

HIV and viral hepatitis coinfection analysis using surveillance data from 15 US states and two cities

Original Paper

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Abstract

Coinfection with human immunodeficiency virus (HIV) and viral hepatitis is associated with high morbidity and mortality in the absence of clinical management, making identification of these cases crucial. We examined characteristics of HIV and viral hepatitis coinfections by using surveillance data from 15 US states and two cities. Each jurisdiction used an automated deterministic matching method to link surveillance data for persons with reported acute and chronic hepatitis B virus (HBV) or hepatitis C virus (HCV) infections, to persons reported with HIV infection. Of the 504 398 persons living with diagnosed HIV infection at the end of 2014, 2.0% were coinfecting with HBV and 6.7% were coinfecting with HCV. Of the 269 884 persons ever reported with HBV, 5.2% were reported with HIV. Of the 1 093 050 persons ever reported with HCV, 4.3% were reported with HIV. A greater proportion of persons coinfecting with HIV and HBV were males and blacks/African Americans, compared with those with HIV monoinfection. Persons who inject drugs represented a greater proportion of those coinfecting with HIV and HCV, compared with those with HIV monoinfection. Matching HIV and viral hepatitis surveillance data highlights epidemiological characteristics of persons coinfecting and can be used to routinely monitor health status and guide state and national public health interventions.

Introduction

Estimates from the USA indicate that 1.2 million residents were living with human immunodeficiency virus (HIV) infection at the end of 2013; >800 000 were infected with hepatitis B virus (HBV); and approximately 4.6 million have ever been infected with hepatitis C virus (HCV) [1–3]. Although effective therapies are available for managing HIV, HBV and HCV infections, these infections sometimes remain undiagnosed because of their often asymptomatic nature [4–6]. Public health efforts to test and link persons with HIV and viral hepatitis infections to care are of crucial importance for mitigating associated morbidity and mortality [7–9].

Because social factors that place persons at risk for acquiring HIV, HBV and HCV are similar and these conditions share some transmission routes, patients can often be coinfecting with

viral hepatitis and HIV. Although the proportion and prevalence of coinfection vary on the basis of disease epidemiology, worldwide estimates report that approximately 10% of persons living with HIV infection are coinfecting with HBV and 25% are coinfecting with HCV [10–13]. HIV infection can increase susceptibility to subsequent infection with HBV or HCV, and concomitant HIV infection can result in an increase in HBV or HCV viraemia, thus accelerating liver damage [14–17]. Coinfected persons are at greater risk for liver and all-cause morbidity and mortality, compared with those who are monoinfected [18–20]. Identifying coinfecting persons and linking them to care and management of both their HIV and viral hepatitis infections is essential. Highly active antiretroviral therapy for HIV, antiviral therapy for HBV and direct-acting antivirals that can cure HCV infection can improve outcomes for coinfecting patients [11, 16, 17].

Communicable disease surveillance data help identify trends and risks associated with infectious agent transmission and guide development and evaluation of public health initiatives [21]. Individual states and cities collect communicable disease data and transmit de-identified records to the Centers for Disease Control and Prevention (CDC) [22]. HIV and viral hepatitis infections are nationally notifiable in the USA but are maintained in disparate surveillance systems within jurisdictions and at CDC. Health departments' surveillance activities for HIV, acute and chronic HBV, and acute and chronic HCV vary by jurisdiction. Although some health departments have used their surveillance data to quantify the number and characteristics of HIV and viral hepatitis coinfections, approaches used for identifying coinfections and analysing results vary greatly [23–27]. Routine linkages of HIV and viral hepatitis surveillance data are necessary to monitor health status, including assessments of the risk for a geographically focused outbreak [28]. This study examined characteristics of HIV and viral hepatitis coinfections by using surveillance data from 15 US states and two cities with a standardised method for matching and analysis.

Methods

Jurisdiction selection

All 65 health departments funded as part of CDC's National HIV Surveillance System were contacted to identify jurisdictions interested in developing a standardised approach for using HIV and viral hepatitis surveillance data for assessing HIV and hepatitis coinfection. Fifteen states (Arizona, Connecticut, Florida, Iowa, Louisiana, Maryland, Massachusetts, Michigan, Minnesota, North Dakota, South Carolina, Texas, Virginia, Washington and Wisconsin) and two independently funded cities (New York City, New York and San Francisco, California) conducted linkages in accordance with their local data security and confidentiality policies and provided de-identified data to CDC. The independently funded city of Houston, Texas, participated in the project, but we limited our analysis to results reported by Texas to avoid duplication of reported cases. We used information collected as part of routine public health surveillance activities classified as non-research; therefore, institutional review board review was not required.

