Prevalence and characterization of occult hepatitis B infection among blood donors in central Saudi Arabia

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ABSTRACT

الاهداف : تحديد النمط الجيني وتقييم مدى انتشار هذه العدوى بين المتبرعين بالدم في الرياض .

الطريقة: أجريت هذه الدراسة المستعرضة في مستشفى الملك خالد الجامعي خلال الفترة من يناير 2011م إلى يناير 2012م. تم خلالها فحص عينات المتبرعين بالدم (العدد=8501) للبروتين السطحي لفيروس الكبد ب والأجسام المضادة للكور بروتين للفيروس. جميع العينات السلبية للبروتين السطحي والايجابية للأجسام المضادة للكور بروتين لفيروس الكبد ب تم اختبارها للكشف عن الأجسام المضادة للبروتين السطحي لفيروس الكبد ب و كذلك تحديد كمية المادة الوراثية و تحليل النمط الجيني.

النتائج: من جميع العينات التي تم اختبارها (8501)، كان هنالك 56 (0.7%) عينة إيجابي و 8445 (99.3%) عينة سلبية للبروتين السطحي لفيروس الكبدُّب. ومن بين العينات السلبية، كان هنالك 198 (2.3%) عينة إيجابية للكور بروتين لفيروس الكبد ب ومن المحتمل أن يكون لديهم عدوى التهاب الكبد ب الغامض. ومن بين العينات الإيجابية للكور بروتين ، كان هنالك 119 (60.1%) عينة ايجابية في حين كان هنالك 79 (39.9%) عينة سُلبية للأجسام المضادة للبروتين السطحي لفيروس الكبد ب. وأظهر التحليل الجزيئي لفيروس الكبد ب في أن 17 (8.6%) عينة من أصل 198 كانت إيجابية (عدوى التهاب الكبد الفيروسي ب الغامض) وكانت جميعها سلبية لاختبار الأجسام المضادة للبروتين السطحي لفيروس الكبد ب . وكانت كمية المادة الوراثية للفيروس منخفضة (186-20> وحدة دولية/مل) في جميع الحالات المكتشفة. و أظهر تحليل النمط الجيني أن 15 (88.2%) عينة من أصل 17 تحتوي النمط الجيني نوع دي، بينما كان 2 (11.8%) عينة تحتوي النمط الجيني إي .

الخاتمة: انتشار عدوى التهاب الكبد الفيروسي ب الغامض بين المتبرعين بالدم في الرياض هو%0.2. ولذلك فمن المستحسن أن يدرج الاختبار الجزيئي لفيروس الكبد ب مع الفحوصات المصلية لفحص المتبرعين بالدم.

Objectives: To evaluate the prevalence of occult hepatitis B viral infections (OBIs) among blood donors considering the clinical and epidemiological importance of identifying OBIs.

Methods: A cross-sectional study conducted at King Khalid University Hospital, Riyadh, Saudi Arabia between January 2011 and January 2012. Blood donors (n=8501) were screened for Hepatitis B virus surface antigen (HBsAg) and hepatitis B core antibodies (HBcAb). All HBsAg-negative and HBcAb-positive samples were tested further for hepatitis B surface antibodies (HBsAb), hepatitis B virus (HBV)-DNA, and HBV genotyping.

Results: Of the 8501 serum samples tested, 56 (0.7%) were positive and 8445 (99.3%) were negative for HBsAg. Among the HBsAg-negative samples, 198 (2.3%) were positive for HBcAb and these patients were suspected to have OBIs. Among the HBcAb-positive samples, 119 (60.1%) were positive while 79 (39.9%) were negative for HBsAb. Analysis of HBV-DNA for the suspected OBIs showed that 17 out of 198 samples (8.6%) yielded positive results, and all of them were HBsAb-negative. The viral load was low (<20-186 IU/mL) in all OBIs. Hepatitis B virus genotyping showed that 15 out of 17 samples (88.2%) were genotype D, and the other 2 samples (11.8%) were genotype E.

Conclusion: The prevalence of OBIs among blood donors in Riyadh was 0.2%. Therefore, it is recommended that HBV molecular testing should be incorporated with serological assays for screening of blood donors.

