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Author(s):	Muhammad Arif Khan Muhammad Waqar Zobaria Rehman Muhammad Idrees Muhammad Wasim Amina Asghar Qazi Shahrukh
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Research article

Hepatitis B virus (HBV) a Severe Health Problem in Mardan, Khyber Pakhtunkhwa: A Molecular Based Study

Muhammad Arif Khan¹, Muhammad Waqar²*, Zobaria Rehman², Muhammad Idrees³, Muhammad Wasim⁴, Amina Asghar⁵, Qazi Shah Rukh⁴

¹DHQ Hospital Mardan, Khyber Paktoonkhwa, Pakistan.

²PCR Section, Genome Centre for Molecular Based Diagnostics & Research Lahore.

³Centre for Applied Molecular Biology, University of the Punjab, Lahore- 53700, Pakistan

⁴Department of Medicine, Khyber Medical Collage Peshawar Khyber Paktoonkhwa, Pakistan

⁵Department of Zoology, University of Education, Lahore, Pakistan

*Corresponding author: Muhammad Waqar

PCR Section, Genome Centre for Molecular Based Diagnostics & Research Lahore, Pakistan, E-mail address: waqarkhan96@gmail.com;

Abstract

Hepatitis B is a severe health issue in both developed and developing countries that infects approximately 3.5 billion individuals globally. HBV infection is a more contagious disease than HCV and HIV infection. The current research work was conducted at D.H.Q hospital, Mardan Khyber Pakhtoon Khwa (KPK). Only HBsAg positive patients (n=750) were selected and for further confirmation of HBV DNA in all patients, Real Time PCR was performed. Out of a total of 750 patients, 34.6 % (n=260) patients were found to be HBV positive using Real Time PCR method. Prevalence of HBV DNA was high in male patients (p= 0.0151) as compared to female patients and also in patients of older age (>60 years). All patients were divided into three groups on the basis of viral load, that is, low viral load 100-10000 IU/ml, intermediate viral load 100,000-10,000,000 IU/ml, and high viral load >10,000,000 IU/ml. The study showed that 61.15% patients were placed in intermediate viral load category. All the HBV DNA positive patients were also screened for HBeAg and 31.53% were found reactive for HBeAg. This study provides a clear spectrum of molecular epidemiology of HBV in the Mardan region of KPK, Pakistan.

Keywords: GCMBDR, HBsAg, HBeAg, KPK

1. Introduction

Hepatitis B infection is a blood borne disease that is caused by partially double-stranded circular DNA virus (1) of the family hepadnaviridae (2). HBV has been reported to cause liver complications. Around 600,000 HBV related deaths occur globally (3), most of them are the result of the chronicity of HBV infection. Approximately 350 million people around the world are chronic carriers of Hepatitis B virus (HBV) (4), while 2 billion are infected with this virus (5-7). Complications like hepatocellular carcinoma, cirrhosis or chronic hepatitis cause deaths every year. Prevalence of HBV is less in developed countries like Australia, Eastern Europe and America, while it is widespread in Africa, Pacific Islands and Asia (8). It has been reported that more than 40 million people in Pakistan are infected with HBV and 12 million more are positive for hepatitis C virus (HCV).

Mucosal or percutaneous exposure to HBV, contaminated blood and/or body fluids can transmit HBV (9). It can be transmitted through body fluids such as saliva, semen, serum or blood. It can even survive in dried blood on razors, syringes, needles or on table surface (10). Common ways of transmitting HBV include blood transfusion (11), mother to child transmission, tattooing (12, 13), use of unsterilized surgical and dental instruments, reuse of disposable syringes and shaving blade by barber, reuse of needles for ear and nose piercing, sharing nail cutters, razors or

⁶Department of Zoology, Kohat University of Science and Technology, Kohat, Khyber Pakhtoonkhwa, Pakistan

toothbrushes belonging to infected people, sharing needles with drug addicts and unsafe sexual practices (14, 15).

