

Intrafamilial spread of hepatitis B: Passage through several generations

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A SZILAGYI, J CHAN-THIN, I FANTUS. Intrafamilial spread of hepatitis B: Passage through several generations. Can J Gastroenterol 1993;7(1):23-27. Uncontrolled spread of hepatitis B virus (HBV) is described in an immigrant Italian family. As many as four generations of offspring appear to have been infected, the source likely originating from a great grandfather. Overt liver disease in four of six members in the second generation as well as a history of hepatitis implicates the virus in the two generations which could not be tested. Five of seven asymptomatic sisters in generation III were found to be carriers, and were both hepatitis B surface antigen (HBsAg)-positive and antihepatitis B antigen antibody-positive. One sister lost HBsAg status shortly after discovery of her carrier status. The overt carrier rate in generation IV was reduced, suggesting later childhood acquisition of HBV infection. This family strongly emphasized the need for investigating patients with a positive family history of liver disease and screening family members of newly discovered HBsAg carriers to allow appropriate measures to be taken to limit the spread of HBV.

Key Words: *Hepatitis familial infection*

Transmission de l'hépatite B entre les générations d'une même famille

RÉSUMÉ: La transmission non maîtrisée du virus de l'hépatite B est décrite au sein d'une famille d'immigrants italiens. Jusqu'à quatre générations semblent avoir été infectées, la source originant probablement d'un arrière-grand-père. Une maladie hépatique évidente chez quatre des six membres de la deuxième

CONSEQUENCES OF CHRONIC HEPATITIS B virus (HBV) infection pose an important economic burden, and cause significant morbidity and mortality. In areas of high or mid-prevalence of HBV, most infections are acquired in the neonatal or early childhood period (1-3). In low prevalence areas, specific high risk lifestyles predispose to infection (1,2). However, a large proportion of cases are related to inapparent sources (4). In such cases the contribution of occult intrafamilial spread of HBV may be overlooked and thus, an opportunity to limit further HBV spread may be missed. Intrafamilial spread both of HBV (5-7) and hepatocellular carcinoma – strongly associated with HBV (8,9) – have been described.

The mechanism of intrafamilial spread has been linked both to vertical transmission (10) and horizontal spread (6) among members of the same or other families. At least one earlier study from North America found little likelihood of horizontal or vertical spread (11). To emphasize the usefulness of current efforts to minimize HBV spread and to shed a historical perspective on

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génération, de même que des antécédents d'hépatite, identifient le virus dans les deux générations qui n'ont pas pu être soumises à l'épreuve. Cinq des sept sœurs asymptomatiques de la génération III se sont révélées porteuses et positives à l'égard de l'antigène de surface de l'hépatite B (AgsHB) et à l'égard d'un anticorps anti-antigène de l'hépatite Be. L'une des sœurs a perdu son statut AgsHB peu de temps après avoir découvert son statut de porteuse. Le taux de porteurs déclarés dans la génération IV a été moindre, donnant à supposer une acquisition plus tardive durant l'enfance de l'infection au HBV. Cette famille souligne nettement la nécessité de pratiquer un dépistage chez les patients qui présentent des antécédents familiaux positifs d'hépatite et le dépistage des membres des familles de porteurs AgsHB nouvellement diagnostiqués afin de promouvoir les mesures appropriées pour limiter la propagation du HBV.

the hazards of unchecked HBV infection, the authors report a family who immigrated from southern Italy with as many as four generations possibly with HBV likely via both horizontal and vertical spread. Such transmission of HBV, spanning several generations, has rarely been reported (12,13).

SUBJECTS AND METHODS

The index case was referred for evaluation of a thyroid nodule and gave a positive strong history of liver disease. This included hepatocellular carcinoma in her father, possible liver disease in a paternal aunt, and blood test ab-

normalities of liver function in a sister. A screen for hepatitis B surface antigen (HBsAg) was positive and the patient was referred for further evaluation. Subsequently, seven sisters, four of their spouses in generation III and nine of their children in generation IV were tested. One additional neonate in generation IV was subsequently vaccinated at birth.

