

Review

Systematic Review of Hepatitis C Virus Prevalence in the WHO Western Pacific Region

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Abstract: Background: This review aimed to identify hepatitis C virus (HCV) prevalence estimates among the general population and six key populations (people who inject drugs, men who have sex with men, sex workers, prisoners/detainees, Indigenous people, and migrants) in the World Health Organization Western Pacific Region (WHO WPR). Methods: Original research articles published between 2016 and 2020 were identified from bibliographic databases. Publications were retrieved, replicas removed, and abstracts screened. Retained full texts were assessed and excluded if inclusion criteria were not met. Methodological quality was assessed using the Johanna Briggs Institute critical appraisal checklist for prevalence data. Data on HCV exposure and active infection were extracted and aggregated and forest plots generated for each population by country. Results: There were no HCV prevalence estimates in any population for more than half of WPR countries and territories. Among the 76 estimates, 97% presented prevalence of exposure and 33% prevalence of active infection. General population viraemic prevalence was 1% or less, except in Mongolia. Results confirm the endemic nature of HCV among people who inject drugs, with estimates of exposure ranging from 30% in Cambodia to 76% in Hong Kong. Conclusions: Countries require detailed knowledge of HCV prevalence in diverse populations to evaluate the impact of efforts to support WHO HCV elimination goals. Results provide baseline estimates from which to monitor and evaluate progress and by which to benchmark future elimination efforts.

Keywords: hepatitis C virus; viral hepatitis; prevalence; systematic review; Western Pacific Region; elimination



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1. Introduction

In 2015, the WHO estimated that worldwide ~71 million people were living with chronic hepatitis C virus (HCV) infection [1], a global prevalence of 1.0% (95% uncertainty interval (UI) 0.8–1.1) [2,3]. In 2019, there were an estimated 542,316 HCV-related deaths, with HCV accounting for 15.3 million (95% UI 13.3–17.5) global DALYs or 0.6% (0.5–0.7) of total global DALYs. Acute hepatitis, cirrhosis, and liver cancer contributed 1.7% (0.9–2.5), 79.5% (76.1–82.7), and 18.9% (15.9–22.2) to DALYs due to hepatitis C, respectively [4].

In 2016, 13.7 million people were estimated to be living with HCV in the WHO WPR (Western Pacific Region) [5] with mortality for HCV (24.1 deaths/100,000) the highest of any WHO region [2]. By 2019, the number of people living with HCV had increased to an estimated 23.5 million or 0.71% prevalence [6] with the region including 25% of the top 20 countries globally for HCV-related deaths [5].

The WHO's 2016 Global Health Sector Strategy on Viral Hepatitis aims to eliminate viral hepatitis as a public health threat by 2030 through a 90% reduction in new infections and a 65% reduction in mortality [1]. HCV surveillance and prevalence data are crucial to monitoring elimination efforts, including estimating the population living with HCV,

documenting the number in need of treatment, and assessing the need for targeted screening and testing in high-prevalence sub-populations.

The study aimed to systematically review and synthesize epidemiological data on HCV prevalence in the general population in 27 WHO WPR countries and 10 territories or areas, and in people who inject drugs, men who have sex with men, sex workers, prisoners/detainees, Indigenous people, and migrants.

2. Materials and Methods

Research articles were retrieved from PubMed/MEDLINE, Embase, and Western Pacific Region Index Medicus (WPRIM) databases in January 2021. The search strategies combined the concepts ‘hepatitis C virus’, ‘prevalence’, and ‘WHO WPR geographic areas’ using controlled and natural vocabulary (Appendix A Table A1). Searches were limited to records published from 1 January 2016 to 31 December 2020. No language restrictions were applied.

All retrieved publications were transferred to a bibliographic data management system (EndNote™X8.2) and spreadsheet (Microsoft Excel 360®). Replica publications were excluded. Country profiles were sent to WHO Country Focal Points with a request for additional publications. The study protocol was registered with PROSPERO (registration number CRD42020223181).

2.1. Eligibility Criteria and Selection

Publications were screened according to the following inclusion criteria: the study was conducted in one of the 37 WHO WPR countries or territories; reported serological or molecular HCV prevalence for a population of interest; included data on the number or frequency of individuals exposed to HCV at a specific time or time-period; data were collected in 2010 or later, and published between 1 January 2016 and 31 December 2020; and the study was conducted in humans. Populations of interest were: (a) the general population, defined as people living in a defined geographic area, excluding children aged < 18 years and high-risk groups; (b) non-remunerated blood donors; (c) pregnant women; (d) people who currently inject or previously injected illicit drugs; (e) men who have sex with men; (f) sex workers, defined as female, male, and transgender adults, over the age of 18, who receive money or goods in exchange for sexual services, either regularly or occasionally, and who may or may not self-identify as sex workers [7]; (g) prisoners/detainees, defined as people detained in closed settings; (h) Indigenous people, defined as distinct ethnic groups with a culture that is associated with a specific geographic region; and (i) migrants, defined as internal or foreign-born migrants living in a WHO WPR country or territory.

Exclusion criteria were that the study was a clinical trial, case–control, qualitative or intervention study, case report, case series, editorial, commentary, letter to editor, author reply, animal study, conference abstract, or review or modelling study that did not provide original HCV prevalence outcomes; had fewer than 15 participants or reported data from the general population, blood donors or pregnant women with a sample < 100; HCV status was self-reported; reported prevalence outcomes for a high-risk patient group (for example people living with HIV or liver disease); or was conducted only among children.

2.2. Selection Process

Identified publications were retrieved and abstracts screened by one reviewer (JI) with 10% checking by a second reviewer (LM). The main reason for exclusion was recorded. Full text publications were independently assessed for relevance by two reviewers (JI/LM) and excluded if they did not meet the inclusion criteria. Citations in systematic reviews, policy-related publications, and modelling studies were manually searched to identify additional publications. WHO Country Focal Points for viral hepatitis in the WPR were contacted and identified an additional 82 studies which were assessed for relevance.