Patterns of HBV vary worldwide. Areas that are known to be highly endemic for HBV infection harbor almost 45% of world population. In such areas, people acquire it during early childhood. About 43% population of the world lives in intermediate endemicity regions where multiple modes of HBV transmission described previously have been observed. In low endemicity regions, HBV infection has been observed in adolescents and adults most likely due to unsafe sex, injecting drugs or blood transfusion. HBV infection is highly endemic and remains a crucial health problem in Pakistan. Almost 9 million people infected with HBV (15, 16) while 3% are its chronic carriers (17, 18). National estimates related to HBV prevalence in Pakistan are not known but data from different selected groups or from different regions have been reported previously. Available data shows its prevalence as 32% anti HBV surface antibodies through natural conservation and 4% carrier rate with a total 38% prevalence (18). HBV assessed by Hepatitis B Surface Antigen (HBsAg) has been observed in 2%-14% blood donors (19, 20) and 7% health professionals (21). Moreover, 2.6% HBsAg positivity was observed in pre-employment screening of healthy individuals in northern Pakistan (22). Furthermore, 78% hepatocellular carcinoma patients and 30-42% chronic liver disease patients were positive for HBsAg (23, 24). Most of these epidemiological studies were restricted to only hospitalized patients. The clearance of HBsAg from the serum of chronic HBV patients have shown a strong association with clinical cure and improved survival. Recent studies have also shown that HBsAg can serve as an effective biomarker for treatment. Likewise, another study shows the significance of quantitated HBeAg and anti-HBe in analyzing the time course of HBsAg positive liver diseases. The serum level of HBeAg was found highest in chronic active hepatitis with lobular distortion and lowest in acute hepatitis in asymptomatic HBsAg carriers.

2. Materials and Methods

2.1. Description of the Study Area

The local population of Mardan, Khyber Pakhtoon Khwa (KPK) was selected for this study. The study was conducted from January 2014 to January 2015. KPK, formerly named as North-West Frontier Province (NWFP), is one of the four provinces of Pakistan with a population of 26.9 million. Peshawar is the largest city and the capital of KPK, followed by Mardan. The current study was conducted at the Department of Medicine, Mardan Medical Complex, Mardan, KPK. Patients from various remote areas of the province who visited Mardan Medical Complex were recruited for the study. Only HBsAg positive patients were enrolled. For further confirmation of HBV DNA by Real Time PCR, serum samples of the patients were sent to Genome Centre, Lahore Pakistan.

2.2. Blood Sample Collection

Samples were received from different collection centers of KPK. All samples were collected according to the standard procedures. Samples were stored at -20°C till further process. All patients were carriers of HBV and were found HBsAg positive by ICT chromatography and ELFA (Enzyme Linked Fluorescent Assay) method.

2.3. HBV DNA Isolation

HBV DNA was extracted from 150 μ l of each sample using Macherey-Nagal nucleic acid extraction kit (USA) which is designed for rapid extraction of highly pure viral nucleic acids from biological samples.

2.4. Real Time PCR

Qualitative and quantitative analysis of the samples was done using Smart Cycler II Real Time PCR (Cepheid, USA). Amplification was done using real time amplification kit prepared by Sacace Biotechnologies, Italy. Total reaction volume was 25 μ l, containing 12.5 μ l HBV amplification mix and 12.5 μ l extracted DNA. For precise checking of the reaction, negative and positive controls were also included with the run. Sensitivity of the assay was 20 copies per ml per blood sample. Specificity of the assay was about 99% (25).

2.5. Statistical Analysis

Statistix 9.0 software was used to analyze and summarize the data. Chi-square test was used for the analysis of categorical variables. A value less than 0.05 was considered as significant (26).

3. Results

A total of 750 HBsAg positive samples comprising 400 (53.33%) males and 350 (46.66%) females were tested for HBV DNA using Real Time PCR. Out of the total 400 male patients, 148 (37%) were found positive for HBV DNA by Real Time PCR, while in females HBV DNA positivity rate was 32% (n=112). Out of the total 750 HBsAg positive samples, HBV DNA was found positive in 260 patients (Table 1). All individuals were categorized into four groups according to their age, that is, 1 to 20 years, 21 to 40 years, 41 to 60 years and above 60 years, respectively. The prevalence of HBsAg was found higher in elderly population.

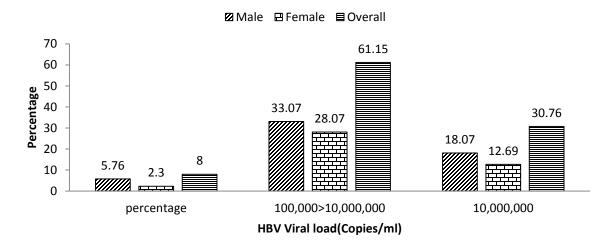
Sr. No.	Gender	Total HBsAg Positive patient	Total HBV DNA positive patients	Percentage	P-value
1	Male	400	148	37	< 0.05
2	Female	350	112	32	
		750	260	34.6	

