

Original Article

Seroprevalence of Human Immunodeficiency Virus (HIV) and Hepatitis C Infection in Hemophilic Patients in Iran

Mohssen Nassiri Toosi¹, Manije Lak², Katayoun Karimi², Mohammadreza Managhchi²,
Katayoun Samimi-rad³, Alireza Abdollahi⁴, Reza Shahsiah⁴

1. Dept. of Internal Medicine, Tehran University of Medical Sciences, Tehran, Iran.

2. Iranian Hemophilia Center, Tehran, Iran.

3. Dept. of Virology, Tehran University of Medical Sciences, Tehran, Iran.

4. Dept. of Pathology, Tehran University of Medical Sciences, Tehran, Iran.

ABSTRACT

Background and Objective: Although transfusion therapy has lead to great improvement in longevity for hemophiliacs, but there have been tragic setbacks especially from transmission of human immunodeficiency virus (HIV) and hepatitis C virus (HCV), HIV was reported to increase the rate of HCV-related liver failure by 4.2 times. In this study, we aimed to determine the seroprevalence of HIV and HCV, the association of HCV with abnormal liver tests, impact of HIV on HCV-related abnormalities and the distribution of HCV genotypes in Iranian hemophiliacs.

Patients and Methods: In a cross-sectional study, we determined virological, clinical and epidemiological characteristics for HIV and HCV infection of 236 hemophiliacs attending our center. Data were analyzed using Chi-square test.

Results: Ten (4.7%) out of 211 patients tested were HIV seropositive and 145 (83.3%) were HCV seropositive. All tested positive HIV patients also had HCV. HCV seroprevalence was significantly higher in patients with hemophilia A and B as compared to other congenital coagulopathies and it was directly related to coagulation severity. HCV seroprevalence was lower in hemophiliacs with positive HBsAg ($p = 0.03$) but it did not differ by HBcAb or HBsAb results. HCV genotype 1a (48.5%) was predominant type and genotype 3a (33.3%) was also common. Frequency of abnormal aspartate aminotransferase and alanine aminotransferase liver enzymes was significantly higher in the HCV positive group ($p = 0.006$).

Conclusion: This study provides evidence that hepatitis c virus infection is a major problem for Iranian hemophiliacs and it has higher prevalence in hemophiliacs with higher age, more severe coagulopathies, abnormal alanine aminotransferase level, and human immunodeficiency virus co-infection.

Key words: Hepatitis C, Hemophilia, Human Immunodeficiency Virus

Received: 8 February 2008

Accepted: 17 March 2008

Address communications to: Dr. Alireza Abdollahi, Department of Pathology, Tehran University of Medical Sciences, Tehran, Iran.

Email: dr_p_abdollahi@yahoo.com

Introduction

The 1940s in the history of hemophilia treatment was the transitional time from primitive, conventional attempts to control bleeding episodes to transfusion therapy as a major accomplishment (1;2). Even though the discovery of cryoprecipitate and factor concentrates and lyophilized concentrates lead to great improvement in both longevity and quality of life for persons with hemophilia, but at the same time a group of tragic setbacks were reported (2). These significant negative consequences were complications resulting from transmission of hepatitis B virus (HBV), hepatitis C virus (hepatitis c virus), and human immunodeficiency virus (human immunodeficiency virus). It was discovered that virtually all hemophilia patients exposed to non-heat-treated factor were hepatitis c virus positive, and over 50% of hemophilia patients in the United States had human immunodeficiency virus seroconvert, and 5-10% of such patients became chronic carriers of hepatitis B (3).

In general, the natural history of hepatitis c virus reveals a slowly progressive disease leading to cirrhosis in approximately 20% of patients after 20 years. The key risk factors that increase the rate of progression of hepatitis c virus are age at infection >40 yrs, alcohol, male gender, obesity, high grade of liver inflammation, hepatic steatosis, immunosuppression, smoking, and co-infection with human immunodeficiency virus and HBV (4-7).

