Prevalence and correlates of HIV and hepatitis B virus coinfection in Northern Alberta

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BACKGROUND: HIV and hepatitis B virus (HBV) share transmission routes, and coinfection is associated with higher morbidity and mortality. To date, no Canadian studies have examined HIV-HBV coinfection, and coinfection is associated with higher morbidity and mortality.

OBJECTIVES: To examine the prevalence and correlates of HIV and HBV coinfections in Northern Alberta.

METHODS: The present study was a retrospective database review of all HIV-infected (HIV+) individuals in Northern Alberta from 1982 to 2010 and a chart review of HBV surface antigen-positive individuals for whom charts were available (46.2%).

RESULTS: Of 2844 HIV+ patients, 2579 (90.7%) had been tested for HBV surface antigen, and 143 (5.5%) of these were HBV coinfected. Coinfected males were primarily Caucasian (70.8%), and coinfected females were primarily black (56.4%) or Aboriginal (31.3%). Cofected individuals were more likely to be male (88.1% versus 71.3%; P<0.001) and to have died (34.3% versus 17.9%; P<0.001).

CONCLUSIONS: The prevalence of coinfection with HBV in HIV-infected patients in Northern Alberta is lower than reported in other developed nations. The pattern of coinfections in Northern Alberta likely follows immigration trends. Recognition and management may be improving with time; however, further research and additional strategies are required to enhance the prevention, identification and management of HBV infection in HIV-infected individuals.

Key Words: Alberta; Canada; Coinfection; Hepatitis B; HIV

HIV and hepatitis B virus (HBV) are major public health concerns. There are >70,000 Canadians living with HIV (1). Approximately 700 new HBV infections occur in Canada annually (2), and >300 million individuals are estimated to be diagnosed with HBV worldwide (3). Undiagnosed reservoir populations of both infections are believed to exist in Canada (1,4).

Canada is a country of low endemicity for both HIV and HBV. While the prevalence of HIV is low in Canada overall, the incidence of infection is increasing among immigrants from endemic countries and men who have sex with men (1,5). The prevalence of HBV declined with the introduction of universal childhood vaccination in 1995 (3). The population of individuals chronically infected with HBV in Canada is largely comprised of individuals from HBV-endemic countries in Asia and Africa, and also includes high rates in Aboriginal and high-risk lifestyle populations (2). Prevalence rates among these populations are comparable with those of endemic areas (4).

As a major centre for the oil industry, Northern Alberta has a unique population. There has been a recent, rapid population expansion, with population growth in Alberta being more than double the national average, in which international immigration is playing an increasing role (6).

HIV and HBV coinfection can occur due to shared routes of transmission (7). In areas of low HBV endemicity, such as North America and Western Europe, HIV and HBV coinfections occur primarily in immigrant populations and adult populations due to sexual and percutaneous transmission (7). The prevalence of chronic HBV coinfection among HIV-infected individuals in areas of low HBV endemicity has been reported to range from 6% to 14% (8,9).

With the use of highly active antiretroviral therapy (HAART), HIV-infected individuals are living longer, and experiencing morbidity and mortality due to other causes. Coinfection with HBV is associated with increased mortality (10) and worse outcome (11) in HIV patients. HBV infections are more likely to be chronic (12,13) and have higher infectivity (14,15), and there is increased loss of HBV immunity in HBV-infected individuals (16,17). Drug-induced toxicity may also play a role in liver disease (18). HIV can hasten the progression of HBV-related liver disease and/or cirrhosis (19,20). HIV-infected individuals with lower CD4 cell counts may also be at greater risk of developing hepatocellular carcinoma (21). Individuals coinfected with HIV-HBV are 17 times more likely to die of liver-related causes compared with those monoinfected with HBV (22).
HIV and HBV coinfection in Northern Alberta

Lamivudine was the only oral agent for antiviral treatment of HBV in Canada from its introduction in 1998 until 2006. While of modest efficacy, the utility of lamivudine is limited due to resistance rates of approximately 70% over four years (23), and it is no longer recommended as monotherapy (24-27). Current Canadian guidelines on the treatment of HIV-HBV coinfection recommend treatment with HAART regimens containing two drugs active against HBV (24). Drugs active against HBV currently used in Canada include interferons, lamivudine, adefovir, entecavir, tafibuvir, tenofovir and emtricitabine (24,28).

