

# Epidemiologic and clinical features of chronic hepatitis B virus infection in 8 Canadian provinces: a descriptive study by the Canadian HBV Network

Carla S. Coffin MD MSc, Alnoor Ramji MD, Curtis L. Cooper MD MSc, David Miles MD, Karen E. Doucette MD MSc, Philip Wong MD, Edward Tam MD, David K. Wong MD, Alexander Wong MD, Sylvester Ukabam MD, Robert J. Bailey MD, Keith Tsoi MD, Brian Conway MD, Lisa Barrett MD PhD, Tomasz I. Michalak MD PhD, Stephen E. Congly MD MSc, Gerald Minuk MD, Kelly Kaita MD, Erin Kelly MD, Hin Hin Ko MD, Harry L.A. Janssen MD PhD, Julia Uhanova PhD, Brendan C. Lethebe MSc, Sarah Haylock-Jacobs PhD, Mang M. Ma MD, Carla Osiowy PhD, Scott K. Fung MD; for the Canadian HBV Network

## Abstract

**Background:** Published Canadian epidemiologic data on hepatitis B virus (HBV) infection include single-centre studies or are focused on Indigenous populations. We performed a study to characterize the demographic and clinical features, liver disease status and treatment of people with chronic hepatitis B in Canada.

**Methods:** In this descriptive, opportunistic, cross-sectional study, available data for people known to be monoinfected with HBV were collected by the Canadian HBV Network from existing clinical databases, with support from the National Microbiology Laboratory, Public Health Agency of Canada. Data were collected in all provinces with the exception of New Brunswick and Prince Edward Island. We analyzed the data using parametric and nonparametric statistical methods, with a significance level of  $p < 0.05$ .

**Results:** In the 9380 unique patient records reviewed, the median age was 48 years, and 5193 patients (55.4%) were male. Ethnicity information was available for 7858 patients, of whom 5803 (73.8%) were Asian, 916 (11.6%) were black and 914 (11.6%) were white. Most of those tested (5556/6796 [81.8%]) were negative for HBV e-antigen, and most of those with fibrosis data (3481/4260 [81.7%]) had minimal liver fibrosis, with more advanced fibrosis noted in older people (> 40 yr). Of the 980 patients with genotype data, 521 (53.2%) had genotype B or C infection. Most of the 9241 patients with known confirmed treatment status received tenofovir disoproxil fumarate (1655 [17.9%]), lamivudine (1434 [15.5%]) or entecavir (548 [5.9%]).

**Interpretation:** Based on available data, Canadian patients with chronic hepatitis B are predominantly Asian and negative for HBV e-antigen, and have genotype B or C infection. Interprovincial variations were noted in antiviral treatment regimen. This multicentre nationwide study provides data regarding patients with chronic hepatitis B and may inform future studies on the epidemiologic features of HBV infection in Canada.

The hepatitis B virus (HBV) is a major human pathogen, with 257 million chronic carriers worldwide — especially in Asia and Sub-Saharan Africa — who remain at risk for cirrhosis, liver failure and hepatocellular carcinoma.<sup>1,2</sup> There are 8 major HBV genotypes, with a specific geographic and ethnic distribution, which have been linked to disease outcomes and response to antiviral therapy.<sup>3</sup> Hepatitis B is reportable to the Canadian Notifiable Disease Surveillance System,<sup>4</sup> yet reliable epidemiologic data are limited owing to the lack of standardized reporting practices and with reporting of only acutely infected people. A Public Health Agency of Canada report on hepatitis B and C in Canada in 2013 lacks data from 2 provinces (Newfoundland and Labrador, Prince Edward Island) and 2 territories

(Northwest Territories, Nunavut).<sup>5</sup> Data from the Public Health Agency of Canada show lower estimates of chronic hepatitis B prevalence and incidence rate in British Columbia than those reported by the BC Centre for Disease Control owing to disparate reporting practices.<sup>6</sup> In a report from the Canadian Liver Foundation, it was estimated that 250 000–

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**Correspondence to:** Carla Coffin, cscoffin@ucalgary.ca

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460 000 Canadians have chronic hepatitis B, with highest rates found in urban centres.<sup>7</sup> The prevalence of acute and chronic HBV infection has declined in Canadian-born people with the advent of universal childhood vaccination, in the mid-1990s. A seroepidemiologic survey of Arctic Indigenous populations, historically considered to be hyperendemic for HBV infection, showed a nonendemic prevalence of 1.2%, yet a substantial number of people were found to have lower than expected vaccine-based immunity, with unknown clinical characteristics.<sup>8</sup> The 2016 Canada long-term census survey showed considerable ethnocultural diversity within Canada, with 1 in 5 Canadians (21.9%) either a landed immigrant or a permanent resident.<sup>9</sup> Importantly, most newcomers were from Asia (61%), with Africa as the second-highest (13.4%) source for recent immigrants, especially highly HBV-endemic countries. Most new Canadians live in established home-country communities in large urban centres (e.g., Toronto, Montréal, Vancouver). Interestingly, there is increasing settlement in less-populous regions (e.g., prairie and Atlantic provinces) owing to economic factors such as job growth in the energy sector or under special provincial nominee programs to encourage population growth (e.g., Manitoba “growing through immigration” [www.immigratemanitoba.com]). This evolving Canadian demographic landscape highlights the need for updated large-scale epidemiological data on HBV infection and the associated disease burden in Canada.