2.3. Data Extraction

The following data elements were entered into the spreadsheet: author(s); year of publication; title; journal; country, region and/or city; study population; study design/sampling method; recruitment year/s; sample size; study response rate; HCV testing methods; numerator: number of HCV antibody positive or HCV RNA positive participants; denominator: number of participants tested for HCV antibody or RNA; and statistical methods used to determine prevalence.

Where multiple publications reported on the same study, only the most comprehensive was retained. Where prevalence estimates were reported for the same population using the same methods across multiple time points, only data for the most recent time point were extracted. Where general population studies included children and data were stratified by age group, prevalence estimates were recalculated with children excluded.

2.4. Assessment of Quality and Risk of Bias

The methodological quality of included studies was assessed using the Joanna Briggs Institute (JBI) critical appraisal checklist for prevalence data [8]. Each publication was assigned a grade using a score of '1' for 'poor', '2' for 'good', and '3' for 'excellent' for each of the nine JBI critical appraisal items, for an aggregate quality score (range 9–27). Two reviewers (JI/LM) independently assessed each study and where there was disagreement, discussion achieved consensus. Publications with an aggregate quality score less than 50% of the maximum score (aggregate quality score < 13.5) were considered poor-quality. Additional analyses were conducted to assess the impact of inclusion of poor-quality publications.

2.5. Data Analysis

Data were aggregated and forest plots with 95% confidence intervals (CIs) were generated by country and each population of interest for HCV exposure (HCV antibody) and active infection (HCV RNA). Where 95% CIs for prevalence measures were not reported in publications, these were calculated using the Clopper–Pearson method [9]. Numerators were also calculated where these were not provided in the text. Where data were either missing or only provided in aggregate form, reviewers contacted authors by email with a request to provide missing or disaggregated data. All statistical analyses were performed with Stata v.14.2 (StataCorp, College Station, TX, USA).

3. Results

A total of 1659 articles were retrieved, including 443 replica records that were removed (Figure 1). Abstracts of the remaining 1216 publications were screened, with 1031 publications excluded on abstract review. Among 185 publications retained for full text review, 63 were retained. An additional study was identified for retention through a search of citations in systematic reviews. From 82 studies identified by Country Focal Points, 2 met the criteria for inclusion and were retained.

A total of 66 publications were retained, resulting in 76 HCV prevalence estimates among sub-populations of interest retrieved from 12 WHO WPR countries and territories (Table 1). No estimates were identified for 25 countries and territories. Among the 76 estimates, 97% (74) were HCV antibody prevalence and 33% (25) were HCV RNA prevalence (Tables A2 and A3). With the exception of pregnant women (two studies, $I^2 = 11\%$), severe heterogeneity ($I^2 > 98\%$) was observed for all populations of interest. Most studies were conducted in sub-national geographic regions. The median aggregate quality score of the 76 included studies was 18 (range 10–24). Three studies were considered poor-quality due to an aggregate quality score < 13.5.

Prevalence of HCV Antibody and RNA

HCV antibody prevalence estimates among the general population or one of the proxy populations were available for 11 countries and territories. The median aggregate quality score of the 31 studies was 19 (range 15–24). Estimates for HCV antibody prevalence

among the general population were typically low at $\leq 1.0\%$, with the highest prevalence observed in Mongolia (13.6%, Figure 2). Eleven estimates of HCV RNA prevalence among the general population or proxy populations were identified; however, HCV RNA was not detected in two of these studies. Where HCV RNA was detected, the prevalence among the general population ranged from $<0.5\%$ in the Hong Kong SAR, Japan, and Malaysia to 11.0% in Mongolia (Figure 3).

