

The global prevalence of hepatitis D virus infection: systematic review and meta-analysis

Alexander J. Stockdale, Benno Kreuels, Marc Y. R. Henrion, Emanuele Giorgi, Irene Kyomuhangi, Catherine de Martel, Yvan Hutin, Anna Maria Geretti

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Supplementary appendix 1. Search Terms for PubMed, EMBASE, and Scopus

PubMed (<https://www.ncbi.nlm.nih.gov/pubmed/>)

#1	Hepatitis D [MeSH] OR Hepatitis Delta Virus [MeSH] or hepatitis D virus [Title/Abstract] OR hepatitis d [Title/Abstract] OR hepatitis delta [Title/Abstract] OR hepatitis delta virus [Title/Abstract] OR HDV [Title/Abstract] OR delta virus [Title/Abstract]
#2	Epidemiology [MeSH] OR Prevalence [MeSH] OR prevalence [Title/Abstract] OR Incidence [MeSH] OR incidence [Title/Abstract] OR epidemiology [Title/Abstract] OR serosurvey [Title/Abstract] OR seroprevalence [Title/Abstract] OR anti-HDV [Title/Abstract] OR HDV RNA [Title/Abstract] OR anti-hepatitis delta virus antibody [Title/Abstract] OR ELISA [Title/Abstract] OR Enzyme-Linked Immunosorbent Assay[MeSH] OR enzyme linked immunosorbent assay [Title/Abstract] OR viraemia [Title/Abstract] OR viremia [Title/Abstract] OR Viremia [MeSH] OR Real-Time Polymerase Chain Reaction [MeSH] OR anti-delta antibody [Title/Abstract] OR "HDV IgG" [Title/Abstract] OR HDV-IgG [Title/Abstract] OR "HDV Ab" [Title/Abstract] OR HDVAb [Title/Abstract] OR HDV-Ab [Title/Abstract] OR "antibodies to HDV" [Title/Abstract] OR anti-HD [Title/Abstract]
#3	#1 AND #2

EMBASE (<https://www.embase.com/login>)

#1	exp DELTAVIRUS/ OR exp "HEPATITIS DELTA VIRUS"/
#2	(hepatitis delta virus OR hdv OR hepatitis d OR delta virus).ti,ab
#3	#1 OR #2
#4	PREVALENCE/ OR EPIDEMIOLOGY/ OR "PREVALENCE STUDY"/ OR "PREVALENCE, SERO"/
#5	(prevalence OR epidemiology OR incidence OR serosurvey OR seroprevalence OR "HDV RNA" OR "hepatitis delta virus ribonucleic acid" OR "anti-HDV" OR "HDV IgG" OR "HDV-IgG" OR "HDV Ab" OR "HDVAb" OR "HDV-Ab" OR "antibodies to HDV" OR "anti-HD" OR "anti-delta antibody" OR "anti-hepatitis delta virus antibody" OR ELISA OR "enzyme linked immunosorbent assay" OR viraemia OR viremia).ti,ab
#6	#4 OR #5
#7	#3 AND #6

SCOPUS (<https://www.scopus.com/home.uri>)

#1	(TITLE-ABS-KEY ("hepatitis d") OR TITLE-ABS-KEY ("hepatitis delta virus") OR TITLE-ABS-KEY ("delta virus") OR TITLE-ABS-KEY ("hepatitis delta") OR TITLE-ABS-KEY (HDV) OR TITLE-ABS-KEY("hepatitis d virus")) AND (TITLE-ABS-KEY (incidence) OR TITLE-ABS-KEY (prevalence) OR TITLE-ABS-KEY (epidemiology) OR TITLE-ABS-KEY (seroprevalence) OR TITLE-ABS-KEY (serosurvey) OR TITLE-ABS-KEY (ELISA) OR TITLE-ABS-KEY ("enzyme linked immunosorbent assay") OR TITLE-ABS-KEY (viraemia) OR TITLE-ABS-KEY (viremia) OR TITLE-ABS-KEY (anti-HDV) OR TITLE-ABS-KEY ("anti HDV") OR TITLE-ABS-KEY ("HDV IgG") OR TITLE-ABS-KEY (HDV-IgG) OR TITLE-ABS-KEY ("HDV Ab") OR TITLE-ABS-KEY ("HDV-Ab") OR TITLE-ABS-KEY ("antibodies to HDV") OR TITLE-ABS-KEY ("anti-HD") OR TITLE-ABS-KEY ("anti-hepatitis delta") OR TITLE-ABS-KEY ("anti-delta antibody") OR TITLE-ABS-KEY ("HDV RNA") OR TITLE-ABS-KEY ("hepatitis delta RNA") OR TITLE-ABS-KEY ("polymerase chain reaction") OR TITLE-ABS-KEY (PCR)) AND PUBYEAR > 1989
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Supplementary appendix 2. Sources of Grey Literature Search

We performed searches of surveys, censuses, vital statistics, and reports using the search terms “hepatitis D” OR “HDV” OR “anti-HDV” OR “hepatitis delta” OR “delta virus”, and screened all articles with the keyword “hepatitis” within the following sources:

1	Global Health Data Exchange http://ghdx.healthdata.org/
2	The Demographic and Health Survey Programmes (https://www.dhsprogram.com/)
3	UNICEF Multiple Indicator Cluster Surveys (http://mics.unicef.org/)
4	World Bank Living Standards Measurement Survey (http://iresearch.worldbank.org/lsm/lsmssurveyFinder.htm)
5	Victoria State (Australia) Health Infectious Diseases Surveillance (https://www2.health.vic.gov.au/public-health/infectious-diseases/infectious-diseases-surveillance)
6	Brazil Health Information (TABNET) (http://tabnet.datasus.gov.br/cgi/deftohtm.exe?sinannet/cnv/hepabr.def)
7	Infectious Disease Epidemiology Annual Report – 2016 Robert Koch Institute (Germany) (https://www.rki.de/DE/Content/Infekt/Jahrbuch/Jahrbuecher/2017.html;jsessionid=055F194F3ED98490F0E98039C03B28A9.1_cid290?nn=2374622)
8	Public Health England (United Kingdom); Sentinel surveillance of blood borne virus testing in England: 2013-2017 (https://www.gov.uk/government/publications/sentinel-surveillance-of-blood-borne-virus-testing-in-england-2017)

Supplementary appendix 3. Included Studies by WHO region: General Populations

African Region (AFR)

- AF1. Ampah KA, Pinho-Nascimento CA, Kerber S, et al. Limited Genetic Diversity of Hepatitis B Virus in the General Population of the Offin River Valley in Ghana. *PLoS One* 2016; 11: e0156864.
- AF2. Andernach IE, Leiss LV, Tarnagda ZS, et al. Characterization of hepatitis delta virus in sub-Saharan Africa. *J Clin Microbiol* 2014; 52: 1629-36.
- AF3. Andersson MI, Maponga TG, Ijaz S, et al. The epidemiology of hepatitis B virus infection in HIV-infected and HIV-uninfected pregnant women in the Western Cape, South Africa. *Vaccine* 2013; 31: 5579-84.
- AF4. Belyhun Y, Liebert UG, Maier M. Clade homogeneity and low rate of delta virus despite hyperendemicity of hepatitis B virus in Ethiopia. *Virology* 2017; 14: 176.
- AF5. Butler EK, Rodgers MA, Collier KE, et al. High prevalence of hepatitis delta virus in Cameroon. *Sci Rep* 2018; 8: 11617.
- AF6. Coffie PA, Tchounga BK, Bado G, et al. Prevalence of hepatitis B and delta according to HIV-type: a multi-country cross-sectional survey in West Africa. *BMC Infect Dis* 2017; 17: 466.
- AF7. Cunha L, Plouzeau C, Ingrand P, et al. Use of replacement blood donors to study the epidemiology of major blood-borne viruses in the general population of Maputo, Mozambique. *J Med Virol* 2007; 79: 1832-40.
- AF8. De Paschale M, Ceriani C, Cerulli T, et al. Prevalence of HBV, HDV, HCV, and HIV infection during pregnancy in northern Benin. *J Med Virol* 2014; 86: 1281-7.
- AF9. Ducancelle A, Abgueguen P, Birguel J, et al. High endemicity and low molecular diversity of hepatitis B virus infections in pregnant women in a rural district of North Cameroon. *PLoS One* 2013; 8: e80346.
- AF10. Francois-Souquiere S, Makuwa M, Bisvigou U, Kazanji M. Epidemiological and molecular features of hepatitis B and hepatitis delta virus transmission in a remote rural community in central Africa. *Infect Genet Evol* 2016; 39: 12-21.
- AF11. Groc S, Abbate JL, Le Gal F, et al. High prevalence and diversity of hepatitis B and hepatitis delta virus in Gabon. *J Viral Hepat* 2019; 26: 170-82.
- AF12. Honge BL, Jespersen S, Medina C, et al. Hepatitis B and delta virus are prevalent but often subclinical co-infections among HIV infected patients in Guinea-Bissau, West Africa: a cross-sectional study. *PLoS One* 2014; 9: e99971.
- AF13. Ifeora IM, Bakarey AS, Adeniji JA, Onyemelukwe FN. Seroprevalence of hepatitis B and delta viruses among HIV-infected population attending anti-retroviral clinic in selected health facilities in Abuja, Nigeria. *J Immunoassay Immunochem* 2017; 38: 608-19.
- AF14. Jaquet A, Wandeler G, Nouaman M, et al. Alcohol use, viral hepatitis and liver fibrosis among HIV-positive persons in West Africa: a cross-sectional study. *J Int AIDS Soc* 2017; 19: 21424.
- AF15. Katwesigye E, Seremba E, Semitala F, Ocamo P. Low sero-prevalence of hepatitis delta antibodies in HIV/ hepatitis B co-infected patients attending an urban HIV clinic in Uganda. *African Health Sciences* 2016; 16: 1089-93.
- AF16. Kommas NP, Ghosh S, Abdou-Chekaraou M, et al. Hepatitis B and hepatitis D virus infections in the Central African Republic, twenty-five years after a fulminant hepatitis outbreak, indicate continuing spread in asymptomatic young adults. *PLoS Negl Trop Dis* 2018; 12: e0006377.
- AF17. Lemoine M, Shimakawa Y, Njie R, et al. Acceptability and feasibility of a screen-and-treat programme for hepatitis B virus infection in The Gambia. *Lancet Glob Health* 2016; 4: e559-67.
- AF18. Makuwa M, Caron M, Souquiere S, Malonga-Mouelet G, Mahe A, Kazanji M. Prevalence and genetic diversity of hepatitis B and delta viruses in pregnant women in Gabon: Molecular evidence that hepatitis delta virus clade 8 originates from and is endemic in central Africa. *J Clin Microbiol* 2008; 46: 754-6.
- AF19. Makuwa M, Mints-Ndong A, Souquiere S, Nkoghe D, Leroy EM, Kazanji M. Prevalence and molecular diversity of hepatitis B virus and hepatitis delta virus in urban and rural populations in northern Gabon in central Africa. *J Clin Microbiol* 2009; 47: 2265-8.

- AF20. Mansour W, Bollahi MA, Hamed CT, et al. Virological and epidemiological features of hepatitis delta infection among blood donors in Nouakchott, Mauritania. *J Clin Virol* 2012; 55: 12-6.
- AF21. Mansour W, Malick FZ, Sidiya A, et al. Prevalence, risk factors, and molecular epidemiology of hepatitis B and hepatitis delta virus in pregnant women and in patients in Mauritania. *J Med Virol* 2012; 84: 1186-98.
- AF22. Matthews PC, Beloukas A, Malik A, et al. Prevalence and characteristics of hepatitis B virus (HBV) coinfection among HIV-positive women in South Africa and Botswana. *PLoS One* 2015; 10: e0134037.
- AF23. Njouom R, Tejiokem MC, Texier G, Fontanet A. Prevalence of hepatitis B, hepatitis C and hepatitis D virus infections in Cameroon: Results from a national population based survey. *J Viral Hepat* 2015; 22: 1-2.
- AF24. Pinho-Nascimento CA, Bratschi MW, Hofer R, et al. Transmission of hepatitis B and D viruses in an African rural community. *mSystems* 2018; 3.
- AF25. Salpini R, Fokam J, Ceccarelli L, et al. High burden of HBV-infection and atypical HBV strains among HIV-infected Cameroonians. *Curr HIV Res* 2016; 14: 165-71.
- AF26. Sanou AM, Benkirane K, Tinto B, et al. Prevalence of hepatitis B virus and hepatitis D virus coinfection in western Burkina Faso and molecular characterization of the detected virus strains. *Int J Infect Dis* 2018; 70: 15-9.
- AF27. Stockdale AJ, Chaponda M, Beloukas A, et al. Prevalence of hepatitis D virus infection in sub-Saharan Africa: a systematic review and meta-analysis. *Lancet Glob Health* 2017; 5: e992-e1003.
- AF28. Torimiro JN, Nanfack A, Takang W, et al. Rates of HBV, HCV, HDV and HIV type 1 among pregnant women and HIV type 1 drug resistance-associated mutations in breastfeeding women on antiretroviral therapy. *BMC Pregnancy Childbirth* 2018; 18: 504.
- AF29. Tuaille E, Kania D, Gordien E, Van de Perre P, Dujols P. Epidemiological data for hepatitis D in Africa. *Lancet Glob Health* 2018; 6: e33.
- AF30. Vray M, Debonne JM, Sire JM, et al. Molecular epidemiology of hepatitis B virus in Dakar, Senegal. *J Med Virol* 2006; 78: 329-34.
- AF31. Wilson P, Jhaveri R, Parr JB, et al. Efficient dried blood spot based determination of hepatitis B seroprevalence from a national survey in the Democratic Republic of the Congo. *J Viral Hepat* 2018; 25: 92-3.
- AF32. Winter A, Letang E, Vedastus Kalinjuma A, et al. Absence of hepatitis delta infection in a large rural HIV cohort in Tanzania. *Int J Infect Dis* 2016; 46: 8-10.

Eastern Mediterranean Region (EMR)

- EM1. Alavian SM, Tabatabaei SV, Ghadimi T, et al. Seroprevalence of HBV infection and its risk factors in the west of Iran: A population-based study. *Int J Prev Med* 2012; 3: 770-5.
- EM2. Alavian SM, Tabatabaei SV, Nourizad S, et al. Seroepidemiology of HBV infection in Kermanshah, west of Iran; a population based study. *Jundishapur J Microbiol* 2012; 5: 564-9.
- EM3. Ali HYM, Yassen SA. Prevalence of hepatitis B & D viral infections among hospital personnel in Mosul Iraq. *Qatar Med J* 2001; 10: 51-3.
- EM4. Al-Nabehi BAH, Saeed WSE, Musa AM, El Hassan AM, Khalil EAG, Al-Shamahy H. Sero-molecular epidemiology and risk factors of viral hepatitis in urban Yemen. *Int J Virol* 2015; 11: 133-8.
- EM5. Al-Traif I, Ali A, Dafalla M, Al-Tamimi W, Qassem L. Prevalence of hepatitis delta antibody among HBsAg carriers in Saudi Arabia. *Ann Saudi Med* 2004; 24: 343-4.
- EM6. Attaran MS, Sharifi Z, Hosseini SM, Samei S, Ataee Z. Prevalence of hepatitis B and hepatitis D coinfection in asymptomatic blood donors in Iran. *Apmis* 2014; 122: 243-7.
- EM7. Djebbi A, Rebai WK, Bahri O, Hogga N, Sadraoui A, Triki H. [serological markers, viral RNA and genotype of hepatitis delta virus in HBs antigen positive Tunisian patients]. *Pathol Biol* 2009; 57: 518-23.
- EM8. Doosti A, Amini-Bavil-Olyaei S, Tajbakhsh E, Adeli A, Mahboudi F. Prevalence of viral hepatitis and molecular analysis of HBV among voluntary blood donors in west Iran. *New Microbiol* 2009; 32: 193-8.