The current study was an opportunistic, cross-sectional descriptive study using available clinical data for people with chronic hepatitis B in Canada. Our study aim was to characterize the demographic and clinical features, virological outcomes and treatment of the more than 9000 patients with hepatitis B followed in diverse academic and community care clinics nationwide. The data collected will facilitate further hypothesis-generating studies to be conducted on hepatitis B in Canada.

## Methods

### Study design

The Canadian HBV Network was formed in 2016 and received official endorsement by the Canadian Association for the Study of the Liver in 2017. It includes 21 academic and community clinical care centres for patients with hepatitis B in 8 Canadian provinces, supported by provincial as well as national reference laboratories (i.e., National Microbiology Laboratory, Public Health Agency of Canada) (Box 1). The network’s mission is to 1) improve treatment and control of hepatitis B in Canada, 2) advance the understanding of HBV infection and disease natural history, 3) promote collaboration on national hepatitis B studies and databases and 4) enhance and inform research and educational opportunities on HBV infection and its consequences.

In this descriptive, opportunistic, cross-sectional study, we collated data for patients positive for serum HBV surface antigen from information submitted by participating sites to the data coordinating centre (University of Calgary). Data

#### Box 1: The 21 sites involved in the Canadian HBV Network

##### Sites that provided clinical data or laboratory data or both to the current study

1. Department of Medicine, University of British Columbia, Vancouver, BC (1323 patients)
2. LAIR Centre, Vancouver, BC (194 patients)
3. Vancouver Infectious Diseases Centre, BC (18 patients)
4. Division of Infectious Diseases, University of Alberta, Edmonton, Alta. (241 patients)
5. Bailey Health Clinic, Edmonton, Alta. (87 patients)
6. Liver Unit, University of Calgary, Calgary, Alta. (1486 patients)
7. Division of Infectious Diseases, University of Saskatchewan, Regina Qu’Appelle Health Region, Sask. (112 patients)
8. Regina General Hospital, Regina, Sask. (67 patients)
9. Department of Internal Medicine, University of Manitoba, Winnipeg, Man. (1051 patients)
10. National Microbiology Laboratory, Public Health Agency of Canada, Winnipeg, Man. (laboratory testing)
11. McMaster University/St. Joseph’s Healthcare, Hamilton, Ont. (80 patients)
12. University of Toronto, Toronto Centre for Liver Disease, Toronto, Ont. (3710 patients)
13. University of Ottawa, Ottawa Hospital Research Institute, Ottawa, Ont. (847 patients)
14. McGill University, Montréal, Que. (132 patients)
15. Dalhousie University, Halifax, NS (19 patients)
16. Memorial University, St. John’s, Nfld. (13 patients)

##### Sites with recent data-sharing agreements or provincial/hospital laboratories that did not contribute data to the current study

1. Providence Health Care (Medical Microbiology)/BC Centre for Disease Control, Vancouver, BC
2. Division of Gastroenterology and Hepatology, University of Alberta, Edmonton, Alta.
3. Western University, London, Ont.
4. Toronto Liver Centre, North York General Hospital, Toronto, Ont.
5. Laval University, Sainte-Foy, Que.

sources included manual review of physician paper charts at local sites and submission to a Web-based portal, or data downloads from existing stand-alone clinical databases (i.e., Microsoft Access or Excel; Vancouver, Calgary, Winnipeg, Toronto, Ottawa). In Alberta, administrative data on demographic characteristics and laboratory data were collected for 3 participating sites from the Alberta Health Services data repository, as previously described.<sup>10</sup> All patients were seen in clinic after January 2012. Cross-sectional data analysis was done on the most recent patient record submitted. Fifteen sites submitted clinical data from 8 provincial health jurisdictions. The data were downloaded and merged into a single Excel spreadsheet or copied manually, verified and analyzed in R 3.5.2 (R Foundation for Statistical Computing) by 2 investigators (S.H.-J. and B.C.L.) for completeness and quality. The Canadian HBV Network coordinator (S.H.-J.) has performed site visits and monitoring checks for

accuracy. All data entry coordinators were trained, and they follow a database guidance document.

### Data elements

We reviewed available records of adult patients with known chronic hepatitis B. Patients were excluded from analysis if they were coinfecting with hepatitis C virus, hepatitis delta virus or HIV. Available data elements included demographic characteristics (age, sex and, in some cases, ethnicity) and most recent tests for HBV DNA, HBV e-antigen (HBeAg), HBV genotype, alanine aminotransferase, liver stiffness measurement via transient elastography (FibroScan, Echosens) and antiviral therapy history. Laboratory testing was performed through provincial diagnostic laboratories as per standard of care. Hepatitis B virus genotyping is not done in routine clinical practice; however, some data were available retrospectively (including a data subset from a previously published study<sup>11</sup>) or were determined prospectively for this study at the National Microbiology Laboratory by in-house direct population sequencing and phylogenetic analysis and/or reverse hybridization assay (Inno-LiPa, Innogenetics).