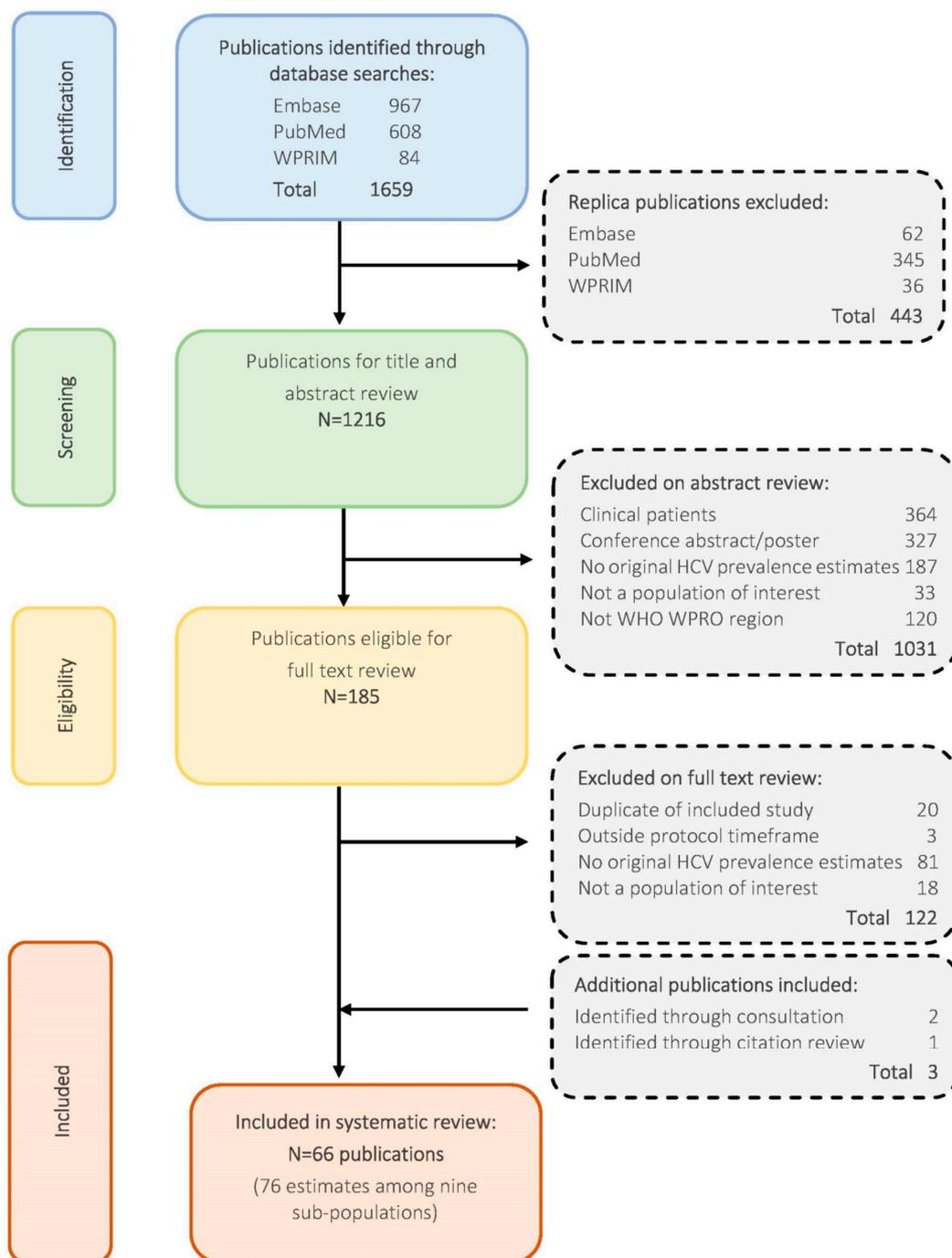


Figure 1. Flow chart of publication identification and selection process.

Table 1. Number of identified estimates for the prevalence of hepatitis C (anti-HCV or RNA) from WHO WPR by population category, 2016–2020 (n = 76).

Country	General Population	Pregnant Women	Blood Donors	People Who Inject Drugs	Men Who have Sex with Men	Sex Workers	Prisoners	Indigenous People	Migrants	Total
Australia	1			4			1		1	7
Cambodia	1			1						2
China	4	1	7	6	10	5		2	3	38
Hong Kong SAR	1		1	1						3
Japan	4	1							1	5
Malaysia	1		1							2
Mongolia	2									2
New Zealand				2						2
Papua New Guinea			1							1
Republic of Korea	3									4
Singapore			1	1						2
Viet Nam			1	3	1	3				8
Total	17	2	12	18	11	8	1	2	5	76

Note: No estimates for any population of interest were available for Brunei Darussalam, Cook Islands, Fiji, Kiribati, Lao People's Democratic Republic, Marshall Islands, Federated States of Micronesia, Nauru, Niue, Palau, Philippines, Samoa, Solomon Islands, Tonga, Tuvalu, Vanuatu, American Samoa, Northern Mariana Islands, French Polynesia, Guam, Macao SAR, New Caledonia, Pitcairn Island, Tokelau, and Wallis and Futuna.

Seventeen estimates for HCV antibody prevalence among people who inject drugs were obtained from seven countries and territories. The median aggregate quality score was 17.5 (range 15–22) and no studies were assessed as poor-quality. HCV antibody prevalence among people who inject drugs ranged from 30% in Cambodia to 76% in the Hong Kong SAR. Seven estimates for HCV RNA prevalence among people who inject drugs were identified from three countries, ranging from 24% in Australia to 51% in New Zealand.

Eleven estimates for HCV antibody prevalence among men who have sex with men were identified from two countries. The median aggregate quality score was 17 (range 13–21). Results did not change when the two studies assessed as poor-quality were excluded. HCV antibody prevalence was 0.6% in China and 26.1% in Viet Nam. Three estimates for HCV RNA prevalence among men who have sex with men were identified from two countries; the prevalence was 0.2% in China and 17.5% in Viet Nam.

Eight estimates for HCV antibody prevalence among sex workers were obtained from two countries. The median aggregate quality score was 18 (range 10–19), including one study assessed as poor-quality. Seven estimates were among female sex workers, where pooled HCV antibody prevalence was 15.4% in Viet Nam and 0.7% in China, noting that prevalence was slightly higher at 0.8% when the poor-quality study was excluded. Two estimates for HCV RNA prevalence among female sex workers were obtained from two countries, where HCV RNA prevalence was 0.04% in China and 8.7% in Viet Nam. One estimate was among male sex workers, where HCV antibody prevalence was 3.7% in Viet Nam (quality score 17). No estimates of HCV RNA prevalence among male sex workers were identified.

Only one estimate for HCV antibody prevalence among prisoners/detainees was identified, where the prevalence was 24% in Australia (quality score 17). No estimates for HCV RNA prevalence among prisoners/detainees were identified. Only one estimate for HCV antibody prevalence among Indigenous people was identified, where the prevalence was 7.0% in a population of Li ethnic minority in Baisha County, China (quality score 18). Similarly, only one estimate for HCV RNA prevalence was identified among Indigenous

people, where the prevalence was 2.7% among Yi people in a Yi autonomous prefecture of southwestern China.

Five estimates for HCV antibody prevalence among migrants were identified from three countries. The median aggregate quality score of these five studies was 17 (range 17–19) and no studies were assessed as poor-quality. HCV antibody prevalence among migrants ranged from 0.4% in China to 1.2% in Australia. One study reported HCV RNA prevalence among migrants in China; however, HCV RNA prevalence was 0%.