- EM9. Elashry A, Gomaa A, Elzohry HA, Badra G, Waked I, Raouf AA. Prevalence and risk-factors of hepatitis virus markers in Egyptian young men: Evidence for decreasing prevalence of hepatitis C. *Hepatology* 2015; 62.
- EM10. Ghadir MR, Belbasi M, Heidari A, et al. Prevalence of hepatitis D virus infection among hepatitis B virus infected patients in Qom province, center of Iran. *Hepat Mon* 2012; 12: 205-8.
- EM11. Gomaa NI, Metwally LA, Nemr N, Younis S. Seroprevalence of HDV infection in HBsAg positive population in Ismailia, Egypt. *Egypt J Immunol* 2013; 20: 23-8.
- EM12. Javanmard D, Abedi F, Namaei MH, Asghari A, Ziaee M, Alavian SM. High prevalence of hepatitis B virus infection in the village of Esfandiar in south Khorasan province, Iran. *Hepat Mon* 2018; 18.
- EM13. Ramia S, El-Zaatari M, Sharara AI, Ramlawi F, Farhat B. Current prevalence of hepatitis delta virus (HDV) infection and the range of HDV genotypes in Lebanon. *Epidemiol Infect* 2007; 135: 959-62.
- EM14. Roshandel G, Semnani S, Abdolahi N, et al. Prevalence of hepatitis D virus infection in hepatitis B surface antigen-positive subjects in Golestan province, northeast Iran. *J Microbiol Immunol Infect* 2008; 41: 227-30.
- EM15. Salehi M, Alavian SM, Tabatabaei SV, et al. Seroepidemiology of HBV infection in south-east of Iran; a population based study. *Iran Red Crescent Med J* 2012; 14: 283-8.
- EM16. Saady N, Sugauchi F, Tanaka Y, et al. Genotypes and phylogenetic characterization of hepatitis B and delta viruses in Egypt. *J Med Virol* 2003; 70: 529-36.
- EM17. Sharifzadeh G, Namaei MH, Ebrahimzadeh A, et al. Prevalence of hepatitis D virus infection and associated factors among HBsAg-positive patients in Birjand, Iran, 2012-2014. *Hepat Mon* 2017; 17.
- EM18. Tajbakhsh E, Abbasian B, Esmaeili A, Behjati M. Seroprevalence of anti-HDV Ab and socioepidemiological characteristics among HBsAg-positive blood donors in Charmahal-O-Bakhtiyari province, Iran. *Hepat Mon* 2011; 11: 130-1.
- EM19. Ur Rahman S, Hafiz A. Study of hepatitis D in high risk population of hepatitis B. *J Coll Physicians Surg Pak*; 10: 29-32.
- EM20. Zaidi G, Idrees M, Malik FA, et al. Prevalence of hepatitis delta virus infection among hepatitis B virus surface antigen positive patients circulating in the largest province of Pakistan. *Virology* 2010; 7: 283.

European Region (EUR)

- EU1. Bissinger AL, Berg CP. Is the HDV seroprevalence in patients admitted to a university hospital representative for all HBV carriers in Germany? *Infection* 2013; 41: 1031-2.
- EU2. Brehar-Ciofleac D, Claici C, Rosiu N, Negrea DA, Moldovan R, Cosnita M. Hepatitis B virus (HBV) and dual HBV-hepatitis delta virus (HDV) infection in apparently healthy persons. *Rom J Virol* 1998; 49: 3-10.
- EU3. Cozzolongo R, Elba S, Petrucci J, et al. Epidemiology of HCV and HBV in the general population: A survey in a southern Italian town. *J Hepatology* 2009; 50.
- EU4. Drobeniuc J, Hutin YJ, Harpaz R, et al. Prevalence of hepatitis B, D and C virus infections among children and pregnant women in Moldova: additional evidence supporting the need for routine hepatitis B vaccination of infants. *Epidemiol Infect* 1999; 123: 463-7.
- EU5. Iliescu EL. Epidemiological study of HBV, HCV, HDV and HEV prevalence in Romania. *J Gastroenterol Hepatology* 2013; 28: 423-4.
- EU6. Kalinin AL, Zhavoronok SV, Mikhailov MI, et al. [Viral hepatitis delta in the republic of Belarus]. *Zh Mikrobiol Epidemiol Immunobiol* 1998; 6: 74-7.
- EU7. Mese S, Nergiz S, Tekes S, Gul K. Seroprevalence of serum HBsAg positivity and hepatitis delta virus infection among blood donors in south-eastern Turkey. *Clin Ter* 2014; 165: 95-8.
- EU8. Quaglio G, Ramadani N, Pattaro C, et al. Prevalence and risk factors for viral hepatitis in the Kosovar population: implications for health policy. *J Med Virol* 2008; 80: 833-40.

- EU9. Rosenberg GK, Lattimore S, Brailsford SR, et al. The diversity of chronic hepatitis B virus infections within blood donors in England and North Wales 2005 through 2010. *Transfusion* 2013; 53: 2467-76.
- EU10. Sellier PO, Maylin S, Brichtler S, et al. Hepatitis B virus-hepatitis D virus mother-to-child co-transmission: A retrospective study in a developed country. *Liver Int* 2018; 38: 611-8.
- EU11. Servant-Delmas A, Le Gal F, Gallian P, Gordien E, Laperche S. Increasing prevalence of HDV/HBV infection over 15 years in France. *J Clin Virol* 2014; 59: 126-8.
- EU12. Tozun N, Ozdogan O, Cakaloglu Y, et al. Seroprevalence of hepatitis B and C virus infections and risk factors in Turkey: A fieldwork TURHEP study. *Clin Microbiol Infect* 2015; 21: 1020-6.
- EU13. Uzun B, Sener AG, Gungor S, Afsar I, Demirci M. Evaluation of hepatitis delta virus (HDV) infection in blood donors in western Turkey. *Transfus Apher Sci* 2014; 50: 388-91.
- EU14. Voiculescu M, Iliescu L, Ionescu C, et al. A cross-sectional epidemiological study of HBV, HCV, HDV and HEV prevalence in the SubCarpathian and South-Eastern regions of Romania. *J Gastrointestin Liver Dis* 2010; 19: 43-8.

Region of the Americas (AMR)

- AM1. Cabezas J, Trujillo O, Balbuena J, Borda A, Fiestas V, Ramos F. Impact of 20 years of implementation of an immunization program for HBV in Huanta, Peru: First clues of elimination of hepatitis delta virus. *Hepatology* 2018; 68.
- AM2. Caetano KAA, Pinheiro RS, Matos MA, et al. Hepatitis B in rural settlement projects in central Brazil: Epidemiology and vaccination. *J Viral Hepat* 2015; 22: 23-4.
- AM3. Crispim MA, Fraiji NA, Campello SC, Schriefer NA, Stefani MM, Kiesslich D. Molecular epidemiology of hepatitis B and hepatitis delta viruses circulating in the western Amazon region, north Brazil. *BMC Infect Dis* 2014; 14: 94.
- AM4. Delfino CM, Gentile EA, Castillo AI, et al. Hepatitis B virus and hepatitis D virus in blood donors from Argentina: Circulation of HBsAg and reverse transcriptase mutants. *Arch Virol* 2014; 159: 1109-17.
- AM5. Njei B, Do A, Lim JK. Prevalence of hepatitis delta infection in the United States: National health and nutrition examination survey, 1999-2012. *Hepatology* 2016; 64: 681-2.
- AM6. Villar LM, Milagres FAP, Lampe E, et al. Determination of hepatitis B, C and D prevalence among urban and Amerindian populations from the eastern Brazilian Amazon: A cross sectional study. *BMC Infect Dis* 2018; 18: 411.

South-East Asia Region (SEAR)

- SE1. Louisirotchanakul S, Myint KS, Srimee B, et al. The prevalence of viral hepatitis among the Hmong people of northern Thailand. *Southeast Asian J Trop Med Public Health* 2002; 33: 837-44.
- SE2. Lusida MI, Surayah, Sakugawa H, et al. Genotype and subtype analyses of hepatitis B virus (HBV) and possible co-infection of HBV and hepatitis C virus (HCV) or hepatitis D virus (HDV) in blood donors, patients with chronic liver disease and patients on hemodialysis in Surabaya, Indonesia. *Microbiol Immunol* 2003; 47: 969-75.
- SE3. Mollah AH, Nahar N, Siddique MA, Anwar KS, Hassan T, Azam MG. Common transfusion-transmitted infectious agents among thalassaemic children in Bangladesh. *J Health Popul Nutr* 2003; 21: 67-71.
- SE4. Theamboonlers A, Hansurabhanon T, Verachai V, Chongsrisawat V, Poovorawan Y. Hepatitis D virus infection in Thailand: HDV genotyping by RT-PCR, RFLP and direct sequencing. *Infection* 2002; 30: 140-4.

Western Pacific Region (WPR)

- WP1. Arakawa Y, Moriyama M, Taira M, et al. Molecular analysis of hepatitis D virus infection in Miyako Island, a small Japanese island. *J Viral Hepat* 2000; 7: 375-81.
- WP2. Chang CJ, Chiang JC, Lu SN, Wang JH. Hepatitis delta virus and GBV-C infection in two neighboring hepatitis B virus and hepatitis C virus-endemic villages in Taiwan. *Chang Gung Med J* 2010; 33: 137-44.

- WP3. Chang SY, Yang CL, Ko WS, et al. Molecular epidemiology of hepatitis D virus infection among injecting drug users with and without human immunodeficiency virus infection in Taiwan. *J Clin Microbiol* 2011; 49: 1083-9.
- WP4. Chen F, Zhang J, Guo F, et al. Hepatitis B, C, and D virus infection showing distinct patterns between injection drug users and the general population. *J Gastroenterol Hepatol* 2017; 32: 515-20.
- WP5. Chen X, Oidovsambuu O, Liu P, et al. A novel quantitative microarray antibody capture assay identifies an extremely high hepatitis delta virus prevalence among hepatitis B virus-infected Mongolians. *Hepatology* 2017; 66: 1739-49.
- WP6. Davaalkham D, Ojima T, Uehara R, et al. Hepatitis delta virus infection in mongolia: analyses of geographic distribution, risk factors, and disease severity. *Am J Trop Med Hyg* 2006; 75: 365-9.
- WP7. Davies J, Tong SY, Davis JS. Hepatitis D is rare or non-existent in hepatitis B virus-infected Indigenous Australians in the Northern Territory. *Aust N Z J Public Health* 2013; 37: 188-9.
- WP8. Dunford L, Carr MJ, Dean J, et al. A multicentre molecular analysis of hepatitis B and blood-borne virus coinfections in Viet Nam. *PLoS One* 2012; 7: e39027.
- WP9. Inoue J, Takahashi M, Nishizawa T, et al. High prevalence of hepatitis delta virus infection detectable by enzyme immunoassay among apparently healthy individuals in Mongolia. *J Med Virol* 2005; 76: 333-40.
- WP10. Lee MH, You SL, Yang HI, et al. Prevalence and risk factors of hepatitis D virus infection: the REVEAL-HBV/HCV/HDV study. *Hepatology Intern* 2014; 8.
- WP11. Li L, He J, Zhao L. [Epidemiologic features of viral hepatitis in Fujian]. *Zhonghua Liu Xing Bing Xue Za Zhi* 1998; 19: 89-92.
- WP12. Liu KSH, Seto WK, Lau EHY, et al. A territory-wide prevalence study on blood-borne and enteric viral hepatitis in Hong Kong. *J Infect Dis* 2019.
- WP13. Lu SN, Chen TM, Lee CM, Wang JH, Tung HD, Wu JC. Molecular epidemiological and clinical aspects of hepatitis D virus in a unique triple hepatitis viruses (B, C, D) endemic community in Taiwan. *J Med Virol* 2003; 70: 74-80.
- WP14. Nguyen VT, McLaws ML, Dore GJ. Highly endemic hepatitis B infection in rural Vietnam. *J Gastroenterol Hepatol* 2007; 22: 2093-100.
- WP15. Tsatsralt-Od B, Takahashi M, Nishizawa T, Inoue J, Ulaankhuu D, Okamoto H. High prevalence of hepatitis B, C and delta virus infections among blood donors in Mongolia. *Arch Virol* 2005; 150: 2513-28.

Supplementary appendix 4. Included Studies by WHO Region: Isolated Populations

European Region (EUR)

EU15. Borresen ML, Olsen OR, Ladefoged K, et al. Hepatitis D outbreak among children in a hepatitis B hyper-endemic settlement in Greenland. *J Viral Hep* 2010; 17: 162-70.

EU16. Karatapanis S, Papastergiou V, Papakonstantinou F, et al. Sixteen years after the implementation of compulsory hepatitis B vaccination in a hepatitis delta endemic area in Greece: Comparison between two population based serosurveys. *J Viral Hep* 2018; 25: 188-9.

EU17. Rex KF, Krarup HB, Laurberg P, Andersen S. Population-based comparative epidemiological survey of hepatitis B, D, and C among Inuit migrated to Denmark and in high endemic Greenland. *Scand J Gastroenterol* 2012; 47: 692-701.

Region of the Americas (AMR)

AM7. Braga WS, Brasil LM, de Souza RA, Castilho Mda C, da Fonseca JC. [The occurrence of hepatitis B and delta virus infection within seven Amerindian ethnic groups in the Brazilian western Amazon]. *Rev Soc Bras Med Trop* 2001; 34: 349-55.

AM8. Braga WS, Castilho Mda C, Borges FG, et al. Hepatitis D virus infection in the Western Brazilian Amazon - far from a vanishing disease. *Rev Soc Bras Med Trop* 2012; 45: 691-5.

AM9. de Paula VS, Arruda ME, Vitral CL, Gaspar AM. Seroprevalence of viral hepatitis in riverine communities from the Western Region of the Brazilian Amazon Basin. *Mem Inst Oswaldo Cruz* 2001; 96: 1123-8.

AM10. di Filippo Villa D, Cortes-Mancera F, Payares E, et al. Hepatitis D virus and hepatitis B virus infection in Amerindian communities of the Amazonas state, Colombia. *Viol J* 2015; 12: 172.

AM11. Duarte MC, Cardona N, Poblete F, et al. A comparative epidemiological study of hepatitis B and hepatitis D virus infections in Yanomami and Piaroa Amerindians of Amazonas State, Venezuela. *Trop Med Int Health* 2010; 15: 924-33.

AM12. Harpaz R, McMahon BJ, Margolis HS, et al. Elimination of new chronic hepatitis B virus infections: results of the Alaska immunization program. *J Infect Dis* 2000; 181: 413-8.

AM13. Leon P, Venegas E, Bengoechea L, et al. [Prevalence of infections by hepatitis B, C, D and E viruses in Bolivia]. *Rev Panam Salud Publica* 1999; 5: 144-51.

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South-East Asia Region (SEAR)

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Supplementary appendix 5. Included Studies by WHO Region: Hepatology Clinic Populations

African Region (AFR)

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- AF36. Anyovi F, Lahaye FM, Rapp C, et al. Molecular epidemiology of hepatitis D virus in Togo. *Trop Med Int Health* 2017; 22: 133-4.
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Eastern Mediterranean Region (EMR)

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- EM23. Ataei B, Yazdani MR, Kalantari H, et al. Hepatitis D virus infection in Isfahan, central Iran: Prevalence and risk factors among chronic HBV infection cases. *Hepat Mon* 2011; 11: 269-72.
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European Region (EUR)

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- EU20. Bahcecioglu IH, Aygun C, Gozel N, Poyrazoglu OK, Bulut Y, Yalniz M. Prevalence of hepatitis delta virus (HDV) infection in chronic hepatitis B patients in eastern Turkey: still a serious problem to consider. *J Viral Hepat* 2011; 18: 518-24.
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- EU1. Bissinger AL, Berg CP. Is the HDV seroprevalence in patients admitted to a university hospital representative for all HBV carriers in Germany? *Infection* 2013; 41: 1031-2.

- EU23. Brancaccio G, Nardi A, Madonia S, et al. The present profile of chronic hepatitis B virus infection highlights future challenges: An analysis of the Multicenter Italian MASTER-B cohort. *Dig Liver Dis* 2018.
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- EU25. Byrne R, Childs K, Sadler MD, et al. Active HDV RNA replication is associated with high HBsAg levels and severe liver damage; a role for HBsAg/HBV DNA ratio to identify those most at risk. *Hepatology* 2016; 64.
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- EU30. De Paschale M, Manco MT, Belvisi L, et al. Epidemiology of hepatitis D virus (HDV) infection in an urban area of northern Italy. *Infection* 2012; 40: 485-91.
- EU31. Degertekin H, Sit D. The prevalence of hepatocellular carcinoma in patients with liver cirrhosis in Southeastern Turkey. *Turkish J Gastroenterol* 1998; 9: 217-20.
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Region of the Americas (AMR)

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AM18. Santos MD, Gomes-Gouvea MS, Nunes JD, et al. The hepatitis delta genotype 8 in Northeast Brazil: The North Atlantic slave trade as the potential route for infection. *Virus Res* 2016; 224: 6-11.

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South-East Asia Region (SEAR)

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SE10. Jat SL, Gupta N, Kumar T, et al. Prevalence of hepatitis D virus infection among hepatitis B virus-infected individuals in India. *Indian J Gastroenterol* 2015; 34: 164-8.

SE11. Jeong SH, Kim JM, Ahn HJ, et al. [The prevalence and clinical characteristics of hepatitis-delta infection in Korea]. *Korean J Hepatol* 2005; 11: 43-50.

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SE14. Regmi K, Shrestha JK, Kc S, et al. Prevalence of Hepatitis D Among Patients with Hepatitis B Viral Infection Attending a Tertiary Care Centre of Nepal. *J Nepal Med Assoc* 2017; 56: 417-20.

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SE16. Zaki H, Darmstadt GL, Baten A, Ahsan CR, Saha SK. Seroepidemiology of hepatitis B and delta virus infections in Bangladesh. *J Trop Pediatr* 2003; 49: 371-4.