### Statistical analysis

We summarized measured variables with mean, count and 95% confidence interval; comparisons were done with *t* tests. We summarized categorical variables using proportion, count and 95% confidence interval; comparisons were done with  $\chi^2$  tests. The comparisons performed were between the target group and the national average, excluding the target group. For example, we conducted the age comparison for BC by comparing the ages in BC to the ages in the 7 other provinces. Missing values were considered as their own group for categorical variables, and the counts of missing data were reported for measured variables. We conducted statistical analysis using R 3.5.2.

### Ethics approval

All data received were anonymous and collected under a protocol approved by the University of Calgary Conjoint Health Research Ethics Board, with subsite ethics and legal agreements for data sharing between sites within the Canadian HBV Network. A waiver of consent was obtained from local research ethics boards or under section 50 of the *Health Information Act* of Alberta, or both, or patients provided signed informed consent to participate.

## Results

### Patient characteristics and genotypes

Clinical and demographic data were submitted for 9380 patients nationwide (Table 1). Most patients (5193 [55.4%]) were men. The median age was 48 years. Ethnicity data were available for 7858 patients, of whom 5803 (73.8%) were Asian, 916 (11.6%) black, 914 (11.6%) white, and 225 (2.9%) other (including Indigenous/First Nations). Just over half (521 [53.2%]) of the 980 patients with available HBV genotype data had genotype B or C infection (Table 2), as expected given the predominantly Asian origin of the patients.

### Hepatitis B virus disease status, fibrosis and antiviral therapy

Most patients with available data (5556/6796 [81.8%]) tested HBeAg negative, and 2319/2820 (82.2%) had HBV DNA levels less than 2000 IU/mL (10 000 virus genome copies/mL) (Table 1). A substantial proportion (779/4260 [18.3%]) had documented stage 2 or greater fibrosis (METAVIR score) based on liver stiffness measurement (FibroScan score > 7.3 kPa, data not shown). The most common antiviral therapy used at any time was tenofovir disoproxil fumarate (1655 patients [17.6%]), followed by lamivudine (1434 [15.3%]) and entecavir (548 [5.8%]). Interprovincial differences were noted in antiviral treatment regimen, with a higher proportion of patients in BC and Ontario than in other provinces receiving first-generation nucleos(t)ide analogs (i.e., lamivudine).

### Clinical characteristics according to hepatitis B virus genotype

Compared to all other genotypes, patients with HBV genotype C were more likely to be HBeAg positive, have more advanced liver fibrosis (> stage 2) and receive nucleos(t)ide antiviral therapy.

### Demographic and clinical characteristics according to sex and age

Analysis of available clinical data for sex- and age-related differences showed that patients aged 40 years or more were more likely than those aged less than 40 to show increased liver fibrosis (> stage 2), whereas those aged less than 40 were more likely than those aged 40 or more to be HBeAg positive (Table 3).

## Interpretation

In this cross-sectional, retrospective descriptive study of 9380 patients from 8 provincial health care jurisdictions, both academic and community clinics, we found that the majority of people with chronic hepatitis B were older, male, Asian and HBeAg-negative, and had HBV genotype B or C infection. Notable proportions had an HBV DNA level greater than 2000 IU/mL and stage 3 or higher fibrosis, which are important indicators regarding the need for antiviral therapy. Many patients received long-term therapy, but there were interprovincial differences between nucleos(t)ide analogue use (i.e., lamivudine v. tenofovir disoproxil/entecavir), likely owing to historic provincial disparities in access to second-generation therapy. Among 980 people with available genotype data, we noted significant differences between HBV genotype C clinical characteristics (higher fibrosis stage, higher alanine aminotransferase levels and higher rates of HBeAg positivity) compared to other genotypes.

These findings are consistent with those from studies of HBV genotype and more severe liver fibrosis in HBV genotype C (v. B) from Asian cohort studies.<sup>12</sup> Our data also provide important comparisons across all reported genotypes in 1 country, which is not feasible in jurisdictions with less ethnic

**Table 1: Demographic characteristics, liver/hepatitis B virus test results, fibrosis and antiviral therapy of Canadians with chronic hepatitis B, by province**