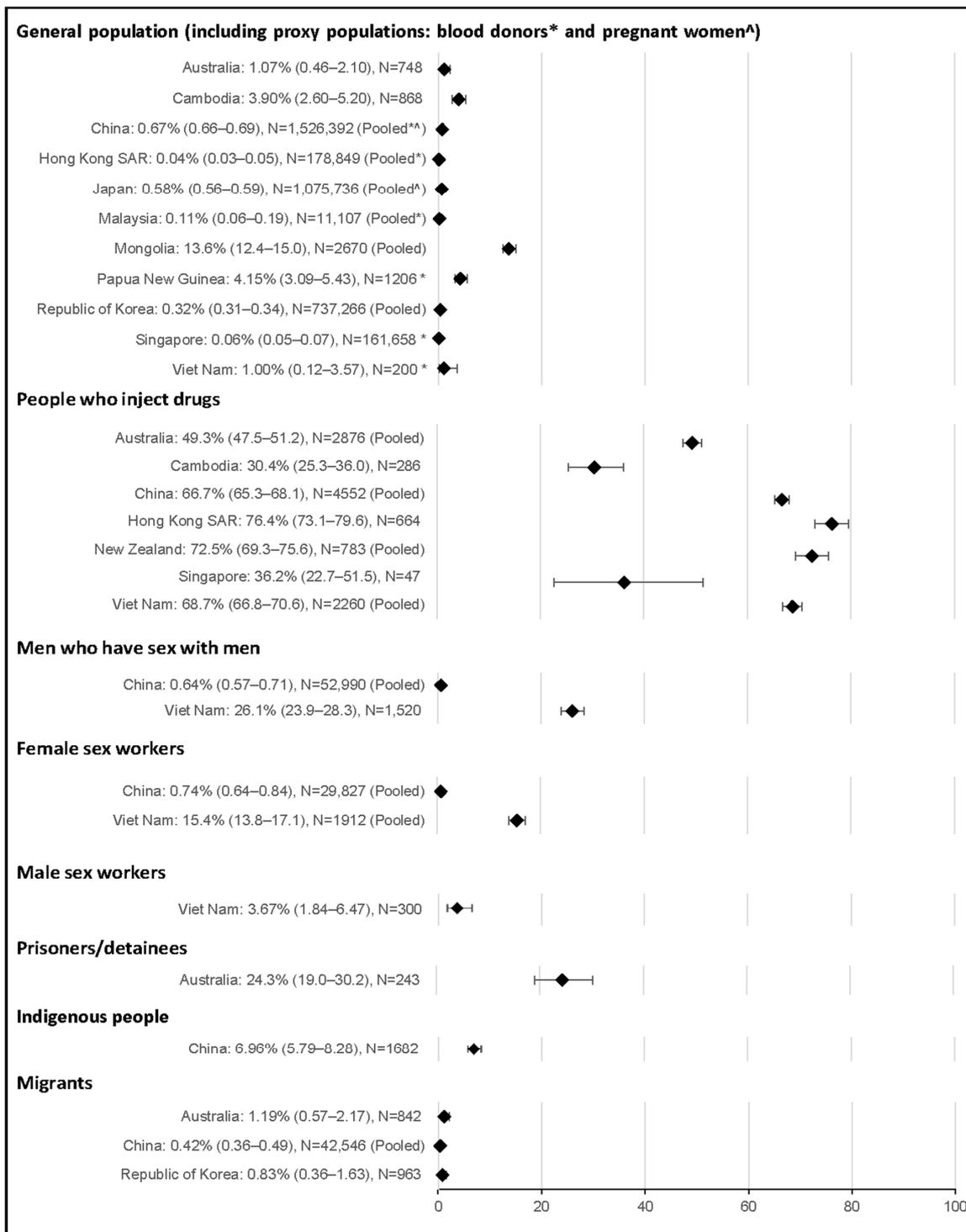


Figure 2. Forest plots HCV antibody prevalence by key population and country.