Table 1. Gender-Wise Distribution of HBV

In the current study, patients were categorized into three groups on the basis of viral load, that is, low viral load (100-10000 cps/ml), intermediate viral load (100,000-10,000,000 cps/ml), and high viral load (>10,000,000 cps/ml), respectively. HBV viral load in both genders is shown in Figure 1. Among genders, 61.15% (n=159) were found with an intermediate viral load followed by a high viral load (n=80) at a rate of 30.76%, while 8.0% (n=21) had a low viral load. Our study also analyzed low, intermediate and high viral loads in various age groups. The current study showed that 54.23% (n=141) of the HBV DNA positive patients of different age groups had an intermediate viral load, followed by 30% of patients with a low viral load (n=78), while 15% of patients had a high viral load. In age group 1-20 years, 3.8% (n=10) had a low viral load, 5%

(n=13) had an intermediate viral load and 2.69% (n=07) had a high viral load. In patients of age group 21-40 years, 16.5% (n=43) had a low viral load, 38.84% (n=101) were carrying an intermediate viral load, while 5% (n=13) had a high viral load. In the age group 41-60, low viral load and intermediate viral load was observed in 8.07% (n=21) and 5.0% patients, respectively.

Figure 1. Distribution of viral loads among both genders of HBV positive patients.



Analysis of HBV in different age groups showed that the prevalence of HBV was higher in elderly population. Distribution of HBV in different age groups is shown in table 2.

S.No	Age	Total HBsAg Positive	Total HBV DNA positive Patients	Percentage	P-value
		Patients			
1	120	81	30	37.03	0.1208
2	2140	503	157	31.21	>0.05 Non- significant
3	4160	130	51	39.23	
4	>60	36	22	61.11	
Total		750	260	34.66	

Table 2. Prevalence of HBV in Various Age Groups

Figure 1 shows the analysis of HBV viral load in both genders. We observed that overall viral titre of HBV was higher in males as compared to females.

Both genders were categorized on the basis of viral load, that is, 20-10000 IU/ml, 100,000 to 10,000,000 IU/ml to 10,000,000 IU/ml, respectively. Viral load among different age groups is analyzed and presented in Figure 2. The majority of patients had an intermediate viral load that was 100,000 to 10,000,000 IU/ml.

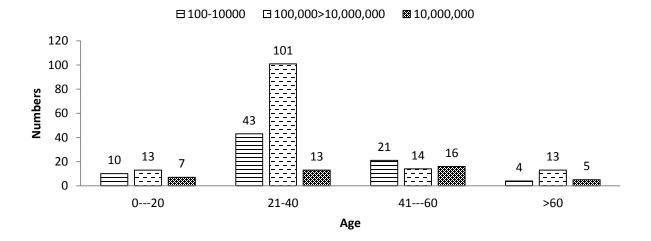


Figure 2. Distribution of HBV viral loads among different age groups.

Table 3 shows HBeAg reactivity in HBV DNA positive patients which shows that 32% of patients with a high viral load, that is >10,000,000, were found reactive for HBeAg. We observed that 82 (31.53%) were HBeAg reactive.

	-				
	HBV Viral load(Copies/ml)				
HBV markers					
	100-10000	100,00>10,000,000	>10,000,000	Total	
HBeAg Reactive	04(5.1%)	42(29.78%)	36(87.80%)	82(31.53%)	
HBeAg Non-Reactive	74(94.87%)	99(70.21%)	05(12.1%)	178 (68.46%)	

Table 3. Hbeag Positivity among HBV DNA Positive Patients

P=0.0000 < 0.05, Significant

4. Discussion

The current study analyzed the prevalence of HBV among the population of KPK. Out of the total 750 samples, 260 were detected for HBV DNA with a prevalence rate of 34.6%. The prevalence of HBV DNA was found higher in males as compared to females. This is in line with several other studies conducted in various areas of Pakistan as well as other countries which also showed that the prevalence of HBV is higher in males as compared to females (25, 27-30). For this study, the patients were categorized in various age groups that are 1-20 years, 21-40 years, 41-60 years, and > 60 years, respectively. High prevalence rate (61.11 %) was found in the age group > 60 years,

followed by age group 41-60 years with a prevalence rate of 39.23%. Patients of age group 21-40 years showed low prevalence rate at 31.21%, while patients of age group 1-20 years showed a prevalence rate of 37.30 %.

According to Munir et al. (2013), overall prevalence of HBV was found 3.0% in Mardan and it was higher in females (3.2%) as compared to males (2%) (31). Likewise, another study by Khan et al. showed the dominance of HBV over HCV in Mardan, KPK Pakistan (32).