Western Pacific Region (WPR)

WP16. Chen X, Xuan M, Yin Y. [Study of HDV infection in Shandong province]. *Zhonghua Liu Xing Bing Xue Za Zhi* 1998; 19: 138-40.

WP17. Lin HH, Lee SS, Yu ML, et al. Changing hepatitis D virus epidemiology in a hepatitis B virus endemic area with a national vaccination program. *Hepatology* 2015; 61: 1870-9.

WP18. Nakasone H, Sakugawa H, Shokita H, et al. Prevalence and clinical features of hepatitis delta virus infection in the Miyako Islands, Okinawa, Japan. *J Gastroenterol* 1998; 33: 850-4.

WP19. Oyunsuren T, Kurbanov F, Tanaka Y, et al. High frequency of hepatocellular carcinoma in Mongolia; association with mono-, or co-infection with hepatitis C, B, and delta viruses. *J Med Virol* 2006; 78: 1688-95.

WP20. Tsatsralt-Od B, Takahashi M, Nishizawa T, Endo K, Inoue J, Okamoto H. High prevalence of dual or triple infection of hepatitis B, C, and delta viruses among patients with chronic liver disease in Mongolia. *J Med Virol* 2005; 77: 491-9.

WP21. Turner S, O'Sullivan T, Wright P, Stuart K. Delta hepatitis testing, epidemiology and management in a large tertiary hospital in Australia. *J Gastroenterol Hepatol* 2016; 31: 84-5.

WP22. Wang Y, Yu F, Wang L, et al. Hepatitis D virus infections among hepatitis B virus (HBV) infected individuals with or without HIV co-infection: A retrospective cross-sectional study in one hospital in Beijing. *Hepatol Intern* 2017; 11.

WP23. Xu PS, Han SY, Sun CY, Zhao J. [The analysis of hepatitis delta virus infective markers in the hepatitis B virus infective people]. *Zhonghua Shi Yan He Lin Chuang Bing Du Xue Za Zhi* 2012; 26: 307-9.

Supplementary appendix 6. Included Studies by WHO Region: Selected Population Groups

African Region (AFR)

- AF4. Belyhun Y, Liebert UG, Maier M. Clade homogeneity and low rate of delta virus despite hyperendemicity of hepatitis B virus in Ethiopia. *Viol J* 2017; 14: 176.
- AF47. Birguel J, Ndong JG, Aurenche C, et al. Viral markers of hepatitis B, C and D and HB vaccination status of a health care team in a rural district of Cameroon. *Med Trop* 2011; 71: 201-2.
- AF6. Coffie PA, Tchounga BK, Bado G, et al. Prevalence of hepatitis B and delta according to HIV-type: a multi-country cross-sectional survey in West Africa. *BMC Infect Dis* 2017; 17: 466.
- AF48. Diop-Ndiaye H, Toure-Kane C, Etard JF, et al. Hepatitis B, C seroprevalence and delta viruses in HIV-1 Senegalese patients at HAART initiation (retrospective study). *J Med Virol* 2008; 80: 1332-6.
- AF14. Jaquet A, Wandeler G, Nouaman M, et al. Alcohol use, viral hepatitis and liver fibrosis among HIV-positive persons in West Africa: a cross-sectional study. *J Int AIDS Soc* 2017; 19: 21424.
- AF49. Ouedraogo HG, Kouanda S, Grosso A, et al. Hepatitis B, C, and D virus and human T-cell leukemia virus types 1 and 2 infections and correlates among men who have sex with men in Ouagadougou, Burkina Faso. *Viol J* 2018; 15: 194.
- AF50. Savassi F, Soares C, Pina De Araujo I, Gomes SA, Motta-Castro AR, Lago BV. Hepatitis B, delta and C viruses detection among HIV-infected individuals from Cape Verde. *J Viral Hepat* 2015; 22: 104-5.

Eastern Mediterranean Region (EMR)

- EM55. Aghasadeghi MR, Mohraz M, Bahramali G, et al. Frequency and genotype of hepatitis D virus infection in patients infected with HIV and those undergoing hemodialysis. *Hepat Mon* 2013; 13: e7481.
- EM56. Bahri F, Kargar Kheirabad A, Ghasemzadeh I, Shoja S, Gouklani H. Hepatitis viruses B and D and Human Immunodeficiency Virus infections in hemodialysis patients in the south of Iran: Prevalence and genotypes. *Hepat Mon* 2016; 16: e32971.
- EM57. Masood Z, Roghiye A, Hasan NM. Prevalence of HCV infection in hemodialysis patients of South Khorasan in comparison with HBV, HDV, HTLV I/II, And HIV infection. *Bangladesh J Med Science* 2014; 13: 36-9.
- EM58. Motamedifar MP, Taheri MM, Lankarani KBM, et al. The Prevalence and Risk Factors of Hepatitis Delta Virus in HIV/HBV Co-Infected Patients in Shiraz, Iran, 2012. *Iran J Med Sci* 2015; 40: 448-53.
- EM59. Njoh J, Zimmo S. Prevalence of antibody to hepatitis D virus among HBsAg-positive drug-dependent patients in Jeddah, Saudi Arabia. *East Afr Med J* 1998; 75: 327-8.
- EM13. Ramia S, El-Zaatari M, Sharara AI, Ramlawi F, Farhat B. Current prevalence of hepatitis delta virus (HDV) infection and the range of HDV genotypes in Lebanon. *Epidemiol Infect* 2007; 135: 959-62.
- EM60. Saeed Ur R, Hafiz A. Seroprevalence of HDV in hemodialysis and drug addicts in Karachi. *J Coll Physicians Surgeons Pakistan* 2000; 10: 470-2.
- EM61. Yadegarynia D, Hatamai H, Roodsari SR, Arab-Mazar Z. Seroprevalence of hepatitis B, C and D viral among hemodialysis patients in Tehran. *Iran J Microbiol* 2017; 9: 195-9.
- EM62. Zahedi MJ, Darvish Moghaddam S, Alavian SM, Dalili M. Seroprevalence of hepatitis viruses B, C, D and HIV infection among hemodialysis patients in Kerman province, south-east Iran. *Hepat Mon* 2012; 12: 339-43.
- EM52. Zaki SA, Abou Khatwa M, Mikail H. Prevalence of infection with delta virus in patients positive for hepatitis B surface antigen. *Hepatol Intern* 2011; 5: 153.

European Region (EUR)

- EU70. Aguilera A, Trastoy R, Barreiro P, et al. Decline and changing profile of hepatitis delta among injection drug users in Spain. *Antivir Ther* 2018; 23: 87-90.
- EU71. Arbune M, Georgescu C. Characteristics of Hepatitis B Co-infection and Disease Evolution in HIV-Positive Paediatric Patients in Romania. *Balkan Med J* 2013; 30: 263-7.
- EU72. Arribas JR, Gonzalez-Garcia JJ, Lorenzo A, et al. Single (B or C), dual (BC or BD) and triple (BCD) viral hepatitis in HIV-infected patients in Madrid, Spain. *AIDS* 2005; 19: 1361-5.

- EU73. Beguelin C, Moradpour D, Sahli R, et al. Hepatitis delta-associated mortality in HIV/HBV-coinfected patients. *J Hepatol* 2017; 66: 297-303.
- EU74. Boyd A, Lacombe K, Miallhes P, et al. Longitudinal evaluation of viral interactions in treated HIV-hepatitis B co-infected patients with additional hepatitis C and D virus. *J Viral Hepat* 2010; 17: 65-76.
- EU75. Catalani C, Giuffreda G, Baldi L, et al. Prevalence of HBV, HDV, HCV infection and alleged risk factors in the Pistoia (Italy) haemodialysis population. *Italian J Allergy Clin Immunol* 2008; 18: 22-9.
- EU76. Childs K, Welz T, Taylor C. Epidemiology and outcomes of hepatitis delta infection in a large, ethnically diverse UK HIV cohort. *HIV Med* 2010; 11: 11.
- EU77. Chircu LV, Barbacini IG, Goldoni E, et al. Evolution of liver damage due to hepatitis viruses (HBV, HDV, HCV), HIV and the development of AIDS. *Mediterranean J Infectious Parasitic Dis* 1998; 13: 21-7.
- EU78. Fernandez-Montero JV, Vispo E, Barreiro P, et al. Hepatitis delta is a major determinant of liver decompensation events and death in HIV-infected patients. *Clin Infect Dis* 2014; 58: 1549-53.
- EU79. Gaeta GB, Milensu S, Cozzi-Lepri A, et al. Prevalence and clinical outcome of hepatitis delta infection in patients with HIV/HBV coinfection. *J Hepatol* 2018; 68.
- EU80. Malliori M, Sypsa V, Psychogiou M, et al. A survey of bloodborne viruses and associated risk behaviours in Greek prisons. *Addiction* 1998; 93: 243-51.
- EU81. Rehak V, Krekulova L. Hidden microepidemic of hepatitis D in the population of injecting drug users in Prague, Czech Republic. *Hepatol Int* 2013; 7.
- EU82. Santana Rodriguez OE, Male Gil ML, HernandezSantana JF, Liminana Canal JM, Martin Sanchez AM. Prevalence of serologic markers of HBV, HDV, HCV and HIV in non-injection drug users compared to injection drug users in Gran Canaria, Spain. *Eur J Epidemiol* 1998; 14: 555-61.
- EU83. Soriano V, Grint D, d'Arminio Monforte A, et al. Hepatitis delta in HIV-infected individuals in Europe. *AIDS* 2011; 25: 1987-92.
- EU84. Stroffolini T, D'Egidio PF, Aceti A, et al. Hepatitis C virus infection among drug addicts in Italy. *J Med Virol* 2012; 84: 1608-12.
- EU14. Voiculescu M, Iliescu L, Ionescu C, et al. A cross-sectional epidemiological study of HBV, HCV, HDV and HEV prevalence in the Subcarpathian and South-Eastern regions of Romania. *J Gastrointest Liver Dis* 2010; 19: 43-8.
- EU85. Zerdali E, Vanli M, Bozkurt M, et al. Hepatitis A, B, C and D seroprevalence among Turkish HIV/AIDS patients. *Hepatol Int* 2018; 12.

Region of the Americas (AMR)

- AM21. Alvarado-Esquivel C, Sablon E, Martinez-Garcia S, Estrada-Martinez S. Hepatitis virus and HIV infections in inmates of a state correctional facility in Mexico. *Epidemiol Infect* 2005; 133: 679-85.
- AM22. Barton EN, King SD, Douglas LL. The seroprevalence of hepatitis and retroviral infection in Jamaican haemodialysis patients. *West Indian Med J* 1998; 47: 105-7.
- AM23. Braga WS, da Costa Castilho M, dos Santos IC, Moura MA, Segurado AC. Low prevalence of hepatitis B virus, hepatitis D virus and hepatitis C virus among patients with human immunodeficiency virus or acquired immunodeficiency syndrome in the Brazilian Amazon basin. *Rev Soc Bras Med Trop* 2006; 39: 519-22.
- AM24. Cocozella DR, Albuquerque MM, Borzi S, et al. [Prevalence of hepatic involvement, alcoholism, hepatitis B, C and HIV in patients with background history of drug use]. *Acta Gastroenterol Latino Am* 2003; 33: 177-81.
- AM25. de Almeida Pereira RA, Mussi AD, de Azevedo e Silva VC, Souto FJ. Hepatitis B Virus infection in HIV-positive population in Brazil: results of a survey in the state of Mato Grosso and a comparative analysis with other regions of Brazil. *BMC Infect Dis* 2006; 6: 34.
- AM26. Fainboim H, Gonzalez J, Fassio E, et al. Prevalence of hepatitis viruses in an anti-human immunodeficiency virus-positive population from Argentina. A multicentre study. *J Viral Hepat* 1999; 6: 53-7.

- AM27. Freitas SZ, Soares CC, Tanaka TS, et al. Prevalence, risk factors and genotypes of hepatitis B infection among HIV-infected patients in the State of MS, Central Brazil. *Braz J Infect Dis* 2014; 18: 473-80.
- AM28. Kucirka LM, Farzadegan H, Feld JJ, et al. Prevalence, correlates, and viral dynamics of hepatitis delta among injection drug users. *J Infect Dis* 2010; 202: 845-52.
- AM29. Leon P, Venegas E, Bengoechea L, et al. [Prevalence of infections by hepatitis B, C, D and E viruses in Bolivia]. *Rev Panam Salud Publica* 1999; 5: 144-51.
- AM30. Mahale P, Aka PV, Chen X, et al. Hepatitis D viremia among injection drug users in San Francisco. *J Infect Dis* 2018; 217: 1902-6.
- AM31. Mendes-Correa MC, Gomes-Gouvea MS, Alvarado-Mora MV, et al. Hepatitis delta in HIV/HBV co-infected patients in Brazil: is it important? *Int J Infect Dis* 2011; 15: e828-32.
- AM32. Oliveira ML, Bastos FI, Telles PR, et al. Prevalence and risk factors for HBV, HCV and HDV infections among injecting drug users from Rio de Janeiro, Brazil. *Braz J Med Biol Res* 1999; 32: 1107-14.
- AM33. Pierre A, Feldner A, Carvalho Filho RJ, et al. Prevalence of hepatitis delta virus among hemodialysis and renal transplant patients. *Int J Artif Organs* 2018; 41: 171-4.

South-East Asia Region (SEAR)

- SE17. Dinesha TR, Boobalan J, Sivamalar S, et al. Occult HBV infection in HIV-infected adults and evaluation of pooled NAT for HBV. *J Viral Hepat* 2018; 25: 718-23.
- SE18. Gibney L, Macaluso M, Kirk K, et al. Prevalence of infectious diseases in Bangladeshi women living adjacent to a truck stand: HIV/STD/hepatitis/genital tract infections. *Sex Transm Infect* 2001; 77: 344-50.
- SE7. Jaiswal SB, Chitnis DS, Salgia P, Sepaha A. Prevalence of hepatitis viruses among chronic renal failure patients on hemodialysis in central India. *Dial Transplant* 2002; 31: 234-40.
- SE2. Lusida MI, Surayah, Sakugawa H, et al. Genotype and subtype analyses of hepatitis B virus (HBV) and possible co-infection of HBV and hepatitis C virus (HCV) or hepatitis D virus (HDV) in blood donors, patients with chronic liver disease and patients on hemodialysis in Surabaya, Indonesia. *Microbiol Immunol* 2003; 47: 969-75.
- SE19. Mittal G, Gupta P, Thakuria B, Mukhiya GK, Mittal M. Profile of hepatitis B virus, hepatitis C virus, hepatitis d virus and human immunodeficiency virus infections in hemodialysis patients of a tertiary care hospital in uttarakhand. *J Clin Exp Hepatol* 2013; 3: 24-8.
- SE20. Prasetyo AA, Ariapramuda R, Kindi EA, et al. Men having sex with men in Surakarta, Indonesia: demographics, behavioral characteristics and prevalence of blood borne pathogens. *Southeast Asian J Trop Med Pub Health* 2014; 45: 1032-47.
- SE21. Prasetyo AA, Dharmawan R, Raharjo I, Hudiyono. Human leukocyte antigen-E alleles are associated with hepatitis C virus, Torque Teno virus, and Toxoplasma vo-infections but are not associated with hepatitis B virus, hepatitis D virus, and GB virus co-infections in human immunodeficiency virus patients. *J Glob Infect Dis* 2016; 8: 75-81.
- SE22. Prasetyo AA, Dirgahayu P, Sari Y, Hudiyono H, Kageyama S. Molecular epidemiology of HIV, HBV, HCV, and HTLV-1/2 in drug abuser inmates in central Javan prisons, Indonesia. *J Infect Dev Ctries* 2013; 7: 453-67.
- SE4. Theamboonlers A, Hansurabhanon T, Verachai V, Chongsrisawat V, Poovorawan Y. Hepatitis D virus infection in Thailand: HDV genotyping by RT-PCR, RFLP and direct sequencing. *Infection* 2002; 30: 140-4.

Western Pacific Region (WPR)

- WP3. Chang SY, Yang CL, Ko WS, et al. Molecular epidemiology of hepatitis D virus infection among injecting drug users with and without human immunodeficiency virus infection in Taiwan. *J Clin Microbiol* 2011; 49: 1083-9.
- WP4. Chen F, Zhang J, Guo F, et al. Hepatitis B, C, and D virus infection showing distinct patterns between injection drug users and the general population. *J Gastroenterol Hepatol* 2017; 32: 515-20.