Iranian Hemophilia Center was established as a main health care system for Iranian hemophilia population in 1965. This reference center provided a good access of hemophilia patients and their family for early diagnosis and treatment. The programs for control of HBV and hepatitis c virus infection in Iran was started in 1993 by routine HBV vaccination and followed by screening for hepatitis c virus in blood products in 1997 (1;4).

In this study, we aimed to determine the prevalence of viral infections, specifically hepatitis c virus, HBV, and human immunodeficiency virus, their contribution to liver tests, the impact of human immunodeficiency virus on hepatitis c virus infection, and the genotypes of hepatitis c virus in Iranian hemophiliacs.

Materials and Methods

In a cross-sectional and prospective study, we enrolled all patients with hemophilia attending the Iranian Hemophilia Center through the year 2003 to determine their virological, clinical and epidemiological characteristics for chronic viral infections. A complete virological, biochemical and epidemiological assessment was done for all hemophiliacs registered at the center. They were tested for anti-hepatitis c virus-ELISA-3, HbsAg, HbsAb, HbcAb and human immunodeficiency virus serology. Also, biochemical liver-related tests including CBC, platelet, liver enzymes (ALT, AST), PT, and albumin were taken. The patients with positive anti-hepatitis c virus were tested for hepatitis c virus-RNA in order to obtain serological confirmation. Genotype was also determined.

Collected clinical information included age, gender, type of hemophilia (A or B or other), severity of hemophilia (classified as severe if factor VIII or IX blood levels were less than 1%, moderate if they were between 1 and 5% and mild if they were greater than 5%). In this study, ALT status was classified normal if ALT levels were below 40 IU/L, defined as the upper limit of normal (ULN). Data were analyzed with Chi-square test.

Results

In total, 236 patients completed the registration. In this population, 73% had hemophilia A, 10% hemophilia B, and 17% other types of congenital coagulopathies like platelet disorders and factor V deficiency. Most of the patients were in the third decade of life. The mean \pm SD of their age was 26.6 ± 12.1 years (Table 1).

On virological examination, more than 80% of patients were anti-hepatitis c virus (ELISA) positive. The frequency of hepatitis c virus-RNA was 73%. Forty four percent of patients with positive HBcAb revealed contact with hepatitis B virus. In this population, only 3% of patients were chronic carrier for hepatitis B virus. About 80% of patients had HbsAb in their blood and showed natural or vaccine immunity to HBV. Only less than 5% of patients were human immunodeficiency virus seroconverted.

No child under the age of 10 was anti-hepatitis c virus, HbsAg, or anti-human immunodeficiency virus (ELISA) positive, in contrast the frequency of these

Tabel-1:Demography of 236 Patients With Hemophilia

	Hemophilia A	Hemophilia B	Others*	Total
Gender				
Female	2	1	18	21 (8.9%)
Male	169	24	22	215 (91.1%)
Total	171 (72.5%)	25 (10.6%)	40 (16.9%)	236 (100%)
Age				
Range (Yr)	4.0-60	6.0-69	9.0-58	4.0-69
Mean±SD	26.1±11.1	32.5±15.9	25.5±12.5	26.6±12.1
0-9	2	2	1	5
10.0-19	47	2	16	65
20-29	70	9	13	92
30-39	29	6	4	39
40-49	15	2	2	19
50-59	7	1	4	12
60 and over	1	3	0	4
Severity				
Severe	128 (74.9%)	16 (64.0%)	(-)	144 (73.5%)
Moderate	24 (14.0%)	3 (12.0%)	(-)	27 (13.8%)
Mild	19 (11.1%)	6 (24.0%)	(-)	25 (12.8%)

* Others= Platelate disorders, Factor V deficiency

infections among older subjects were higher. The age distribution of positive anti-hepatitis c virus and HBcAb differs statistically among the hemophiliacs.