Hepatitis case selection

Jurisdictions varied by viral hepatitis conditions that were reportable and by when each condition became reportable (Table 1). Data were extracted from surveillance systems used to maintain viral hepatitis data in each jurisdiction and input into SAS® (SAS

Institute, Inc., Cary, North Carolina, USA) datasets. Datasets included acute HBV, acute HCV, chronic HBV and chronic HCV conditions, with case classifications consistent with applicable CDC/Council of State and Territorial Epidemiologists case definitions [29]. Each jurisdiction was responsible for assigning case classifications to viral hepatitis cases by using the applicable case definition. Chronic HBV and chronic HCV are not reportable in Texas; therefore, standard definitions in alignment with the chronic HCV case definition were applied to HCV laboratory data reported electronically to identify cases in Texas. Hepatitis event date was determined for each hepatitis case by a CDC-developed hierarchy of dates associated with the condition [30]. Each jurisdiction determined the earliest event date and conditions to be included on the basis of the jurisdiction's hepatitis surveillance practices (Table 1). Health departments de-duplicated their viral hepatitis data to create a unique identifier for each person across all reported conditions or to create an identifier for each person separately by HBV and HCV conditions.

HIV case selection

All jurisdictions have reported HIV infection stage 3 (AIDS) since the beginning of the epidemic in the early 1980s. However, HIV infection reporting was implemented at different times across US jurisdictions (Table 1). Data were extracted from the HIV surveillance system within each jurisdiction by using a standardised SAS program and input into a SAS dataset. All jurisdictions had routine quality-assurance procedures in place, including a requirement to de-duplicate HIV cases on a monthly basis. Datasets included all persons with HIV infection reported to health departments and meeting data completeness eligibility criteria for transfer to CDC (unpublished data CDC, 2017).

Data matching

All jurisdictions used an automated hierarchical deterministic matching method to link HIV and hepatitis datasets to reduce the matching time and to minimise variation in manual adjudication. A SAS program was developed for matching data on 14 keys (i.e., character string of values from a variable or combination of variables) (Table 2) and was similar to the method previously described by New York City [26]. Six jurisdictions validated the deterministic matching method against their existing matching methods that included a probabilistic matching component. Manual review was required only when multiple records in one dataset matched to a single record in the other dataset on the same lowest key number.

Analysis

All jurisdictions used a standardised SAS program to summarise results from the matched datasets. Aggregate data from each jurisdiction were combined. Coinfections were defined as both HIV and viral hepatitis (HBV or HCV) infections in the same person. We examined characteristics of coinfections within three cohorts: (1) persons living with diagnosed HIV as of 31 December 2014; (2) persons ever reported with HBV; and (3) persons ever reported with HCV. When assessing coinfections among persons living with diagnosed HIV infection, HIV cases were restricted to those among persons meeting the following criteria: (1) HIV infection diagnosis date on or before 31 December 2014; (2) alive as of 31 December 2014; and (3) most recent known address

Table 1. Comparison of the earliest year^a included in the analysis and the year registry started, HIV and hepatitis surveillance registries, 15 US states and two cities

Jurisdiction	Earliest year ^a /year registry started					
	HIV	Stage 3 (AIDS)	Hepatitis B, acute	Hepatitis B, chronic	Hepatitis C, acute	Hepatitis C, chronic
State						
Arizona	1968/1987	1981/1987	1933/1990	1975/1990	–/1997	1998/1997
Connecticut	1980/1981	1980/1981	2004/1992	–/–	2004/1994	2004/1994
Florida	1973/1997	1979/1981	2001/1999	1944/1999	2009/1999	1943/1999
Iowa	1979/1998	1979/1983	1976/1990	1980/1990	2001/1990	1951/1990
Louisiana	1979/1984	1979/1984	–/1990	2009/1990	–/1990	2009/1990
Maryland	1976/1981	1979/1981	2006/1989	1981/2003	2006/1989	1949/2003
Massachusetts	1976/1999	1979/1983	2007/1985	2007/1985	2007/1992	2007/1992
Michigan	1980/1983	1981/1983	2004/2000	2004/2000	2004/2000	2004/2000
Minnesota	1982/1982	1982/1982	2005/2005	1971/1987	2001/2005	1941/1998
North Dakota	1983/1983	1983/1983	2000/1976	2000/1976	2000/1994	1991/1994
South Carolina	1964/1986	1977/1986	2004/2004	2004/2004	2004/2004	2004/2004
Texas	1980/1999	1980/1983	2004/2000	–/–	2004/2000	2001/–
Virginia	1950/1993	1963/1986	1974/1996	1961/2005	2005/1996	1952/2005
Washington	1980/1987	1982/1984	2004/1965	1969/2000	1990/1981	1981/2000
Wisconsin	1981/1985	1981/1982	2000/1987	2000/1987	2000/2000	2000/2000
City						
New York City	1972/2000	1977/1981	2005/2001	2005/2001	2005/2002	2005/2002
San Francisco	1978/2002	1979/1981	2007/2004	2007/1984	2008/2004	2007/2001

AIDS, acquired immunodeficiency syndrome; HIV, human immunodeficiency virus; *MMWR*, *Morbidity and Mortality Weekly Report*.