HIV-HBV coinfection is a public health concern with distinct management issues, and high morbidity and mortality. There are currently no Canadian studies on the prevalence, correlates and mortality of HIV-HBV coinfections. We conducted a retrospective review on a cohort of HIV-positive patients followed between 1982 and 2010 in Northern Alberta. Our objectives were to examine the prevalence of HIV-HBV coinfection and to describe the correlates associated with coinfection in this region.

METHODS

Data sources
The Northern Alberta HIV Program (NAP) Database was created in 2003 and contains information on all HIV-positive patients from a population of nearly two million individuals in Northern Alberta and parts of the Northwest Territories. All patients who test positive for HIV in this catchment area are referred to the NAP, and confirmed positive patients are entered into the database. The NAP database was created for clinical and research purposes, and has been used in several recent studies (29-31). The NAP database is built on the Oracle platform and each patient file contains a unique identifier as well as demographic and selected clinical information. The NAP data are obtained from charts, laboratory results and Alberta NetCare electronic health records, and are entered by clerical staff. Data entry and modification is restricted by a user profile, and it is not double entered or checked. The data are updated after patient visits, with new laboratory results and when the NAP is made aware of changes in patient status. A database analyst ensures that the database is functional, runs database queries as requested and manages database fields.

Data for the present study were extracted from the NAP database inclusive of cases entered from the index case of HIV in Alberta in 1982 to May 2010. These data were used to identify patients who were reported as having tested hepatitis B surface antigen positive (HBsAg+). Information on sex, ethnicity, age, results and dates of microbial testing, and risk factor information was extracted for both HBsAg+ and HBsAg-negative (HBsAg−) patients from the NAP database.

The four NAP clinical sites located in Edmonton (Alberta) were visited and available charts of confirmed patients were reviewed for further information. Every attempt was made to locate all records, including from potential storage locations. Both medications and dates were recorded for HAART regimens and information was based on chart notes, prescription records and medication profiles. Laboratory investigation results specific to HBV infection and HBV-related liver disease were also recorded based on chart notes and hard-copy laboratory results filed in the chart.

Testing methods
All microbiological testing was performed at the Provincial Laboratory for Public Health (Edmonton and Calgary, Alberta), and analyzed and reported as per their protocols. Testing methodologies have changed over time; however, all HIV screening during this time period was conducted with an enzyme immunoassay (EIA). All screen-positive tests were followed by a second EIA and confirmation by Western blot. HBsAg testing involved EIA screening followed by a confirmatory EIA. Other hepatitis B serology, including hepatitis B surface antibody, hepatitis B core antibody (anti-HBc), hepatitis B e antigen (HBeAg), and antibody to HBeAg, were tested using a single EIA. HBV DNA samples were analyzed using quantitative DNA polymerase chain reaction (PCR) and were available in Alberta from December 2003.

Hepatitis C virus (HCV) PCR samples were analyzed using qualitative RNA PCR. HCV antibody testing began in 1992; methods have varied during this time span, but involve EIA for screening and confirmation with or without a third EIA or immunoblot test.

Statistical analysis
Variables were compared between HBsAg+ and HBsAg− individuals using χ² or Fisher exact tests for categorical variables and Mann-Whitney tests for continuous variables. Multivariable logistic regression was used to identify independent demographic and risk factor correlates for HBsAg positivity and assess the impact of possible confounders such as age at HIV and HCV diagnosis. Any variables identified by univariate analysis to be associated with HBV infection at a significance level of ≤0.10 were included in multivariable analyses, as well as any potential confounding variables. If inclusion of variables did not result in meaningful changes to the coefficients or model, variables were not included in the final model. Statistical analyses were performed using Stata version 10 (StataCorp, USA).

Ethics
Ethics approval for this project was obtained from the Health Research Ethics Board at the University of Alberta (Edmonton, Alberta).