Variable	Province; no. (%) of patients* (95% CI)								
	Total n = 9380	BC n = 1535	Alta. n = 1814	Sask. n = 179	Man. n = 1051	Ont. n = 4637	Que. n = 132	NS n = 19	NL n = 13
Age, yr, mean (median)	49.1 (48) (48.80–49.36)	52.7‡ (53) (52.01–53.35)	45.6‡ (44) (45.01–46.21)	44.3 (41) (42.22–46.42)	46.6‡ (46) (45.76–47.37)	50.0‡ (50) (49.62–50.44)	49.6 (47) (47.28–51.84)	40.9‡ (36) (34.47–47.32)	44.5 (50) (37.73–51.20)
Male sex	5193 (55.4) (54.35–56.37)	864 (56.3) (53.76–58.78)	919 (50.7)‡ (48.33–52.99)	106 (59.2) (51.62–66.42)	558 (53.1) (50.02–56.14)	2642 (57.0)‡ (55.54–58.41)	81 (61.4) (52.47–69.59)	13 (68.4) (43.50–86.44)	10 (76.9) (45.98–93.84)
Born in Canada	80 (0.8) (0.68–1.07)	5 (0.3)‡ (0.12–0.81)	4 (0.2)‡ (0.07–0.60)	3 (1.7)‡ (0.43–5.21)	0 (0.0)‡	68 (1.5)‡ (1.15–1.87)	0 (0.0)‡	0 (0.0)‡	0 (0.0)
Immigrant status missing	7063 (75.3) (74.41–76.17)	939 (61.2)‡ (58.68–63.61)	1398 (77.1)‡ (75.05–78.97)	84 (46.9)‡ (39.49–54.50)	693 (65.9)‡ (62.97–68.79)	3797 (81.9)‡ (80.74–82.98)	132 (100.0)‡ (96.48–100.00)	7 (36.8)‡ (17.23–61.37)	13 (100.0) (71.66–100.00)
Asian	5803 (61.9) (60.87–62.85)	1120 (73.0)‡ (70.65–75.16)	1109 (61.1)‡ (58.84–63.38)	97 (54.2)‡ (46.60–61.59)	252 (24.0)‡ (21.45–26.70)	3198 (69.0)‡ (67.61–70.29)	13 (9.8)‡ (5.56–16.57)	11 (57.9) (33.97–78.88)	3 (23.1)‡ (6.16–54.02)
Black	916 (9.8) (9.18–10.39)	16 (1.0)‡ (0.62–1.73)	272 (15.0)‡ (13.40–16.74)	34 (19.0)‡ (13.68–25.67)	76 (7.2)‡ (5.77–9.01)	498 (10.7)‡ (9.87–11.67)	17 (12.9)‡ (7.90–20.09)	2 (10.5) (1.84–34.54)	1 (7.7)‡ (0.40–37.91)
White	914 (9.7) (9.16–10.37)	32 (2.1)‡ (1.45–2.97)	94 (5.2)‡ (4.23–6.33)	24 (13.4)‡ (8.95–19.49)	21 (2.0)‡ (1.27–3.09)	664 (14.3)‡ (13.33–15.37)	79 (59.8)‡ (50.94–68.17)	0 (0.0)	0 (0.0)‡
Ethnicity missing	1522 (16.2) (15.49–16.99)	343 (22.3)‡ (20.30–24.53)	305 (16.8)‡ (15.14–18.63)	16 (8.9)‡ (5.36–14.35)	690 (65.6)‡ (62.68–68.51)	144 (3.1)‡ (2.63–3.66)	12 (9.1)‡ (5.00–15.67)	4 (21.0) (6.97–46.10)	8 (61.5)‡ (32.28–84.87)
HBeAg-positive	1240 (13.2) (12.54–13.93)	232 (15.1)‡ (13.38–17.03)	217 (12.0)‡ (10.52–13.57)	30 (16.8)‡ (11.76–23.23)	27 (2.6)‡ (1.73–3.77)	697 (15.0)‡ (14.02–16.10)	30 (22.7)‡ (16.08–31.00)	3 (15.8) (4.17–40.49)	4 (30.8) (10.36–61.12)
HBeAg status missing	2584 (27.5) (26.65–28.47)	713 (46.4)‡ (43.93–48.98)	680 (37.5)‡ (35.26–39.77)	14 (7.8)‡ (4.50–13.03)	745 (70.9)‡ (68.02–73.60)	421 (9.1)‡ (8.28–9.95)	2 (1.5)‡ (0.26–5.92)	4 (21.0) (6.97–46.10)	5 (38.5) (15.13–67.72)
HBV DNA level < 2000 IU/mL	2319 (24.7) (23.85–25.61)	881 (57.4)‡ (54.87–59.88)	582 (32.1)‡ (29.95–34.3)	87 (48.6)‡ (41.12–56.15)	480 (45.7)‡ (42.63–48.74)	252 (5.4)‡ (4.81–6.14)	27 (20.4)‡ (14.13–28.54)	10 (52.6)‡ (29.50–74.79)	0 (0.0)
HBV DNA level missing	6560 (69.9) (68.99–70.86)	654 (42.6)‡ (40.12–45.13)	731 (40.3)‡ (38.04–42.60)	92 (51.4)‡ (43.85–58.88)	571 (54.3)‡ (51.26–57.37)	4385 (94.6)‡ (93.86–95.19)	105 (79.5)‡ (71.46–85.87)	9 (47.4)‡ (25.21–70.50)	13 (100.0) (71.66–100.00)
Alanine transaminase level, U/L, mean (median)†	40.1 (27) (37.43–42.86)	33.4 (25)‡ (30.67–36.10)	38.0 (28) (33.11–42.81)	36.9 (27) (27.00–46.89)	51.3‡ (27) (42.32–60.32)	39.2 (30) (36.34–41.98)	43.8 (28) (33.43–54.10)	30.1‡ (26) (23.66–36.58)	NA
Stage F3–4 fibrosis	324 (3.4) (3.10–3.85)	103 (6.7)‡ (5.53–8.11)	113 (6.2)‡ (5.18–7.47)	7 (3.9) (1.72–8.21)	4 (0.4)‡ (0.12–1.04)	96 (2.1)‡ (1.69–2.53)	1 (0.8)‡ (0.04–4.77)	0 (0.0)‡	0 (0.0)‡
Fibrosis data missing	5120 (54.6) (53.57–55.60)	102 (6.6)‡ (5.47–8.04)	962 (53.0)‡ (50.70–55.35)	93 (52.0) (44.40–59.43)	1004 (95.5)‡ (94.05–96.66)	2807 (60.5)‡ (59.11–61.94)	120 (90.9)‡ (84.33–95.00)	19 (100.0)‡ (79.08–100.00)	13 (100.0)‡ (71.66–100.00)
Lamivudine prescription (at any time)	1434 (15.3) (14.57–16.04)	386 (25.1)‡ (23.01–27.41)	71 (3.9)‡ (3.09–4.94)	1 (0.6)‡ (0.03–3.55)	14 (1.3)‡ (0.76–2.28)	958 (20.6)‡ (19.51–21.86)	2 (1.5)‡ (0.26–5.92)	1 (5.3)‡ (0.28–28.11)	1 (7.7) (0.40–37.91)
Tenofovir disoproxil prescription (at any time)	1655 (17.6) (16.88–18.43)	376 (24.5)‡ (22.38–26.74)	408 (22.5)‡ (20.60–24.50)	17 (9.5)‡ (5.80–15.01)	35 (3.3)‡ (2.36–4.65)	788 (17.0) (15.93–18.11)	30 (22.7) (16.08–31.00)	1 (5.3)‡ (0.28–28.11)	0 (0.0)
Entecavir prescription (at any time)	548 (5.8) (5.38–6.34)	80 (5.2)‡ (4.18–6.48)	199 (11.0)‡ (9.59–12.52)	0 (0.0)‡	9 (0.8) (0.42–1.68)	259 (5.6) (4.95–6.30)	1 (0.8)‡ (0.04–4.77)	0 (0.0)‡	0 (0.0)
Treatment data missing	139 (1.5) (1.25–1.75)	0 (0.0)‡	0 (0.0)‡	66 (36.9)‡ (29.89–44.43)	0 (0.0)‡	70 (1.5) (1.19–1.91)	0 (0.0)‡	3 (15.8)‡ (4.17–40.49)	0 (0.0)