- WP24. Chu FY, Chiang SC, Su FH, Chang YY, Cheng SH. Prevalence of human immunodeficiency virus and its association with hepatitis B, C, and D virus infections among incarcerated male substance abusers in Taiwan. *J Med Virol* 2009; 81: 973-8.
- WP8. Dunford L, Carr MJ, Dean J, et al. A multicentre molecular analysis of hepatitis B and blood-borne virus coinfections in Viet Nam. *PLoS One* 2012; 7: e39027.
- WP25. Hall N, Thuy LN, Diem Tdo T, et al. High prevalence of hepatitis delta virus among persons who inject drugs, Vietnam. *Emerg Infect Dis* 2015; 21: 540-3.
- WP26. Hung CC, Wu SM, Lin PH, et al. Increasing incidence of recent hepatitis D virus infection in HIV-infected patients in an area hyperendemic for hepatitis B virus infection. *Clin Infect Dis* 2014; 58: 1625-33.
- WP27. Huo TI, Wu JC, Wu SI, et al. Changing seroepidemiology of hepatitis B, C, and D virus infections in high-risk populations. *J Med Virol* 2004; 72: 41-5.
- WP28. Kao JH, Chen PJ, Lai MY, Chen DS. Hepatitis D virus genotypes in intravenous drug users in taiwan: decreasing prevalence and lack of correlation with hepatitis B virus genotypes. *J Clin Microbiol* 2002; 40: 3047-9.
- WP29. Kemp R, Miller J, Lungley S, Baker M. Injecting behaviours and prevalence of hepatitis B, C and D markers in New Zealand injecting drug user populations. *N Z Med J* 1998; 111: 50-3.
- WP17. Lin HH, Lee SS, Yu ML, et al. Changing hepatitis D virus epidemiology in a hepatitis B virus endemic area with a national vaccination program. *Hepatology* 2015; 61: 1870-9.
- WP30. Wang LJ, Lin SK, Chiang SC, Su LW, Chen CK. Risk factors for HIV, viral hepatitis, and syphilis among heroin users in northern Taiwan. *Subst Use Misuse* 2013; 48: 89-98.
- WP22. Wang Y, Yu F, Wang L, et al. Hepatitis D virus infections among hepatitis B virus (HBV) infected individuals with or without HIV co-infection: A retrospective cross-sectional study in one hospital in Beijing. *Hepatol Int* 2017; 11.

Supplementary appendix 7. Quality Assessment Checklist

1. Are the characteristics of included participants and the study setting described in detail?

Yes: The authors report the total number, age, and gender distribution of the participants, recruitment setting, clinical setting (if relevant), and selected population group (if relevant).

Partially reported: Relevant characteristics not completely reported.

No: None of the relevant characteristics of the participants are reported.

2. Are the eligibility criteria (inclusion and exclusion criteria) for the study explicit and appropriate?

Yes: The eligibility criteria are clearly stated and replicable. A statement on serological testing entry criteria, and any relevant important inclusion or exclusion criteria are clearly reported.

No: The eligibility criteria are not clearly stated.

3. Is the method of recruitment adequately described and appropriate?

Yes: There is a clear statement of the method of selection of participants.

Unclear: The method used to recruit participants is not clearly stated or no information is provided about the method used to recruit participants in the study.

No: The participants were recruited based on other criteria, such as access to intervention or availability of resources, that could lead to a significant selection bias.

4. Is the sample representative of the target population? Consider which of these target populations is represented:

A. General populations (e.g., community samples, antenatal, blood donors).

B. Selected population groups (e.g., injecting drug users, haemodialysis recipients, men who have sex with men, commercial sex workers, people with HCV or HIV).

C. Liver disease populations (e.g., people attending hepatology clinics, patients with cirrhosis or HCC).

Yes: The recruited population represents the underlying source population.

Unclear: It is not clear whether the recruitment population is representative.

No: The recruited population has material differences from the source population; it is not a representative sample, or there are significant inclusion or exclusion criteria that reduce applicability to the target population.

5. Detail any significant non-representative factors

a) Was this a previously known high prevalence community?

b) Was this a selective sub group identified as high risk e.g., sex workers

c) Were there significant exclusions (e.g., criteria based on ALT results or HCV results)

6. Is there significant bias arising from recall for follow up or retrospective data?

Yes: Only a subset of patients returned for analysis; there is incomplete analysis of the eligible participants (<80%).

Partially/unclear: There is incomplete analysis of eligible participants (<100%); or it is unclear whether all eligible individuals were tested.

No: All eligible individuals or a random sample of eligible individuals who consented to participate were tested.

7. Is the HBsAg laboratory method clearly described in the methods section?

Yes: The HBsAg method is reproducible from the description in the methods.

No: The HBsAg method is not adequately described to permit replication.

8. Is the anti-HDV and/or HDV RNA laboratory method clearly described in the methods section?

Yes: The method is reproducible from the description in the methods/referenced article.

No: The method is not adequately described to permit replication.

9. Is the response rate reported?

Yes: The number or proportion of patients who refused to participate is reported.

Unclear: It is not clear from the information provided how many participants who were eligible who did not participate.

No: The number or proportion of patients who did not participate is not reported.

10. Are both competing interest and source of support for the study reported?

Yes: Both competing interest and source of support (financial or other) received for the study are reported, or the absence of any competing interest and source of support is acknowledged.

Partially reported: Only one of these elements is reported.

No: Either there is no information available about competing interests and sources of support, or only one of these elements is reported.

Supplementary appendix 8. Binomial Mixed Model and Parametric Bootstrapping Procedure

A binomial mixed model was developed to estimate the probability of anti-HDV detection among HBsAg-positive people in general and hepatology clinic populations. Here we describe details of the statistical model and how prevalence estimates have been obtained.

Let Y_i denote the number of HBsAg-positive people who tested positive for anti-HDV out of n_i for the i -th survey, for $i = 1, \dots, n$. We then assumed that, conditionally on independent and identically distributed Gaussian random effects Z_i with mean zero and variance σ^2 , the Y_i was mutually independent. Binomial distribution with probability of a positive HDV test p_i , such that

$$\log \left\{ \frac{p_i}{1-p_i} \right\} = \beta_0 + \beta_1 LD_i + Z_i, \quad (1)$$

where LD_i is a dummy variable taking value 1 if the i -th survey was conducted in a hepatology clinic and 0 otherwise.

To estimate the unknown regression coefficients (β_0 and β_1) and random effects variance (σ^2), we accounted for data-quality variation as follows:

Let $[X]$ be a shorthand notation for “the distribution of the random variable X .” We then defined a weighted likelihood function for $\theta = (\beta_0, \beta_1, \sigma^2)$ as

$$L(\theta) = \prod_{i=1}^n L_i(\theta)^{w_i}, \quad (2)$$

where

$$L_i(\theta) = \int_{-\infty}^{+\infty} [Z_i][Y_i | Z_i] dZ_i \quad (3)$$

and $w_i = e^{q_i}$, with q_i denoting a quality score obtained via the principal component analysis run using answers to questions detailed in Supplementary appendix 7. Finally, we maximised (2) by approximating the integral in (3) with a Quasi Monte Carlo method.

Let C denote the set of surveys falling within a predefined geographical area (e.g., country or WHO region) and for a given target population (general population or hepatology clinic). To predict anti-HDV prevalence among HBsAg-positive people for C , we took $B = 10,000$ samples, say $p_i^{(j)}$, from the distribution of p_i conditioned on the data, also known as the predictive distribution for p_i , to obtain

$$\hat{p}_C = \frac{1}{B} \sum_{j=1}^B \frac{\sum_{i \in C} N_i w_i p_i^{(j)}}{\sum_{i \in C} N_i w_i}, \quad (4)$$

where N_i is the size of the target population for the i -th survey.

To predict anti-HDV prevalence without conditioning on HBsAg-positive status, we modified equation (4) as

$$\hat{p}_C = \frac{1}{B} \sum_{j=1}^B \frac{\sum_{i \in C} N_i w_i r_i^{(j)} p_i^{(j)}}{\sum_{i \in C} N_i w_i}, \quad (5)$$

where $r_i^{(j)}$ corresponds the j -th sample drawn from a prior distribution for the probability of carrying HBsAg. The prior distributions were obtained by matching the lower and upper bound of the reported estimate of HBsAg prevalence to the 0.025 and 0.975 quantiles of a Beta distribution.

The number of people with anti-HDV was computed by multiplying the regional HBV prevalence estimate (from the WHO Global Hepatitis Report 2017, (Available at: <http://www.whohbsagdashboard.com/#global-strategies>) and HBsAg conditional anti-HDV prevalence estimates and multiplying the resulting anti-HDV prevalence with the United National Population Division (UNPD) population estimates for 2018 (World Population Prospects 2019 (Available at: <https://population.un.org/wpp/>), with countries grouped by WHO regions.

To derive 95% confidence intervals (CIs) we used a parametric bootstrap procedure. From the 95% CIs for the population level HBV prevalence and the HBsAg conditional anti-HDV prevalence estimates, we found the beta distributions for the 2.5th and 97.5th percentiles which matched the 95% CI bounds for HBV prevalence and anti-HDV, conditional on HBsAg prevalence respectively. We generated N=10,000,000 prevalence samples from each distribution and multiplied these. This was used to obtain a distribution of N estimates of the population anti-HDV prevalence.

We then took the empirical 2.5th and 97.5th quantiles of this sample to be the lower and upper bound of the 95% CI for the population level anti-HDV estimate. To take the product of prevalences, we had to assume both prevalence estimates (HBV, HDV conditional on HBsAg) to be independent. As they were obtained from separate datasets, this was a reasonable assumption. Finally, the CI for the number of people with anti-HDV was derived by multiplying the 95% CI for the anti-HDV prevalence with the UNPD 2018 population estimates.

Statistical source code for the binomial mixed model and the parametric bootstrapping procedure are available on request from the authors.

Supplementary appendix 9. Data Sources for Estimations of Population Sizes

Country / Group	Source	URL
Albania	Instituti i Statistikës	http://www.instat.gov.al/
Algeria	Office National des Statistiques	http://www.ons.dz/
Argentina	Instituto Nacional de Estadística y Censos	http://www.indec.gov.ar/
Benin	Institut National de la Statistique	http://www.insae-bj.org/
Bolivia	Instituto Nacional de Estadística	http://www.ine.gob.bo/
Botswana	Central Statistics Office	http://www.cso.gov.bw/
Brazil	Instituto Brasileiro de Geografia e Estatística	http://www.igbe.gov.br/
Bulgaria	National Statistical Institute	http://www.nsi.bg/
Burkina Faso	Institut National de la Statistique et de la Demographie	http://www.insd.bf/
Cameroon	Bureau Central des Recensements et des Etudes de Population	http://www.bucrep.cm/
Cameroon	Statistics Cameroon	http://www.statistics-cameroon.org/
CAR	Division des Statistiques, des Etudes Economiques et Sociales	http://www.stat-centrafrique.com/
Colombia	Departamento Administrativo Nacional de Estadística	http://www.dane.gov.co
Cote d'Ivoire	Institut National de la Statistique	http://www.ins.ci/
Czechia	Czech Statistical Office	http://www.czso.cz/
Denmark	Denmark Statistik (StatBank)	http://www.statbank.dk
Egypt	Central Agency for Public Mobilization and Statistics	http://www.capmas.gov.eg/
Ethiopia	Central Statistical Agency	http://www.csa.gov.et/
Europe	Eurostat	https://ec.europa.eu/eurostat/en/
France	Institut National de la Statistique et des Études Économiques	http://www.insee.fr/
Gabon	Direction Générale des Statistiques	http://www.statgabon.ga/
Gambia	Gambia Bureau of Statistics	http://www.gbos.gov.gm
Germany	Statistisches Bundesamt	http://www.destatis.de/
Germany	Statistik Hessen	https://statistik.hessen.de/
Ghana	Ghana Statistical Service	http://www.statsghana.gov.gh/
Global	United Nations/ UNAIDS AIDS Info	http://aidsinfo.unaids.org/
Guinea-Bissau	Instituto Nacional de Estadística	http://www.stat-guineebissau.com/
Iran	Statistical Centre	http://www.amar.org.ir
Iraq	Central Organization for Statistics and Information Technology	http://www.cosit.gov.iq/
Italy	Istituto Nazionale di Statistica	http://www.istat.it
Jamaica	Jamaica Information Service	https://jis.gov.jm/
Kosovo	Kosovo Agency of Statistics	http://www.ask.rks-gov.net/
Latin America	Ethnologue: Languages of the World	http://www.ethnologue.com/
Libya	Bureau of Statistics and Census	http://www.bsc.ly/
Mali	Institut National de la Statistiques	http://www.instat.gov.ml/
Mauritius	Office National de la Statistique	http://www.ons.mr/
Mexico	Instituto Nacional de Estadística y Geografía	http://www.inegi.org.mx/
Republic of Moldova	Departamentul Statistica si Sociologie	http://www.statistica.md/

Mozambique	Instituto Nacional de Estatística	http://www.ine.gov.mz/
Nigeria	National Bureau of Statistics	http://www.nigerianstat.gov.ng/
Pakistan	Pakistan Bureau of Statistics	http://www.pbs.gov.pk/
Peru	Instituto Nacional de Estadística e Informática	http://www.inei.gob.pe
Romania	Romania National Institute of Statistics	http://www.insse.ro/
Russian Federation	Federal State Statistics Service	http://www.gks.ru/
Saudi Arabia	Central Department of Statistics & Information	http://www.cdsi.gov.sa/
Senegal	Agence Nationale de la Statistique et de la Démographie	http://www.ansd.sn/
South Africa	Statistics South Africa	http://www.statssa.gov.za/
Spain	Instituto Nacional de Estadística	http://www.ine.es/
Tajikistan	State Statistical Committee	http://www.stat.tj/
Togo	Direction Générale de la Statistique et de la Comptabilité Nationale	http://www.stat-togo.org
Tunisia	Institut National de la Statistique	http://www.ins.tn/
Turkey	State Institute of Statistics	http://www.turkstat.gov.tr
Uganda	Uganda Bureau of Statistics	http://www.ubos.org
United Kingdom	Office for National Statistics	https://www.ons.gov.uk/
USA	US Census Bureau	http://www.census.gov/
Uzbekistan	State Committee of Uzbekistan on Statistics	http://www.stat.uz/
Venezuela	Instituto Nacional de Estadística	http://www.pib.socioambiental.org
Yemen	Central Statistical Organisation	http://www.cso-yemen.org/
Country-level population estimates	United Nations Population Division. World Population Prospects: 2019 Revision; Census reports and other statistical publications from national statistical offices; Eurostat: Demographic Statistics; United Nations Statistical Division: Population and Vital Statistics Report; US Census Bureau: International Database; Secretariat of the Pacific Community: Statistics and Demography Programme	https://population.un.org/wpp/ ; https://ec.europa.eu/eurostat/web/population-demography-migration-projections/data ; https://unstats.un.org/unsd/demographic-social/products/vitstats/index.cshtm ; https://www.census.gov/programs-surveys/international-programs/about/idb.html ; https://prism.spc.int/
PWID	UNAIDS estimates	http://aidsinfo.unaids.org/ ; Ref ^a
MSM	UNAIDS estimates	http://aidsinfo.unaids.org/
HCV-positive	Polaris Observatory HCV Collaborators	Ref ^b
HIV-positive	UNAIDS estimates	http://aidsinfo.unaids.org/

^aDegenhardt L, Peacock A, Colledge S, et al. Global prevalence of injecting drug use and sociodemographic characteristics and prevalence of HIV, HBV, and HCV in people who inject drugs: a multistage systematic review. *Lancet Glob Health* 2017; 5: e1192-e1207.

^bGlobal prevalence and genotype distribution of hepatitis C virus infection in 2015: a modelling study. *Lancet Gastroenterol Hepatol* 2017; 2: 161-76.

Abbreviations: CAR, Central African Republic; USA, United States of America; PWID, People who inject drugs; MSM, Men who have sex with men; HCV, Hepatitis C virus.

Supplementary appendix 10. Included Populations by WHO Region

	AFR	EMR	EUR	AMR	SEAR	WPR	Total
Overall population ^a , millions	1,0152.8	715.4	928.5	1,006.5	1,982.2	1,945.7	7,631.1
General population							
Population represented ^b , millions	158.9	308.7	313.2	360.0	148.4	236.9	1,526.1
Number tested for anti-HDV	7,940	2,455	6,924	417	170	6,118	24,024
Community survey	4,400	801	259	210	29	6,043	11,742
Blood donors	1,244	1,645	5,545	207	141	30	8,812
Antenatal clinic	1,098	0	1,118	0	0	0	2,216
HIV clinic ^c	1,088	0	0	0	0	0	1,088
Student / occupational setting	110	9	2	0	0	45	166
Hepatology clinic populations							
Population represented ^b , millions	96.4	365.0	527.2	19.5	304.0	245.5	1,557.6
Number tested for anti-HDV	3,526	15,194	58,958	1,951	2,364	6,208	88,201
Selected populations groups^d							
Population represented ^b , millions	1.7	1.2	6.9	7.2	1.0	0.9	19.1
Number tested for anti-HDV	314	423	2,870	813	216	3,432	8,068
People who inject drugs	0	112	713	166	67	1,696	2,754
Haemodialysis recipients	17	105	19	119	121	41	422
Men who have sex with men	68	0	29	0	14	0	111
Commercial sex workers	0	0	0	1	14	424	439
HCV-positive	0	19	891	257	0	432	1,599
HIV-positive	229	187	1,218	270	0	839	2,743

^aUnited Nations Population Division 2018 population estimates (World Population Prospects 2019).