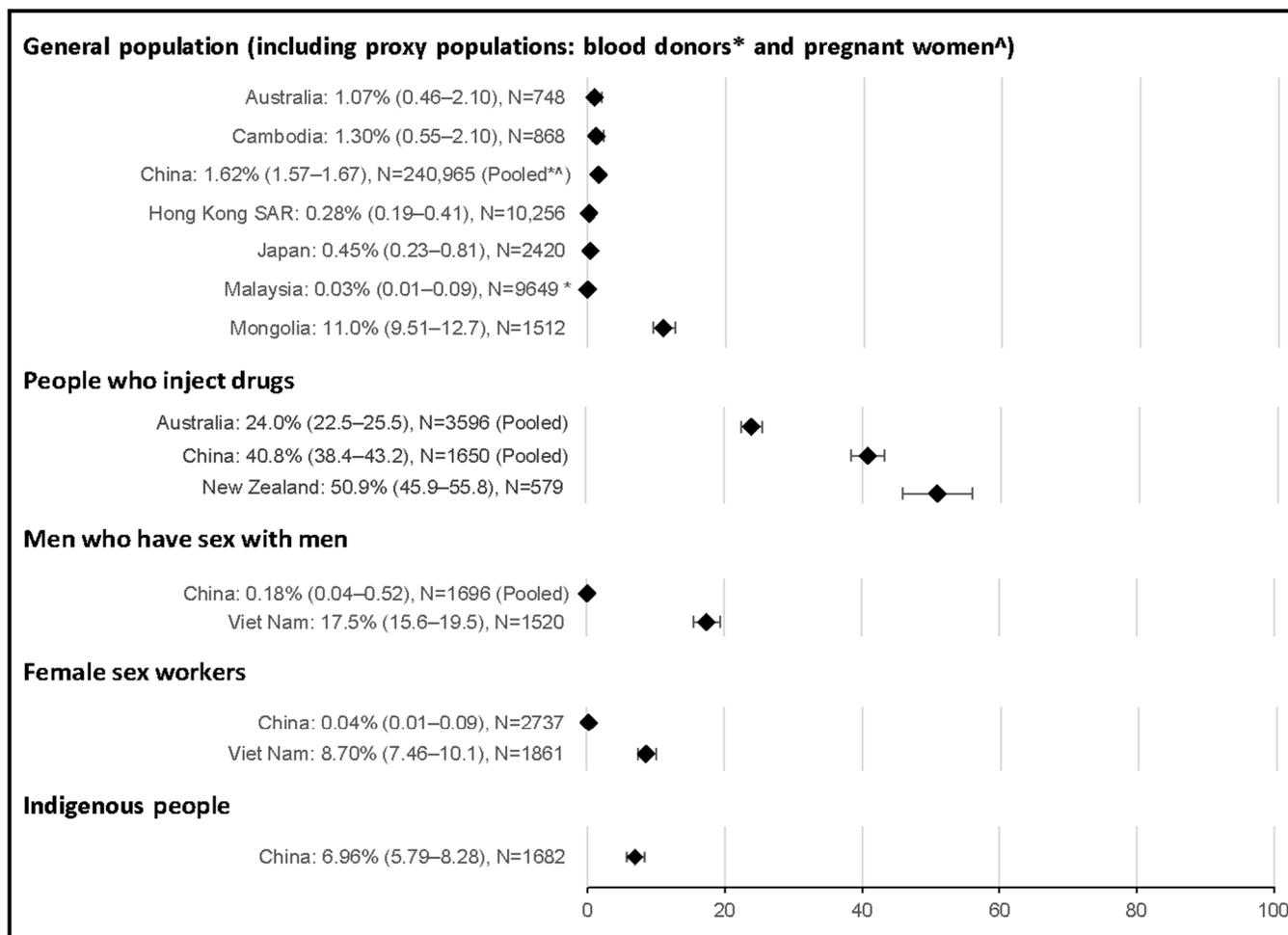


Figure 3. Forest plots HCV RNA prevalence by key population and country.

4. Discussion

Globally, an estimated 1% of people (62.5 million) are living with hepatitis C infection. Viraemic prevalence of HCV in the WHO WPR is estimated at 1% (14 million people). In this review, viraemic prevalence of HCV infection among the general population was $\leq 1\%$ in most WHO WPR countries where data were identified (Australia, China, Japan, Malaysia, Republic of Korea, Singapore, Viet Nam, and Hong Kong SAR). While HCV antibody prevalence was comparatively high in Papua New Guinea at 4.2%, only one small study among blood donors ($n = 1206$) was identified [10] and this estimate may not be representative of the general population. HCV antibody prevalence was also comparatively high at 3.9% in Cambodia where, similarly, only one small study ($n = 868$) was identified [11], However, hepatitis C viraemic prevalence in this setting was substantially lower at 1.3%. Mongolia was a notable exception, with this review confirming the endemic nature of HCV in this setting [12].

HCV infection is endemic among people who inject drugs, with an estimated 52% of people who inject drugs exposed to HCV globally [13] and 39% living with HCV [14]. In this review, estimates of exposure to HCV infection among people who inject drugs were comparatively high in the Hong Kong SAR (76%), New Zealand (73%), Viet Nam (69%), and China (67%). In the remaining three countries with available data, exposure to HCV among people who inject drugs was $< 50\%$ (Australia, 49%; Singapore, 36%; Cambodia, 30%). Among the three countries with available estimates for HCV viraemic prevalence among people who inject drugs, approximately one-third had cleared the virus in New Zealand (73% exposed vs. 51% with active infection) and China (67% exposed vs. 41% with active infection), with half clearing the virus in Australia (49% exposed vs. 24% with active infection).

The global prevalence of HCV among men who have sex with men is estimated at 3.4%, with considerable geographic variation likely due to variation in the prevalence of both injection drug use and HIV among this population [15]. Estimates for the prevalence of HCV infection among men who have sex with men were identified from only two countries. In China, the prevalence of HCV infection among men who have sex with men (0.6% exposed) was comparable to the general population (0.7% exposed). In Viet Nam, the prevalence of HCV infection in men who have sex with men was substantially higher than in the general population (1% exposed), with one in four (26%) men who have sex with men estimated to be exposed to HCV. This was likely due to overlapping risk factors, with 19% of the Vietnamese sample reporting illicit drug use, 6% injection drug use, and 15% living with HIV [16].

Although sex workers may face elevated risk of HCV infection through either sex or drug use, there is limited information on the global prevalence of HCV in this group. This review identified prevalence estimates for female sex workers in only two countries (China and Viet Nam) and only one estimate among male sex workers (Viet Nam). In China, the prevalence among female sex workers (0.7% exposed) was comparable to the general population (0.7% exposed), with no estimate identified among male sex workers. In Viet Nam, the prevalence of exposure among female sex workers was substantially higher at 15.4% and 3.7% among male sex workers, likely due to overlapping risk factors with 8% of the female Vietnamese sample reporting injection drug use and 14% living with HIV [17].

Very few studies reported HCV prevalence estimates among prisoners/detainees (one study in Australia) or Indigenous people (two studies in China). In Australia, one in four prison entrants were exposed to HCV, likely due to overlapping risk factors [18]. In China, Li and Yi ethnic minorities appear to have a higher prevalence of HCV (7.0% exposed and 2.7% with active infection respectively) compared to the general population (0.7% exposed). Tattooing has been hypothesized as a potential route of transmission among Li people [19], while injection drug use is a probable route of transmission among the Yi ethnic minority [20]. Five studies reported HCV prevalence among migrants (Australia, the Republic of Korea, and three in China). In all three countries, HCV antibody prevalence among migrants was $\leq 1\%$ and comparable to the general population.

There were significant gaps in the evidence, with no estimates of HCV prevalence among the general population, proxy populations, or key populations in 60% (16/27) of WHO WPR countries and 90% (9/10) of territories. There were no countries with available evidence of HCV prevalence among all populations of interest and only six countries and one member state had evidence for the prevalence of HCV among people who inject drugs, the group at highest risk of HCV infection and transmission. Gaps in the evidence for all populations of interest were particularly evident among the Pacific Islands and Territories.

While a strength of this review was the assessment of both exposure to HCV and viraemic prevalence, only one-third (25/76) of studies reported on viraemic prevalence. Among the estimates, there was significant heterogeneity for all populations of interest except for pregnant women, limiting comparability of results by country. Country-level estimates were pooled using source publication numerators and denominators, rather than prevalence. More weight was therefore given to studies with a larger sample size, further limiting the comparability of results between countries.