There was no prominent difference in anti-hepatitis c virus (ELISA) positivity between male and female hemophiliacs. The anti-hepatitis c virus (ELISA) positivity significantly increased in hemophilia A and B as compared to other congenital coagulopathies (platelate disorders and factor V deficiency).

We did anti-hepatitis c virus (ELISA) test in 8 patients with positive human immunodeficiency virus serology. All of them were anti-hepatitis c virus positive (100%). This relationship due to low number of patients in this group was not statistically significant. There was a lower frequency of anti-hepatitis c virus (ELISA) positive test in hemophilia patients with HBsAg positivity. The anti-hepatitis c virus test (ELISA) was positive in 2 out of 5 HBsAg positive patients with hemophilia in contrast to 134 out of 160 HBsAg negative hemophilics. ($p = 0.03$). We did not find any significant difference in anti-hepatitis c virus (ELISA) positivity by HBcAb (contact to HBV) or HBsAb (immune to HBV) positive and negative.

Hepatitis c virus RNA test by reverse transcriptase-polymerase chain reaction (RT-PCR) was positive in 80.2% of anti-hepatitis c virus (ELISA) positive hemophilia patients. It means that viremia was found out in 80% of anti-hepatitis c virus (ELISA) positive patients. Non-viremic infection (liver cell or mononuclear cell sequestration), transient absence (or fall) of viremia, recovered infection, viremia below limit of assay, and false positive (or non-specific) ELISA results are possible explanations for the rest of them.

In a study based on the results of PCR positivity, hepatitis c virus genotyping was carried out using genotype specific probes and the dot blot hybridization assay. Genotype 1a (48.5%) was predominant type, and genotype 3a (33.3%) was common. Four patients were found infected with type 1b (12.1%), 1 patient with type 2b (3%), and one patient with type 4b (3%). The double infection with two hepatitis c virus types was not tried in this study.

In registration of laboratory findings in 236 hemophiliacs, frequency of abnormal AST liver enzyme was significantly higher in anti-hepatitis c

virus (ELISA) positive group. ($p = 0.015$). There was no difference in frequency of abnormal AST between human immunodeficiency virus positive and negative groups. The frequency of abnormal ALT liver enzyme was significantly higher in anti-hepatitis c virus (ELISA) positive group. ($p = 0.006$). There was also no difference in frequency of abnormal ALT between human immunodeficiency virus positive and negative groups.

Patients with hepatitis c virus and human immunodeficiency virus seropositive status had a higher ratio of prolonged prothrombin time in respect of hepatitis c virus or human immunodeficiency virus negative patients ($p =$ non-significant). Patients with factor V deficiency were excluded in this analysis. But patients with anti-hepatitis c virus (ELISA) and human immunodeficiency virus positive tests had a higher frequency of platelet lower than 150000/ μ l. In analysis of serum albumin results among hemophilic patients, there was only a significant difference between human immunodeficiency virus positive and negative groups ($p = 0.008$). The human immunodeficiency virus seropositive patients had a lower serum albumin level.

Discussion

It should be noted that Iranian Hemophilia Center started to diagnose and treat hemophilic population (1965) before inactivation of blood products for transfusion-transmitted viral infections (8). In this center, hemophiliacs were treated with imported lyophilized concentrates and locally produced cryoprecipitate and fresh frozen plasma without viral inactivation treatment before 1997. This could lead to early and ready exposure of this population to contaminated clotting factor products. The high prevalence of hepatitis c virus infection observed in this hemophilic population (80%) indicates that hepatitis C largely contributes to the morbidity of persons with hemophilia in Iran. This high prevalence is comparable to developed countries (8-10) with a good access of patients to products and the widespread use of clotting factor concentrates in replacement therapy before the era of inactivated products.