^aFor HIV and stage 3 (AIDS), based on the earliest diagnosis date included in the analysis; the HIV surveillance system automatically calculated the diagnosis date by using recorded laboratory and clinical (non-laboratory) evidence. For hepatitis, based on the earliest event date included in the analysis as selected by each jurisdiction. The event date was determined for each hepatitis case by a hierarchy of dates associated with the condition (i.e., onset date, diagnosis date, laboratory report date or first report to the public health system, state or *MMWR* date).

on or before 31 December 2014 was in the jurisdiction. When assessing coinfections among persons with a viral hepatitis condition, HIV cases were restricted to persons with HIV infection diagnosed on or before 31 December 2014 who were reported to the jurisdiction regardless of vital status and residence. When assessing coinfections among all three cohorts described previously, viral hepatitis cases were restricted to those with a condition event date on or before 31 December 2014 reported to the jurisdiction regardless of residence or vital status. Among persons with multiple reported HBV conditions (e.g., reported with both an acute and a chronic condition), the HBV condition with the earliest event date was used when summarising the coinfection; the same method was used among persons with multiple reported HCV conditions. When assessing coinfections among persons living with diagnosed HIV as of 31 December 2014, we included persons ever diagnosed with a viral hepatitis condition and reported with a condition event date on or before 31 December 2014; due to limitations of viral hepatitis surveillance data we could not determine whether individuals had cleared their viral hepatitis infections before 31 December 2014. Because the number of persons coinfecting with HIV, HBV and HCV was expected to be low, our analysis was not designed to identify these coinfections. If a person was coinfecting with all three conditions, both the HIV and HBV coinfection information and the HIV and HCV coinfection information would be summarised.

Age group was based on age at diagnosis of HIV or viral hepatitis infection; age for coinfections was based on age at diagnosis

of the second reported virus. Transmission category was selected from the most likely route of transmission of HIV on the basis of a hierarchy of reported risk information [1]. Among coinfecting persons, sex and race/ethnicity were first derived from the HIV dataset, and supplemented with information from the hepatitis dataset if missing from the HIV dataset. For HIV infection, sex indicated sex at birth. For viral hepatitis cases, sex was not uniformly defined across all jurisdictions and indicated sex at birth, sex at the time of viral hepatitis event or current sex at the time the data were extracted depending on the jurisdiction. Among coinfecting persons, the timing of when coinfection became known was determined by comparing the HIV diagnosis date and hepatitis event date. This represented the earliest known date associated with each virus but might not reflect the true order of infection.

Results

The earliest year included in the analysis and the year the registry started for viral hepatitis and HIV data varied across the 15 states and two cities (Table 1). Of 504 398 persons living with diagnosed HIV infection as of 31 December 2014 in 17 total jurisdictions, 10 216 (2.0%; range: 0.1–4.5%) were coinfecting with HBV, and 33 993 (6.7%; range: 0–11.3%) were coinfecting with HCV (Table 3). Of 269 884 persons ever reported with HBV, 14 117 (5.2%; range: 2.6–12.2%) were coinfecting with HIV. Of 1 093 050

Table 2. Matching keys used by 15 US states and two cities for the deterministic matching method^a

Key	Description
1	Full LAST NAME + first six letters of FIRST NAME + full DOB
2	First letter of LAST NAME + letters 3–10 of LAST NAME + letters 2–9 of FIRST NAME + full DOB
3	Letters 2–7 of LAST NAME + first six letters of FIRST NAME + full DOB
4	First two letters of LAST NAME + first three letters of FIRST NAME + full SSN + full DOB
5	Full LAST NAME + first three letters of FIRST NAME + full DOB
6	Letters 3–5 of LAST NAME + first three letters of FIRST NAME + full DOB
7	First four letters of LAST NAME + first four letters of FIRST NAME + full DOB
8 ^b	First letter of LAST NAME + letters 3–10 of LAST NAME + letters 2–9 of FIRST NAME + month and year of DOB
9 ^b	First letter of LAST NAME + letters 3–10 of LAST NAME + letters 2–9 of FIRST NAME + day and year of DOB
10 ^b	Full SSN
11 ^b	First five letters of LAST NAME + first four letters of FIRST NAME + month and year of DOB
12 ^b	First letter of LAST NAME + letters 3–10 of LAST NAME + letters 2–9 of FIRST NAME + month and year of DOB, switching the first and last name in one dataset
13 ^b	First letter of LAST NAME + letters 3–10 of LAST NAME + letters 2–9 of FIRST NAME + day and year of DOB, switching the first and last name in one dataset
14 ^b	First five letters of LAST NAME + first four letters of FIRST NAME + month and year of DOB, switching the first and last name in one dataset

DOB, date of birth; HIV, human immunodeficiency virus; SSN, social security number.