Note: CI = confidence interval, HBeAg = hepatitis B e-antigen, HBV = hepatitis B virus, NA = not available.  
 \*Except where noted otherwise.  
 †Data missing for 5487 patients overall: 112 from British Columbia, 1378 from Alberta, 74 from Saskatchewan, 32 from Manitoba, 3875 from Ontario, 14 from Quebec and 2 from Nova Scotia.  
 ‡p < 0.05 for difference between the value for a given province, and the national total minus the value for that province.

diversity. Most patients tested in the current study were HBeAg negative. Being negative for HBeAg may be due to the presence of precore and basal core promoter mutations in the viral genome and is associated with the development of cirrhosis or hepatocellular carcinoma, or both, in Asian cohort studies.<sup>13</sup> Data on HBV included results of standard clinical tests (HBV DNA level and HBeAg status),<sup>11,13</sup> as analysis for muta-

tions associated with HBeAg-negative disease is not available or usually needed in routine clinical practice to assess need for treatment. Most of the data were contributed for patients residing in urban centres and from tertiary referral clinics in regions with high rates of immigration.<sup>11</sup> Universal childhood vaccination in Nunavut has led to a significant decline in the prevalence of chronic infection in Inuit populations.<sup>8</sup> This

**Table 2: Demographic characteristics, liver/hepatitis B virus test results, fibrosis and treatment of Canadians with chronic hepatitis B, by hepatitis B virus genotype**

