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Hepatitis B and C Viruses incidence, the Risk Factors of Hepatocellular Carcinoma, is low in Aseer Region, Saudi Arabia

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ABSTRACT

Aim: Risks of hepatocellular carcinoma (HCC) following hepatitis B (HBV) and/or hepatitis C virus (HCV) infection are well known. The aim of this study was to determine the prevalence and risk factors of hepatitis B and C virus infections in blood donors at Aseer Region, Saudi Arabia

Methods: The study was conducted on random blood samples collected from healthy blood donor volunteers, who were referred to Blood Transfusion Centers found at Aseer region, during the period March 2012 to February 2013. All the collected blood units were screened for hepatitis B surface antigen (HBsAg), anti-HBc, HCV, human immunodeficiency virus (HIV) 1 and 2, human T-cell lymphotropic virus (HTLV) I/II, venereal disease research laboratory (VDRL) and malaria. All donated blood were checked for HBV-DNA, HCV-RNA and HIV-RNA by nucleic acid test (NAT) technology.

Results: Of 7267 (26 (0.36%) females and 7241 (99.64) males) blood donors screened, with median age of 28 (female) and 30 years (males), 71 (0.98%) were HBsAg positive of them 66 (0.91) were positive to HBV-DNA, 449 (6.18%) were anti-HBc positive of them 78 (1.07%) were positive to HBV-DNA. Cases positive to both HBsAg and HBc-Ab were 69 (0.95%) all of them were positive to HBV-DNA. There were 5 (0.069%) cases positive for HCV, none of them showed mixed infection with HVB. All positive cases for HBsAg, HBc-Ab and HCV were shown to be among male volunteers.

Conclusion: Prevalence of HCV in Asser region is very low. In the time that HBsAg is low, prevalence of HBcAb is relatively moderate. Expected incidence of heptatocellular carcinoma due to infection with HCV is much lower than HBV.

Key words: Blood donors, HBV, HBsAg, HCV, HCC, CLD, Saudi Arabia

INTRODUCTION

Chronic liver disease is responsible for over 1.4 million deaths annually and is characterized by permanent inflammatory processes that predispose to liver cancer and in particular hepatocellular carcinoma (HCC) (Ayele and Gebre-Selassie, 2013). In healthy liver, inflammatory processes stimulate growth and repair and restore normal liver architecture. Chronic liver disease

(CLD) results from an inflammatory injury to the liver, which has persisted for six or more months without complete resolution. Chronic liver disease comprises of a spectrum of disease such as chronic hepatitis, liver cirrhosis, and HCC (Laraba *et al.*, 2010). In CLD, the balance of damage versus regeneration in the liver is disrupted and can lead to the formation of excessive scar tissue, termed fibrosis. In the long-term, an

exacerbation of fibrosis will lead to cirrhosis, which is characterized by abnormal liver architecture and function and is associated with a significant reduction in overall health wellbeing. At cirrhotic stages, liver damage is often irreversible or difficult to treat. Cirrhosis leads frequently to death from liver failure or to HCC (Bartosch, 2010). Indeed, HCC is the first cause of death in cirrhotic patients (Parkin et al., 2001), and is a tumor with poor prognosis, ranking third in terms of death by cancer. Furthermore, it is the fifth most prevalent cancer worldwide, with 800,000 new cases per year in the world (Parkin et al., 2001, El-Serag, 2007).

Hepatitis B virus (HBV) and hepatitis C virus (HCV) infections are primarily hepatropic. Chronic infection with these viruses causes progressive liver disease and HCC (Beasley et al., 1981; Simonetti et al., 1989; Colombo et al., 1989; Bruix et al., 1989; Lauer and Walker, 2001; Ferrari et al., 2003), and HBV/HCV co-infection further increases the risk of HCC (Shi et al., 2005). Also HBV and HCV have been shown to be lymphotropic (Pasquinelli et al., Ferri 1993; et al., Pawlotsky 1995). Some studies have examined the role of infections in non-Hodgkin's these lymphoma (NHL) but the findings were inconclusive (Kuniyoshi et al., 2001; Negri et al., 2001). HCV infection has been shown to be associated with other malignant diseases such as multiple myeloma and thyroid cancer (Montella et al., 2001).