^aAutomated SAS[®] (SAS Institute, Inc., Cary, North Carolina, USA) program used to match records on 14 keys. Manual review was required only when multiple records from one dataset matched to a single record in the other dataset on the same lowest key value.

^bIf matched on this key, the following three additional criteria had to be met to be considered a match:

- (1) Value of sex had to be same in both datasets or the full date of birth and digits one through four and six through nine of the social security number had to be the same in both datasets.
- (2) First name in the HIV dataset was not among the 20 most common names in the HIV dataset for the jurisdiction.
- (3) Last name in the HIV dataset was not among the 20 most common names in the HIV dataset for the jurisdiction.

persons ever reported with HCV, 47 240 (4.3%; range: 0.2–13.3%) were coinfecting with HIV.

Persons living with diagnosed HIV infection with or without HBV infection

Among persons living with diagnosed HIV infection, a greater proportion of those coinfecting with HBV were black/African American (53.9%), and a lower proportion were Hispanic (14.2%), compared with persons living with diagnosed HIV infection without HBV (44.9% and 22.2%, respectively) (Table 4). The largest proportion of HIV/HBV coinfecting persons were aged 40–49 years at the time of their second diagnosis (35.8%). A greater proportion of persons living with diagnosed HIV infection were male among those with HBV (82.9%), compared with those without HBV (74.0%). Among persons living with diagnosed HIV infection, a greater proportion of those with HBV were males with HIV infection attributed to male-to-male sexual contact (49.8%), compared with those without HBV (44.4%). A lower proportion of persons living with diagnosed HIV infection and coinfecting with HBV were females with HIV infection attributed to heterosexual contact (8.6%), compared with those without HBV (13.8%). Among 74.4% of HIV/HBV coinfecting persons, HIV diagnosis year preceded the HBV event year.

Persons living with diagnosed HIV infection with or without HCV infection

No differences were identified in the distribution of race/ethnicity by >5.0 percentage points among persons living with diagnosed

HIV infection with and without HCV (Table 4). A greater proportion of persons coinfecting with HIV and HCV were aged ≥ 50 years (37.2%), compared with those coinfecting with HIV and HBV (24.0%). Distributions by sex among persons living with diagnosed HIV infection with and without HCV were similar. Males and females with HIV infection attributed to injection drug use (IDU) (24.3% and 13.6%, respectively) represented a greater proportion of persons living with diagnosed HIV infection and HCV, compared with those without HCV (5.7 and 3.4%, respectively). Males with HIV infection attributed to male-to-male sexual contact and IDU (12.7%) represented a greater proportion of persons living with diagnosed HIV infection and HCV, compared with those without HCV (3.7%). In contrast, males with HIV infection attributed to male-to-male sexual contact (25.1%) and females with HIV infection attributed to heterosexual contact (7.4%) represented a lower proportion of persons living with diagnosed HIV infection and HCV, compared with those without HCV (46.0% and 14.2%, respectively). As with HIV and HBV coinfections, HIV diagnosis year preceded HCV event year among the majority (83.6%) of persons coinfecting with HCV and HIV.

Persons ever receiving a diagnosis of viral hepatitis with and without HIV infection

Race/ethnicity was unknown for the majority of HBV mono-infected persons (53.1%), and comparisons with HBV/HIV-coinfecting persons should be avoided (Table 5). The largest proportion of HBV/HIV-coinfecting persons was those aged 40–49 years at the

Table 3. Number and percentage of HIV and hepatitis coinfections among persons living with diagnosed HIV infection and among persons with hepatitis infection, 15 US states and two cities