HBsAg, anti-HBsAg antibody, hepatitis Be antigen (HBeAg) anti-HBeAg antibody, immunoglobulin (Ig) G anti-hepatitis Bc antigen (HBcAg) antibody were measured by standard assay (Enzyme Immunoassay, Boehring diagnos-

tics and Microparticle Enzyme Immunoassay, Abbott). All positive HBsAg were confirmed by reverse passive hemagglutination (Abbott Diagnostics, Illinois). Anti-HBcAg antibody were measured by Enzyme Immunoassay (Ortho Diagnostics). Wilson's disease, hemochromatosis and α 1 antitrypsin deficiency were ruled out by standard methods.

RESULTS

Affected and probably affected members are shown in Figure 1. Historically, the great grandfather (generation I [Gen II]) migrated from Valsini in southern Italy to Turin in Maternal province. He was reported to have died in his early 50s with ascites and jaundice. In Gen II, four of six siblings were reported to have had liver disease. The grandfather (Gen II₁) immigrated to Montreal and was reported to have died in his 50s with ascites and probable liver cancer. The brother (Gen II₂) apparently had cirrhosis, but was thought to drink alcohol excessively. Two other sisters in Italy have known liver disease and at least one recalled being told hepatitis B was the cause. Unfortunately, this branch of the family declined to be tested. There was no history of liver disease on the great grandmother or grandmother's side.

Gen III, seven females, were born in Italy and immigrated in the early 1960s to Montreal with their parents. Their mean age at present is 40.7 years. Two additional siblings died as neonates. None of the sisters is symptomatic, and only one has mildly abnormal liver biochemistries (Gen III₆). Their abdominal ultrasound examinations were normal. Because of no direct serological evidence of HBV infection in Gen I and II, serum ceruloplasmin, serum iron studies and α 1 antitrypsin were measured in six of seven sisters; these studies were uniformly negative. In addition, serology for anti-HBc was negative in five sisters of Gen III and two of their spouses.

The serological findings in the seven sisters, four of their spouses and nine children are listed in Table 1. Tests were repeated at least three times during a 24-month observation period.