Hepatocellular carcinoma is the sixth most common cancer and the third most common cause of mortality from cancer worldwide (Parkin et al., 2002). **Hepatitis** В virus and **HCV** monoinfections are well-known major risk factors for HCC, and the relative importance varies worldwide changes over time (Lu et al., 2006). Because of their shared modes of

transmission, coinfection of HBV and HCV is not uncommon, particularly in countries with a high prevalence of HBV or HCV. Hepatitis B virus and HCV coinfection results in more severe liver disease (Sato *et al.*, 1994) and in an increased risk of HCC (Zarski *et al.*, 1998) than monoinfection.

The epidemiology of liver cancer is linked to the incidence and mortality rates from liver cirrhosis, since a large proportion of HCCs develop from cirrhotic liver (London and McGlynn, 2006). The improved survival and reduced mortality from cirrhosis, due to improvements in the prevention and treatment of this condition, have in fact increased the possibility of developing HCC in cirrhotic patients. Of some importance are also the improvements in diagnosis, mainly due to widespread use of ultrasound and measurement of alphafetoprotein since the early 1980s, which led to more frequent detection of neoplastic liver in cirrhotic patients (Shu-Chun et al., 2009).

HCV is a single-stranded RNA virus of the flavivirus family, about 9.5 kb in length. HCV does not integrate into the host genome, and the means by which it establishes chronic infection are unknown. The association of chronic HCV infection with development of HCC is well-established; this nearly always occurs in the presence cirrhosis. It is estimated that approximately 80% of persons infected with HCV will develop a chronic infection, and 15-20% will develop serious liver disease, often many decades later. Among those who develop cirrhosis, 1-4% per year will develop HCC. Several important cofactors, such as alcohol consumption and older age at infection, are known to affect the probability of developing HCC among persons chronically infected with HCV (Ikeda et al., 1993; Tsukuma et al., 1993; Lauer and Walker, 2001).

HBV is a small, partially doublestranded DNA virus of the hepadnavirus family, which replicates through a reverse transcription phase. Unlike HCV, HBV integrates into the host hepatocyte genome early in infection, usually at sites of DNA damage (Bill and Summers, 2004). The virus carries no known oncogenes, and sites of integration are not consistent from cell to cell. But the presence of multiple integrated viral genes is thought to create genomic instability in the host and may lead to loss of heterozygosity for suppressor genes. Moreover, hepatitis B x antigen acts as a transactivator that may lead to increased transcriptional activity of cellular oncogenes as well interference with the function of tumor suppressor genes, such as p53 (Feitelson et al., 2002). In populations where HBV infection occurs in early childhood, the lifetime risk of HCC in an HBV-infected person has been estimated at 27% for males and 4% for females. (Dickinson et al., 2002). Among endemic populations, however, there may be substantial disparities in HCC risk due to differences in the natural history of chronic HBV infection between populations (Evans et al., 1998).

MATERIALS AND METHODS I. Study population

The study was conducted on random blood samples collected from healthy blood donor volunteers, who were referred to Blood Transfusion Centers found at Aseer region (Southern part of KSA), during the period March 2012 to January 2013. According to routine practice, volunteer blood donors were interviewed (history of intravenous drug abuse, jaundice, admission to fever hospital and history of HBV vaccination) and medically examined before donation. Those with high-risk behaviors including intravenous drugs abusers, history of promiscuous sexual relationships, homosexuals, homeless or those with any

medical problem especially jaundice or hospitalization at fever hospitals, bleeding disorders necessitating component transfusion, pregnancy or recent delivery less than 12 weeks were rejected.

Sera of the 7267 (26 females and 7261 males) collected samples were separated and kept at -80°C till use.

II- Viral Markers screening: Hepatitis B virus surface antigen (HBsAg) test:

Detection of HBsAg was done using commercially available MonolisaTM HBsAg ULTRA ELISA kit (BIO-RAD, Mames-la-Coquette, France) for the detection of HBsAg in serum and plasma.

Confirmatory test for HBsAg positive units:

Blood units which were shown to be HBsAg positive or at border line were retested using HBsAg confirmation kit (DIA.PRO Diagnostic, Milano, Italy), a set of reagents for the confirmation of HBsAg positivity in human sera or plasma.