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Tepatitis B infection is a major health problem worldwide.^{1,2} It is caused by hepatitis B virus (HBV), which primarily targets the liver and causes acute and chronic hepatitis that may progress to cirrhosis, hepatocellular carcinoma, or in rare conditions, fulminant hepatitis.^{1,2} Hepatitis B virus is an enveloped virus with a double-stranded circular DNA genome. It belongs to the family of Hepadnaviridae and has 8 different genotypes, namely HBV genotypes A-H.¹ The HBV genotype is a variable that can potentially influence the outcome of chronic hepatitis B and the success of antiviral therapy. Hepatitis B virus has 3 antigens, surface antigen (HBsAg), e antigen (HBeAg), and core antigen (HBcAg), and therefore, after infection 3 antibodies are generated by the immune system: HBsAb, HBeAb, and HBcAb. These HBV markers along with the viral load are particularly important in defining the stage of HBV infections and in their management.¹ Hepatitis B virus is transmitted parentally by different routes including exposure to infectious blood or body fluids, organ transplant, or by sexual contact.² Evidence has shown that HBV transmission can occur during the serological window period when the HBsAg is absent and the viral DNA is present in the circulation of infected carriers.³ It has been estimated that more than 2 billion people have been infected with HBV worldwide.⁴ This includes 350 million chronic carriers of the virus. The disease has caused epidemics in different parts of the world including Asia and Africa.⁴ Moreover, it is endemic in China, Thailand, Vietnam, Brazil, Peru, Venezuela, and southern African countries.⁵ Despite the introduction of the HBV subunit vaccine in Saudi Arabia in 1989, HBV infection was the predominant type of viral hepatitis identified among Saudi adults for the 2000-2010 period comprising 49.3-53% of all cases.⁶⁻⁸ However, a subgroup of HBV-infected individuals are characterized by the persistent presence of HBV-DNA and absence of HBsAg in their circulation; this condition is known as an occult HBV infection (OBI).^{9,10} The prevalence of OBIs among HBcAb-positive blood donors has been reported as 12.2% in Iran, 2.8% in Lebanon, and 2.9% in Pakistan.¹¹⁻¹³ Previous study revealed that OBIs may persist for years in infected individuals without apparent symptoms of overt HBV infection; however, it can be still transmitted to others.¹⁴ Other reports have shown that co-infection, drug abuse, or immunosuppressants

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can trigger an increase in HBV-DNA levels without an increase in HBsAg.^{15,16} The advent of highly sensitive polymerase chain reaction (PCR) assays has made it possible to identify HBV carriers who are negative for HBsAg, but who harbour low levels of HBV-DNA in serum or liver tissue, especially among blood and organ donors. Considering the clinical and epidemiological importance of detecting OBIs, we aimed to utilize molecular and the serological testing for the characterization and evaluation of the prevalence of OBIs among blood donors in Riyadh, the central region of Saudi Arabia.

Methods. A cross-sectional study performed at King Khalid University Hospital (KKUH) between January 2011 and January 2012. A total of 8501 volunteer donors were enrolled in this study including 8161 males (96%) and 340 females (4%). The mean age of the participants was 40 years (range, 17-63 years). All participants were eligible to donate blood and were selected based on physical examination and medical history, including a modified version of the general blood bank questionnaire form from the American Association of Blood Banks (AABB). Serum samples were received by the Virology and Molecular Biology laboratories for routine blood screening and molecular testing at KKUH. All samples that were positive for hepatitis C virus antibodies (HCV Ab), human immunodeficiency virus antigen and antibodies (HIV Ag/Ab), and human T lymphocytic leukaemia virus I and II antibodies (HTLV I and II Ab) were excluded. This study was approved by and performed according to the guidelines of KKUH and the college of medicine institutional review board committee. All procedures performed in were in accordance with the ethical standards of the institutional, or national research committee and with the 1964 Helsinki declaration and its later amendments or comparable ethical standards. Informed consent was obtained from all participants included in the study.

Detection of HBsAg, HBcAb, and HBsAb. Serum samples were tested for HBsAg, HBcAb (total antibodies IgM and IgG), and HBsAb based on chemiluminescent microparticle immunoassays (CMIAs). These tests were performed using CE-approved kits (ARCHITECT[®] HBsAg, ARCHITECT[®] anti-HBc, ARCHITECT[®] anti-HBs) and the Architect i2000 analyzer (Abbott, Lake Forest, IL, USA).