Jurisdiction	Among persons living with diagnosed HIV infection, 2014 ^a			Among cumulative HBV cases through 2014 ^b		Among cumulative HCV cases through 2014 ^b	
	Living HIV	% HIV/HBV	% HIV/HCV	Cum HBV	% HIV/HBV	Cum HCV	% HIV/HCV
State							
Arizona	16 664	3.8	7.9	18 904	5.5	108 608	2.1
Connecticut	10 478	0.1	9.3	339	4.7	30 325	4.9
Florida	110 145	2.1	4.6	25 317	10.9	137 172	4.4
Iowa	2496	3.4	0	3122	4.1	1118	0.2
Louisiana	20 231	1.8	2.1	5467	8.4	17 634	3.2
Maryland	35 000	2.9	9.8	14 989	8.8	51 305	8.5
Massachusetts	21 243	1.6	7.4	15 190	2.6	67 767	2.8
Michigan	15 257	4.0	5.6	17 033	5.6	81 289	1.7
Minnesota	8140	4.5	6.6	23 340	3.0	41 198	2.7
North Dakota	353	3.7	7.4	660	2.7	7669	0.7
South Carolina	18 238	3.0	7.4	6822	12.2	40 374	5.1
Texas ^c	79 733	0.3	7.7	4472	6.7	211 117	3.8
Virginia	24 631	2.0	4.8	13 151	4.9	54 307	3.1
Washington	12 805	2.7	8.1	16 839	3.1	78 988	2.2
Wisconsin	6677	2.3	8.7	4391	4.8	42 846	2.2
City							
New York City	108 723	2.3	7.3	89 717	3.9	101 980	10.9
San Francisco	13 584	1.3	11.3	10 131	3.3	19 353	13.3
Total	504 398	2.0	6.7	269 884	5.2	1 093 050	4.3

Cum, cumulative; HBV, hepatitis B virus (acute or chronic); HCV, hepatitis C virus (acute or chronic); HIV, human immunodeficiency virus.

^aIncludes persons living with diagnosed HIV infection, regardless of stage at disease diagnosis, whose most recently known address through 31 December 2014, was within the jurisdiction and who were presumed to be alive as of 31 December 2014. Coinfections refer to persons living with diagnosed HIV infection matched with a hepatitis infection event occurring through 31 December 2014.

^bIncludes persons reported with the hepatitis infection (acute or chronic) from the earliest event date included in the analysis for each jurisdiction (see Table 1) through 31 December 2014. Coinfections refer to persons reported with the hepatitis infection event date through 31 December 2014, matched with an HIV infection diagnosis through 31 December 2014.

^cHouston, Texas (USA), independently reported to CDC the following coinfection information for this project:

- (1) Among 23 272 persons living with diagnosed HIV in Houston, 396 (1.7%) were coinfecting with HBV and 1126 were coinfecting with HCV (4.8%).
- (2) Among the 7884 cumulative persons with HBV in Houston, 573 (7.3%) were coinfecting with HIV.
- (3) Among the 27 769 cumulative persons with HCV in Houston, 1722 (6.2%) were coinfecting with HIV.
- (4) Variation in coinfection data is caused by differences in the hepatitis information reported to the Houston Health Department and the Texas Department of State Health Services. Houston HBV data include acute and chronic conditions, but Texas HBV data include only acute conditions.

time of second diagnosis (35.9%). The proportion of males was higher among the HBV/HIV coinfecting cohort, compared with the HBV mono-infected (83.4% vs. 53.8%). Among HBV/HIV-coinfecting persons, the largest proportion was among persons with HIV infection attributed to male-to-male sexual contact (48.4%). Among the HBV/HIV coinfecting population, HIV diagnosis year preceded HBV event year in 75.8% of all cases.

Similar to HBV/HIV coinfections, the greatest proportion of persons coinfecting with HCV and HIV were black/African American (42.3%) (Table 5). The proportion of HCV/HIV-coinfecting persons aged ≥ 50 years at the time of the second diagnosis was 39.2%. A greater proportion of HCV/HIV coinfecting patients were male than those only infected with HCV (75.1% vs. 61.1%). Among HCV/HIV-coinfecting persons, the largest proportion was among persons with HIV infection

attributed, at least in part, to IDU (53.6%). Among the HCV/HIV coinfecting population, HIV diagnosis year preceded HCV event year in 84.1% of cases.

Discussion

We report here on a multijurisdictional HIV and viral hepatitis coinfection match conducted by using routinely collected nationally notifiable disease surveillance data in the USA. The project summarised results from >500 000 persons living with diagnosed HIV infection, >250 000 persons reported with HBV, and >1 million persons reported with HCV from 15 states and two cities. Overall, among persons living with diagnosed HIV infection, we determined that the proportion coinfecting with HBV was 2.0% and HCV was 6.7%. Among persons ever reported to be infected

Table 4. Number and percentage of HIV and hepatitis coinfections among persons living with diagnosed HIV infection, by selected characteristics, 15 US states and two cities, 2014