Anti-Hepatitis B core antigen antibodies (HBcAb) test:

Detection of HBcAb was done using commercially available MonolisaTM Anti-HBc PLUS ELISA kit (BIO-RAD), a detection kit for antibodies to nucleocapsid antigen (core) of the HBV in human serum or plasma by enzyme immunoassay.

Confirmatory test for HBcAb positive units:

Blood units which were positive or at border line to HBcAb were retested using anti-HBc detection kit (total, DIA.PRO) which is Enzyme immunoassay for the detection of antibodies to hepatitis B core antigen (anti-HBc) in human serum or plasma.

Anti-Hepatitis C virus antibodies (HCV-Ab) test:

Detection HCV-Ab was done using the commercially available Murex anti-HCV4th generation (DiaSorin, S. P. A.

UK Branch, Central road, Dartfod DA1 5Lr, UK), an enzyme immunoassay for the detection of antibodies to hepatitis C virus in human serum or plasma.

Confirmatory test for Anti-HCV-Ab:

Positive and borderline HCV-Ab units were confirmed using the commercially available HCV confirmation kit (DIA.PRO), an enzyme immunoassay for the confirmation of HCV Ab positivity in human sera or plasma.

Nucleic acid test (NAT)

All samples were tested for the presence of HBV, HCV and HIV nucleic acids by NAT using Roche COBAS® TaqScreen MPX Test which is a qualitative multiplex test that enables simultaneous screening of HIV-1 Group M and Group O RNA, HIV-2 RNA, HCV RNA and HBV DNA in pooled and individual plasma donations.

Statistical analysis

The biochemical data recorded were expressed as mean±SD and statistical and correlation analyses were undertaken using the One-way ANOVA followed by a post-hoc LSD (Least Significant Difference) test. A P value < 0.05 was statistically significant. A Statistical analysis was performed with the Statistical Package for the Social Sciences for Windows (SPSS, version 10.0, Chicago, IL, USA).

RESULTS

I. Donor selection and sample Collection:

Both donor selection and blood testing were done according to WHO and Saudi Ministry of Health recommendations.

The studied samples included 7267 (26 (0.36% females; 7241 (99.64%) randomly selected blood donations from donors with a median age of 28 and 30 years respectively (Table 1).

Table 1: Gender distribution among blood donors accepted for donation.

Gender	Number	%	Median age
Female	26	0.35778	28
Male	7241	99.6422	30
Total	7267	100	30

Donors of ages between 21 and proportion (50.59%, $P \le 0.001$) with a 30 years constituted the largest median age of 26 years (Table 2).

Table 2: Number and age rage distribution among blood donors accepted for donation.

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Age range	Number	%	Median	Variance
18-20	429	5.903	20	0.649317
21-30	3676	50.585	26	7.722881
31-40	2203	30.315	35	7.967831
41-50	786	10.816	45	8.186748
51-60	173	2.381	55	8.828577
Total	7267	100	30	

Table 3 lists the distribution of the nationalities of the study participants. Blood donors were mostly Saudi nationals (95.13%). Non-Saudi donors were Afghani, Bengali, Egyptians, Eritrean, Indians, Filipinos, Jordanians, Lebanese, Pakistanis, Palestinians, Sudanese, Turkish and Yemenis. The higher non-Saudi proportion was Yemenis (1.58%), followed by Egyptians (1.32%), Sudanese (0.37%), Pakistani (0.35%), Indians (0.30%), Syrian (0.29%) and then Jordanians (0.27%).

Nationality	Number	%
Afghani	2	0.0275
Bengali	5	0.0688
Egyptian	96	1.3210
Erytrian	3	0.0413
Indian	22	0.3027
Jordon	20	0.2752
Lebanese	1	0.0138
Pakistani	26	0.3578
Philippine	5	0.0688
Palastine	7	0.0963
Saudi	6913	95.1287
Sudanese	27	0.3715
Syrian	21	0.2891
Tyrkey	4	0.0550
Yemani	115	1.5825
Total	7267	100.0000

Table 3: Nationality distribution blood donors accepted for donation.

II- Routine screening:

Hepatitis B virus surface antigen (HBsAg) test:

Detection of hepatitis B surface antigen (HBsAg) was done using commercially available ELISA kit. Of the 7267 donated blood, 71 (0.098%)

case were positive to HBsAg (Table 4). There were no infection detected among volunteers in age range 18-20. The highest infection rate inside groups was in age ranges 41-50 (3.05%) and 51-60 (2.89%) and was lowest in age range 21-30 (0.52%).