Assessment of HBV viral load and genotyping. Serum from HBsAg-negative and HBcAb-positive volunteer blood donors were processed for HBV-DNA quantification and genotyping. The HBV viral load was measured for individual donors using the COBAS[®] TaqMan[®] analyzer and COBAS[®] AmpliPrep/COBAS[®] TaqMan[®] HBV Quantities test, version 2.0 (Roche Diagnostics, Germany) as previously described.¹⁷ The limit of detection of the assay was 16.4 IU/mL and the linear range of quantification was 20 IU/mL as stated by the manufacturer kit insert. Serum samples containing HBV-DNA were subjected to HBV genotyping using nested PCR and an INNO-LiPA[®] HBV Genotyping kit (INNOGENETICS[®] N.V, Belgium) as described previously.¹⁸

Statistical analysis. Data were collected and entered into a Microsoft Office Excel file for statistical and descriptive analysis. The percentage and the mean values were used where applicable.

Results. Of the 8501 blood donors samples screened for HBsAg, 56 samples (0.7%) were positive, confirming current HBV infection, whereas 8445 samples (99.3%) were negative (Figure 1). Among the HBsAg-negative samples, 198 (2.3%) were positive for HBcAb and 8247 (97.7%) were negative. All HBcAb-positive samples were subjected to HBsAb and HBV viral load testing. Of the 198 HBcAb-positive samples, 119 (60.1%) were positive for HBsAb, but all of them

were negative for HBV-DNA (Figure 1). Moreover, 79 out of 198 samples (39.9%) were negative for HBsAb, but among those, 17 samples (21.5%) were HBV-DNA positive, confirming OBIs, and 62 samples (78.5%) were HBV-DNA negative (Figure 1 and Table 1). The viral load of HBV-DNA-positive samples was low, ranging between 20 and 186 IU/mL (Table 1). Analysis of HBV genotyping for those samples showed that 15 out of 17 samples (88.2%) were genotype D, and the other 2 samples (11.8%) were genotype E (Figure 1 & Table 1).

Discussion. This is the first report that characterizes and defines the frequency of OBIs among blood donors in Riyadh. The prevalence of HBV infection among blood donors was 0.9%. This includes a 0.7% HBsAg-positive and a 0.2% HBV-DNA-positive, but HBsAg-negative population (OBIs). In agreement with these results, a recent report has shown that the prevalence of HBsAg among blood donors was 1.5%.¹⁹ Other studies have revealed that the HBsAg positivity rates in blood donors was 4.3% in Egypt and 2.21% in Pakistan in 2006.²⁰ On the other hand, the prevalence of OBI among blood donors was 0.04-0.2% in Italy, 0.02-0.15% in India, 0.016% in Korea, and 1.18% in Egypt.^{14,21,22}

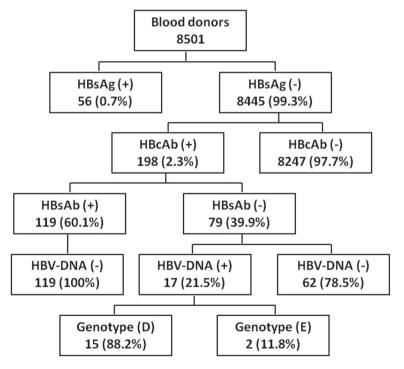


Figure 1 - Prevalence of serological and molecular markers of hepatitis B virus (HBV) infection among blood donors in central, Saudi Arabia. HBsAb - hepatitis B surface antibodies

In this study, 198 out of 8501 samples (2.3%) were HBsAg-negative, but HBcAb-positive and these donors were suspected to have an OBI. This result was markedly lower than that of a previous study carried out in 1997, which reported the prevalence of HBcAb among Saudi blood donors as 16.4%.23 However, our data were consistent with other recent reports which have observed an overall reduction in HBV infection over the last ten years because of the introduction of the HBV vaccine.²⁴ Previous published studies revealed that the prevalence of HBcAb within HBsAg-negative blood donors was 0.56% in the United Kingdom, 0.8% in the United State of America, 1.4% in Germany, 2.1% in Iran, 3.1% in Brazil, 4.9% in Italy, 13.5 in Korea, 15% in Greece, and 76% in Ghana.^{21,22,25-30} In the current study, 198 samples were suspected with an OBI and 17 samples were confirmed by using the real-time PCR. However, 181 samples were positive for HBcAb but negative for HBV-DNA, and those were still considered as suspected HBV infection cases. The failure of detecting HBsAg and HBV-DNA could be due to viral mutation, or due to the limitation of the serological and molecular assays used. It has been suggested that

such conditions may be observed during infection with mutant HBV, or during the period when the levels of circulating HBsAg in the infected individuals are below the detection threshold.^{1,10}

In our study, all HBV-DNA-positive cases had low viral load (<186 IU/mL) and were negative for HBsAb. Thus, these cases may have recovered from previous infections but had persistently low levels of HBV. In agreement with this result, several studies have reported low HBV-DNA levels (<200 IU/ml) in subjects with OBI.^{14,22,27,31} Also, a study carried out in Iran has demonstrated that all OBI cases were individuals negative for HBsAb.¹¹ Other studies have shown that there was no significant difference in HBV-DNA positivity between HBsAb-positive and HBsAb-negative samples.^{31,32} These results suggested that blood obtained from donors positive for HBsAb and HBcAb markers was not necessarily safer than that from donors positive HBcAb alone.