Variable	Genotype; no. (%) of patients* (95% CI)					
	A-E n = 980	A n = 218	B n = 278	C n = 243	D n = 157	E n = 84
Age, yr, mean (median)	46.2 (44) (45.36–47.01)	44.8 (43) (43.00–46.70)	48.3‡ (47) (46.81–49.77)	49.2‡ (49) (47.61–50.81)	44.1‡ (42) (41.90–46.27)	37.9‡ (37.5) (36.14–39.67)
Male sex	533 (54.4) (51.21–57.53)	121 (55.5) (48.64–62.17)	143 (51.4) (45.41–57.43)	127 (52.3) (45.79–58.66)	95 (60.5) (52.38–68.12)	47 (56.0) (44.72–66.63)
Born in Canada	3 (0.3) (0.08–0.97)	0 (0.0)‡	1 (0.4) (0.02–2.30)	2 (0.8)‡ (0.14–3.26)	0 (0.0)	0 (0.0)
Immigrant status missing	678 (69.2) (66.17–72.04)	120 (55.0)‡ (48.18–61.73)	208 (74.8) (69.21–79.73)	194 (79.8)‡ (74.12–84.58)	105 (66.9) (58.86–74.05)	51 (60.7) (49.43–71.01)
Asian	389 (39.7) (36.63–42.84)	78 (35.8)‡ (29.50–42.57)	161 (57.9)‡ (51.86–63.74)	119 (49.0)‡ (42.55–55.43)	28 (17.8)‡ (12.37–24.92)	3 (3.6)‡ (0.93–10.81)
Black/African/Caribbean	128 (13.1) (11.05–15.37)	43 (19.7)‡ (14.78–25.76)	1 (0.4)‡ (0.02–2.30)	5 (2.0)‡ (0.76–5.00)	20 (12.7)‡ (8.14–19.22)	59 (70.2)‡ (59.13–79.47)
White	104 (10.6) (8.79–12.75)	6 (2.8)‡ (1.12–6.18)	36 (12.9)‡ (9.35–17.61)	38 (15.6)‡ (11.43–20.96)	22 (14.0)‡ (9.18–20.66)	2 (2.4)‡ (0.41–9.14)
Ethnicity missing	333 (34.0) (31.03–37.05)	91 (41.7)‡ (35.17–48.61)	80 (28.8)‡ (23.61–34.55)	81 (33.3)‡ (27.51–39.69)	63 (40.1)‡ (32.48–48.26)	18 (21.4)‡ (13.52–32.00)
HBeAg positive	140 (14.3) (12.19–16.67)	23 (10.6)‡ (6.95–15.60)	33 (11.9) (8.43–16.40)	61 (25.1)‡ (19.88–31.13)	12 (7.6)‡ (4.19–13.27)	11 (13.1) (7.03–22.64)
HBeAg status missing	233 (23.8) (21.17–26.59)	64 (29.4)‡ (23.50–35.96)	62 (22.3) (17.64–27.75)	53 (21.8)‡ (16.89–27.64)	33 (21.0)‡ (15.10–28.39)	21 (25.0) (16.47–35.85)
HBV DNA level < 2000 IU/mL	341 (34.8) (31.83–37.88)	91 (41.7)‡ (35.17–48.61)	80 (28.8)‡ (23.61–34.55)	76 (31.3) (25.58–37.57)	62 (39.5) (31.88–47.62)	32 (38.1) (27.91–49.38)
HBV DNA level missing	554 (56.5) (53.36–59.65)	121 (55.5)‡ (48.64–62.17)	168 (60.4)‡ (54.40–66.17)	140 (57.6) (51.12–63.86)	80 (51.0) (42.90–58.97)	45 (53.6) (42.41–64.41)
Alanine transaminase level, U/L, mean (median)†	38.7 (28) (33.09–44.23)	34.6 (27) (29.84–39.37)	32.0‡ (28) (28.80–35.16)	52.7 (28) (31.06–74.31)	28.6‡ (24) (25.42–31.73)	51.8 (31) (30.80–72.73)
Stage F3–4 fibrosis	27 (2.8) (1.86–4.04)	3 (1.4)‡ (0.36–4.30)	8 (2.9)‡ (1.34–5.81)	11 (4.5)‡ (2.40–8.18)	2 (1.3) (0.22–5.00)	3 (3.6) (0.93–10.81)
Fibrosis data missing	636 (64.9) (61.81–67.87)	166 (76.1)‡ (69.82–81.53)	154 (55.4)‡ (49.34–61.30)	154 (63.4)‡ (56.94–69.38)	105 (66.9) (58.86–74.05)	57 (67.8) (56.66–77.40)
Lamivudine prescription (at any time)	28 (2.8) (1.94–4.16)	1 (0.4)‡ (0.02–2.92)	11 (4.0) (2.09–7.17)	11 (4.5) (2.40–8.18)	5 (3.2) (1.18–7.66)	0 (0.0)
Tenofovir disoproxil prescription (at any time)	155 (15.8) (13.62–18.29)	18 (8.2)‡ (5.11–12.94)	36 (12.9) (9.35–17.61)	68 (28.0)‡ (22.53–34.15)	25 (15.9) (10.76–22.81)	8 (9.5) (4.50–18.41)
Entecavir prescription (at any time)	35 (3.6) (2.54–4.99)	2 (0.9)‡ (0.16–3.63)	15 (5.4) (3.16–8.93)	12 (4.9) (2.70–8.68)	5 (3.2) (1.18–7.66)	1 (1.2) (0.06–7.37)
Treatment data missing	13 (1.3) (0.74–2.32)	0 (0.0)‡	5 (1.8) (0.66–4.39)	5 (2.0) (0.76–5.00)	2 (1.3) (0.22–5.00)	1 (1.2) (0.06–7.37)

Note: CI = confidence interval, HBeAg = hepatitis B e-antigen, HBV = hepatitis B virus.  
 †Data missing for 381 patients: 70 with genotype A, 105 with genotype B, 103 with genotype C, 70 with genotype D and 33 with genotype E.  
 ‡p < 0.05 for difference between a given genotype, and the total for all other genotypes minus the value for that genotype.

situation is mirrored in the current study, which did not show a substantial number of people with chronic hepatitis B in Canadian Indigenous populations (< 1% reported, data not shown). The baseline demographic and clinical data reported

in the present study are similar to those reported by the Hepatitis B Research Network, a cooperative network of investigators from the United States and Canada, which found that 50% of adults were men, the median age was 42 years, and

**Table 3: Demographic characteristics, liver/hepatitis B virus test results, fibrosis and treatment of Canadians with chronic hepatitis B, by sex and age**