Table 4: Age range of HBsAg positive among blood donors accepted for donation.

Age range	Number	% in total population	HBsAg positive (n)	% infection in total population	% infection inside the age group
18-20	429	5.903	0	0	0
21-30	3676	50.585	19	0.2615	0.5169
31-40	2203	30.315	23	0.3165	1.0440
41-50	786	10.816	24	0.3303	3.0534
51-60	173	2.381	5	0.0689	2.8902
Total	7267	100	71	0.9770	0.9770

Of the 69 HBsAg positive donors there were one case coinfected with HIV, 70 (0.96%) of positive to antibody to

core antigen (HBcAb) and 66 (0.91%) were positive to HBV-DNA (Table 5).

Table 5: Other markers associated with HBsAg positive blood donors accepted for donation.

Positive to	HBsAg	HBcAb	HCV	HIV	RPR	HTLV	NAT
HBsAg	71	70	0	1	0	0	66

Anti-Hepatitis B core antigen antibodies (HBcAb):

Detection of HBcAb was done using commercially available ELISA kit for the detection of antibodies to nucleocapsid antigen (core) of the HBV in human serum or plasma by enzyme

immunoassay. Screening resulted in 449 positive donations (Table 6). There was no infection detected among volunteers in age range 18-20. The highest infection rate inside groups was in age ranges 51-60 (21.96%) and age range 21-30 (3.67%) was the lowest.

Table 6: Age range	of HRCAh pocitive	among blood donor	s accepted for donation.
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Age range	Number	% in total population	HBcAb positive (n)	% infection in total population	% infection inside the age group
18-20	408	6.091	0	0	0
21-30	3384	50.523	135	1.8577	3.6725
31-40	2021	30.173	157	2.1605	7.1267
41-50	725	10.824	119	1.6375	15.1399
51-60	160	2.389	38	0.5229	21.9653
Total	6698	100	449	6.1786	6.1786

Of these positive cases there were 70 cases positive for HBsAg. Cases positive to HBc-Ab and positive to HBV-

DNA in the same time were 78. One case was coinfected with HCV (Table 7).

Table 7: Other markers associated with HBcAb positive blood donors accepted for donation.

Positive to	HBsAg	HBcAb	HCV	HIV	RPR	HTLV	NAT
HBcAb	70	449	1	0	0	0	78

Anti-Hepatitis C virus antibodies (HCV-Ab) test:

A detection hepatitis C virus antibody (HCV-Ab) was done using the commercially available 4th generation kit.

Five cases were found to be positive for antibodies against HCV (Table 8). All cases were Saudi national aged 23, 36, 38, 45 and 50 years old.

Table 8: Age range of HCV-Ab positive among blood donors accepted for donation.

Age range	Number	% in total population	HBsAg positive (n)	% infection in total population	% infection inside the group
18-20	408	6.09137	0	0	0
21-30	3384	50.5225	1	0.0138	0.0272
31-40	2021	30.1732	2	0.0275	0.0908
41-50	725	10.8241	2	0.0275	0.2544
51-60	160	2.38877	0	0	0
Total	6698	100	5	0.0688	0.0688

Of these positive cases there was one case (45 years old) positive for HBcAb and HBV-DNA in the same time

and one case positive for HBV-DNA (Table 9).

Table 9: Other markers associated with HCV positive blood donors accepted for donation.

Positive to	HBsAg	HBcAb	HCV	HIV	RPR	HTLV	NAT
HCV	0	1	5	0	0	0	2

DISCUSSION

The group of viruses (hepatitis A, B, C, D and E) that cause acute and/or chronic infection and inflammation of the liver gives rise to a major public health problem globally. Hepatitis B and C viruses are major causes of severe illness and death. The global burden of disease due to acute hepatitis B and C and to

cancer and cirrhosis of the liver is high (about 2.7% of all deaths) and is forecast to become a higher ranked cause of death over the next two decades. (WHO, 2010).