Characteristic ^a	HIV without HBV event	HIV/HBV coinfections	HIV without HCV event	HIV/HCV coinfections
	No. (column %)	No. (column %)	No. (column %)	No. (column %)
Race/ethnicity				
American Indian/Alaska Native	1522 (0.3)	31 (0.3)	1383 (0.3)	170 (0.5)
Asian ^b	6049 (1.2)	227 (2.2)	5961 (1.3)	315 (0.9)
Black/African American	221 923 (44.9)	5511 (53.9)	213 112 (45.3)	14 322 (42.1)
Hispanic/Latino ^c	109 518 (22.2)	1452 (14.2)	102 807 (21.9)	8164 (24.0)
Multiple races	8702 (1.8)	209 (2.1)	8075 (1.7)	836 (2.5)
Native Hawaiian/Other Pacific Islander	305 (0.1)	8 (0.1)	302 (0.1)	11 (0)
Unknown	465 (0.1)	0 (0)	458 (0.1)	4 (0)
White	145 698 (29.5)	2778 (27.2)	138 307 (29.4)	10 171 (29.9)
Age group^d (years)				
0–12	7240 (1.5)	16 (0.2)	7149 (1.5)	27 (0.1)
13–29	162 833 (33.0)	1292 (12.7)	158 076 (33.6)	2480 (7.3)
30–39	169 121 (34.2)	2798 (27.4)	160 540 (34.1)	6298 (18.5)
40–49	107 135 (21.7)	3653 (35.8)	100 004 (21.3)	12 530 (36.9)
50–64	43 852 (8.9)	2251 (22.0)	40 736 (8.7)	11 940 (35.1)
≥65	3973 (0.8)	206 (2.0)	3872 (0.8)	718 (2.1)
Unknown	28 (0)	0 (0)	28 (0)	0 (0)
Sex^e				
Male	365 602 (74.0)	8467 (82.9)	348 614 (74.1)	25 455 (74.9)
Female	128 579 (26.0)	1749 (17.1)	121 790 (25.9)	8538 (25.1)
Unknown	1 (0)	0 (0)	1 (0)	0 (0)
Sex^e and HIV transmission category^f				
Male				
IDU	34 184 (6.9)	695 (6.8)	26 636 (5.7)	8243 (24.3)
Male-to-male sexual contact	219 593 (44.4)	5089 (49.8)	216 139 (46.0)	8543 (25.1)
Male-to-male sexual contact and IDU	21 161 (4.3)	644 (6.3)	17 475 (3.7)	4330 (12.7)
Heterosexual contact ^g	34 661 (7.0)	836 (8.2)	33 858 (7.2)	1639 (4.8)
Other/unknown ^h	56 003 (11.3)	1203 (11.8)	54 506 (11.6)	2700 (7.9)
Female				
IDU	20 124 (4.1)	390 (3.8)	15 880 (3.4)	4634 (13.6)

(Continued)

Table 4. (Continued.)

Characteristic ^a	HIV without HBV event No. (column %)	HIV/HBV coinfections No. (column %)	HIV without HCV event No. (column %)	HIV/HCV coinfections No. (column %)
Heterosexual contact ^b	68 187 (13.8)	882 (8.6)	66 561 (14.2)	2508 (7.4)
Other/unknown ^b	40 268 (8.2)	477 (4.7)	39 349 (8.4)	1396 (4.1)
Timing of coinfection				
HIV diagnosis year before the year of hepatitis event	N/A	7601 (74.4)	N/A	28 419 (83.6)
Same HIV diagnosis year and year of hepatitis event	N/A	1979 (19.4)	N/A	4042 (11.9)
HIV diagnosis year after year of hepatitis event	N/A	636 (6.2)	N/A	1532 (4.5)

HBV, hepatitis B virus (acute or chronic); HCV, hepatitis C virus (acute or chronic); HIV, human immunodeficiency virus; N/A, not applicable; IDU, injection drug use.

^aFor persons with coinfection, information comes first from the HIV surveillance system. If information was missing in the HIV surveillance system, information from the hepatitis surveillance system was used.

^bIncludes persons for whom the surveillance system did not differentiate between Asian and Native Hawaiian/Other Pacific Islander.

^cHispanics/Latinos can be of any race.

^dFor HIV cases without a hepatitis event, based on age at diagnosis of HIV. For coinfection cases, based on age at coinfection or second reported virus infection to the health department.

^eFrom HIV surveillance system, sex indicates sex at birth. From hepatitis surveillance system, sex might indicate sex at birth, sex at the time of hepatitis event, or current sex at the time the data were extracted.

^fData have not been statistically adjusted to account for unknown transmission categories.

^gHeterosexual contact with a person known to have, or to be at high risk for, HIV infection.

^hIncludes haemophilia, blood transfusion, or perinatal exposure, and persons with an unknown transmission category.

with HBV, 5.2% were ever reported to be infected with HIV, whereas among persons ever reported to be infected with HCV, 4.3% were ever reported to be infected with HIV. Differences in the number of coinfections between the two analytic methods are the result of differences in the inclusion of decedents and those with an out-of-jurisdiction residency between the two methods. These proportions represent reported coinfections among participating jurisdictions. Infected persons who were never tested for HIV or viral hepatitis or who were identified as infected but never reported to public health are not represented in these data. Because HIV and viral hepatitis might be undiagnosed, estimates of viral hepatitis coinfection among persons with HIV are often higher than reported here [10–13].

