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Associations between the prevalence of chronic hepatitis B among people who inject drugs and country-level characteristics: an ecological analysis

Short running title: PWID HBV global ecological analysis

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49 **Abstract**

50

51 **Background**

52 Globally, hepatitis B virus (HBV) is a leading cause of liver disease. People who inject drugs (PWID) are at greater
53 risk than the general population of contracting HBV. This risk could depend on societal factors in different
54 countries. We investigated the associations between country-level chronic HBV prevalence in PWID with
55 national indicators of development and prevalence of HIV and hepatitis C virus (HCV).

56

57 **Methods**

58 We used global systematic review data on chronic HBV prevalence (hepatitis B Surface antigen-positive) among
59 PWID and country-level sociodemographic characteristics from online databases. National random-effects
60 meta-analysis estimates of HBV prevalence were the outcome in linear regression models testing for
61 associations with country-level characteristics.

62

63 **Results**

64 The study included 131,710 PWID from 304 estimates in 55 countries: the pooled HBV prevalence among PWID
65 in the countries analysed was 4.5% (95%CI 3.9-5.1), the highest regional pooled prevalence was in East and
66 Southeast Asia (17.6% (13.3-22.3)), and the lowest was in Western Europe (1.7% (1.4-2.1)). In multivariable
67 models, no indicators of development were associated with HBV prevalence, but there was evidence of positive
68 associations between HBV prevalence in the general population and among PWID, and evidence of HIV and HCV
69 prevalence in PWID being associated with HBV prevalence in PWID: multivariable coefficients 0.03 (95%CI 0.01-
70 0.04); $p < 0.001$, and 0.01 (95%CI 0.00-0.03); $p = 0.01$, respectively.

71

72 **Discussion and Conclusions**

73 HBV prevalence among PWID was associated with HIV and HCV prevalence among PWID and background HBV
74 prevalence in the general population, highlighting the need for improving harm reduction in PWID and
75 implementation of HBV vaccination, especially where HBV is endemic.

76

77 **Keywords:** Hepatitis B virus, people who inject drugs, prevalence, ecological

78 Introduction

79

80 Hepatitis B virus (HBV), a blood-borne virus that causes liver disease, primarily hepatocellular carcinoma and
81 cirrhosis, results in substantial morbidity and mortality(1). Worldwide, chronic HBV is the most common cause
82 of hepatocellular carcinoma, with attributable cases ranging from 50% in regions with low HBV endemicity and
83 up to 70-80% in highly endemic areas(2). Chronic infection is indicated by the sustained presence of the
84 serological marker, hepatitis B surface antigen (HBsAg), for at least six months(3). The risk of progression from
85 acute to chronic disease is inversely proportional to age; approximately 5% of adults exposed to HBV develop
86 chronic infection. Most of the global population (~88%) live in countries where HBV prevalence is intermediate
87 (2-8%) or high ($\geq 8\%$), as classified by the World Health Organisation(4).

88

89 HBV is transmitted by parental or mucosal exposure to HBsAg-positive bodily fluids, particularly blood. Lifelong
90 protection is provided by a safe and effective vaccine delivered through infant programmes. Whilst
91 immunisation coverage has increased considerably in recent years from 30% in 2000 to 85% in 2019, rates vary
92 radically across the globe(3). Direct, percutaneous inoculation by sharing contaminated injecting equipment,
93 such as needles and syringes, is an important mode of HBV transmission(5). There are an estimated 15.6 million
94 (95% Uncertainty Interval 10.2, 23.7 million) people who inject drugs (PWID) across the globe with a higher risk
95 of contracting HBV than the general population of their respective countries(5). For example, in 2015, 30.3% of
96 newly HBV-infected individuals in the United States reported injecting drug use (IDU) as a critical risk factor(4),
97 indicating the importance for further work to address prevention and treatment of HBV infection among PWID.
98 The World Health Organization has set targets for the elimination of hepatitis by 2030 as a public health threat
99 and PWID are a key population to consider if these goals are to be achieved(6).

100

101 Recent estimates suggest that 9.1% (95% uncertainty interval: 5.1-13.2%) of PWID suffer from HBV, globally,
102 equating to roughly 1.4 million (0.7-2.4 million) people(5). 22.6% of PWID have evidence of past infection(7).
103 East and Southeast Asia accounts for more than half of all HBsAg-positive PWID worldwide(5) and 51% of the
104 global IDU-attributable HBV burden(8), yet the estimated 4.0 million (95% Uncertainty Interval 3.0, 5.0 million)
105 PWID living there only represent 25% of the global population of PWID(5). The burden of the HBV epidemic
106 among PWID in different parts of the world may be influenced by country-level factors.