Similar to the general hemophilic population, hemophilia A was the predominant type (73%). The risk of infection with each of these viruses is directly related to the severity of the hemophilia. The predominance of severe hemophilia was expected in a population infected by hepatitis c virus with such a

high exposure. Individuals with more severe disease bleed more frequently and require more clotting factor, thus increasing their risk of blood-borne infections. This shows that the likelihood of infection increases with bleeding frequency and amount of factor used, indicating that these are the key risk factors of interest.

Another important problem is multiple infections, especially the combination of hepatitis C and human immunodeficiency virus. As expected, given their similar routes of transmission, such co-infection is common among persons with hemophilia (9;11-13). We detected a high proportion of past exposure to HBV (44%), with 3% of positive HBsAg, and important frequency of co-infection with human immunodeficiency virus (5%). This shows higher exposure of this population to transfusion-transmitted viral infections as compared to Iranian general population. We registered lower prevalence of human immunodeficiency virus infection in our hemophilic population as compared to developed countries (13;14). Based on our anecdotal observations in our center, lower survival of human immunodeficiency virus positive patients could affect this lower number in our registry. However, co-infection especially with human immunodeficiency virus is growing in importance because these patients are at a higher risk of progression to chronic liver disease than those infected with hepatitis c virus alone (2;14;15). This is illustrated by our surveillance data, which indicated that among human immunodeficiency virus infected subjects, 100% were hepatitis c virus positive and these patients had a lower serum albumin level and a higher ratio of prolonged prothrombin time.

Children who were born after 1993 have no evidence of hepatitis c virus, HBV, and human immunodeficiency virus infection. Others have observed this decline in risk as well (16-18). Although the occurrence of new infections with HBV, hepatitis c virus, and human immunodeficiency virus has substantially decreased in recent years, chronic liver disease by these viruses continues to be of significant concern to patients with hemophilia. The impact of human immunodeficiency virus infection has been devastating and despite recent advances in treatment, it is likely to be felt for some time into the future.

The observation that most anti-hepatitis c virus-positive hemophiliacs (57%) did not show abnormal ALT levels at the time of study was interesting, demonstrating the lack of a relationship between the degree of abnormality of serum transaminase levels

and seropositivity for hepatitis c virus. Also, in this regard absence of a history of alcoholism in our patients needs to be considered.

The association between hepatitis c virus viremia and raised ALT levels ($p = 0.006$) was also reported in hemophiliac populations by other authors (2;19). This information can be useful for clinicians who treat positive anti-hepatitis c virus hemophiliacs but do not have any molecular method (i.e. PCR) available to determine the persistence of hepatitis c virus infection. Raised ALT can be a clue of hepatitis c virus viremia, an important piece of information for hemophiliacs, for whom there has been an obvious reluctance to perform liver biopsies due to the risk of bleeding complications (20-22). The protective effect of chronic HBV infection (positive HBsAg) found in this study ($p = 0.03$) has also been described by other authors (23-25). It has been attributed to viral interference between HBV and hepatitis c virus, with the mechanism still not elucidated.

The distribution of hepatitis c virus genotypes in the studied population did not significantly differ from the published data on hemophilic and non-hemophilic populations in Iran, where genotype 1a infection is predominant (26-28). Actually, the distribution of hepatitis c virus genotype is similar with England, European countries and USA, where usually reflects the origin of the blood donors used in the manufacture of imported pooled factor VIII and IX concentrates (29;30). In Iran, the hepatitis c virus genotype distribution would be more a reflex of indigenous and some intermixing of strains with strains from other parts of the world (23).

Conclusion

In summary, this study provides evidence that hepatitis c virus infection is a major problem for Iranian hemophiliacs and it has a higher prevalence in patients with higher age, more severe coagulopathies, abnormal ALT level, and uman immunodeficiency virus co-infection. Transmission of viruses via blood products has been a significant source of morbidity for persons with hemophilia. Fortunately, a number of effective strategies including viral inactivation procedures have been implemented in order to prevent such transmission in the future. The most recent innovation in hemophilia treatment has been the introduction of recombinant factor therapy, which holds great promise for safety as well as efficacy.

