T lymphocytes into the liver cells is critical for the pathogenesis of both HBV infection and hepatocellular carcinoma (Wong, 2014). On the other hand, detection of HBsAg as the first serologic marker of the disease is done routinely, the persistence of which beyond six months signifies chronic infection, while its disappearance from the circulation marks closure to cure outcome of HBV infection. The quantitative measurement of HBsAg gained momentum in recent times due to its importance as a predictor of early virologic response to antiviral therapy as well as helping to distinguish disease status in chronic HBV patients. Furthermore, quantitative measurement of serum HBsAg is indicated as a surrogate marker for viral covalently closed circular DNA and intra hepatic HBV DNA, which are believed to maintain the replicative form of HBV DNA, serum HBsAg levels, and the numeric status of circulating immune competent cells in Indian chronic HBV patients are far from adequate, the present study explores the possible relationship between these parameters in Indian subjects with chronic HBV infection.

# MATERIALS AND METHODS

We manually searched epidemiologic studies that Thirty-one subjects who attended the outpatient's department of the F H Medical College & Hospital, Firozabad, during the period May 2014 to August 2014, were positive for HBsAg for more than six months, and had clinical features of chronic HBV, were prospectively included as study group patients. Twenty-one voluntary blood donors devoid of HBsAg in their sera served as controls for the lymphocyte estimation study. Informed consent was obtained from both the control and study group subjects, and the study was approved by the Institutional Ethics Committee.

#### Serologic and Biochemical Parameters

The serum HBeAg and anti-HBe status of the subjects were determined by commercial enzyme-linked immunosorbent assay (ELISA) kits (Amar-EASE, Taiwan) as per the manufacturer's instructions. Serum aspartate transferase and alanine transferase levels were ascertained by an automated clinical biochemistry analysis system (Randox Oceanside, CA).

# Quantitation of Serum HBV: DNA and Genotyping

Extraction of viral DNA from patient's sera was done using the High Pure System Viral Nucleic Acid kit (Roche Molecular Systems, Belleville, NJ) as per the manufacturer's instructions. Amplification and subsequent quantification of extracted DNA were performed by the COBAS® TaqMan®HBV test using a COBAS TaqMan 48 Analyzer (Roche Molecular Systems) which has a lower limit detection of 6 IU (35 copies) of HBV DNA/mL. HBV genotypes were determined using an in house nested polymerase chain reaction assay involving type-specific primers to generate respective genotype-specific amplicons. (Naito H, 2001).

# Quantitation of Serum HBsAg

Serum concentration of HBsAg was measured by a sandwich ELISA kit (Alpha Diagnostic International, SanAntonio, TX) as per the manufacturer's instructions. This kit meets the requirements for a thirdgeneration test upon testing against the Food and Drug Administration reference panel. Performance of this kit was compared with a commercial test kit (Abbot Auszyme II, Abbott Laboratories, Abbott Park, IL) and assumed to detect all possible combinations of HBsAg subtypes (adw, adr, ayw, ayr), having a lower limit of sensitivity of 0.3 ng/mL, as per the manufacturer's declaration. For convenience, HBsAg concentrations measured in ng/mL were converted to IU/mL, considering that 0.15 ng/mL of HBsAg is equivalent to 350 IU/mL (Biswas, 2003).

# Quantitation of Peripheral Blood Lymphocytes

The following antibodies were divided in two panels for fluorescence-activated cell sorter (FACS) analysis. A three-color direct immunofluorescence reagent TriTEST CD4 fluorescein isothiocyanate (FITC)/CD8 phycoerythrin (PE)/CD3 peridinin chlorophyll protein (PerCP) was used. Lymphocytes and

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HBsAg in HBV infection to identify and determine the absolute counts of mature human T lymphocytes (CD3+), helper/inducer (CD3+CD4+) T lymphocytes and suppressor/cytotoxic (CD3+CD8+)T lymphocytes, and a four-color direct immunofluorescence reagent MultiTEST CD3 FITC/CD16+CD56 PE/CD45 PerCP/CD19 allophycocyanin was employed to identify and determine the percentages and absolute counts of mature human T (CD3+), natural killer (CD3-CD16+ CD56+), and B (CD3-CD19+) lymphocytes in erythrocyte-lyzed whole blood (Fisicaro, 2012; Larrubia, 2013). The reagents employ fluorescence triggering, allowing direct fluorescence gating of the lymphocyte population to reduce contamination by unlyzed or nucleated red blood cells in the gate. Matched isotype antibodies were used for control staining. All reagents were obtained from Becton Dickinson (San Jose, CA) and used as per the manufacturer's instructions. Three- and four-color analysis was performed by gating 10,000  $\pm$  20,000 events on acquisition of respective cellular expression for each analysis using a FACS Caliberor FAC Scan (Becton Dickinson) flow cytometer.