^bRefers to the sum of the catchment populations represented by included studies, excluding overlapping or repeated samples. For hepatology clinic populations, this refers to the catchment area of the clinical service. For selected population groups, this was based on estimates of the size of the population at risk.

^cIncluded in general populations in countries with adult HIV prevalence >1%.

^dData from selected population groups partially overlap with hepatology clinic and/or general population samples.

Abbreviations: AFR, African Region; EMR, Eastern Mediterranean Region; EUR, European Region; AMR, Region of the Americas; SEAR, South-East Asian Region; WPR, Western Pacific Region; HCV, Hepatitis C virus.

Supplementary appendix 11. Isolated Populations Tested for Anti-HDV

Country Reference	Population and Region	Year	Tested for HBsAg Number	HBsAg prevalence % (95% CI)	HBsAg-positive tested for anti-HDV Number	Anti-HDV prevalence % (95% CI)
Bolivia AM13	Yurakare, Yuki and Trinitario ethnicities, Bolivian Amazonia	1996	366	1.6 (0.8, 3.5)	6	16.7 (3.0, 56.4)
Bolivia AM13	Rural Andean Highlands	1996	187	0.5 (0.1, 3.0)	1	0.0 (0.0, 79.3)
Brazil AM7	Jurua, Purus and Wood Valleys, Amazon State	<2001	688	9.7 (7.7, 12.2)	67	13.4 (7.2, 23.6)
Brazil AM8	Labrea, South West Amazonas	2005-2006	787	11.8 (9.7, 14.3)	93	41.9 (32.4, 52.1)
Brazil AM9	Acre and Purus Rivers, States of Acre and Amazonas	1997	349	5.2 (3.3, 8.0)	18	66.7 (43.7, 83.7)
Brazil AM14	Parakana Indians Apakterewa and Xingu villages	1995-2003-2005	196-258	5.6 (3.2, 9.8)-3.9 (2.1, 7.0)	11-10	0.0 (0.0, 25.9)-0.0 (0.0, 27.8)
Brazil AM15	Acre State	2002	2656	3.4 (2.7, 4.1)	89	21.3 (14.1, 31.0)
Brazil AM6	Eastern Amazon	2011-2017	603	0.5 (0.1, 1.5)	3	0.0 (0.0, 56.1)
Colombia AM10	Amazonas State	2011-2013	862	2.7 (1.8, 4.0)	23	26.1 (12.5, 46.5)
Venezuela AM11	Piaroan communities	2002-2004	412	5.1 (3.4, 7.7)	21	19.0 (7.7, 40.0)
Venezuela AM11	Yanomami communities	2002-2004	231	14.3 (10.4, 19.4)	33	6.1 (1.7, 19.6)
USA AM12	Alaska Native Population, Bristol Bay	1993-1994	603	8.8 (6.8, 11.3)	53	0.0 (0.0, 6.8)
Denmark EU15	Itilleq, Greenland	2005-2007	115	27.0 (19.7, 35.7)	31	51.6 (34.8, 68.0)
Denmark EU15	Sarfannuaq, Greenland	2005-2007	72	5.6 (2.2, 13.4)	4	0.0 (0.0, 49.0)
Denmark EU17	West Greenland	<2012	153	2.6 (1.0, 6.5)	4	0.0 (0.0, 49.0)
Denmark EU17	East Greenland	<2012	288	28.8 (23.9, 34.3)	483	1.2 (0.2, 6.5)
Greece EU16	Archangelos, Rhodes	1997	1938	11.5 (10.2, 13.0)	223	25.1 (19.9, 31.2)
Greece EU16	Archangelos, Rhodes	2013	1076	3.3 (2.3, 4.5)	35	20.0 (10.0, 35.9)
India SE6	Andaman and Nicobar Islands	2002	NS	NS	223	3.6 (1.8, 6.9)
India SE5	Upper Dibang Valley, Uttar Pradesh	2005	NS	NS	93	35.5 (26.5, 45.6)

Abbreviations: HBsAg, Hepatitis B surface antigen; CI, Confidence interval; USA, The United States of America; NS, Not stated

Supplementary appendix 12. List of anti-HDV assays used in included studies

Method and Manufacturer ^a	N	%
ELISA, Abbott, USA	71	18.0
ELISA, ETI-AB-DELTAK-2, Diasorin, Italy	66	16.8
Method not specified	51	12.9
ELISA, Dia.Pro Diagnostic Bioprobes, Italy	50	12.7
ELISA, Diasorin, Italy	39	9.9
ELISA, manufacturer not specified	39	9.9
ELISA, Murex anti-delta, Wellcozyme, United Kingdom	9	2.3
ELISA, Radim Diagnostics, Germany	8	2.0
ELISA, Diagnostic Automation, USA	6	1.5
ELISA, Wantai, China	6	1.5
Architect chemiluminescence assay, Abbott, USA	4	1.0
ELISA, In-house assay	4	1.0
RIA, Abbott, USA	4	1.0
ELISA, DRG Diagnostics, Germany	3	0.8
ELISA, Hepanostika, Biomeriux, France	3	0.8
ELISA, Biovendor, Germany	3	0.8
ELISA, Biokit, Spain	2	0.5
RIA, Diasorin, Italy	2	0.5
ELISA, Globe Diagnostics, Italy	2	0.5
ELISA, Institute of Chinese Academy of Preventive Medicine, China	2	0.5
ELISA, RPC Diagnostics Systems, Russian Federation	2	0.5
ELISA, Sorin Biomedica, Italy	2	0.5
Western blot	2	0.5
ELISA, MBS, Italy	1	0.3
ELISA, Beijing Bell Bioengineering, China	1	0.3
Cobas e601 chemiluminescence assay, Roche, Switzerland	1	0.3
ELISA, Creative Diagnostics, USA	1	0.3
ELISA, Deltassay, Sanofi Pasteur Diagnostic, France	1	0.3
ELISA, Detect, Biochem Immunosystems, Italy	1	0.3
ELISA, Equipar, Italy	1	0.3
ELISA, Genway Biotech, USA	1	0.3
ELISA, InGen, France	1	0.3
ELISA, Ortho Diagnostics, USA	1	0.3
RIA, manufacturer not specified	1	0.3
RIA, Dianabot, Japan	1	0.3
ELISA, SmarTest Diagnostics, Israel	1	0.3
ELISA, TKA 4HD, Teknolabo, Italy	1	0.3

^aSome studies used more than one assay.

Abbreviations: ELISA Enzyme linked immunosorbent assay; USA, The United States of America; RIA Radioimmunoassay.

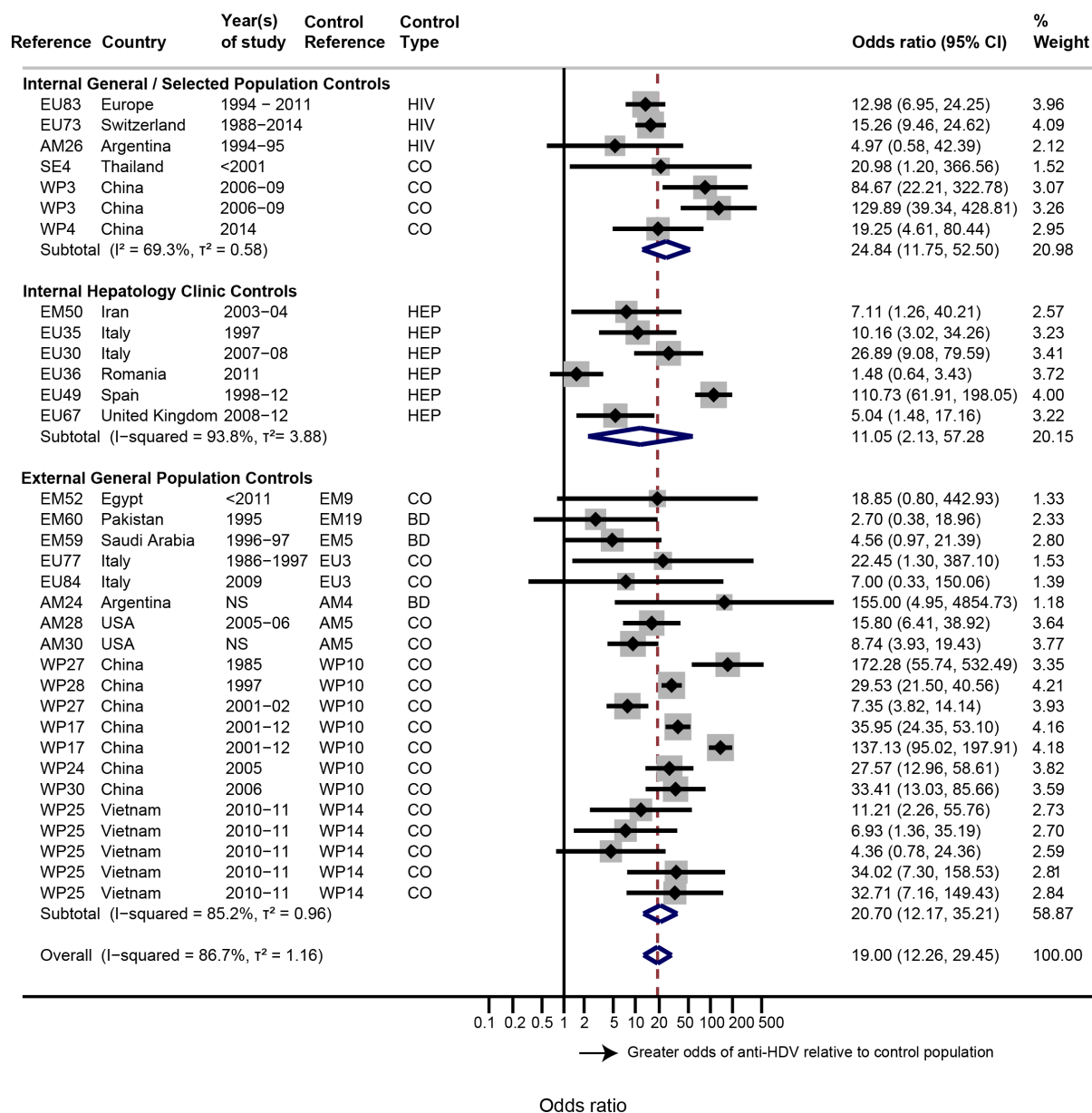
Supplementary appendix 13. Country-Level Anti-HDV Prevalence Estimates in General Populations (HBsAg-Positive and Total) and Hepatology Clinic Populations (HBsAg-Positive)

WHO Region	Country	General Populations % (95% CI)		Hepatology Clinic Populations % (95% CI)
		HBsAg-positive	Total	HBsAg-positive
AFR	Algeria	-	-	5.6 (2.2, 10.3)
AFR	Benin	10.5 (3.7, 20.4)	1.7 (0.6, 3.4)	-
AFR	Botswana	3.7 (0.2, 15.6)	0.1 (0.01, 0.7)	-
AFR	Burkina Faso	1.7 (1.1, 2.4)	0.2 (0.1, 0.2)	3.5 (0.2, 13.3)
AFR	Cameroon	11.1 (8.9, 14.9)	1.0 (0.9, 1.2)	28.7 (22.2, 35.4)
AFR	CAR	11.2 (6.9, 16.2)	0.9 (0.6, 1.4)	43.6 (21.2, 67.7)
AFR	Cote d'Ivoire	14.7 (6.2, 26.0)	1.2 (0.5, 2.3)	-
AFR	Ethiopia	3.4 (0.9, 7.5)	0.3 (0.1, 0.6)	9.7 (5.6, 14.8)
AFR	Gabon	22.0 (16.1, 29.2)	2.0 (1.5, 2.8)	-
AFR	Gambia	1.4 (0.7, 2.3)	0.1 (0.05, 0.1)	11.5 (7.4, 16.3)
AFR	Ghana	5.6 (3.1, 8.9)	0.5 (0.3, 0.8)	11.8 (6.3, 18.8)
AFR	Guinea-Bissau	23.9 (15.1, 34.2)	3.9 (2.4, 5.9)	-
AFR	Malawi	1.8 (0.3, 4.6)	0.2 (0.05, 0.5)	-
AFR	Mali	6.5 (0.2, 31.9)	0.8 (0.02, 3.9)	-
AFR	Mauritania	19.4 (16.1, 22.9)	2.9 (2.4, 3.5)	32.3 (27.1, 37.5)
AFR	Mozambique	0.7 (0.1, 2.2)	0.03 (0.003, 0.1)	-
AFR	Nigeria	10.6 (5.1, 18.0)	1.6 (0.7, 3.1)	4.7 (3.2, 7.8)
AFR	Senegal	3.0 (1.1, 5.8)	0.2 (0.1, 0.5)	13.8 (5.6, 26)
AFR	South Africa	1.6 (0.4, 4.0)	0.1 (0.02, 0.2)	-
AFR	United Republic of Tanzania	5.0 (2.6, 8.3)	0.2 (0.1, 0.4)	-
AFR	Togo	18.5 (4.5, 40.0)	1.1 (0.2, 2.6)	14.0 (5.7, 26.5)
AFR	Uganda	3.3 (1.3, 6.1)	0.2 (0.1, 0.4)	-
EMR	Egypt	3.8 (1.9, 6.6)	0.1 (0.03, 0.1)	7.4 (2.6, 16.2)
EMR	Iran	3.2 (1.9, 5.3)	0.1 (0.04, 0.1)	11.5 (9.7, 13.3)
EMR	Iraq	3.7 (0.1, 16.1)	0.2 (0.01, 0.8)	-
EMR	Lebanon	1.4 (0.2, 4.1)	0.02 (0.003, 0.05)	3.0 (0.9, 7.0)
EMR	Libya	-	-	2.6 (1.2, 4.6)
EMR	Pakistan	9.0 (1.6, 22.7)	0.6 (0.1, 1.7)	20.5 (11.8, 33.2)
EMR	Saudi Arabia	3.6 (0.7, 8.9)	0.1 (0.01, 0.2)	6.7 (4.5, 9.6)
EMR	Tunisia	4.7 (2.1, 8.1)	0.2 (0.1, 0.4)	5.7 (2.5, 12.4)
EMR	Yemen	2.9 (0.1, 11.8)	0.1 (0.004, 0.4)	-
EUR	Albania	-	-	10.6 (6.4, 15.7)
EUR	Belarus	3.4 (1.8, 5.4)	0.1 (0.05, 0.1)	21.7 (15.5, 28.5)
EUR	Bulgaria	-	-	10.7 (6.0, 16.2)
EUR	France	2.0 (1.7, 2.4)	0.01 (0.004, 0.01)	4.7 (4.1, 5.4)
EUR	Germany	2.6 (1.1, 4.8)	0.02 (0.01, 0.03)	6.6 (5.9, 7.3)
EUR	Israel	-	-	9.7 (3.5, 18.5)
EUR	Italy	3.2 (0.2, 13.2)	0.02 (0, 0.1)	9.3 (8.4, 10.2)
EUR	Kosovo	4.0 (0.4, 12.1)	0.1 (0.01, 0.3)	-
EUR	Republic of Moldova	15.0 (9.4, 22.6)	1.3 (0.8, 1.9)	46.4 (33.5, 59.5)
EUR	Poland	-	-	5.1 (1.6, 10.6)

EUR	Romania	7.0 (1.0, 27.2)	0.4 (0.1, 1.5)	32.0 (29.7, 34.3)
EUR	Russian Federation	-	-	19.7 (16.5, 23.0)
EUR	Slovenia	-	-	0.4 (0.1, 0.7)
EUR	Spain	-	-	10.8 (8.4, 13.5)
EUR	Tajikistan	-	-	24.0 (13.1, 36.3)
EUR	Turkey	3.1 (1.5, 5.4)	0.1 (0.1, 0.2)	32.7 (26.8, 39.1)
EUR	United Kingdom	2.1 (0.9, 3.9)	0.02 (0.001, 0.1)	5.3 (4.7, 6.0)
EUR	Uzbekistan	-	-	17.3 (10.4, 25.6)
AMR	Argentina	2.7 (0.5, 7.6)	0.004 (0.001, 0.01)	-
AMR	Brazil	7.2 (1.1, 30.5)	0.05 (0.01, 0.2)	11.0 (8.7, 13.7)
AMR	Canada	-	-	2.6 (1.8, 3.6)
AMR	Peru	5.7 (0.2, 26.1)	0.1 (0.001, 0.3)	-
AMR	USA	5.9 (3.0, 9.8)	0.02 (0.01, 0.03)	2.6 (1.8, 3.6)
SEAR	Bangladesh	4.1 (0.2, 17.5)	0.3 (0.01, 1.4)	15.4 (11.3, 20.6)
SEAR	India	-	-	4.9 (3.5, 6.6)
SEAR	Indonesia	0.8 (0.1, 2.8)	0.02 (0.002, 0.07)	0.7 (0.1, 2.0)
SEAR	Nepal	-	-	2.0 (0.1, 6.8)
SEAR	Republic of Korea	-	-	2.0 (1.2, 3.1)
SEAR	Thailand	1.6 (0.1, 6.1)	0.02 (0.002, 0.1)	-
WPR	Australia	1.3 (0.1, 4.5)	0.01 (0.0, 0.02)	4.4 (3.1, 6.0)
WPR	China	2.1 (1.6, 2.7)	0.4 (0.3, 0.5)	4.5 (4.0, 5.1)
WPR	Japan	8.5 (5.9, 11.4)	0.7 (0.5, 0.9)	21.3 (17.4, 25.5)
WPR	Mongolia	36.9 (31.3, 43.1)	4.0 (3.2, 4.8)	74.7 (69.3, 79.8)
WPR	Vietnam	10.8 (5.2, 18.2)	1.4 (0.7, 2.3)	-