The demography of the cohort of coinfecting persons in our study matched that of other US studies regarding race and sex [23–27]. HIV transmission categories were correlated with the most common viral hepatitis transmission risks in the USA (sexual transmission for HBV and IDU for HCV) [3, 31, 32]. Identified coinfections are not necessarily recent infections, but rather new diagnoses, at least some of which must be of historical acquisition. HIV diagnosis often preceded the viral hepatitis event date in our study. Because the timing of coinfection in our analysis is based on surveillance data, HIV diagnosis preceding the viral hepatitis event date does not necessarily reflect the order in which each infection was acquired, but rather the timing of the diagnoses. Recommendations for testing persons living with HIV infection for HBV and HCV might explain the substantial proportion with an HIV diagnosis year before the hepatitis event year [33]. A public health need exists for screening all persons at risk for viral hepatitis infection, in addition to those with diagnosed HIV.

Our results are subject to certain limitations. First, viral hepatitis and HIV are chronic and often asymptomatic infections, and event year might not be consistent with the year of exposure or infection. Because our results were ascertained from surveillance data, persons with undiagnosed infection or diagnosed infection not reported to public health are not included in our analysis. Underreporting of viral hepatitis cases has been documented and might vary by jurisdiction or over time [34, 35]. Participating jurisdictions included 15 states and two cities, and therefore, our results might not be representative of the entire USA. Data from the various jurisdictions were not homogenous, particularly with regard to viral hepatitis. Although HIV surveillance is fairly similar across jurisdictions, interjurisdictional viral hepatitis surveillance activities, de-duplication efforts and data quality differ, and these differences might have confounded estimates of proportions of coinfecting persons. Moreover, each jurisdiction sets its own priorities for viral hepatitis surveillance on the basis of state or local funding, regulations and resources. National definitions for viral hepatitis case surveillance have evolved and implementation of these definitions has not necessarily been uniform across jurisdictions [29]. Jurisdictions were encouraged to include data that they believed were reasonably valid; therefore, conditions and timeframe for which data were included varied by location. National surveillance for viral hepatitis infections is founded on an incident disease surveillance paradigm. The majority of jurisdictions do not track viral hepatitis cases prospectively, and therefore, cumulative viral hepatitis cases might include persons who cleared infection spontaneously (HBV or HCV) or through treatment (HCV). Finally, minor inaccuracies might have occurred during the matching process, affecting the results.

Table 5. Number and percentage of HIV and hepatitis coinfections among persons with hepatitis B infection and hepatitis C infection, by selected characteristics, 15 US states and two cities, cumulative through 2014

Characteristic ^a	HBV without HIV diagnosis	HIV/HBV coinfections	HCV without HIV diagnosis	HIV/HCV coinfections
	No. (column %)	No. (column %)	No. (column %)	No. (column %)
Race/ethnicity				
American Indian/Alaska Native	1655 (0.7)	52 (0.4)	8440 (0.8)	294 (0.6)
Asian ^b	51 190 (20)	288 (2.0)	7771 (0.7)	392 (0.8)
Black/African American	29 378 (11.5)	7464 (52.9)	89 531 (8.6)	19 987 (42.3)
Hispanic/Latino ^c	7090 (2.8)	1931 (13.7)	46 369 (4.4)	11 035 (23.4)
Multiple races	3923 (1.5)	300 (2.1)	10 612 (1.0)	1209 (2.6)
Native Hawaiian/Other Pacific Islander	2122 (0.8)	9 (0.1)	1012 (0.1)	17 (0)
Unknown	135 791 (53.1)	0 (0)	601 105 (57.5)	5 (0)
White	24 618 (9.6)	4073 (28.9)	280 970 (26.9)	14 301 (30.3)
Age group^d (years)				
0–12	4318 (1.7)	19 (0.1)	4795 (0.5)	29 (0.1)
13–29	58 497 (22.9)	1657 (11.7)	117 648 (11.3)	3006 (6.4)
30–39	59 172 (23.1)	3830 (27.1)	140 563 (13.4)	8347 (17.7)
40–49	47 965 (18.8)	5064 (35.9)	247 300 (23.7)	17 344 (36.7)
50–64	49 176 (19.2)	3218 (22.8)	396 480 (37.9)	17 348 (36.7)
≥65	18 245 (7.1)	329 (2.3)	77 640 (7.4)	1166 (2.5)
Unknown	18 394 (7.2)	0 (0)	61 384 (5.9)	0 (0)
Sex^e				
Male	137 710 (53.8)	11 769 (83.4)	639 195 (61.1)	35 478 (75.1)
Female	112 798 (44.1)	2348 (16.6)	393 997 (37.7)	11 762 (24.9)
Unknown	5259 (2.1)	0 (0)	12 618 (1.2)	0 (0)
Sex^e and HIV transmission category^f				
Male				
IDU	N/A	1195 (8.5)	N/A	12 337 (26.1)
Male-to-male sexual contact	N/A	6827 (48.4)	N/A	10 966 (23.2)
Male-to-male sexual contact and IDU	N/A	979 (6.9)	N/A	6283 (13.3)
Heterosexual contact ^g	N/A	1088 (7.7)	N/A	2103 (4.5)
Other/unknown ^h	N/A	1680 (11.9)	N/A	3789 (8.0)
Female				
IDU	N/A	602 (4.3)	N/A	6696 (14.2)