Hepatitis B is a potentially lifethreatening liver infection caused by the hepatitis B virus. It is a major global health problem and the most serious type of viral hepatitis. It can cause chronic liver disease and puts people at high risk of death from cirrhosis of the liver and liver cancer. Worldwide, an estimated two billion people have been infected with the hepatitis B virus and more than 240 million have chronic (long-term) liver infections. About 600,000 people die every year due to the acute or chronic consequences of hepatitis B. A vaccine against hepatitis B has been available since 1982. Hepatitis B vaccine is 95% effective in preventing infection and its chronic consequences, and is the first vaccine against a major human cancer. Hepatitis B virus can cause an acute illness with symptoms that last several weeks, including yellowing of the skin and eyes (jaundice), dark urine, extreme fatigue, nausea, vomiting and abdominal pain. Hepatitis B is endemic in China and other parts of Asia. Most people in this region become infected with the hepatitis B virus during childhood and 8-10% of the adult population is chronically infected. Liver cancer caused by hepatitis B is among the first three causes of death from cancer in men, and a major cause of cancer in women in this region (WHO, Fact sheet N°204, 2012).

In the present study, age range 21-30 years old constituted the largest population among blood donors. In addition majority of the donors were male and there were very little number of female donors during the period of this study. The percentage of non-Saudi donors was low. A similar study conducted on blood donors in Saudi Arabia by El-Hazmi (2004) also showed that the largest groups of donors were those at age range 20-29 years old and female donors were as low as 1.2% at year 2000 and declined to reach 0.7% by year 2002. Also El-Hamzi (El-Hazmi 2004) showed that the percentage of non-Saudi donors declined from 17.2% at year 2000 to reach 14.8 by year 2002. Also Ankra-Badu et al. (2001) previously showed that the proportion of Saudi

blood donors increased with the decrease in the non-Saudis blood donors.

In the present study blood donors were screened for the presence of HBsAg, anti-HBc, HCV-Ab, HCV-RNA and HBV-DNA. We found large number of anti-HBc carrier with or without HBsAg positivity. According to De Villa et al. (2003) HBcAb positivity with HBsAg negative status can reflect a number of situations: (1) it may indicate a false-positive result, so in the present study positive cases were confirmed using different detection kit; (2) it may represent past and currently healed infection, and this why we in the present screened units for HBsAg positivity; and (3) it may constitute the sole marker of occult HBV infection, which is thus potentially transmissible, as has been demonstrated by contagion occurring through blood transfusion from donors who are only HBcAb (+) (Hoofnagle, Seeff et al. 1978).

In the present study it was found that HBsAg positive cases were low while HBc-Ab positive cases were relatively high. Similar work done by Panhotra et al. (2005) on blood donors and showed that 1.9% were HBsAg positive alone, 3.2% were anti-HBc positive alone and 10.1% were both anti-HBc and anti-HBsAg positive. In the current study we found only one case (0.02%) positive for HBsAg alone which means that the presence of low HBsAg infection in Aseer region. It was previously shown shat there is a decline in hepatitis B viral infection in South-Western Saudi Arabia and it was attributed to the effectiveness efficacy of the integration of hepatitis B vaccination into the extended program of immunization in KSA. The significant decline of HBV markers among unvaccinated Saudi adults indicated an indirect effect of other factors (for example health education and socioeconomic progress) on the prevalence and transmission of HBV. In areas of

high endemicity, the epidemiological characteristics HBV are modified significantly by the combination of HBV vaccination and other complimentary control strategies(Ayoola, Tobaigy *et al.* 2003).

Occurrence of HBsAg positivity among subpopulation was the lowest in 50-60 group, while it was zero in young population (group 18-20 years old). Vaccination against HBV was introduced 1989 for all infants at birth and in 1990 for school children (Al-Faleh 2003). This may be the most important factor responsible for the decline in HBV infection (Al-Faleh 2003).