A systematic analysis of HBV prevalence in Pakistan conducted in 2011 revealed that the percentage of HBV infection was 4.3318%, among healthcare persons (3.25%), among military recruits (4.276%), among pregnant women (5.872%), among healthy blood donors (3.93%), among patients with liver diseases (27.54% \pm 6.385%), among patients with cirrhosis (28.87%), among patients with hepatitis (15.896% \pm 14.824%), among patients with HCC (22% \pm 2.645%), among multiple transfused patients (6.223%), among users of injectable drugs (14.95%), among opthalmic patients (3.89%), among surgical patients (7.397%), among prisoners (5.75%), among opthalmic patients (3.89%), and among users of injectable drugs (14.95%) (33).

The next most important aspect of the current study is that all the HBV DNA positive patients were checked for HBeAg reactivity. About 88% of patients with a high viral load were found reactive for HBeAg, followed by patients with an intermediate viral load, that is, 42 (29.78%), while 04 (5.1%) of patients with a low viral load were found reactive for HBeAg.

5. Conclusion

It is concluded that HBV DNA positivity rate was higher in males as compared to females. It is also confirmed that HBV DNA positivity rate is high in age group >60 as compared to other age groups and patients with a high viral load were mostly reactive for HBeAg.

Competing interest

None.

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References

- 1. Barker LF, Shulman NR, Murray R, Hirschman RJ, Ratner F, Diefenbach WC, et al. Transmission of serum hepatitis. *JAMA*. 1970;211(9):1509–12.
- 2. Pungpapong S, Kim WR, Poterucha JJ, eds. Natural history of hepatitis B virus infection: An update for clinicians. *Mayo Clin Proc.* 2007;82(8):967–975.
- 3. Guirgis BS, Abbas RO, Azzazy HM. Hepatitis B virus genotyping: Current methods and clinical implications. *Int J Infect Dis*. 2010;14(11):e941–53.
- 4. Marín JAM, Anduiza CIA, Calderón GM, Rodríguez SC, Allen JLF, Flores RA, et al. Prevalence and resistance pattern of genotype G and H in chronic hepatitis B and HIV co-infected patients in Mexico. *Ann Hepatol.* 2012;11(1):47–51.

- 5. Ali H, Memon M. Prevalence of hepatitis B infection in pregnant women in a tertiary care hospital Karachi. *Infect Dis J Pak.* 2007;16(2):35–38.
- 6. Zhu C-T, Dong C-L. Characteristics of general distribution of hepatitis B virus genotypes in China. *Hepatobiliary Pancreatic Dis Int*. 2009;8(4):397–401.
- 7. Li G, Li W, Guo F, Xu S, Zhao N, Chen S, et al. A novel real-time PCR assay for determination of viral loads in person infected with hepatitis B virus. *J Virol Methods*. 2010;165(1):9–14.
- 8. Torbenson M, Thomas DL. Occult hepatitis B. Lancet Infect Dis. 2002;2(8):479-86.
- 9. Shepard CW, Simard EP, Finelli L, Fiore AE, Bell BP. Hepatitis B virus infection: Epidemiology and vaccination. *Epidemiol Rev.* 2006;28(1):112–25.
- Workowski KA, Berman SM. CDC sexually transmitted diseases treatment guidelines. *Clin Infect Dis.* 2002;35(Supplement-2):S135–7.
- 11. Francisci D, Antonelli S, Preziosi R, Mecozzi F, Stagni G, Pauluzzi S. Risk factors for acute parenterally transmitted viral hepatitis: A 20-year study. *Eur J Epidemiol*. 1993;9(6):625–8.
- 12. Gibb D, Goodall R, Dunn D, Healy M, Neave P, Cafferkey M, et al. Mother-to-child transmission of hepatitis C virus: Evidence for preventable peripartum transmission. *Lancet*. 2000;356(9233):904–7.
- 13. Wahid B, Saleem K, Rasool N, Rafique S, Ali A, Waqar M, et al. Tattooing trend: Major cause of HCV transmission among youngsters. *Int J Infect Dis.* 2018:1–3.
- 14. Haley RW, Fischer RP. Commercial tattooing as a potentially important source of hepatitis C infection: Clinical epidemiology of 626 consecutive patients unaware of their hepatitis C serologic status. *Medicine*. 2001;80(2):134–51.
- 15. Hakim S, Kazmi S, Bagasra O. Seroprevalence of hepatitis B and C genotypes among young apparently healthy females of Karachi, Pakistan. *Libyan J Med*. 2008;3(2):66–70.
- 16. Hakeem K, Khan MS, Abdullah M, Rehman MA, Hashmi MI. Prevalence of HBs Ag and Anti HCV in pregnant ladies attending antenatal clinic at Shaikh Zayed Medical Complex, Rahim Yar Khan. *Esculapio J Service Inst Med Sci.* 2006;2(3):6–8.
- 17. Noorali S, Hakim ST, McLean D, Kazmi SU, Bagasra O. Prevalence of hepatitis B virus genotype D in females in Karachi, Pakistan. *J Infect Dev Countries*. 2008;2(05):373–8.
- 18. Khan F, Shams S, Qureshi ID, Israr M, Khan H, Sarwar MT, et al. Hepatitis B virus infection among different sex and age groups in Pakistani Punjab. *Virol J*. 2011;8(1):225.
- 19. Chaudhry NT, Khan SJ, Khan TA, Saeed M, Syed M, Iqbal J, et al. Prevalence of hepatitis B carriers and blood group frequency in voluntary blood donors. *J Ayub Med Coll*. 1996;8(2):29–32.
- 20. Rehman K, Khan A, Haider Z, Shahzad A, Iqbal J, Khan R, et al. Prevalence of sero markers of HBV and HCV in health care personnel and apparently healthy blood donors. *JPMA*. 1996;46(7):152–4.
- 21. Sheikh M, Shamsh K. Prevalence of HBV markers in health care personals Vs matched control. *J Coll Phys Surg Pak.* 1995;5:19–21.
- 22. Khokhar N, Gill ML, Malik GJ. General sero prevalence of hepatitis C and hepatitis B virus infections in population. Journal of the College of Physicians and Surgeons--Pakistan: *J Coll Phys Surg Pak*. 2004;14(9):534–6.