RESULTS
Of 2844 HIV patients in the NAP as of May 2010, 265 (9.3%) were of unknown HBsAg status. Of 2579 patients with HBV test results, 143 (5.5%) were HBsAg+.

Compared with individuals for whom HBsAg testing results were documented, the 265 patients with unknown HBsAg status had a significantly earlier year of HIV diagnosis [1992 [interquartile range (IQR) 1988 to 2000] versus 1999 [IQR 1993 to 2004]; P=0.002], were younger (median age 31 years [IQR 25 to 38 years] versus 34 years [IQR 28 to 41 years]; P=0.001), were more likely to have died (33.2% versus 18.8%; P=0.001), had fewer years of life after their HIV diagnosis (three years [IQR one to five years] versus five years [IQR three to eight years]; P<0.001), and were more likely to be Caucasian (64.0% versus 54.5%; P=0.003) and less likely to be black (8.4% versus 14.5%; P=0.007).

There were no statistically significant differences in age at time of HIV diagnosis, age at death and years of life after HIV diagnosis between the HBsAg+ and HBsAg− groups (Table 1). In univariate analyses, the HBsAg+ group was significantly more likely to have an earlier median date of HIV diagnosis [1996 [IQR 1999 to 2001] versus 1999 [IQR 1993 to 2004]; P=0.001], be male (88.1% versus 71.3%; P=0.001), more likely to have died (34.3% versus 17.9%; P<0.001) and less likely to be Aboriginal (14.0% versus 27.7%; P=0.006). HBsAg+ patients were less likely to be viremic with HCV (15.8% versus 63.7%; P=0.002) despite comparable anti-HCV seropositivity rates (37.0% versus 40.0%; P=0.6). No differences were observed in serological results for cytomegalovirus, toxoplasmosis, varicella zoster virus and syphilis between HBsAg+ and HBsAg− groups (data not shown).

The prevalence of HBV coinfection decreased with year of HIV detection; patients with HIV infection identified between 1982 and 1987 had an HBsAg prevalence of 13.0%, which those diagnosed after 1996 had a rate of 4.2% (P=0.001). Figure 1 presents the HIV cases shown (median age 31 years [IQR 25 to 38 years] versus 34 years [IQR 28 to 41 years]; P=0.001), were more likely to be Caucasian (64.0% versus 54.5%; P=0.003) and less likely to be black (8.4% versus 14.5%; P=0.007).

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HBsAg prevalence among males was significantly higher than among females (6.8% versus 2.4%; P=0.001.) Among males, a significant correlate of HBV coinfection on univariate analysis was being a man who has sex with men (OR=1.50; P=0.03; Table 1). Being a man who has sex with men did not remain significant in multivariable analysis.

When patterns of ethnicity were examined, Caucasians represented the majority of infected male patients (n=87 [70.8%]), followed by blacks (n=16 [13.0%]) and Aboriginals (n=15 [12.2%]).
majority of coinfected females were of black ethnicity (nine of 16 [56.2%]), followed by Aboriginal ethnicity (n=5 [31.3%]). Except for one patient, all black females were from HIV-endemic countries, as defined by the Public Health Agency of Canada for surveillance purposes (32). Black ethnicity was highly associated with HBV infection in females (56.2% versus 25.1%; P<0.01).

In multivariable analyses, factors identified as independent correlates of HBsAg+ status in this HIV population were male sex (adjusted OR [aOR] 2.56 [95% CI 1.43 to 4.56]), being deceased (aOR 1.84 [95% CI 1.21 to 2.80]), earlier year of HIV detection (aOR 0.95 [95% CI 0.93 to 0.98]) and black ethnicity (referent: Caucasian; aOR 2.22 [95% CI 1.28 to 3.88]).

Charts were located for less than one-half (n=66 [46.2%]) of HBsAg+ patients. These patients had a later diagnosis of HIV infection (2000 versus 1991; P=0.0001), were less likely to have died (21% versus 45%; P=0.001), were less likely to be male (81% versus 93.5%; P=0.04) and were more likely to be of black ethnicity (31.8% versus 5.5%; P=0.0001) than the remaining HBsAg+ patients for whom charts were not available.