Countries and territories require robust, timely, and detailed knowledge of HCV prevalence and incidence in both the general population and key sub-populations to optimize and evaluate the impact of HCV prevention and treatment activities developed to support WHO HCV elimination goals. This review found limited studies with recent data on prevalence, with data only available for the general population, proxy populations, or key populations from 11 WHO WPR countries and 1 of 10 territories. HCV elimination efforts in the region, and especially in the 16 countries and 9 member states for which no estimates were identified, would be enhanced by implementing standardized seroprevalence surveys. Increased and improved data are particularly important for populations most at risk of infection and transmission where there are likely to be significant treatment benefits, such

as people who inject drugs [21]. With the exception of one sub-population in one country (people who inject drugs in Australia), this review did not identify evidence of progress towards HCV elimination through hepatitis C treatment.

Systematic reviews of prevalence are becoming more conventional and have the potential to inform burden of disease policy and practice, including elimination efforts. However, their utility is limited by the availability and quality of existing data. Empirical estimates of HCV prevalence using direct methods are unavailable in many countries. Of available estimates, many are not recent, nationwide, or inclusive of key sub-populations. Along with improved efforts to develop systems for monitoring the epidemic and tracking progress towards elimination, the use of indirect methods is recommended to monitor prevalence and incidence and to validate elimination goals. Our results highlight the diversity of HCV infection among specific sub-populations in the WHO WPR, with a high prevalence of exposure to HCV among people who inject drugs and specific sub-populations where injection drug use is prevalent. Findings provide baseline prevalence estimates from which to monitor and evaluate progress and by which to benchmark ongoing elimination efforts, as well as a template for future reviews.

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Conflicts of Interest: The authors declare no conflict of interest.

Appendix A

Table A1. Search strategy.

All Searches: Language Limit: No Limits, Date Limits: from 2016 to 2020		Items
Embase: Search Date 27/01/2021		
#4	#1 AND #2 AND #3	967
#3	(Australia or Brunei Darussalam or Brunei or Cambodia or China or Macao or Hong Kong or Cook Islands or Fiji or Japan or Kiribati or Lao People’s Democratic Republic or Laos or Malaysia or Marshall Islands or Federated states of Micronesia or Micronesia or Mongolia or Nauru or New Zealand or Niue or Palau or Papua New Guinea or Philippines or Republic of Korea or Korea or Samoa or Singapore or Solomon Islands or Tonga or Tuvalu or Vanuatu or Viet Nam or Vietnam or Polynesia or Guam or New Caledonia or Mariana or Pitcairn or Tokelau or Wallis or Futuna).ab,hw,kw,ti	342,058
#2	(prevalence or seroepidemiology or disease surveillance or sero epidemiology or sero epidemiological or sero epidemiologic or seroepidemiolog * or surveillance or serolog * or serosurvey * orprevalence * or population surveillance).ab,hw,kw,ti	512,912
#1	(hepatitis C or hepacivirus or hcv).ab,hw,kw,ti	49,421

Table A1. Cont.

All Searches: Language Limit: No Limits, Date Limits: from 2016 to 2020		Items
PubMed: Search date 27/01/2021		
#4	#1 AND #2 AND #3	608
#3	Search Australia[ad] OR Australia[tiab] OR "Brunei Darussalam"[ad] OR Brunei[ad] OR "Brunei Darussalam"[tiab] OR Brunei[tiab] OR Cambodia[ad] OR Cambodia[tiab] OR China[ad] OR China[tiab] OR "Hong Kong"[ad] OR "Hong Kong"[tiab] OR Macao[ad] OR Macao[tiab] OR "Cook Islands"[ad] OR "Cook Islands"[tiab] OR Fiji[ad] OR Fiji[tiab] OR Japan[ad] OR Japan[tiab] OR Kiribati[ad] OR Kiribati[tiab] OR "Lao People's Democratic Republic"[ad] OR Laos[ad] OR "Lao People's Democratic Republic"[tiab] OR Laos[tiab] OR Malaysia[ad] OR Malaysia[tiab] OR "Marshall Islands"[ad] OR "Marshall Islands"[tiab] OR "Federated states of Micronesia"[ad] OR Micronesia[ad] OR "Federated states of Micronesia"[tiab] OR Micronesia[tiab] OR Mongolia[ad] OR Mongolia[tiab] OR Nauru[ad] OR Nauru[tiab] OR "New Zealand"[ad] OR "New Zealand"[tiab] OR Niue[ad] OR Niue[tiab] OR Palau[ad] OR Palau[tiab] OR "Papua New Guinea"[ad] OR "Papua New Guinea"[tiab] OR Philippines[ad] OR Philippines[tiab] OR "Republic of Korea"[ad] OR Korea[ad] OR "Republic of Korea"[tiab] OR Korea[tiab] OR Samoa[ad] OR Samoa[tiab] OR Singapore[ad] OR Singapore[tiab] OR "Solomon Islands"[ad] OR "Solomon Islands"[tiab] OR Tonga[ad] OR Tonga[tiab] OR Tuvalu[ad] OR Tuvalu[tiab] OR Vanuatu[ad] OR Vanuatu[tiab] OR "Viet Nam"[ad] OR Vietnam[ad] OR "Viet Nam"[tiab] OR Vietnam[tiab] OR Polynesia[ad] OR Polynesia[tiab] OR Guam[ad] OR Guam[tiab] OR "New Caledonia"[ad] OR "New Caledonia"[tiab] OR Mariana[ad] OR Mariana[tiab] OR Pitcairn[ad] OR Pitcairn[tiab] OR Tokelau[ad] OR Tokelau[tiab] OR "Wallis and Futuna"[ad] OR "Wallis and Futuna"[tiab]	1,625,027
#2	Search "Prevalence"[Mesh] OR prevalence*[TIAB] OR "Population Surveillance"[Mesh] OR "Seroepidemiologic Studies"[Mesh:NoExp] OR prevalence*[TIAB] OR seroepidemiolog*[TIAB] OR "sero epidemiologic"[TIAB] OR "sero epidemiological"[TIAB] OR "sero epidemiology"[TIAB] OR serosurvey*[TIAB] OR serolog*[TI] OR epidemiolog*[TI] OR surveillance[TI]	254,647
#1	Search "hepatitis c"[MeSH Terms] OR "hepacivirus"[Mesh] OR "hepatitis c"[TIAB] OR hepaciviru*[TIAB] OR "hcv"[TIAB] OR "hepatitis c antibodies"[MeSH Terms] OR "Hepatitis C Antigens"[Mesh]	21,825
Western Pacific Region Index Medicus: Search date 14/01/2021 (publications beyond 2019 were not indexed)		
#3	#1 AND #2	84
#2	ArticleTitle:prevalence OR Abstract:prevalence OR ArticleTitle:seroep*\ OR Abstract:seroep*\ OR ArticleTitle:surveillance OR Abstract:surveillance	7597
#1	ArticleTitle:hepatitis C OR ArticleTitle:hcv OR Abstract:hepatitis C OR Abstract:hcv OR Keywords:hepatitis C OR Keywords:hcv OR MeSH:hepacivirus OR MeSH:hepatitis c	897