Variable	No. (%) of patients* (95% CI)			
	Men n = 5193	Women n = 4187	Age < 40 yr n = 2621	Age ≥ 40 yr n = 6759
Age, yr, mean (median)	49.9‡ (50) (49.56–50.30)	48.0‡ (47) (47.59–48.46)	32.7§ (33) (32.48–32.85)	55.4§ (54) (55.19–55.7)
Male sex	5193 (100.0)‡ (99.91–100.00)	–‡	1260 (48.1)§ (46.15–50.01)	3933 (58.2)§ (57.00–59.37)
Born in Canada	47 (0.9) (0.67–1.21)	33 (0.8) (0.55–1.12)	22 (0.8)§ (0.54–1.29)	58 (0.8)§ (0.66–1.12)
Immigrant status missing	3943 (75.9) (74.74–77.08)	3120 (74.5) (73.16–75.83)	1922 (73.3)§ (71.59–75.01)	5141 (76.1)§ (75.02–77.07)
Asian	3147 (60.6)‡ (59.25–61.93)	2656 (63.4)‡ (61.95–64.89)	1510 (57.6)§ (55.69–59.51)	4293 (63.5)§ (62.35–64.66)
Black/African/Caribbean	516 (9.9)‡ (9.14–10.79)	400 (9.6)‡ (8.69–10.49)	433 (16.5)§ (15.13–18.01)	483 (7.1)§ (6.55–7.79)
White	562 (10.8)‡ (10.00–11.71)	352 (8.4)‡ (7.59–9.30)	168 (6.4)§ (5.52–7.43)	746 (11.0)§ (10.30–11.81)
Ethnicity missing	834 (16.1)‡ (15.08–17.09)	688 (16.4)‡ (15.33–17.60)	427 (16.3)§ (14.91–17.77)	1095 (16.2)§ (15.33–17.11)
HBeAg positive	659 (12.7)‡ (11.80–13.63)	581 (13.9)‡ (12.85–14.97)	578 (22.0)§ (20.49–23.70)	662 (9.8)§ (9.10–10.53)
HBeAg status missing	1384 (26.6)‡ (25.46–27.88)	1200 (28.7)‡ (27.30–30.06)	747 (28.5)§ (26.79–30.28)	1837 (27.2)§ (26.12–28.26)
HBV DNA level < 2000 IU/mL	1235 (23.8) (22.63–24.97)	1084 (25.9) (24.57–27.25)	681 (26.0)§ (24.32–27.72)	1638 (24.2)§ (23.22–25.28)
HBV DNA level missing	3673 (70.7) (69.47–71.96)	2887 (69.0) (67.52–70.35)	1776 (67.8)§ (65.93–69.54)	4784 (70.8)§ (69.68–71.86)
Alanine transaminase level, U/L, mean (median)†	48.2‡ (31) (43.45–52.86)	30.5‡ (22) (28.66–32.43)	42.5 (27) (36.27–48.67)	39.3 (27) (36.34–42.18)
Stage F3–4 fibrosis	232 (4.5) (3.93–5.07)	92 (2.2) (1.78–2.70)	37 (1.4)§ (1.01–1.96)	287 (4.2)§ (3.78–4.76)
Fibrosis data missing	2753 (53.0) (51.64–54.38)	2367 (56.5) (55.01–58.04)	1601 (61.1)§ (59.18–62.95)	3519 (52.1)§ (50.86–53.26)
Lamivudine prescription (at any time)	926 (17.8)‡ (16.81–18.91)	508 (12.1)‡ (11.17–13.17)	217 (8.3)§ (7.27–9.42)	1217 (18.0)§ (17.10–18.95)
Tenofovir disoproxil prescription (at any time)	1087 (20.9)‡ (19.84–22.07)	568 (13.6)‡ (12.55–14.65)	350 (13.4)§ (12.09–14.73)	1305 (19.3)§ (18.38–20.27)
Entecavir prescription (at any time)	350 (6.7)‡ (6.08–7.46)	198 (4.7)‡ (4.12–5.43)	93 (3.5)§ (2.89–4.35)	455 (6.7)§ (6.15–7.36)
Treatment data missing	90 (1.7)‡ (1.40–2.14)	49 (1.2)‡ (0.88–1.56)	35 (1.3)§ (0.95–1.87)	104 (1.5)§ (1.26–1.87)

Note: CI = confidence interval, HBeAg = hepatitis B e-antigen, HBV = hepatitis B virus.  
 \*Except where noted otherwise.  
 †Data missing for 3078 men, 2422 women, 1556 patients aged less than 40 years and 3944 patients aged 40 years or more.  
 ‡p < 0.05 for difference between men and women.  
 §p < 0.05 for difference between patients aged less than 40 years and those aged 40 years or more.

74% of the cohort were HBeAg-negative.<sup>14</sup> In addition, our HBV genotype data are similar to those reported by the Hepatitis B Research Network (mainly B [39%] and C [33%], followed by A [18%], D [8%] and E [3%]), which also

reflected the largely foreign-born population of Asian and Africans.<sup>14</sup> However, the report did not include treatment information, and all fibrosis data were based on liver histopathological examination or serum-based markers of liver

fibrosis (FibroTest or aspartate-aminotransferase-to-platelet ratio index [APRI]), as transient elastography was approved in the United States only relatively recently.