Hepatitis B virus genotype D infection is the most prevalent genotype in Saudi Arabia comprising an approximately 81.4% of HBV infections, followed by genotype E with 5.7%.²⁴ It has been reported

Sample	Gender	Age	HBsAg	HBcAb	HBsAb	HBV-DNA IU/mL	Genotype
D3	М	29	-	+	-	82	D
D4	М	28	-	+	-	83	D
D44	М	22	-	+	-	45	D
D47	F	36	-	+	-	< 20	D
D70	М	27	-	+	-	186	E
D71	F	35	-	+	-	66	D
D75	М	47	-	+	-	77	D
D76	М	28	-	+	-	133	D
D84	М	48	-	+	-	< 20	Е
D92	М	52	-	+	-	< 20	D
D93	М	27	-	+	-	51	D
D96	М	37	-	+	-	99	D
D100	М	40	-	+	-	< 20	D
D102	М	34	-	+	-	< 20	D
D107	М	21	-	+	-	54	D
D111	М	39	-	+	-	< 20	D
D135	М	23	-	+	-	27	D
*/ **	15M/2F*	33.7**	0*	17*	0*	82.1**	15D/2E*
Percentage	0.20/0.02	NA	0	0.20	0	NA	0.20/0.02

Table 1 - Demographic and serological profile of occult HBV infections.

that treatment of HBV genotype D infections in the absence of HBeAg with peginterferon alpha-2a for 72 weeks reduced the viral load to <400 copies/mL in only 23% of the patients.³³ Compared with this observation therapy with the nucleoside analogues such as entecavir, or tenofovir for 48 weeks has been shown to reduce the viral load to undetectable levels in 58.4-66% of the HBV infected patients with negative HBeAg.^{34,35} In the current study, the majority of OBIs identified (88.2%) were genotype D. However, the result obtained was not surprising since the vast majority of HBV infections in Saudi Arabia are genotype D.²⁴ Similar to our study, a study conducted on blood donors from Italy and Poland showed that the prevalence of HBV genotype D among OBIs was significantly higher than that of other genotypes.³⁶ Another recent report from Colombia concluded that HBV genotype F is the most prevalent OBI among blood donors.³⁷

In this study, the HBV-DNA detection assay was conducted only on samples from blood donors who tested positive for HBcAb. This is because HBcAb is considered a pathognomonic marker of HBV infection especially in the absence of HBsAg. It has been suggested that HBcAb screening has the advantage of excluding the majority of OBIs, leaving only a few cases of undetectable OBIs.³⁸ However, this approach is not practical in areas where the prevalence of HBcAb is more than 10%, because many donor rejections will negatively affect the blood supply. In the current study, we found that the prevalence of HBcAb in the Riyadh region was low (2.3%). Thus, the positive HBcAb blood units can either be directly discarded to cut costs, or screened for HBV-DNA and discarded only when positive for HBV-DNA, thus saving more blood units. By contrast, in other countries such as Korea, Greece, and Ghana where the prevalence of HBcAb is high (>10%),^{21,26,28} it would be mandatory to screen all blood units for HBV-DNA and discard only the positive samples in order to maintain adequate blood supply. Maximal prevention of HBV blood transmission can be achieved by developing tests that are more sensitive for HBsAg and HBcAb detection. Moreover, high-sensitivity HBV-DNA testing can be performed on blood donors either in individual samples using real-time PCR, which has a sensitivity of 12-20 IU/mL,16 or in minipool samples using nucleic acid testing (NAT), which has sensitivity of 3-8 IU/mL,³⁹ to exclude all blood units containing HBV-DNA.^{10,38}

Study limitation. One of the limitations of this study was that low numbers of OBIs were identified,

and thus we could not correlate the findings with the patients' demographic data.

In conclusion, all identified OBIs had a low viral load (<186 IU/mL) and mostly were genotype D. Thus, screening of blood and organ donors with more sensitive methods, including nucleic acid testing along with serological assays, especially in endemic areas, can potentially reduce the incidence of HBV blood transmission.

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