Table A2. Estimates for the prevalence of HCV antibodies (n = 31) and HCV RNA (n = 11) among the general population.

Population Group	Country	Author (Publication Year)	Geographic Location	Sampling Period	Study Design	Sample Size	Quality Score	HCV Antibody Estimate (95% CI)	HCV RNA Estimate (95% CI)
General population	Australia	Glenister et al. (2020)	Victoria	2016–2018	CSS	748	23	1.07% (0.46–2.10)	1.07% (0.46–2.10)
	Cambodia	Yamamoto et al. (2020)	Siem Reap	2010–2014	CSS	868	18	3.90% (2.60–5.20)	1.30% (0.55–2.10)
	China	Chen et al. (2017)	Hunan	2014	CSS	672	17	1.34% (0.61–2.53)	0.89% (0.33–1.93)
		Jiao et al. (2016)	Beijing	2010–2011	CSS	8374	18	0.37% (0.25–0.53)	0.13% (0.07–0.23)
		Wang et al. (2018)	Jilin	2016	CSS	3132	24	0.38% (0.20–0.67)	
		Zhang et al. (2016)	Jilin	2010–2013	CSS	227,808	18	2.98% (2.91–3.05)	1.70% (1.65–1.76)
	Hong Kong SAR	Liu et al. (2019)	Hong Kong SAR	2015–2016	CSS	10,256	23	0.50% (0.30–0.60)	0.28% (0.19–0.41)
	Japan	Akao et al. (2019)	Ehime	2018	CSS	305	18	0.98% (0.20–2.85)	
		Akita et al. (2020)	National	2014	CSS	742,783	18	0.68% (0.66–0.70)	
		Sugiyama et al. (2018)	Hiroshima	2011–2016	CSS	2420	20	1.00% (0.63–1.44)	0.45% (0.23–0.81)
		Tatemichi et al. (2020)	National	2016–2017	CSS	315,193	19	0.30% (0.33–0.37)	
	Malaysia	Muhamad et al. (2020)	National	2006–2012	PCS	1458	18	0.30% (0.10–0.70)	
	Mongolia	Baatarxhuu et al. (2017)	National	2012–2014	CSS	1512	16	15.6% (13.8–17.5)	11.0% (9.51–12.7)
		Dashtseren et al. (2017)	National	2013	CSS	1158	22	11.1% (9.31–13.0)	
Republic of Korea	Hong et al. (2017)	Seoul and Suwon	2002–2013	CSS	438,924	18	0.18% (0.17–0.19)		
	Jang et al. (2019)	National	2015	CSS	268,422	20	0.60% (0.57–0.63)		
	Kim et al. (2020)	National	2012–2016	CSS	29,920	20	0.80% (0.56–0.78)		
Blood donors	China	Deng et al. (2020)	Liaoning	2013–2018	N/A	411,216	20	0.37% (0.35–0.39)	
		Ding et al. (2019)	31 provinces	2016–2017	N/A	115,841	15	0.38% (0.23–0.53)	
		Fu et al. (2019)	5 provinces	2013–2016	N/A	648,607	20	0.17% (0.16–0.18)	
		Wu et al. (2017)	Shanghai	2014	N/A	38,952	20	0.14% (0.03–0.42)	
		Wu et al. (2016)	Ningxia Hui	2013	N/A	2099	20	0.46% (0.39–0.53)	0% (N/A)
		Xu et al. (2019)	Sichuan	2015	N/A	19,988	20	0.40% (0.33–0.51)	
		Yang et al. (2016)	Hubei	2014	N/A	47,691	21	0.25% (0.21–0.29)	
	Hong Kong SAR	Wong et al. (2018)	Hong Kong SAR	2016	N/A	168,593	19	0.01% (0.01–0.02)	
	Malaysia	Ramli et al. (2020)	Kelantan	2017–2018	N/A	9649	20	0.07% (0.03–0.15)	0.03% (0.01–0.09)
	Papua New Guinea	Varpit et al. (2020)	East New Britain	2018	N/A	1206	19	4.15% (3.09–5.43)	
	Singapore	Soh et al. (2019)	Singapore	2011–2015	N/A	161,658	20	0.06% (0.05–0.07)	
Viet Nam	Ishizaki (2017)	Hai Phong	2012	N/A	200	18	1.00% (0.12–3.57)		
Pregnant women	China	Wu et al. (2016)	Ningxia Hui	2013	CSS	2012	19	0.10% (0.01–0.36)	0% (N/A)
	Japan	Sugiyama et al. (2017)	Hiroshima	2010–2011	CSS	15,035	18	0.30% (0.17–0.33)	

Table A3. Estimates for the prevalence of HCV antibodies (n = 43) and HCV RNA (n = 14) among key population groups.