Abbreviations: HBsAg, Hepatitis B surface antigen; CI, confidence interval; AFR, African Region; EMR, Eastern Mediterranean Region; EUR, European Region; AMR, Region of the Americas; SEAR, South-East Asian Region; WPR, Western Pacific Region; CAR, Central African Republic; USA, The United States of America

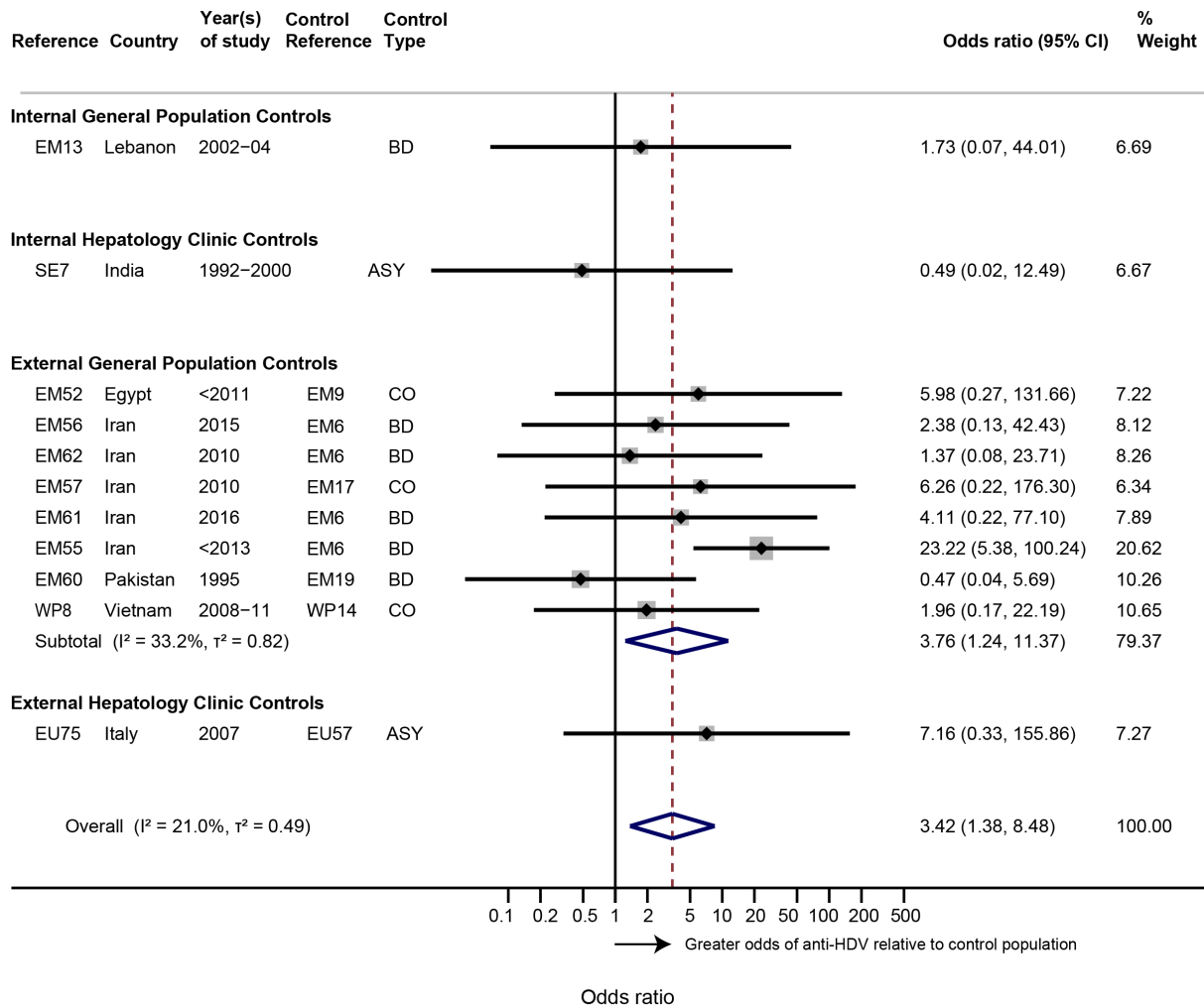
Supplementary appendix 14. Forest Plot of Anti-HDV Prevalence Among People Who Inject Drugs Relative to Comparator Populations^a



^aComparator populations: HIV clinic attendees (HIV); Community (CO), Hepatology clinic population (HEP), Blood donors (BD); ^bEurope includes 34 European countries and Argentina.

Abbreviations: CI, confidence interval; USA, The United States of America.

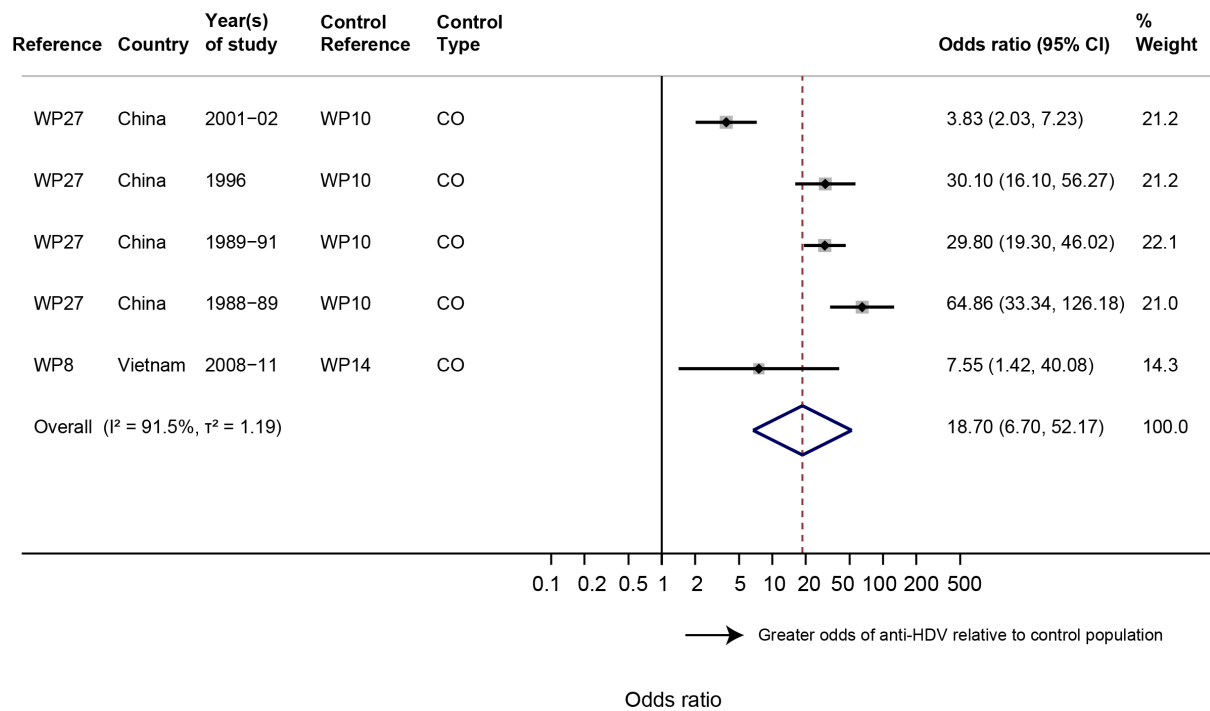
Supplementary appendix 15. Forest Plot of Anti-HDV Prevalence Among Haemodialysis Recipients Relative to Comparator Populations^a



^aComparator populations: Blood donors (BD), Asymptomatic HBsAg-positive people (ASY), Community (CO).

Abbreviations: CI, confidence interval.

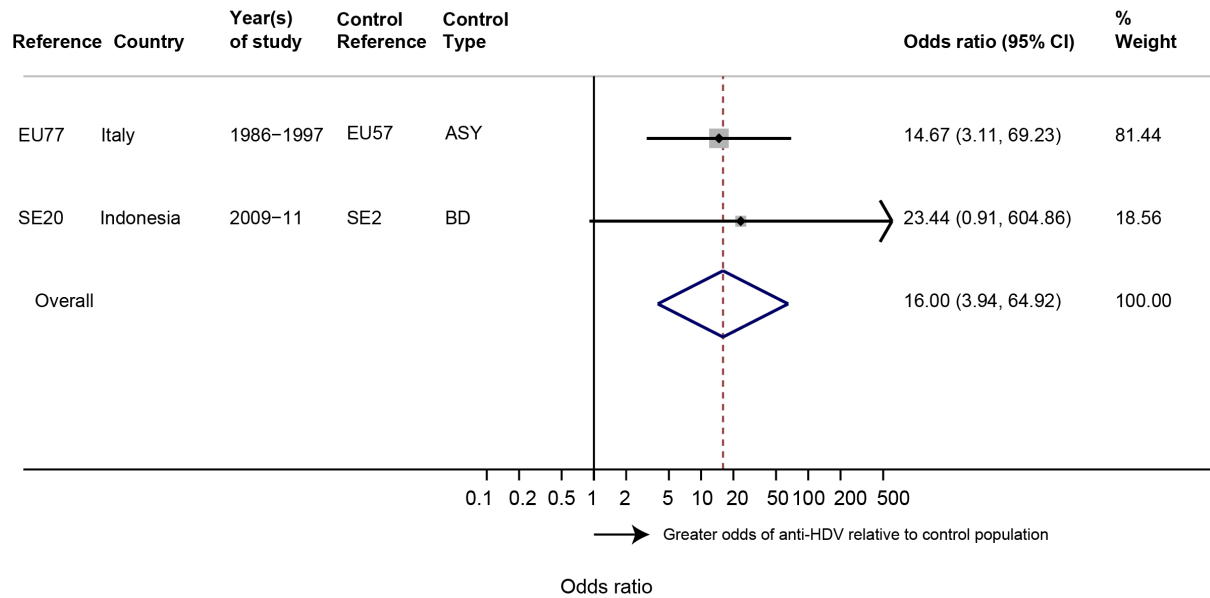
Supplementary appendix 16. Forest Plot of Anti-HDV Prevalence among Commercial Sex Workers Relative to Comparator Populations^a



^aComparator populations: All were external community (CO) populations from the same geographic region.

Abbreviations: CI, confidence interval.

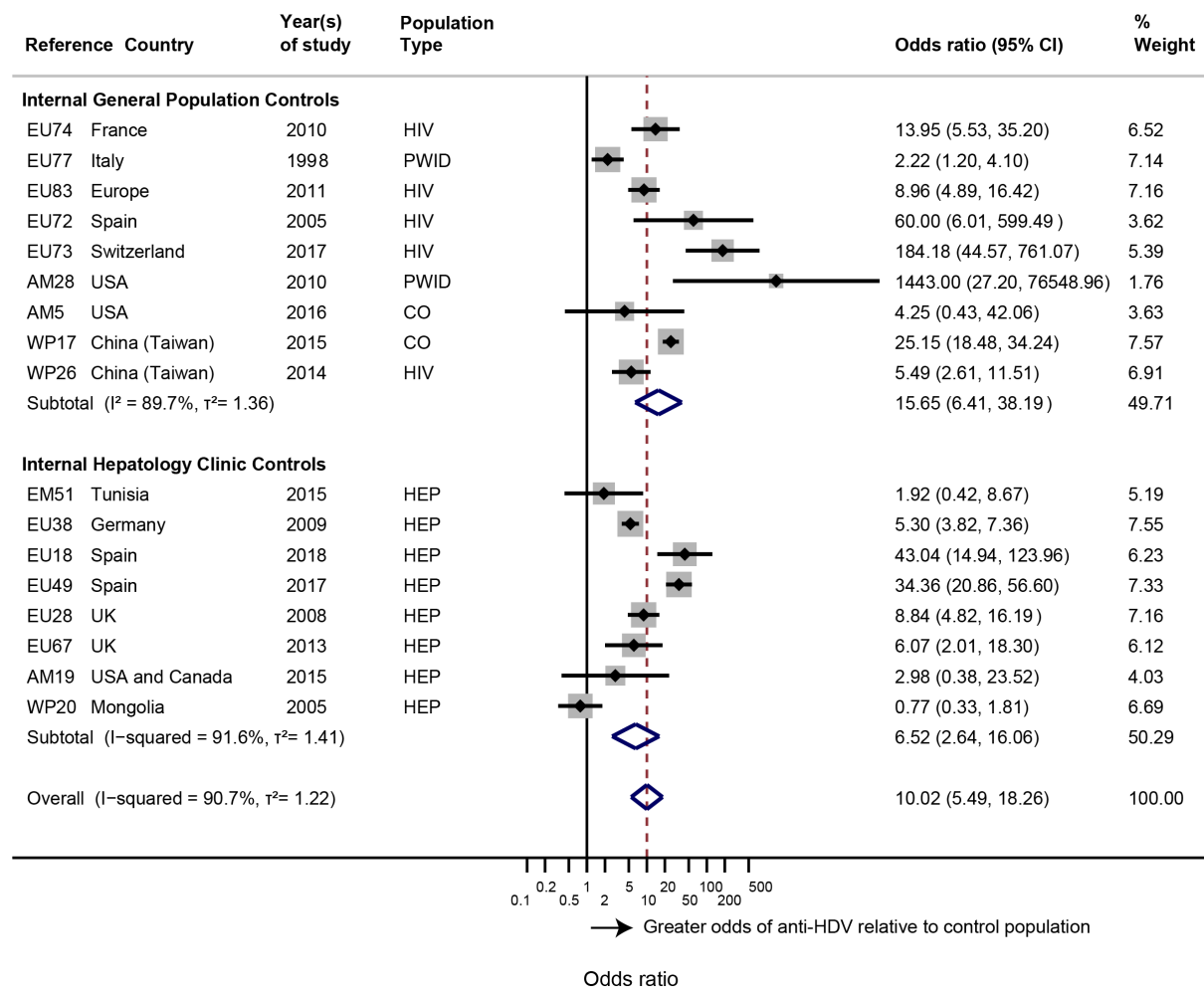
Supplementary appendix 17. Forest Plot of Anti-HDV Prevalence Among Men Who Have Sex with Men Relative to Comparator Populations^a



^aComparator populations: Asymptomatic HBsAg-positive people (ASY), Blood donors (BD).

Abbreviations: CI, confidence interval.