A study done by El-Hazmi (El-Hazmi 1989) conducted on male and female population in different provinces of Saudi Arabia. The overall prevalence of hepatitis B surface antigen (HBsAg) was high (16.7%) and no significant difference was encountered between the rate in males and females. Different regions of Saudi Arabia showed a significantly variable prevalence HBsAg. The eastern province had a prevalence of about 9% compared to the southwestern province where the prevalence was 25% in Jizan. The antibodies anti-HBs and anti-HBc were encountered in 30-67% of the individuals in different provinces, suggesting that a significant number of Saudis were already immune to HBsAg before they reached adulthood.

of **HBV-DNA** Presence HBsAg positive samples was in nearly all cases except five samples. This indicates to how much these blood units are highly infective. Also, HBV-DNA was found in high percentage (17.37%) of HBc-Ab positive cases. This indicated positive blood units for HBc-Ab should be discarded as it carries high possibility of infectivity. Current infection showing HBsAg and HBc-Ab in the same time was high (15.59%). In the time HBc-Ab positive cases with HBV-DNA negative is more than those cases with HBV-DNA positive, it still not secure to use these units in blood transfusion. The virus may be found in polymorhonuclear cells or other places other than serum or plasma used for the detection of HBV-DNA (Catterall, Murray-Lyon *et al.* 1994).

Previous studies since 1980s showed a high prevalence of HBV infection in Saudi Arabia ranging between 5-10% of the population varying from one region to another (Arya et al., 1985; Parande et al., 1986; Al-Faleh, 1988). The prevalence of HBsAg in children with ages of 1 to 10 years was about 7% in 1989, just before adding the HBV vaccine as the seventh primary immunogen of the Extended Program of Immunization (EPI) (Al-Faleh et al., 1992). This followed by vaccination of all children entering school in 1990 as a program launched by MOH (Al-Faleh et al., 1999).

In 1997, the efficacy of the vaccination hepatitis program evaluated by means of a stratified cluster sampling technique in all 13 regions of This community-based study detected a decline of HBV infection among Saudi children up to the age of 12 years—from 7% in 1989 to 0.3% in 1997 (Al-Faleh et al., 1999). Children positive for HBV were found to have an HBVinfected father, mother, sisters brothers. This indicates a major role for familial horizontal transmission of HBV in KSA. In the 1980s, a study of MOH blood donors showed that HBV infection averaged 5% to 10% (Al-Faleh, 1988). Screening of blood donors by Ministry of health showed that HBV infection in the year 2000 was at a prevalence rate of 3.2% and 5.9% among persons between 18-44 years old and over 50 years old respectively with an average of 3.25%. The prevalence of HBV infection declined from 3.7% in year 1987 to 1.7% in year 2000 (Al-Faleh, 2003).

Hepatitis C is a contagious liver disease that results from infection with the hepatitis C virus. It can range in

severity from a mild illness lasting a few weeks to a serious, lifelong illness. The hepatitis C virus is usually spread when blood from an infected person enters the body of a susceptible person. It is among the most common viruses that infect the liver. Every year, 3–4 million people are infected with the hepatitis C virus. About 150 million people are chronically infected and at risk of developing liver cirrhosis and/or liver cancer. More than 350 000 people die from hepatitis C-related liver diseases every year (WHO, Fact sheet N°164, 2012).

The current study showed that HCV infection is very low (0.067%) with complete absence of infection in age ranges 18-20 and 51-60 years old with cases having **HCV-RNA** 2 positivity. Since the discovery of HCV in 1988 and introduction of diagnostic tests for this virus, data on HCV prevalence has accumulated in Saudi Arabia. In 1989, a baseline community randomized study in Jizan was done on children from the age of 1 to 10 years for prevalence of HCV. The average prevalence was calculated to be 0.87% (Al-Faleh et al., 1991). Another study in the same region in 1991 but on the adult population over the age of 10 years was done. The average prevalence was only 1.8%, which increased with age, reaching 3.5% in persons over 50 years (Al-Faleh et al., 1995). A study done on children aged 1-12 years in 13 regions of KSA in 1997, showed that the prevalence of HCV was only 0.04% . Al-Faleh (2003) also analyzed Saudi blood donors from 1996 to 2001 at KKUH and demonstrated that the prevalence of HCV decreased steadily from 0.58% in 1996 to 0.28% in 2001, which is a decline among blood donors of more than 50% over 4 years.

CONCLUSIONS

Prevalence of HBsAg in Asser region is very low. The rate of HBc-Ab in units of blood donation is relatively high. The presence of HBV-DNA in

HBc-Ab positive donations make it risky for use. Vaccination program against HBV decreased the rate of HBV transmission. Infection rate of HCV is very low. Expected incidence of heptatocellular carcinoma due to infection with HCV is much lower than HBV.

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