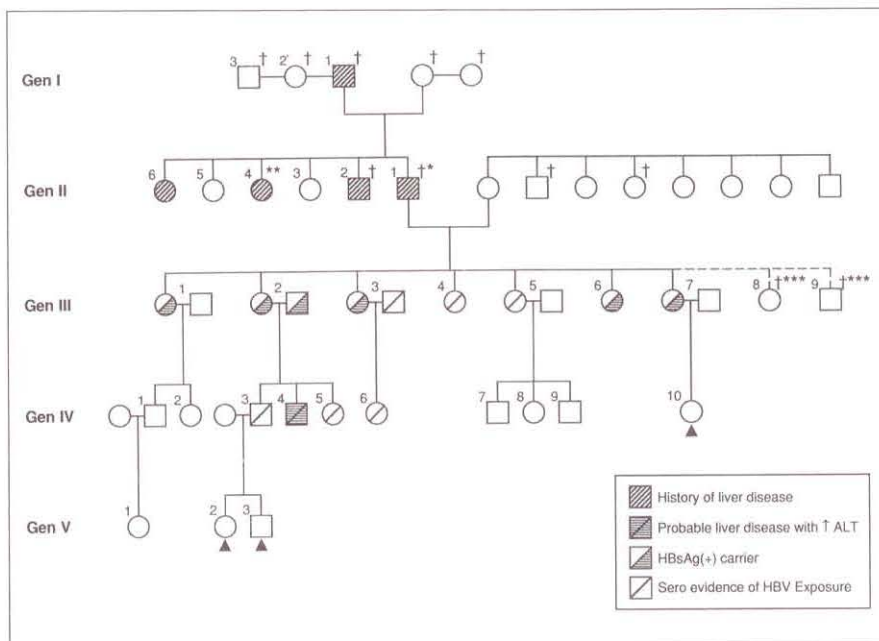


Figure 1 Intrafamilial spread of hepatitis B is depicted through four consecutive generations. Hepatitis B virus (HBV) serology was measured only in generations III and IV. The presence of frequent liver disease and a historical relationship to HBV in generation II implicates the virus in the pathogenesis of the liver disease in generations I and II. *Probable hepatocellular carcinoma; **History of positive HBV serology; ***Neonatal death; † Died

All seven sisters were found to have evidence of exposure to HBV. Five were initially HBsAg-positive and anti-HBeAg-positive. Within eight months of the initial tests, III₃ converted from HBsAg-positive to anti-HBsAg antibody-positive and lost HBsAg. This event was verified by the finding of borderline negative HBsAg on repeat testing. Two other sisters had only antibodies; III₅ had anti-HBsAg only and III₄ had only positive IgG anti-HBcAg antibody found on all three tests. One of four spouses was HBsAg-positive and anti-HBeAg-positive while one was positive only for anti-HBsAg.

In Gen IV (mean age 15.7 years) three children from III₅ (IV 7,8,9) were negative for HBV-related serology. Of the remaining six, IV₄ was positive for both HBsAg and HBeAg. He also had mild aspartate aminotransferase and alanine aminotransferase elevations. This subject has no known either risk factors for HBV infection. The brother and sister of IV₄ (IV₃ and IV₅, respectively) both have anti-HBsAg and anti-HBeAg but are otherwise asymptomatic. One cousin, IV₆, was also anti-HBsAg-positive. In addition, the neonate (Gen IV₁₀) was vaccinated against HBsAg, and Gen V₂ and V₃ were vaccinated because of close contact with Gen III₂.

DISCUSSION

Familial clustering of HBV infection or hepatoma has been previously described (5-9,12,13). What is interesting about this family is that probably four successive generations were affected by HBV. Also of interest is the observation that the likely source of infection in Gen I and II could be traced to the father's side. Finally, it is noteworthy that the successive unchecked passage of HBV led to an apparent natural reduction in the number of HBsAg carrier children by the fourth generation.

Infection with HBV can occur by several routes. Blood transfusion (now rare) (14), shared needles and sexual contact are well described (15-17) but do not represent the majority of traced infections in North America (4). With the exception of stool, HBV has been

TABLE 1
Hepatitis B serology in generations III and IV and their spouses

Gen/spouse	Age (years)	HBsAg	anti-HBsAg	HBeAg	anti-HBeAg	anti-IgGHbCAg
Gen III ₁	49	+	-	-	+	+
Spouse	56	ND	ND	ND	ND	ND
Gen IV ₁	27	-	-	-	-	-
Gen IV ₂	23	-	-	-	-	-
Gen III ₂	47	+	-	-	+	+
Spouse	51	+	-	-	+	ND
Gen IV ₃	26	-	+	-	+	ND
Gen IV ₄	24	+	-	+	-	+
Gen IV ₅	28	-	+	-	+	ND
Gen III ₃ *	43	+	-	-	+	+
Spouse	49	-	+	-	+	+
Gen IV ₆	25	-	+	-	-	+
Gen III ₄ †	40	-	-	-	-	+
Gen III ₅	38	-	+	-	-	+
Spouse	36	-	-	-	-	-
Gen IV ₇	8	-	-	-	-	-
Gen IV ₈	6	-	-	-	-	-
Gen IV ₉	4	-	-	-	-	-
Gen III ₆	35	+	-	-	+	+
Gen III ₇	32	+	-	-	+	+
Spouse	33	-	-	-	-	-
Gen IV ₁₀	0.5	Vaccinated				