References

1. Remesar M, Gamba C, Kuperman S, Marcosa MA, Miguez G, Caldarola S, et al. Antibodies to hepatitis C and other viral markers in multi-transfused patients from Argentina. *J Clin Virol* 2005 Dec;34 Suppl 2:S20-6.:S20-S26.
2. Telfer P, Sabin C, Devereux H, Scott F, Dusheiko G, Lee C. The progression of HCV-associated liver disease in a cohort of haemophilic patients. *Br J Haematol* 1994 Jul;87(3):555-61.
3. Chambost H, Gerolami V, Halfon P, Thuret I, Michel G, Sicardi F, et al. Persistent hepatitis C virus RNA replication in haemophiliacs: role of co-infection with human immunodeficiency virus. *Br J Haematol* 1995 Nov;91(3):703-7.
4. Moylan CA, Muir AJ. Treatment of hepatitis C in special populations. *Clin Liver Dis* 2005 Nov;9(4):567-77.
5. Fattovich G, Giustina G, Degos F, Tremolada F, Diodati G, Almasio P, et al. Morbidity and mortality in compensated cirrhosis type C: a retrospective follow-up study of 384 patients. *Gastroenterology* 1997 Feb;112(2):463-72.
6. Darby SC, Ewart DW, Giangrande PL, Spooner RJ, Rizza CR, Dusheiko GM, et al. Mortality from liver cancer and liver disease in haemophilic men and boys in UK given blood products contaminated with hepatitis C. UK Haemophilia Centre Directors' Organisation. *Lancet* 1997 Nov 15;350(9089):1425-31.
7. Lefkowitz JB, Monroe DM, Kasper CK, Roberts HR. Comparison of the behavior of normal factor IX and the factor IX Bm variant Hilo in the prothrombin time test using tissue factors from bovine, human, and rabbit sources. *Am J Hematol* 1993 Jul;43(3):177-82.
8. Giangrande PL. Hepatitis in haemophilia. *Br J Haematol* 1998 Oct;103(1):1-9.
9. Brettler DB, Alter HJ, Dienstag JL, Forsberg AD, Levine PH. Prevalence of hepatitis C virus antibody in a cohort of hemophilia patients. *Blood* 1990 Jul 1;76(1):254-6.
10. Troisi CL, Hollinger FB, Hoots WK, Contant C, Gill J, Ragni M, et al. A multicenter study of viral hepatitis in a United States hemophilic population. *Blood* 1993 Jan 15;81(2):412-8.
11. Ghany MG, Leissinger C, Lagier R, Sanchez-Pescador R, Lok AS. Effect of human immunodeficiency virus infection on hepatitis C virus infection in hemophiliacs. *Dig Dis Sci* 1996 Jun;41(6):1265-72.