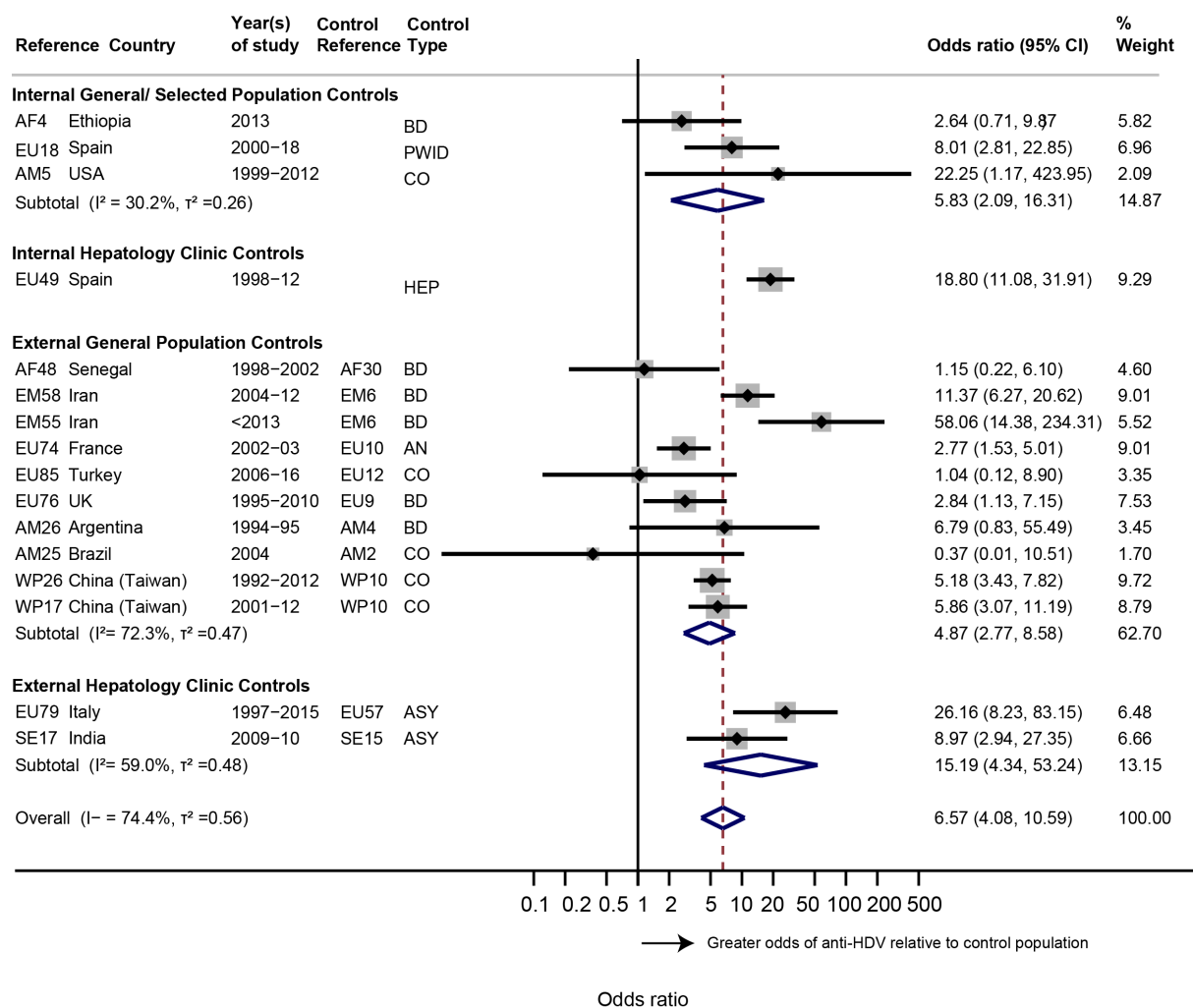
Supplementary appendix 18. Forest Plot of Anti-HDV Prevalence Among Hepatitis C Virus (HCV)-Antibody Positive People Relative to Comparator Populations^a



^aComparator populations: HIV clinic attendees (HIV), people who inject drugs (PWID), Community (CO), Hepatology clinic attendees (all, analysed by HCV status) (HEP); ^bEurope includes 34 European countries and Argentina.

Abbreviations: CI, confidence interval; USA, The United States of America.

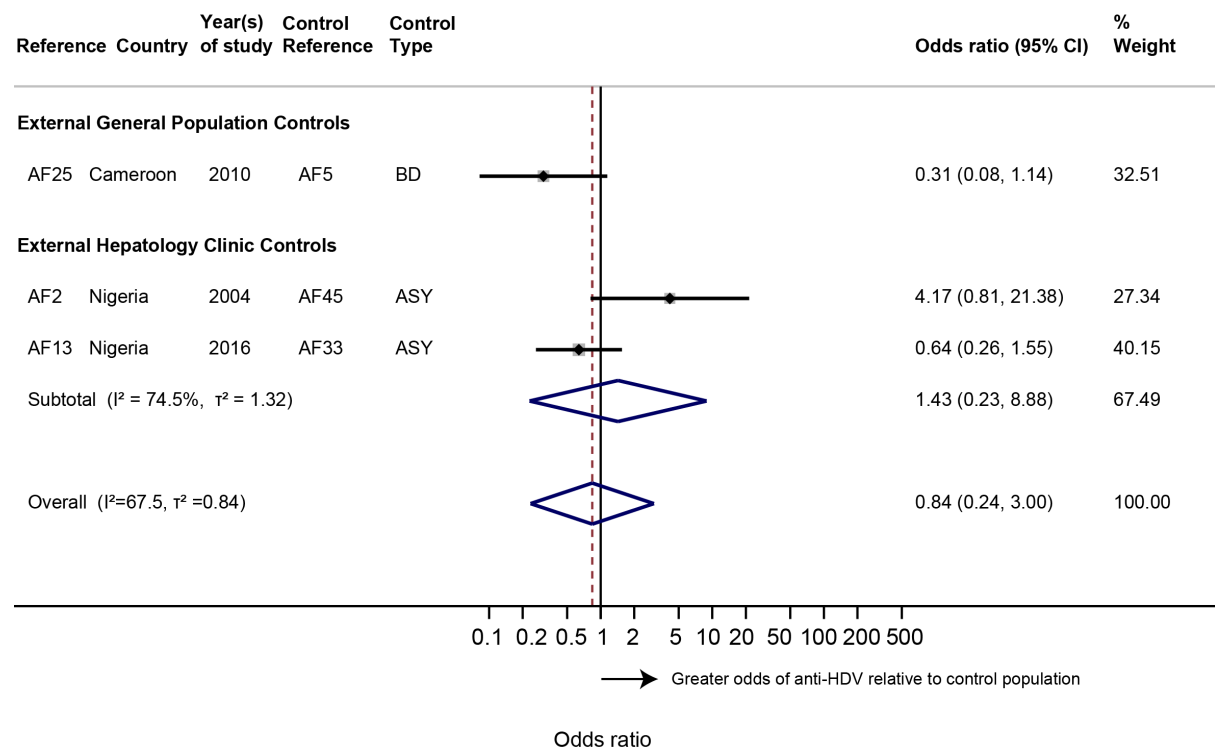
Supplementary appendix 19. Forest Plot of Anti-HDV Prevalence Among People with HIV Relative to Comparator Populations^a in Countries Without Generalised HIV Epidemics^b



^aComparator populations: Blood donors (BD), HIV-positive and HIV-negative people who inject drugs (PWID), Community (CO), Hepatology clinic population (all, analysed by HIV status) (HEP), Antenatal population (AN), Asymptomatic HBsAg-positive people (ASY); ^bGeneralised HIV epidemics were defined by an adult HIV prevalence >1%.

Abbreviations: CI, confidence interval; USA, The United States of America.

Supplementary appendix 20. Forest Plot of Anti-HDV Prevalence Among People with HIV Relative to Comparator Populations^a in Countries with Generalised HIV Epidemics^b



^aComparator populations: Blood donors (BD), Asymptomatic HBsAg-positive people (ASY); Generalised HIV epidemics were defined by an adult HIV prevalence >1%.

Abbreviations: CI, confidence interval.

Supplementary appendix 21. Meta-Regression Analysis of Characteristics Associated with HDV RNA Detection Among People with Anti-HDV

Characteristic	HDV RNA prevalence % (95% CI)	Co-efficient ^a (95% CI)	P value
Overall	58.5 (52.4 - 64.5)		
WHO Region			
Europe	64.1 (54.3 – 73.3)	Reference	
Africa	41.3 (31.8 – 51.1)	-0.13 (-0.28 – 0.02)	0.09
Americas	64.2 (21.5 – 98.0)	0 (-0.29 – 0.30)	0.98
Eastern Mediterranean	49.4 (30.1 – 68.7)	-0.09 (-0.27 – 0.09)	0.32
South-East Asia	50.1 (31.4 – 70.3)	-0.09 (-0.47 – 0.29)	0.65
Western Pacific	73.3 (57.8 – 86.7)	0.07 (-0.07 – 0.22)	0.33
Type			
Hepatology clinic	65.7 (58.3 – 72.7)	Reference	
General population	38.5 (25.9 – 51.8)	-0.20 (-0.33 – -0.07)	0.003
Isolated population	62.7 (26.2 – 93.8)	-0.02 (-0.48 – 0.44)	0.95
Selected population group	63.8 (49.6 – 77.0)	-0.02 (-0.16 – 0.11)	0.73
HDV RNA detection method			
qRT-PCR	50.1 (41.6 – 58.6)	Reference	
End point PCR	63.3 (53.8 – 72.4)	0.05 (-0.06 – 0.17)	0.33
Not specified	76.6 (52.4 – 64.5)		
Anti-HDV prevalence			
Per 10% increase		0.03 (0.02 – 0.06)	<0.001

^aThe co-efficient refers to the change in HDV RNA prevalence per unit change in the reported characteristics/explanatory variables, where for example a co-efficient of 0.1 refers to an increase of 10% in HDV RNA detection.

Abbreviations: CI, confidence interval; qRT-PCR, quantitative real time reverse transcriptase polymerase chain reaction.

Supplementary appendix 22. List of Included Studies Reporting HDV Genotype Data by WHO Region

African Region (AFR)

- AF2. Andernach IE, Leiss LV, Tarnagda ZS, et al. Characterization of hepatitis delta virus in sub-Saharan Africa. *J Clin Microbiol* 2014; 52: 1629-36.
- AF4. Belyhun Y, Liebert UG, Maier M. Clade homogeneity and low rate of delta virus despite hyperendemicity of hepatitis B virus in Ethiopia. *Virology* 2017; 14: 176.
- AF5. Butler EK, Rodgers MA, Collier KE, et al. High prevalence of hepatitis delta virus in Cameroon. *Sci Rep* 2018; 8: 11617.
- AF9. Ducancelle A, Abgueguen P, Birguel J, et al. High endemicity and low molecular diversity of hepatitis B virus infections in pregnant women in a rural district of North Cameroon. *PLoS One* 2013; 8: e80346.
- AF11. Groc S, Abbate JL, Le Gal F, et al. High prevalence and diversity of hepatitis B and hepatitis delta virus in Gabon. *J Viral Hepat* 2019; 26: 170-82.
- AF16. Komas NP, Ghosh S, Abdou-Chekaraou M, et al. Hepatitis B and hepatitis D virus infections in the Central African Republic, twenty-five years after a fulminant hepatitis outbreak, indicate continuing spread in asymptomatic young adults. *PLoS Negl Trop Dis* 2018; 12: e0006377.
- AF18. Makuwa M, Caron M, Souquiere S, Malonga-Mouelet G, Mahe A, Kazanji M. Prevalence and genetic diversity of hepatitis B and delta viruses in pregnant women in Gabon: molecular evidence that hepatitis delta virus clade 8 originates from and is endemic in central Africa. *J Clin Microbiol* 2008; 46: 754-6.
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Supplementary appendix 23. HDV Genotype Geographic Distribution Data

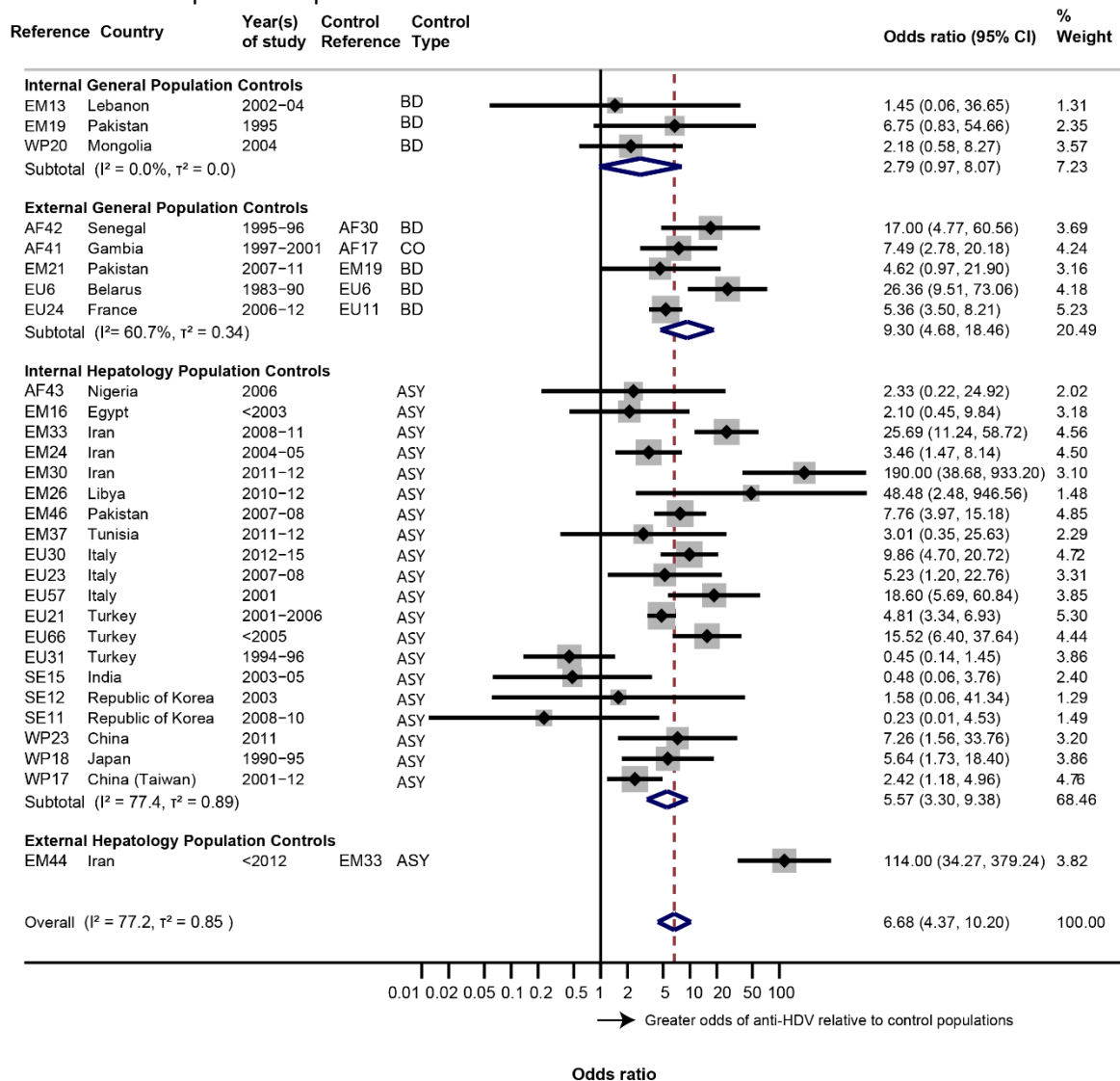
Ref	Year	Country	Region	N	Genotype (n)
EU101	1996	Sweden	Malmö	7	G1 (7)
EH9	2013	United Kingdom	National	2	G1 (2)
EU33	2015	United Kingdom	London	25	G1 (19) G5 (6)
EU93	2010	Germany	Berlin, Essen	42	G1 (42)
EU95	2019	Germany	Munich, Hannover	72	G1 (72)
EU73	2017	Switzerland	National	70	G1 (66) G5 (4)
EU11	2014	France	National	14	G1 (12) G6 (1) G7 (1)
EU97	2017	France	National	2152	G1 (1734) G2 (7) G3 (7) G5 (291) G6 (29) G7 (60) G8 (24)
EU90	1998	Spain	Catalonia	37	G1 (37)
EU95	2019	Spain	Madrid	12	G1 (12)
EU98	1997	Italy	Foggia	46	G1 (46)
EU95	2019	Italy	Turin	18	G1 (18)
EH81	2013	Czechia	Remedis	7	G1 (7)
EU22	2006	Poland	Gdansk	4	G1 (4)
EU91	2000	Russian Federation	Samara	4	G1 (4)
EU94	2001	Russian Federation	Yakutia	29	G1 (14) G2 (15)
EU92	2018	Republic of Moldova	Kishinev	2	G1 (2)
EU54	2013	Romania	Bucharest	38	G1 (38)
EU99	1997	Greece	Achangelos	12	G1 (12)
EU99	1997	Greece	Mainland	15	G1 (15)
EU87	2004	Turkey	Ankara	13	G1 (13)
EU86	2007	Turkey	Izmir	26	G1 (26)
EU89	2011	Turkey	Ankara	9	G1 (9)
EU96	2012	Turkey	Central Eastern Turkey	34	G1 (34)
EU88	2014	Turkey	Elazig	40	G1 (40)
EM13	2007	Lebanon	National	1	G1 (1)
EU100	2018	Israel	National	55	G1 (55)
EM64	2004	Iran	Tehran	1	G1 (1)
EM68	2008	Iran	Tehran	22	G1 (22)
EM66	2009	Iran	Tehran	26	G1 (26)
EM55	2013	Iran	Tehran	3	G1 (3)
EM70	2015	Iran	Tehran	38	G1 (38)
EM67	2019	Iran	Tehran	14	G1 (14)
EM32	2017	Saudi Arabia	Jeddah	1	G1 (1)
EU42	2008	Tajikistan	Dushanbe	10	G1 (10)
EM69	2012	Pakistan	Karachi	22	G1 (22)
EM65	2014	Pakistan	Lahore	21	G1 (21)
EM63	2018	Pakistan	Lahore	17	G1 (17)
WP39	2004	Mongolia	Ulaanbaatur	20	G1 (20)
WP41	2006	Mongolia	Ulaanbaatur	32	G1 (32)
WP15	2005	Mongolia	Ulaanbaatur	117	G1 (117)
WP40	2007	Mongolia	Ulaanbaatur	13	G1 (13)
WP32	2007	Mongolia	Ulaanbaatur	24	G1 (24)