- 23. Tong C, Khan R, Beeching N, Tariq W, Hart C, Ahmad N, et al. The occurrence of hepatitis B and C viruses in Pakistani patients with chronic liver disease and heaptocellula carcinoma. *Epidemiol Infect*. 1996;117(2):327–32.
- 24. Khan TS, Rizvi F, Rashid A. Hepatitis C sero positivity among chronic liver disease patients in Hazara, Pakistan. *J Ayub Med Coll*. 2003;15(2):53–5.
- 25. Naz Z, Wahid B, Usman S, Saleem K, Rafique S, Ali A, et al. Expression of SOCS1 and SOCS3 genes in interferon-treated and direct-acting antiviral drugs-treated hepatitis C patients. *J Interferon Cytokine Res.* 2018;38(6):255–60.
- 26. Wahid B, Waqar M, Rasool N, Wasim M, Khalid I, Idrees M. Prevalence of thyroid stimulating hormone dysfunction among sofosbuvir- treated HCV infected patients: A real- world clinical experience. *J Med Virol*. 2019;91(3)514–517.
- 27. Shazi L, Abbas Z. Comparison of risk factors for hepatitis B and C in patients visiting a gastroenterology clinic. *J Coll Phys Surg Pak*. 2006;16(2):104–7.
- 28. Mahtab M-A, Rahman S, Karim MF, Khan M, Foster G, Solaiman S, et al. Epidemiology of hepatitis B virus in Bangladeshi general population. *Hepatobiliary Pancreatic Dis Int.* 2008;7(6):595–600.
- 29. Wahid B, Waqar M, Rasool N, Rehman Z, Saeed J, Wasim M, et al. Recent trends in molecular epidemiology of Hepatitis C virus in Mardan, KPK, Pakistan. *Infect Genet Evol*. 2018;66:66–71.
- 30. Bhatti S, Quraishi M, Mahmood C, Javaid K. Sero prevalence of HBs Ag and HCV antibodies in healthy individuals of high socio-economic status. *Biomedica*. 2007;23:131–3.
- Munir M, Shams S, Lodhi MA, Parveen Z, Ullah N. Prevalence of hepatitis B in the students, and employees of Abdul Wali Khan University, Mardan Shankar Campus. Pakhtunkhwa. J Life Sci. 2013;3:120–9.
- 32. Khan AJ, Luby SP, Fikree F, Karim A, Obaid S, Dellawala S, et al. Unsafe injections and the transmission of hepatitis B and C in a pre-urban community in Pakistan. *Bull W H O*. 2000;78:956–63.
- 33. Ali M, Idrees M, Ali L, Hussain A, Rehman IU, Saleem S, et al. Hepatitis B virus in Pakistan: A systematic review of prevalence, risk factors, awareness status and genotypes. *Virol J*. 2011;8(1):102.