12. Eyster ME, Fried MW, Di Bisceglie AM, Goedert JJ. Increasing hepatitis C virus RNA levels in hemophiliacs: relationship to human immunodeficiency virus infection and liver disease. Multicenter Hemophilia Cohort Study. *Blood* 1994 Aug 15;84(4):1020-3.
13. Hill HA, Stein SF. Viral infections among patients with hemophilia in the state of Georgia. *Am J Hematol* 1998 Sep;59(1):36-41.
14. Eyster ME, Diamondstone LS, Lien JM, Ehmann WC, Quan S, Goedert JJ. Natural history of hepatitis C virus infection in multitransfused hemophiliacs: effect of coinfection with human immunodeficiency virus. The Multicenter Hemophilia Cohort Study. *J Acquir Immune Defic Syndr* 1993 Jun;6(6):602-10.
15. Sabin CA, Telfer P, Phillips AN, Bhagani S, Lee CA. The association between hepatitis C virus genotype and human immunodeficiency virus disease progression in a cohort of hemophilic men. *J Infect Dis* 1997 Jan;175(1):164-8.
16. Morfini M, Mannucci PM, Ciavarella N, Schiavoni M, Gringeri A, Rafanelli D, et al. Prevalence of infection with the hepatitis C virus among Italian hemophiliacs before and after the introduction of virally inactivated clotting factor concentrates: a retrospective evaluation. *Vox Sang* 1994;67(2):178-82.
17. Pistello M, Ceccherini-Nelli L, Cecconi N, Bendinelli M, Panicucci F. Hepatitis C virus seroprevalence in Italian haemophiliacs injected with virus-inactivated concentrates: five year follow-up and correlation with antibodies to other viruses. *J Med Virol* 1991 Jan;33(1):43-6.
18. Schramm W, Roggendorf M, Rommel F, Kammerer R, Pohlmann H, Rasshofer R, et al. Prevalence of antibodies to hepatitis C virus (HCV) in haemophiliacs. *Blut* 1989 Oct;59(4):390-2.
19. Mauser-Bunschoten EP, Bresters D, van Drimmelen AA, Roosendaal G, Cuyper HT, Reesink HW, et al. Hepatitis C infection and viremia in Dutch hemophilia patients. *J Med Virol* 1995 Mar;45(3):241-6.
20. Aledort LM, Levine PH, Hilgartner M, Blatt P, Spero JA, Goldberg JD, et al. A study of liver biopsies and liver disease among hemophiliacs. *Blood* 1985 Aug;66(2):367-72.
21. Hanley JP, Dolan G, Day S, Skidmore SJ, Irving WL. Interaction of hepatitis B and hepatitis C infection in haemophilia. *Br J Haematol* 1993 Nov;85(3):611-2.
22. Mimms LT, Mosley JW, Hollinger FB, Aach RD, Stevens CE, Cunningham M, et al. Effect of concurrent acute infection with hepatitis C virus on acute hepatitis B virus infection. *BMJ* 1993 Oct 30;307(6912):1095-7.
23. Samimi-Rad K, Nategh R, Malekzadeh R, Norder H, Magnus L. Molecular epidemiology of hepatitis C virus in Iran as reflected by phylogenetic analysis of the NS5B region. *J Med Virol* 2004 Oct;74(2):246-52.
24. Mellor J, Holmes EC, Jarvis LM, Yap PL, Simmonds P. Investigation of the pattern of hepatitis C virus sequence diversity in different geographical regions: implications for virus classification. The International HCV Collaborative Study Group. *J Gen Virol* 1995 Oct;76(Pt 10):2493-507.
25. McCaw R, Moaven L, Locarnini SA, Bowden DS. Hepatitis C virus genotypes in Australia. *J Viral Hepat* 1997 Sep;4(5):351-7.
26. Harris KA, Gilham C, Mortimer PP, Teo CG. The most prevalent hepatitis C virus genotypes in England and Wales are 3a and 1a. *J Med Virol* 1999 Jun;58(2):127-31.
27. Vinelli E, Lorenzana I. Transfusion-transmitted infections in multi-transfused patients in Honduras. *J Clin Virol* 2005 Dec;34 Suppl 2:S53-60.:S53-S60.
28. Lichterfeld M, Schmeisser N, Qurishi N, Vogel M, Brackmann HH, Spengler U, et al. Clinical outcomes of HIV-HCV co-infection in a large cohort of hemophiliac patients. *J Infect* 2005 Apr;50(3):221-8.
29. Shen F, Huang Q, Sun HQ, Ghildyal R. Significance of blood analysis in hemophiliacs co-infected with human immunodeficiency virus and hepatitis viruses. *World J Gastroenterol* 2007 Mar 28;13(12):1862-6.
30. Koziel MJ, Peters MG. Viral hepatitis in HIV infection. *N Engl J Med* 2007 Apr 5;356(14):1445-54.