HBV DNA levels were reported for 33 of 120 (27.5%) HBsAg+ patients in the database and 32 of 58 (55%) patients for whom charts were available, with 17 of 32 (53.1%) having detectable HBV DNA (note that the cut off for detection varied over time). Anti-HBc results were reported for 31 of 143 (21.6%) HBsAg+ patients in the database and 30 of 66 (45.5%) patient charts, with 19 of 30 (63.3%) testing positive. HBV DNA testing was not routinely available before that time, and denominators exclude patients who had died before December 2003. HIV diagnosis was made more than four weeks before HBV diagnosis in 41 of 66 patients (62.1%), within four weeks of HIV diagnosis in nine of 66 (13.6%) and more than four weeks after HBV diagnosis in six of 66 patients (9.1%).

### Table 1

<table>
<thead>
<tr>
<th>Demographic characteristic</th>
<th>Positive (n=143)</th>
<th>Negative (n=2436)</th>
<th>P‡</th>
</tr>
</thead>
<tbody>
<tr>
<td>Age at diagnosis, years</td>
<td>32 (28–40)</td>
<td>34 (28–41)</td>
<td>0.2</td>
</tr>
<tr>
<td>Age at death, years</td>
<td>41 (35–48)</td>
<td>49 (33–48)</td>
<td>0.8</td>
</tr>
<tr>
<td>Years of life after HIV diagnosis</td>
<td>5 (3–8)</td>
<td>5 (3–8)</td>
<td>0.9</td>
</tr>
<tr>
<td>Male sex, n (%)</td>
<td>126 (88.1)</td>
<td>1736 (71.3)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Deceased, n (%)</td>
<td>49 (34.3)</td>
<td>437 (17.9)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Incarceration, n (%)</td>
<td>26 (19.3)</td>
<td>504 (24.4)</td>
<td>0.2</td>
</tr>
<tr>
<td>Ethnicity, n (%)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Caucasian</td>
<td>88 (61.5)</td>
<td>1314 (53.9)</td>
<td>0.004</td>
</tr>
<tr>
<td>Aboriginal</td>
<td>20 (14.0)</td>
<td>674 (27.7)</td>
<td></td>
</tr>
<tr>
<td>Black</td>
<td>25 (17.5)</td>
<td>351 (14.4)</td>
<td></td>
</tr>
<tr>
<td>Other</td>
<td>6 (4.2)</td>
<td>97 (4.0)</td>
<td></td>
</tr>
<tr>
<td>Risk factors§, n (%)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Needle stick</td>
<td>0 (0.0)</td>
<td>11 (0.45)</td>
<td>1.0</td>
</tr>
<tr>
<td>Transfusion</td>
<td>3 (2.1)</td>
<td>56 (2.3)</td>
<td>1.0</td>
</tr>
<tr>
<td>Vertical</td>
<td>1 (0.7)</td>
<td>13 (0.53)</td>
<td>0.6</td>
</tr>
<tr>
<td>Bisexual</td>
<td>20 (14.0)</td>
<td>245 (10.1)</td>
<td>0.1</td>
</tr>
<tr>
<td>Men who have sex with men (males only)</td>
<td>75 (59.5)</td>
<td>859 (49.5)</td>
<td>0.03</td>
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<tr>
<td>Injection drug use</td>
<td>39 (27.3)</td>
<td>833 (54.2)</td>
<td>0.09</td>
</tr>
<tr>
<td>Unknown</td>
<td>3 (2.1)</td>
<td>23 (9.4)</td>
<td>0.2</td>
</tr>
<tr>
<td>Serology, n (%)</td>
<td></td>
<td></td>
<td></td>
</tr>
<tr>
<td>Anti-hepatitis B surface antigen positive</td>
<td>22 (17.7)</td>
<td>1167 (50.4)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Anti-hepatitis B core antibody negative</td>
<td>6 (9.8)</td>
<td>773 (61.6)</td>
<td>&lt;0.001</td>
</tr>
<tr>
<td>Anti-hepatitis C virus positive</td>
<td>37 (37.0)</td>
<td>801 (40.0)</td>
<td>0.6</td>
</tr>
<tr>
<td>Hepatitis C virus RNA positive</td>
<td>3 (15.8)</td>
<td>261 (63.7)</td>
<td>0.002</td>
</tr>
<tr>
<td>Anti-hepatitis A virus positive</td>
<td>43 (70.4)</td>
<td>1035 (69.9)</td>
<td>0.9</td>
</tr>
</tbody>
</table>