Population Group	Country	Author (Publication Year)	Geographic Location	Sampling Period	Study Design	Sample Size	Quality Score	HCV Antibody Estimate (95% CI)	HCV RNA Estimate (95% CI)
People who inject drugs	Australia	Iversen et al. (2020)	National	2019	CSS	2531	20	45.5% (43.5–47.4)	18.0% (16.4–19.7)
		Palmer et al. (2020)	Victoria	2008–2012	PCS	720	18		25.0% (21.9–28.3)
		Peach et al. (2018)	Victoria	2014	CSS	127	17	92.9% (87.0–96.7)	63.0% (54.0–71.4)
		Spelman et al. (2019)	Victoria	2005–2010	PCS	218	21	68.8% (62.2–74.9)	56.0% (49.1–62.7)
	Cambodia	Saing et al. (2020)	National	2017	CSS	286	20	30.4% (25.3–36.0)	
	China	Chen et al. (2017)	Hunan	2014	CSS	1049	18	45.9% (42.9–49.0)	39.7% (36.7–42.7)
		Jiang et al. (2020)	Chongqing	2016–2017	CSS	1716	17	79.7% (77.7–81.5)	
		Liu et al. (2017)	Guangdong	2013	CSS	371	15	79.2% (74.8–83.3)	
		Su et al. (2018)	Yunnan	2015	CSS	610	16	73.9% (70.3–77.4)	
		Wu et al. (2016)	Ningxia Hui	2013	CSS	601	19	47.3% (43.2–51.3)	42.7% (38.8–46.8)
		Zhang et al. (2020)	Guangdong	2011–2017	CSS	205	17	78.0% (71.8–83.5)	
		Hong Kong SAR	Chan et al. (2017)	Hong Kong SAR	2013–2014	CSS	664	17	76.4% (73.1–79.6)
	New Zealand	Noller et al. (2020)	South Island	2017	CSS	204	18	63.2% (56.2–69.9)	
		O'Connor et al. (2016)	North Island	2015	CSS	579	17	75.8% (72.1–79.3)	50.9% (45.9–55.8)
Singapore	Soh et al. (2019)	Singapore	2009–2014	CSS	47	16	36.2% (22.7–51.5)		
Viet Nam	Des Jarlais et al. (2016)	HaiPhong	2014	RDS	603	22	66.8% (62.9–70.6)		
	Des Jarlais et al. (2020)	HaiPhong	2019	RDS	1268	22	72.8% (70.3–75.2)		
	Ishizaki et al. (2017)	HaiPhong	2012	CSS	389	15	58.4% (53.3–63.3)		
Men who have sex with men	China	An et al. (2020)	Guangdong	2015	CSS	267	21	0% (N/A)	
		Bai et al. (2019)	Tianjin	2016–2018	CSS	1200	13	0.75% (0.34–1.42)	
		Guanghua et al. (2018)	10 Provinces	2015	CSS	1697	19	0.41% (0.17–0.85)	
		Han et al. (2020)	Shandong	2015–2017	CSS	1300	13	0.31% (0.08–0.79)	
		Hu et al. (2017)	Shandong	2014	CSS	3060	17	0.69% (0.43–1.05)	
		Jiao et al. (2016)	Beijing	2010–2011	CSS	1296	20	0.46% (0.17–1.05)	0.23% (0.05–0.67)
		Lu et al. (2020)	Chongqing	2018	CSS	578	17	0.35% (0.04–1.24)	
		Qin et al. (2016)	National	2013	CSS	42,535	20	0.67% (0.60–0.75)	
		Shen et al. (2017)	Jiangsu	2012–2013	CSS	657	16	0.61% (0.17–1.55)	
		Wu et al. (2016)	Ningxia Hui	2013	CSS	400	19	0% (N/A)	0% (N/A)
Viet Nam	Nadol et al. (2016)	Hanoi, HaiPhong, Can Tho, Ho Chi Minh City	2009–2010	CSS	1520	17	26.1% (23.9–28.3)	17.5% (15.6–19.5)	

Table A3. Cont.

Population Group	Country	Author (Publication Year)	Geographic Location	Sampling Period	Study Design	Sample Size	Quality Score	HCV Antibody Estimate (95% CI)	HCV RNA Estimate (95% CI)
Sex workers	China	Chen et al. (2016)	Guangxi	2015	CSS	22,459	19	0.63% (0.53–0.74)	
		Hong et al. (2016)	Zhejiang	Unstated	CSS	800	10	0.25% (0.03–0.90)	
		Hu et al. (2020)	Chongqing	2018	CSS	2805	19	1.14% (0.78–1.61)	
		Wu et al. (2016)	Ningxia Hui	2013	CSS	2737	19	0.62% (0.36–0.99)	0.40% (0.20–0.72)
	Yu et al. (2016)	Guangxi	2010–2014	CSS	1026	19	2.63% (1.74–3.81)		
	Viet Nam	Colby et al. (2016)	Ho Chi Minh City	2010	CSS	300	17	3.67% (1.84–6.47)	
		Ishizaki et al. (2017)	HaiPhong	2012	CSS	51	15	35.3% (22.4–49.9)	
		Le et al. (2019)	HaiPhong, Ho Chi Minh City, Hanoi	2013	CSS	1861	17	14.9% (13.3–16.6)	8.70% (7.46–10.1)
Prisoners	Australia	Kerslake et al. (2020)	4 states/2 territories	2016	CSS	243	17	24.3% (19.0–30.2)	
Indigenous people	China	Xu et al. (2019)	Hainan Island	2014–2015	CSS	1682	18	6.96% (5.79–8.28)	
		Shi et al. (2020)	Yi	2014–2015	CSS	2608	16		2.70% (2.10–3.40)
Migrants	Australia	Ngo et al. (2018)	Sydney	2013–2014	CSS	842	17	1.19% (0.57–2.17)	
	China	Chimungu et al. (2020)	Guangdong	2010–2017	CSS	40,935	17	0.42% (0.36–0.49)	

Note: CCS: Cross-sectional survey, PCS: Prospective cohort study, RDS: Respondent driven sampling. N/A: not applicable.

Refs. [10,11,16–20,22–80] included studies.

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