## Statistical Analysis

Descriptive statistics (mean, median, standard deviations, and interquartile ranges [IQR]), Student's t-test, and Fisher's Exact tests were performed as and where applicable using SPSS software, version 13.0 (SPSS, Inc., Chicago, IL). A value of P < 0.05 was considered statistically significant.

## **RESULTS AND DISCUSSION**

Of 31 mostly male (96%) patients, 93.5% were negative for HBeAg and positive for anti-HBe. Genotype D was found to be predominant (83.8%), followed by genotype A (16.2%). A significantly lowered total T cell (P = 0.02) and cytotoxic T cell population (P = 0.018) was evident in HBV-infected subjects in comparison with the healthy controls. The median values and IQR of total T cells for control and study subjects observed were 69.1 (IQR = 9.1) and 66.9 (IQR = 7.55), respectively, while the median values of cytotoxic T cells for control and study subjects were 27.2 (IOR = 13.1) and 24.6 (IOR = 0.2) respectively. No significant alteration of cytotoxic to helper T cell ratios was apparent upon comparing the control (median = 1.07; IOR = 0.39) and patient (median = 1.11; IQR = 0.68) groups. The median viral load (log copies/mL) and HBsAg (log IU/mL) levels noted in the patient group were 3.91 (IQR = 2.10) and 4.59(IQR = 1.04). As depicted in Table 1, upon grouping of patients on the basis of viral load (<2000 IU/mL) in comparison with controls, while the natural killer cell status remain unaltered, a significant drop in the total T cell population was observed in the group of patients having a viral load, (< 2000 IU/mL)(P =0.02), as well as in the group of patients having a viral load (> 2000 IU/mL) (P = 0.05). Interestingly, compared with control subjects, the population of cytotoxic T cells reduced significantly (P = 0.04) in patient saving a higher viral load (>2000 IU/mL) than in the patients having a lower viral load (>2000 IU/mL), while no alteration in the helper T cell population was encountered in either group (Table 1). No significant differences were found in immune cell profile and HBsAg levels between the patients having lower or higher viral loads, but both alanine transferase and aspartate transferase levels increased significantly (P < 0.02) with increasing viral load (Table 2).

The types of cell-mediated responses expressed in the early stages of HBV infection can influence the subsequent outcome. In chronic HBV infection, the HBV-specific T cell responses weak in peripheral blood (Stoop, 2005). Mainly under the influence of high viral and antigen load, T cells are attracted into the infected liver where they are diluted among virus-nonspecific T cells and non-T cells that are the predominant cell population of the intrahepatic infiltrate, resulting in the T cell hyporesponsiveness typical of chronic patients (Maini, 2000; Tseng, 2013). Furthermore, suppression of T cell responses is more profound in highly viremic patients (Von, 2007) which is reported to be overcome to some extent by enhanced levels of peripheral blood CD4-mediated T cell reactivity to HBV nucleocapsid antigens.20 In the present study, we attempted to evaluate the influence of serum viral load (HBV DNA) and serum HBsAg load on the status of peripheral immune competent cells in chronic HBV infection. We found suppression of the total peripheral T cell population, but not of natural killer cells, in our chronic HBV patients in comparison with healthy controls, indicating HBV-induced suppression of adaptive immunity, but non induction of innate immunity in the peripheral lymphocyte pool. Considering the importance of

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the cytotoxic T cell mediated response for elimination and suppression of HBV of replication, 19 selective depletion of cytotoxic T cells, but not the population helper T cells, as evident in our patients having a higher viral load (.2000 IU/mL), is indicative of persistent chronicity of HBV infection. However, our data, to some extent, differ from an earlier study that showed decreases in the percentages of CD3+, CD4+ cells, and the ratio of CD4+cells/CD8+ cells, with an increase in CD8+ cells in chronic HBV infection (Xu, 2006) Apart from these, although aspartate transferase and alanine transferase values correlated with serum HBV DNA levels, the concentration of serum HBsAg, viral load, and peripheral immune cell status appeared independent of each other in our study subjects (Martinot-Peignoux, 2013).