*Subject converted to antihepatitis B surface antigen (HBsAg) within eight months of first test; †Retest done 12 months later revealed same pattern; Gen Generation; HBeAg Hepatitis Be surface antigen; IgGHbCAg Immunoglobulin G hepatitis Bc antigen; ND Not done

isolated from all bodily secretions (18). However, because up to 70% of infections with HBV are subclinical (4), tracing sources may be quite difficult. While 5 to 10% of infected adults become carriers, as many as 90% of infected neonates born to HBeAg-positive mothers will be carriers (19). Older children have an intermediate rate of carrier status. The age dependency on outcome of HBsAg carrier rate has a major impact on geographical epidemiology of HBV carriage. For example, in the Orient where a large exposure to HBV occurs through vertical transmission, the HBsAg carrier rate is the highest (20). In the Mediterranean, infectivity is believed to occur in older children largely through horizontal spread and their carriage rates are much lower (21). In North America, HBV exposure occurs mainly in minority immigrants or among high risk adult populations leading to a much smaller HBV carriage rate (21). Immigrants from endemic areas retain their carrier rates (22).

Early reports on intrafamilial spread

of HBV or clustering of hepatoma focused on possible genetic predisposition to either carriage or carcinogenesis (23,24). Figure 1 certainly could be interpreted in the light of a dominant mode of inheritance of HBV with incomplete penetrance in Gen IV. However, early reports did not take into consideration the biological behaviour of HBV and a genetic linkage was not verified (25).

Intrafamilial spread traced to the paternal side is reported much less often than maternal sources (26). The mechanism of spread under these circumstances relates either to subclinical infection to the mother (consequently in whom high replicative HBeAg-positivity occurs) or more likely to horizontal transmission through close familial contact (27).

The apparent passage of HBV through multiple generations has been reported only occasionally. Three successive generations affected by hepatoma were reported by Lok et al (12) supporting the notion of multigeneration passage of HBV (12). Although

historical inference imply that Gen I and II were infected, the elimination of other genetic liver disease and the uniform positive HBV serology in sisters of Gen III coupled with the high penetrance of liver disease in Gen II strongly implicate HBV as being pathogenic in both Gen I and II. Such a generational spread of HBV was also documented in a study of another Italian American family in which two-thirds of five generations had evidence of positive HBV serology (13).

The authors noted that the carrier rate of HBsAg decreased by the fourth generation (five of seven, 71% Gen III versus one of six, 17% Gen IV). There is no clear explanation for this apparent reduction. Large population-based studies from both Italy and Japan also noted a carrier rate reduction in

children of HBsAg-positive mothers (28,29). Plausible explanations include an improvement in socioeconomic conditions and decreased family size (which limits sibling spread). Nevertheless, HBV infectivity remained high in Gen IV since four of six children had HBV markers. Therefore, one wonders why three of four children cleared HBsAg in Gen IV and not in Gen III (since the rate of HBsAg clearance in carriers is no more than 1 to 2% per year [30]). It is possible that in Gen IV the children acquired subclinical hepatitis horizontally, in which case the clearance is higher and carrier rate is lower. The observation that passage of HBV infection is much less efficient in anti-HBeAg-positive mothers supports this notion (4,19). Although the mothers in Gen III began delivering in their early and late 20s, we are not cer-

tain that mothers in this generation were anti-HBeAg-positive at delivery. These results also differ from Sampliner *et al* (13) whose work indicated that 13 of 18 children born to HBsAg mothers became carriers. Further study of this discrepancy is required.

CONCLUSIONS

In summary, the present authors report a family in which at least four generations may have been infected with HBV. The combination of serological findings suggest that both vertical and horizontal spread were responsible for dissemination of the infection. This family study re-emphasizes the need for close historical and medical scrutiny of newly discovered patients with HBV. Appropriate medical intervention will then limit further spread of this potentially lethal infection.

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