Our data highlight regional variations in HBV disease management that can inform policy-makers of areas where best practice can be implemented. Under the *Canada Health Act*, universal health care is publicly funded, and all Canadians should have reasonable access to medically necessary services.<sup>15</sup> However, “medically necessary” is not specifically defined in the act, and health care spending and services are implemented by provincial/territorial government health care boards. This has led to substantial variation in public access to prescription medications and even in childhood HBV vaccination schedules. For example, Alberta and Quebec have included second-generation nucleos(t)ide analogues as first-line therapy for HBV for almost a decade, yet these drugs were not approved on public drug benefits list in Ontario until recently and are still not covered by BC pharmacare unless patients have advanced fibrosis. The World Health Organization recommends that all infants receive their first dose of HBV vaccine as soon as possible after birth, as the greatest risk of chronic hepatitis B is within the first 5 years of life.<sup>1</sup> Only 5 provinces (BC, Alberta [since September 2018], Quebec, New Brunswick and Prince Edward Island) and the territories offer universal infant HBV vaccination. Universal HBV screening during pregnancy is mandatory in many provinces but is only “strongly recommended” in Ontario, although it is widely implemented as good clinical practice.<sup>16</sup> Although there has been considerable public health focus on hepatitis C and HIV infection in Canada, hepatitis B remains underappreciated and underfunded, and there is a lack of awareness of disease burden and complexity of management, and a need for better treatments.<sup>17</sup> According to the Ontario Burden of Infectious Diseases Study, HBV was the fifth-ranked pathogen causing substantial health-adjusted life-years lost.<sup>18</sup> Modelling data showed that immigrants with hepatitis B in Canada lost an average of 4.6 life-years, with higher lifetime risk of end-stage liver disease.<sup>19</sup> Current immigration trends and aging of the Canadian population mean that it is imperative that uniform, structural programs be developed for identification (screening), education and access to care for Canadians at risk for HBV infection and possible life-threatening consequences. The Canadian HBV Network is committed to conducting longitudinal, prospective and rigorous epidemiological studies on HBV infection in Canada. Our network is working to establish global partnerships such as with the International Coalition to Eliminate HBV (<https://www.ice-hbv.org>), which includes more than 50 scientists from 21 countries.

### Limitations

The data may not represent patients who are not engaged in care. In addition, to date, there are limited genotype data from participating sites in BC, Ontario and Quebec, which have large populations of patients with hepatitis B. However, ongoing robust prospective data collection by Canadian

HBV Network investigators will help address these data gaps in future studies. The data completeness may be affected by underreporting and referral bias (i.e., patients with symptoms of advanced liver disease are more likely to be referred for treatment by their primary care provider than asymptomatic patients); however, patients with cirrhosis and hepatocellular carcinoma are often asymptomatic, and many people are referred too late or present with nonspecific symptoms.<sup>20</sup> All study investigators highlight consistent or increasing number of patients in their clinical practice needing treatment.

### Conclusion

The current study provides data on the demographic and clinical characteristics, virological outcomes and antiviral treatment in people with chronic hepatitis B in Canada. Hepatitis B is an ongoing and important cause of morbidity and mortality in Canada. Country-specific epidemiological data will inform the development of programs to eliminate hepatitis B as a public health threat and ultimately achieve a cure for the disease. The data from this study will inform global research on the impact of HBV, even in wealthy countries such as Canada.

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**Affiliations:** Cumming School of Medicine (Coffin, Lethebe, Congly, Haylock-Jacobs), University of Calgary, Calgary, Alta.; Division of Gastroenterology (Ramji, Ko), Department of Medicine, Faculty of Medicine, University of British Columbia, Vancouver, BC; Division of Infectious Diseases (Cooper), University of Ottawa, Ottawa Hospital Research Institute; Division of Gastroenterology (Kelly), Department of Medicine, University of Ottawa, Ottawa Hospital Research Institute, Ottawa, Ont.; Department of Medicine (Doucette, Ma), University of Alberta, Edmonton, Alta.; Division of Gastroenterology (P. Wong), Department of Medicine, McGill University, Royal Victoria Hospital, Montréal, Que.; LAIR Centre (Tam), Vancouver, BC; Department of Medicine (D. Wong, Janssen, Fung), University of Toronto, Toronto Centre for Liver Disease, Toronto, Ont.; Department of Medicine (A. Wong), University of Saskatchewan; Regina General Hospital (Ukabam), Regina, Sask.; Bailey Health Clinic (Bailey), Edmonton, Alta.; Department of Medicine (Tsoi), McMaster University, St. Joseph's Healthcare, Hamilton, Ont.; Vancouver Infectious Diseases Centre (Conway), Vancouver, BC; Division of Infectious Diseases (Barrett), Dalhousie University, Halifax, NS; Faculty of Medicine (Michalak), Memorial University of Newfoundland, St. John's, Nfld.; Department of Internal Medicine (Minuk, Uhanova, Miles, Kaita), University of Manitoba; National Microbiology Laboratory (Osiowy), Public Health Agency of Canada, Winnipeg, Man.

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