(Continued)

Table 5. (Continued.)

Characteristic ^a	HBV without HIV diagnosis No. (column %)	HIV/HBV coinfections No. (column %)	HCV without HIV diagnosis No. (column %)	HIV/HCV coinfections No. (column %)
Heterosexual contact ^b	N/A	1114 (7.9)	N/A	3220 (6.8)
Other/unknown ^h	N/A	632 (4.5)	N/A	1846 (3.9)
Timing of coinfection				
HIV diagnosis year before the year of hepatitis event	N/A	10 695 (75.8)	N/A	39 717 (84.1)
Same HIV diagnosis year and year of hepatitis event	N/A	2600 (18.4)	N/A	5521 (11.7)
HIV diagnosis year after year of hepatitis event	N/A	822 (5.8)	N/A	2002 (4.2)

HBV, hepatitis B virus (acute or chronic); HCV, hepatitis C virus (acute or chronic); HIV, human immunodeficiency virus; N/A, not applicable; IDU, injection drug use.

^aFor coinfecting cases, information comes first from the HIV surveillance system. If information was missing in the HIV surveillance system, information from the hepatitis surveillance system was used.

^bIncludes persons for whom the surveillance system did not differentiate between Asian and Native Hawaiian/Other Pacific Islander.

^cHispanics/Latinos can be of any race.

^dFor hepatitis cases without an HIV diagnosis, based on age at diagnosis of hepatitis. For coinfecting cases, based on age at coinfection or second reported virus infection to the health department.

^eFrom HIV surveillance system, sex indicates sex at birth. From hepatitis surveillance system, sex might indicate sex at birth, sex at the time of hepatitis event, or current sex at the time the data were extracted.

^fData have not been statistically adjusted to account for unknown transmission categories.

^gHeterosexual contact with a person known to have, or to be at high risk for, HIV infection.

^hIncludes haemophilia, blood transfusion, or perinatal exposure and persons with an unknown transmission category.

Our findings highlight key public health opportunities. Racial disparities exist with regard to the populations affected by HIV and viral hepatitis. Blacks/African Americans comprise approximately 12% of the US population, but in our analysis represented >50% of persons coinfecting with HIV/HBV and 42% of persons with HIV/HCV coinfection. Male-to-male sexual contact was the predominant risk factor for HIV and HBV coinfections, whereas IDU was more common among persons coinfecting with HIV and HCV. Efforts to reduce coinfections (e.g., safe sex, preexposure prophylaxis and syringe service programmes) should target gay, bisexual and other men who have sex with men and persons who inject drugs, respectively. National guidelines recommend that, at an entry to care, all HIV-infected persons be tested for HBV, vaccinated for HBV if susceptible and screened for HCV infection with annual retesting of HCV-uninfected persons thereafter [33]. Automated electronic medical record orders can provide testing reminders in accordance with published guidelines and help remove barriers to patient screening, testing and vaccination. Health departments might consider potential benefits of co-locating and integrating HIV and viral hepatitis testing and prevention services, which can help patients navigate care for HIV or viral hepatitis infection or both.

Shared social factors that place persons at risk for acquiring HIV and viral hepatitis along with some shared transmission routes for these conditions make coinfections more likely. Assessing coinfection trends provides important information about clinical care needs (e.g., linkage to care and treatment) and for public health intervention (e.g., preexposure prophylaxis or syringe service programmes). Using surveillance data to assess coinfections is crucial for monitoring health status and measuring benchmarks to eliminate HIV and viral hepatitis infections [28, 34, 36]. Our analysis demonstrated that a standardised approach for assessing coinfections can be applied to surveillance data from different systems and jurisdictions. However, limitations of the surveillance systems might have affected the results of this analysis and resulted in an underestimation of coinfections. The ultimate goal of identification is early intervention to decrease morbidity and mortality associated with these conditions, improve clinical outcomes and limit viral transmission to susceptible persons [28, 37].

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