ABSTRACT

Three major hepatitis B virus (HBV) antigens include HBcAg, HBeAg and HBsAg. HBeAg is the extracellular form of HBcAg, and is seen almost exclusively in people who have circulating serum HBV DNA. Presence of HBsAg in serum indicates that the individual has contracted HBV infection. Chronic hepatitis HBeAg-negative/anti-HBe-positive is known as an important form of chronic hepatitis B in the Mediterranean region. In this report, we used Real-Time PCR and ELISA for detection of HBV and HBeAg/HBsAg, respectively. In our investigation on 4243 HBV cases referred to the Mahdieh Clinical Laboratory between 2008 and 2016, we found a 53-year-old man with clinical symptoms of hepatitis and abnormal molecular and serological features. Despite the presence of clinical symptoms and high viral load ($128 \times 10^5$ iu/ml), the patient was HBsAg-positive and HBeAg-negative. Identifying this type of HBV could indicate spread of this type of hepatitis in Isfahan, Iran.

Keywords: Hepatitis B, HBsAg, HBeAg.

Detection of Mediterranean Hepatitis B in a 45 Years Old Man in Mahdieh Clinical Laboratory, Isfahan, Iran

Majid Komijani (PhD)
Department of Biology, Faculty of Science, Arak University, Arak 38156-8-8349, Iran

Mohammad Taghi Kardi (PhD)
Department of Microbiology, Mahdieh Laboratory, Isfahan, Iran

Khashayar Shahin (MSc)
State Key Laboratory Cultivation Base of Most, Institute of Food Safety and Nutrition, Jiangusa Academy of Agricultural Sciences, Nanjing, China

Mahsa Yazdi (MSc)
Department of Biology, Faculty of Sciences, University of Isfahan, Isfahan, Iran

Corresponding author: Mohammad Taghi Kardi
E-mail: m.kardi@gmail.com
Tel: +989133094626
Address: Department of Microbiology, Mahdieh Laboratory, Isfahan, Iran

Received: 28 Jun 2017
Revised: 1 Nov 2017
Accepted: 04 Nov 2017
A mutation in the HBV precore region (G to A conversion at position 1896) leads to inhibition of HbeAg production. Moreover, the levels of viral replication, viral genotype and probably the genetic variability of the virus influence the expression of viral antigens [4-6]. In the early 1980s, a mutant form of CHB was reported with replication ability and negative HBeAg [7-9]. We hereby report for the first time, a case with anti-HBe-positive but HBeAg-negative CHB infection in Isfahan, Iran.

CASE REPORT

We examined 5800 suspected hepatitis B cases referred to the Mahdieh Clinical Laboratory in Isfahan (Iran) from 2008 to 2016. Diagnosis was made based on the number of HBV DNA copies and presence of HBsAg and HBeAg in the serum of individuals. Viral DNA was extracted using QIAamp DNA Mini Kit (Cat No: 51304, Qiagen, Germany). Real-Time PCR was performed using Rotor gene 6000 Corbett thermal cycler (Australia) and Artus HBV RG PCR Kit (Cat No: 4506265, Qiagen, Germany). ELISA was performed using Autobio kit (Cat No: E0315, China) to detect HBeAg and HBsAg.

According to the results, 4243 cases were HBV-positive. Among these cases, there was a 53-year-old man with primary clinical symptoms of hepatitis (vomiting, jaundice, tiredness and pain) and high viral load \((128 \times 10^5)\). Interestingly, the HBsAg and HBeAg of this case were positive and negative, respectively. To ensure the accuracy of the results, all experiments were repeated several times.
DISCUSSION

Approximately thirty years ago, CHB patients with replicating HBV were reported from the Mediterranean region, while having positive anti-HBe and negative HBeAg [10]. Molecular investigations on HBeAg-negative HBV led to the detection of CHB. To date, eight HBV genotypes (A–H) have been identified [2]. Genotype D has been associated with anti-HBe-positive CHB infection in the Mediterranean region [11]. HBeAg-negative CHB indicates that the patient was exposed to HBV strains that are not able to produce HBeAg [10]. In clinical practice, the term HBeAg-negative CHB is appropriate for patients with chronic HBV infection who are HBeAg-negative but usually anti-HBe-positive and have elevated serum HBV DNA levels. The most important and reliable methods of detecting this antigen is hybridization methods, while PCR is a more sensitive technique [3]. Therefore, choosing the most suitable and effective therapy relies on the detection of HBV DNA. Given the rising prevalence of CHB, it is recommended to inform laboratories and education systems about CHB cases similar to our case in order to maintain quality in detection and therapeutic approaches as well as monitoring the disease and its distribution. Studies have shown that the prevalence of this type of hepatitis is higher in some countries compared to Iran [10]. In addition, other studies in Iran have shown that this type of hepatitis is more common in Mashhad and Tehran (12, 13).

CONCLUSIONS

Considering the detection of this case in Isfahan, it seems that the prevalence of this type of hepatitis in Isfahan is increasing. Therefore, it is highly recommended to provide a comprehensive plan to prevent the spread of this type of hepatitis.

ACKNOWLEDGMENTS

The authors would like to thank staff of the Mahdieh Laboratory for their helpful and constructive comments.