WP33	2015	China	Guandong	1	G1 (1)
WP34	2015	China	Beijing	3	G1 (3)
WP4	2017	China	Hunan Province	48	G2 (48)
WP43	1995	China (Taiwan)	Taipei	14	G1 (8) G2(6)
WP13	2003	China (Taiwan)	Tzukuan Township	4	G2 (4)
WP37	2006	China (Taiwan)	Tapei	61	G1 (15) G2 (41) G4 (5)
WP26	2014	China (Taiwan)	Tapei	28	G2 (9) G4 (19)
WP2	2011	China (Taiwan)	Yunlin County	100	G1 (5) G2 (42) G4 (53)
SE4	2002	Thailand	Prathumtani and Songkla Province	8	G1 (8)
WP14	2007	Vietnam	Thai Binh province	25	G1 (5) G2 (20)
WP38	2014	Vietnam	Hanoi	21	G1 (19) G2 (2)
WP25	2015	Vietnam	Ha Noi	17	G1 (12) G2 (5)
WP31	2018	Vietnam	Hanoi	57	G1 (52) G2 (5)
SE12	2011	Republic of Korea	Kildong, Kangdong-gu, Seoul	2	G1 (2)
WP36	1999	Japan	Mikayo Island	6	G4 (6)
WP42	2003	Japan	Mikayo Island	33	G1 (2) G2 (1) G4 (30)
WP35	2003	Japan	Okinawa	1	G2 (1)
EM16	2003	Egypt	Mansoura	7	G1 (9)
EM70	2009	Tunisia	Tunis	1	G1 (1)
EM51	2015	Tunisia	Bizerte, Tunis, Nabeul, Sousse, Monastir	11	G1 (11)
AF39	2018	Algeria	Algiers	1	G1 (1)
AF20	2012	Mauritania	Nouakchott	28	G1 (25) G5 (3)
AF21	2012	Mauritania	Nouakchott	31	G1(28) G5 (3)
AF40	2013	Mauritania	Nouakchott	47	G1 (40) G5 (7)
AF41	2019	Gambia	Banjul, Fajara, Bansang	17	G1 (5) G5 (12)
AF27	2017	Ghana	Kumasi	2	G1 (1) G5 (1)
AF2	2014	Nigeria	Ibadan	1	G1 (1)
AF2	2014	Nigeria	Lagos	2	G1 (1) G6 (1)
AF46	2016	Nigeria	Osogbo, Ogbomoso, Ile-Ife, Ibadan	14	G1 (14)
AF38	2011	Cameroon	Yaoundé	25	G1 (22) G5 (1) G6 (1) G7 (1)
AF9	2013	Cameroon	Tokombéré	9	G1 (9)
AF51	2018	Cameroon	Yaoundé, Douala	211	G1 (138) G7 (61) G6 (11) G8 (1)
AF24	2018	Cameroon	Bankim District	3	G1 (3)
AF18	2008	Gabon	Libreville	3	G1 (1) G8 (2)
AF19	2009	Gabon	Woleu-Ntem (rural)	17	G1 (10) G7 (2) G8 (5)
AF11	2019	Gabon	Estuaire	40	G1 (21) G7 (7) G8 (12)
AF2	2014	CAR	Bangui	5	G1 (5)
AF16	2018	CAR	Bangui	18	G1 (18)
AF51	2017	DRC	Orientale Equateur Provinces	12	G1 (12)
AF34	2017	Ethiopia	Addis Ababa	12	G1 (12)
AF4	2017	Ethiopia	Gondar	6	G1 (6)
AM44	2000	Canada	Quebec	1	G1 (1)
AM46	1997	USA	Maryland	22	G1 (22)
AM41	2019	USA	Bethesda, MD	28	G1 (28)
AM42	2001	Venezuela	NorthWest	3	G3 (3)

AM45	2001	Venezuela	Sierra de Perija	6	G1 (6)
AM45	2001	Venezuela	Amazonas	4	G3 (4)
AM34	2007	Colombia	Leiticia	5	G3 (5)
AF10	2015	Colombia	Amazonas State	8	G1 (1) G3 (7)
AM36	1993	Peru	Yavari River	2	G3 (2)
AM43	2006	Brazil	Acre and Rodondo	40	G1 (22) G3 (18)
AM39	2008	Brazil	East Amazon	14	G3 (14)
AM38	2009	Brazil	Western Amazon	14	G3 (14)
AF16	2011	Brazil	Maranhão	3	G3 (1) G8 (2)
AF31	2011	Brazil	Sao Paulo	1	G1 (1)
AM3	2014	Brazil	Manaus City	38	G3 (38)
AM3	2014	Brazil	North Brazilian Amazonas	12	G3 (12)
AM3	2014	Brazil	Eirenepe City	20	G3 (20)
AM40	2014	Brazil	Porto Velho, Rio Branco, Cruzeiro do Sul	90	G3 (90)
AM35	2015	Brazil	Rondonia	52	G1 (4) G3 (48)
AF18	2016	Brazil	Maranhao Province	4	G8 (4)
AM37	2016	Brazil	Amazon	46	G3 (46)
AF4	2014	Argentina	Buenos Aires (Central Region)	1	G1 (1)

Abbreviations: DRC, Democratic Republic of the Congo; CAR, Central African Republic; USA, The United States of America.

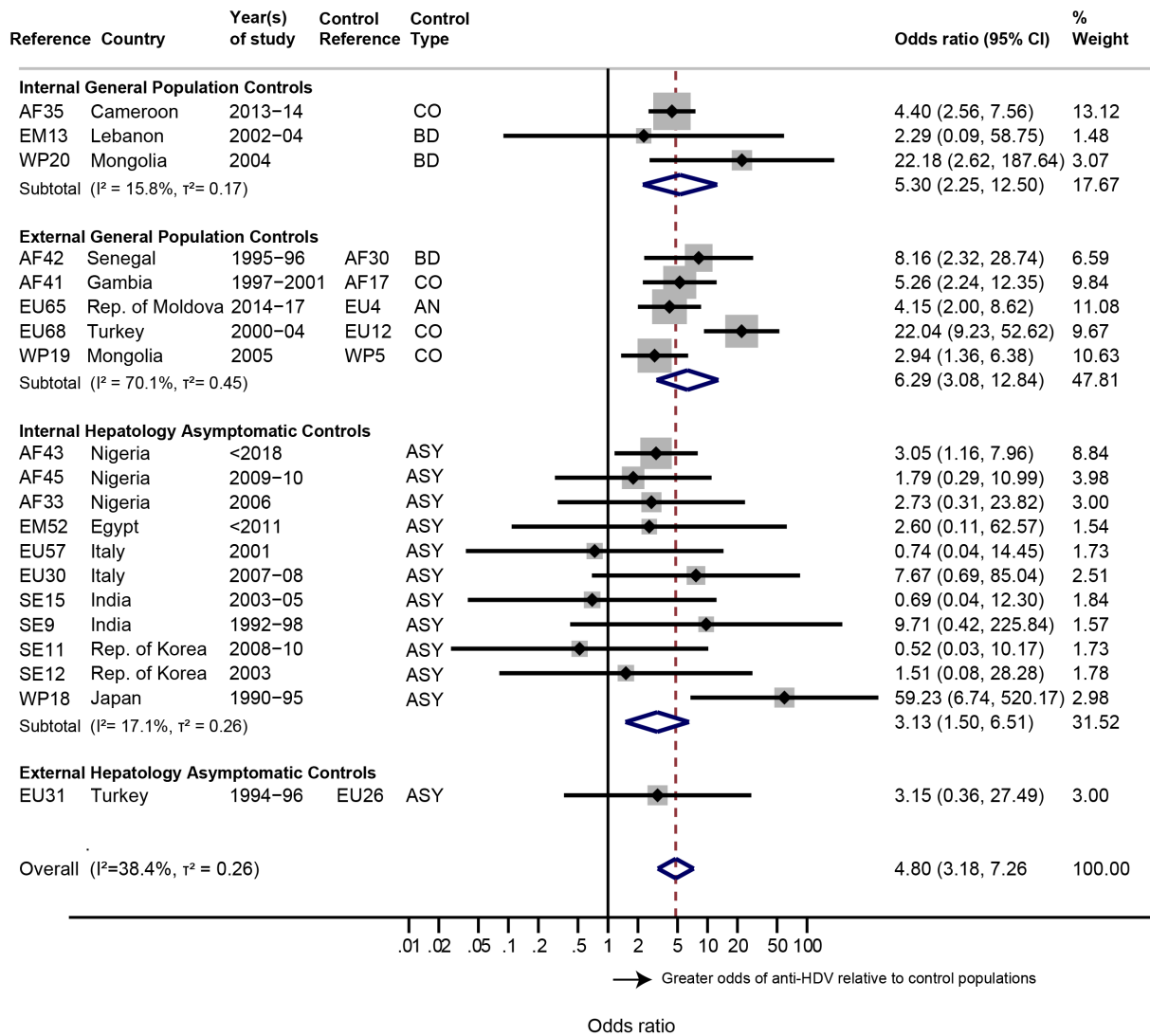
Supplementary appendix 24. Forest Plot of Anti-HDV Prevalence Among People with Cirrhosis Relative to Comparator Populations^a



^aComparator populations: Blood donors (BD), Community (CO), Asymptomatic HBsAg positive people (ASY).

Abbreviations: CI, confidence interval

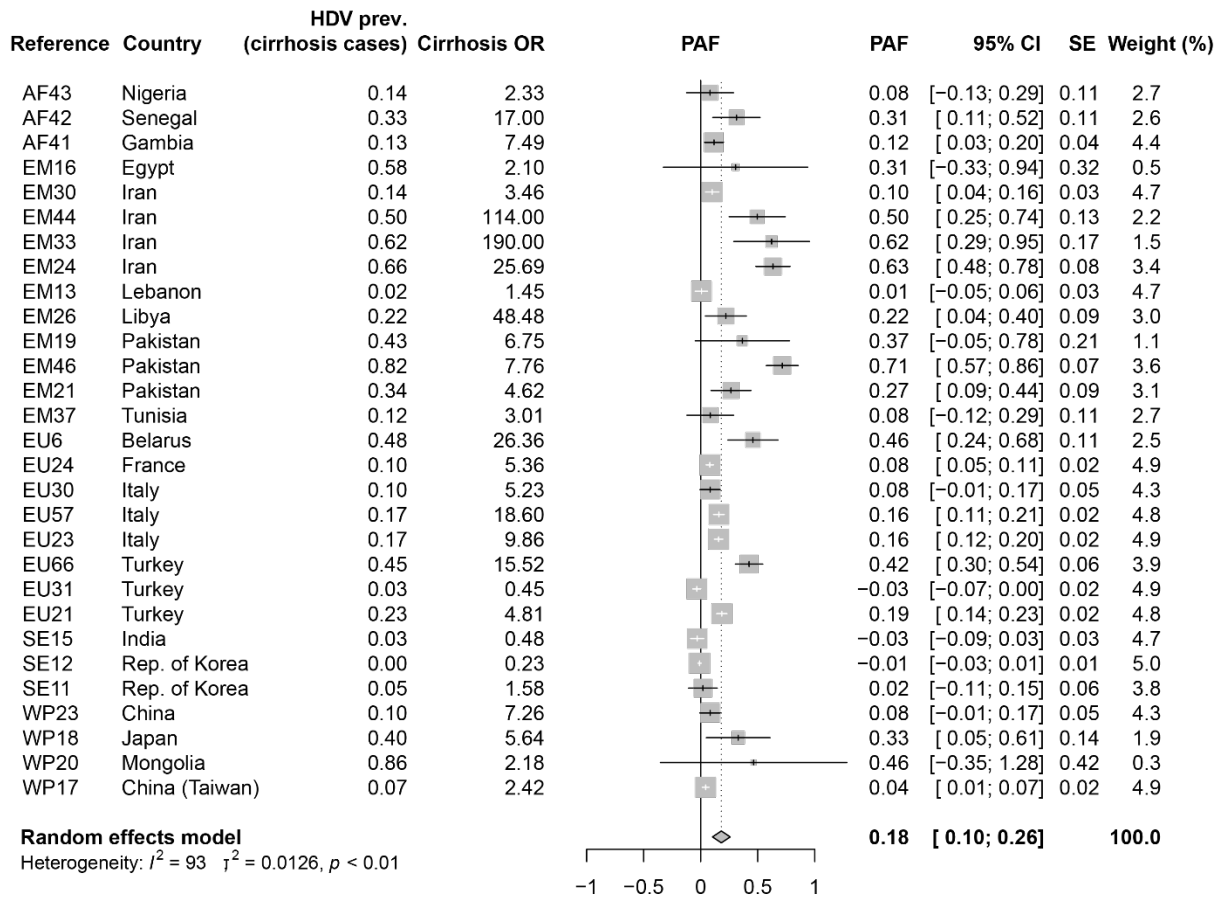
Supplementary appendix 25. Forest plot of Anti-HDV Prevalence Among People with Hepatocellular Carcinoma Relative to Comparator Populations^a



^aComparator populations: Community (CO), Blood donors (BD), Antenatal (AN), Asymptomatic HBsAg positive people (ASY).

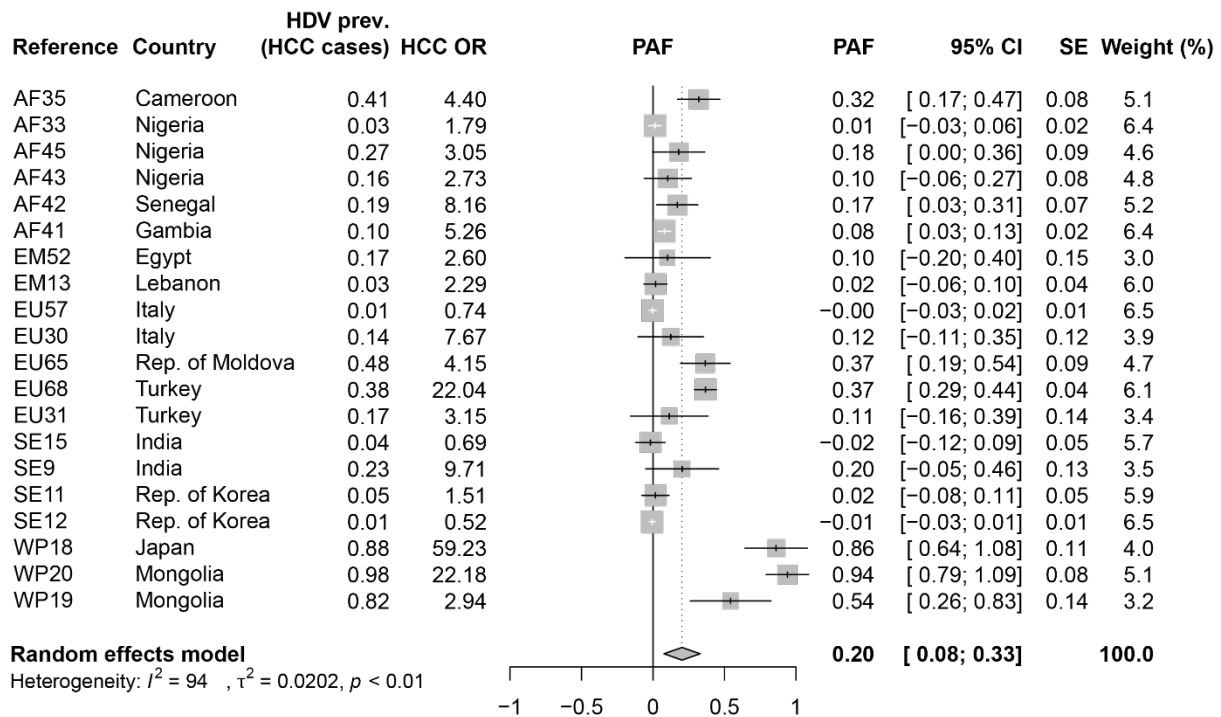
Abbreviations: CI, Confidence interval; Rep, Republic.

Supplementary appendix 26. Population Attributable Fraction of HDV to the Detection of Cirrhosis Among HBsAg-Positive People



Abbreviations: HBsAg, Hepatitis B surface antigen, Prev., prevalence; OR, odds ratio; PAF, population attributable fraction; CI, confidence interval (normal approximation using standard error calculated via parametric bootstrap); SE standard error (calculated via parametric bootstrap); Rep. Republic.

Supplementary appendix 27. Population Attributable Fraction of HDV to the Detection of Hepatocellular Carcinoma Among HBsAg-Positive People



Abbreviations: HBsAg, Hepatitis B surface antigen; Prev., prevalence; HCC, Hepatocellular carcinoma; RR, risk ratio; PAF, population attributable fraction; CI, confidence interval (normal approximation using standard error calculated via parametric bootstrap); SE standard error (calculated via parametric bootstrap); Rep., Republic.