Cell types%	Groups					
	Controls (n = 21)	Patients-Gr1 (n = 12)	P value	Controls (n = 21)	Patients-Gr2 (n = 19)	P value
T Cells	$68.98 \pm 5.74*$	$63.5 \pm 6.21*$	0.02*	$68.98 \pm 5.74*$	64.6 ± 7.31*	0.05*
B Cells	$14.82\pm4.87$	$16.9\pm4.54$	0.27	$14.82\pm4.87$	$16.7\pm7.15$	0.54
NK Cells	$17.19 \pm 4.45$	$16.82\pm4.35$	0.83	$17.19 \pm 4.45$	$16.73\pm6.06$	0.79
Tc Cells	31.69±12.65*	$26.22\pm5.67$	0.204	31.69±12.65*	$24.41\pm7.05*$	0.046*
TH Cells	$30.9\pm5.28$	$33 \pm 4.93$	0.30	$30.9\pm5.28$	$30.40\pm5.34$	0.781
TH: Tc	$1.114\pm0.413$	$1.32\pm0.463$	0.208	$1.114\pm0.413$	$1.34\pm0.44$	0.111

 Table 1: Comparison of peripheral immune cells profile of control and two groups of HBV-infected subjects at baseline

**Notes:** \*Significant (Student's unpaired t-test); ¶Viral load, 2000 IU/mL; §Viral load. 2000 IU/mL **Abbreviations:** NK, natural killer; HBV, hepatitis B virus; Tc, cytotoxic T cells; TH, helper T cells; Gr1, low viral load group; Gr2, high viral load group

Parameters	Viral load, 2000	Viral load. 2000 IU/mL (n =	P value
	IU/mL (n = 12)	19)	
T cell	$63.52 \pm 6.2$	$64.6 \pm 7.3$	0.701
B cell	$16.89 \pm 4.54$	16.14 ± TC 7.39	0.745
NK cell	$16.82\pm4.35$	$16.73\pm6.06$	0.968
cell	$26.22\pm5.67$	$24.41\pm7.05$	0.501
TH cell	$33\pm4.93$	$30.40\pm5.34$	0.227
TH/TC	$1.326\pm0.46$	$1.345 \pm 0.44$	0.919
Viral load (log DNA copies)	2.57 ± 1.13*	$4.68 \pm 2.6^{*}$	0.0237*
HBsAg (log IU/mL)	$4.80 \pm 4.62$	$4.61\pm4.59$	0.17
ALT (IU/mL)	$24.1 \pm 10.3*$	$75.8 \pm 54.1*$	0.01*
AST (IU/mL)	$23.57 \pm 7.13*$	$39.12 \pm 16.75*$	0.011*

 Table 2: Comparative cellular, virological, serogical and biochemical parameters of two HBV

 disease groups based on viral load at baseline

**Note:** \*Statistically significant (Student's unpaired t-test). **Abbreviations:** HBV, hepatitis B virus; NK, natural killer; ALT, alanine transaminase; AST, aspartate transaminase; HBsAg, hepatitis B surface antigen; Tc, cytotoxic T cells; TH, helper T cells

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In a previous study, peripheral natural killer cell levels in chronic HBV have been shown to be lower than in controls and, in comparison between patients grouped on the basis of viral load, the level of natural killer cells was significantly lower in patients having a higher viral load (6.81 log copies/mL) than those having a lower viral load (4.81 log copies/mL) (Karageorgopoulos, 2014). In contrast, in our study involving lesser viral load, while no alteration was found in the level of natural killer cells compared between patients having lower (2.57 log copies/mL) and higher (4.68 log copies/mL) viral loads and controls, a significant reduction in cytotoxic T cells was evident in patients having a higher viral load. Although the immunologic cellular mechanism by which the body clears HBV involves specific and nonspecific T lymphocyte responses, the HBV-specific cellular immune response mounted by the body against the virus has been considered to be an important factor related to viral clearance. In this respect, one of the limitations of the present study is that we did not determine the percentage of HBV-specific peripheral lymphocytes and this, along with the small sample size, prevents the drawing of a conclusion, and further study along this line is warranted. Quantitative measurement of serum HBsAg based on ELISA and chemiluminescence micro particle assay has been used (Efe and Wahlin, 2013) which recently gained importance providing level of HBsAg as a potential marker for monitoring therapeutic responses. Furthermore, the role of serum HBsAg quantification in distinguishing inactive carriers from subjects having the active form of the disease and harboring genotype D has been implicated in chronic HBV. Genotype D was predominant in our study subjects, which is in accordance with our previous findings from this part of the country. The present study is the first of its kind evaluating the relationship between serum HBsAg level, HBV DNA, and peripheral immune cell populations in Indian patients with chronic HBV who are supposed to harbor genotype D (Bertoletti, 2012). Despite the aforementioned limitations, the present study evidenced a non association of serum HBsAg concentration with serum HBV DNA, as reported earlier. The study further indicates that, irrespective of serum HBsAg level, HBV suppresses the total T cell population in the peripheral circulation, and specifically the cytotoxic T cell population, which was more pronounced in patients having comparatively higher viral load, without altering the status of other cell types or T cell subsets, reflecting the chronicity of the disease.

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