Data presented as median (interquartile range) unless otherwise indicated. *Data collection ended in May 2010; ‡Denominators may not equal column totals due to missing data or serology testing not being performed; §Calculated using Mann-Whitney test for continuous variables and χ² or Fisher’s exact test for categorical variables; †Risk factor information is not mutually exclusive.

**Figure 1** HIV patients tested for hepatitis B surface antigen (HBsAg) and proportion testing HBsAg positive according to year of HIV diagnosis, Northern Alberta, 1982 to 2010. HBsAg test result represents lifetime prevalence rates because testing could have occurred at any point until May 2010.
For HBsAg+ patients with available charts, the HAART regimen was divided into three time periods: 1989 to 1997 (pre-lamivudine); 1998 to 2006 (lamivudine available as of 1998); and 2006 to 2010 (newer HBV agents available) (Figure 2). From 1989 to 1997, the majority of HAART regimens initiated in coinfected patients did not contain any drugs active against HBV, although 23 HAART regimens were initiated in the cohort that contained lamivudine before its formal approval on the Canadian drug market, and two trial HAART regimens consisted of both lamivudine and tenofovir. After lamivudine was formally approved in 1998, the number of HAART regimens containing at least one HBV-active antiretroviral doubled. The number of regimens initiated containing at least two HBV-active antiretrovirals increased by almost 40% after 2007 and, concurrently, the number of regimens containing only one HBV active drug decreased fivefold (Figure 2).

Liver ultrasound results were documented in 24 of 66 HBsAg+ patient charts, with 12 of 24 showing no abnormalities, four showing fatty infiltration, two showing hepatomegaly, and four showing cirrhosis. Liver biopsy results were available in nine of 66 HBsAg+ patient charts, with one of nine having cirrhosis, one having acute drug-induced hepatitis, four having HBV-related liver disease (chronic hepatitis with fibrosis) and three having unrelated liver disease (granuloma, steatohepatitis).

**DISCUSSION**

We found a prevalence of HBV coinfection of 5.5% among a cohort of HIV infected individuals in Northern Alberta. This is lower than the coinfection rate of 6% to 14% previously reported in other areas of low endemicity for both HIV and HBV (8,9) but higher than estimates of 0.1% to 0.5% prevalence of HBV infection in the general population in Canada (2,4).

The majority of coinfected males were Caucasian, while coinfected females were more likely to be of black ethnicity. The strong association with black ethnicity in females may, in part, reflect demographic changes in Alberta’s HIV-positive population. Women from HIV- and HBV-endemic countries in sub-Saharan Africa represent an increasing proportion of HIV cases in Alberta (32,33). A recent large systematic review found an HBsAg prevalence of 15% among those both infected with HIV and from sub-Saharan Africa (34).

Limited access to HBV vaccination remains a significant problem in developing countries, although this is improving (35-37). As of the end of 2007, HBV vaccine had been introduced to 171 countries and an estimated 65% of the world’s population, compared with 3% in 1992 (38). Coverage is highest in the Americas at 88%, while it is 69% in Africa and 30% in Southeast Asia (38). The decline in prevalence of HBV infection from 13.0% between 1982 and 1987 to 4.2% after 1996 may, in part, be attributed to the introduction of immunization programs in Canada. While the HBV vaccine is of limited use in individuals already infected with HIV due to poor immune response, completion of the vaccine series before HIV infection is an important strategy for preventing HBV coinfections (39,40) and is best accomplished through universal vaccination at an early age. In addition, vaccination strategies for preventing HBV coinfection in HIV-infected individuals are being investigated, and benefits may be observed with alterations in vaccination dosing and scheduling (41-43).

Despite similar HCV seroprevalence among HIV patients with and without chronic HBV coinfection, we found that the rates of persistent HCV viremia were significantly higher among HIV patients without chronic HBV infection. Thus, chronic HCV infection appeared to be less likely in HIV patients also chronically infected with HBV. This pattern may reflect a “viral interference” phenomenon whereby one viral infection may predominate in cases of dual- or triple-infected patients (44).

Seventy-five per cent of all deaths in both the HBV-infected and -noninfected groups occurred in those diagnosed with HIV infection before 1997. However, coinfected patients exhibited a higher proportion of death throughout the study period, particularly in the post-HAART era, when HBsAg+ patients were more than two times more likely to die than those who were HBsAg- (23.5% versus 9.5%; P<0.001). No conclusions on liver-related outcomes or causes of mortality can be made due to data limitations.

Data regarding HAART regimens was limited but suggestive of an increased use of two HBV active agents in our cohort after the introduction of the Canadian guidelines in 2007 (Figure 2). This may reflect greater adherence to guidelines for treatment of coinfections but may also reflect the recommendation in HIV treatment guidelines (27) to use Truvada (tenofovir/emtricitabine; Gilead Sciences Inc, USA) as the preferred backbone for HIV therapy in recent years. Analysis of differences in HAART regimens between HIV-HBV coinfected and HIV monoinfected groups was not possible with our data set and is an area for future research.

Our study had several limitations, including limited results of testing for HBV infection. Of our cohort, the 9.3% of patients of unknown HBsAg status were likely never tested for hepatitis B; therefore, unidentified coinfections may exist in our population. The very low overall rate of HBsAg and HBV DNA testing may reflect lack of documentation of these HBV markers in either the hard-copy charts (where available) or in the electronic database. These issues are similar to those reported in a Unites States study, which found that screening of HCV and HBV increased from <20% to >60% from 1998 to 2004, but that screening and vaccination were not universally conducted or, if conducted, not documented despite care guidelines promoting screening and vaccination for HBV (45).

Due to the retrospective nature of the study and the method of data entry at the clinic, serial HBsAg+ results were not available, thus making it impossible to definitively characterize the HBV infection as chronic based on a definition of persistent HBsAg positivity over a minimum six-month period. Although less likely, it is possible that some of the patients who tested HBsAg+ had acute infections that subsequently cleared. Correlation with liver enzyme results would have been helpful to further evaluate this possibility; however, limitations to the data preclude this. In addition, because anti-HBe test results were inconsistently available, HBsAg+/anti-HBc+ individuals were not included in our cohort as coinfected individuals. Serial HBV test results and results of other coinfections (eg, hepatitis D) would also have allowed us to characterize the infectivity of HBV in our population, and to control for potential confounders. The retrospective nature of our study also precluded estimation of HBV incidence rates; this is a potential area for future study.

We were unable to evaluate outcomes, such as complications (eg, cirrhosis) or causes of death, due to limited information in the NPRD.
database and available charts. Patients whose charts were reviewed represented a later cohort of HIV-HBV infected patients and, thus, our observations on the treatment and management of these patients may not be generalizable to the entire cohort of coinfected patients in Alberta.

CONCLUSION

Our data reveal a prevalence of 5.5% of HBV coinfection in Northern Alberta HIV-infected patients, slightly lower than that reported in HIV-infected populations in other developed countries but significantly higher than rates reported in the general population in Canada. Coinfected persons were more likely to be male, and more likely to be deceased. The majority of coinfected males were Caucasian, while coinfected females were more likely to be of black ethnicity, many of whom were from HIV-endemic countries. While our observations suggest that both HBV-specific laboratory testing and treatment improved with time, it also showed a need for enhanced patient follow-up and treatment for HBV in accordance with national guidelines as well as further research regarding the outcomes of HIV-HBV coinfections.

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REFERENCES

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