107

108 Whilst previous research has highlighted that country-level income is associated with HBV prevalence in the
109 general population, there has been limited research into country-level associations with HBV prevalence among
110 PWID. This ecological analysis used country-level indicators of a nation's socioeconomic and development status
111 and epidemiological data from our previously published global systematic review to explore whether country-
112 level measures of disadvantage were associated with chronic HBV prevalence in PWID. Our hypothesis was that
113 countries with higher levels of inequality and poverty would have higher prevalence of HBV among PWID when
114 controlling for general population HBV prevalence, a relationship that would be acting through the unequal
115 distribution of vaccinations and harm-reduction measures.

116

117 **Methods**

118

119 **Data Sources**

120 The data were from a previous, multistage, global systematic review on population sizes and prevalence of HIV,
121 HBV, and hepatitis C (HCV) in PWID conducted by Degenhardt et al(5) (PROSPERO registration numbers
122 CRD42016052858 and CRD42016052853), combined with country-level sociodemographic characteristics from
123 various sources, described below. The review was completed in 2017 as an update to the last available estimates
124 for viral hepatitis prevalence from 2011(9). Full methods have been published elsewhere(5). The review included
125 journal articles indexed in Medline, Embase, and PsycINFO and online reports from government,
126 intergovernmental, and non-governmental organisations published between January 2008 to June 2017, as well
127 as data via expert requests. Some of the main inclusion criteria were: PWID samples greater than 40 individuals,
128 samples that represented the whole PWID population and not just sub-populations (e.g. prison populations),
129 and, where multiple estimates were available, literature with a higher quality grade and increased geographic
130 coverage took precedence over the recency of the estimate(5). The inclusion flow diagram of the review by
131 Degenhardt et al is included as supplementary figure 1, whilst a GATHER checklist for this current manuscript is
132 included as supplementary table 1.

133

134 **Variables**

135 As this analysis focused on chronic HBV, only studies with from the Degenhardt et al review(5) with data on
136 HBsAg were included. Studies were excluded that looked exclusively at alternative serological markers (such as
137 anti-HBc, a marker that only indicates past or current infection and not whether the infection is acute or chronic).
138 Most studies reported a percentage of the PWID sample infected with HBsAg. Where this information was
139 omitted, we calculated the percentage using the sample size and the number of HBsAg-positive individuals.

140

141 The values for the country-level characteristics were obtained from multiple sources. Country-level HIV and HCV
142 (antibodies) prevalence in PWID were as in the review by Degenhardt et al(5). The rest of the indicators: HBV
143 prevalence in the whole population, Gender Inequality Index (GII), Gini coefficient, Gross Domestic Product
144 (GDP) (Billion US\$), Gross National Income (GNI) (US\$), Human Development Index (HDI), HepB3 (3-dose)
145 immunisation coverage among 1-year-olds, labour force participation rate (women), prisoners per 100,000 of
146 the whole population, Sociodemographic Index (SDI), urbanisation, women with secondary education and both
147 male and female youth unemployment, were obtained from global online databases. We imputed any missing
148 values using similar sources. For full details on the indicators, source, values, and additional notes, see
149 supplementary tables 2 and 3.

150

151 **Meta-analysis**

152 Taiwan and the Occupied Palestinian Territory had large amounts of missing country-level characteristic data
153 and were thus omitted from further analysis. A random-effects meta-analysis was conducted using the study
154 sample size and the number of individuals infected with HBsAg to generate a single pooled chronic HBV
155 prevalence estimate per country. This was to allow for studying between-country variation as opposed to
156 between-study. A secondary meta-analysis was undertaken to stratify the sample by UNAIDS region instead of
157 country to generate regional average percentages of those infected by HBsAg.

158

159 **Country-level regression**

160 We used general linear models to test for associations between the national estimate of chronic HBV prevalence
161 among PWID (converted to a proportion and logit transformed: $[\log p/(1-p)](10)$) and the country-level
162 characteristics as the independent variables. Scatter plots of HBV prevalence and country-level characteristics
163 were generated and superimposed by linear and quadratic curves to visually inspect trends. For each
164 independent variable, both a linear and quadratic univariable model were run and then compared for their fit
165 using the Bayesian Information Criterion (BIC) and a likelihood ratio test (LRT). If the BIC values were highly

166 similar, the LRT were used to solve any discrepancies. The model with the best fit was subsequently used moving
167 forward. Data were analysed using Stata (version 16.0); source code for this analysis can be shared upon request
168 with the corresponding author.

169

170 If HBV prevalence is highly endemic in a particular nation, it is likely that the chronic HBV prevalence in PWID
171 would also be higher as a result. Thus, we included HBV prevalence in the whole population, in addition to the
172 country-level characteristics, in the multivariable models.

173 **Results**

174

175 For this analysis, data on chronic HBV prevalence in PWID was reported in 55 countries, accounting for ~63% of
176 the global population. The study included 131,710 PWID from 304 estimates in total. Greece had the highest
177 number of estimates (n=70, partially due to having separate estimates for different cities/regions), and there
178 were 20 countries with only 1 estimate. All studies from which estimates were generated were conducted
179 between 2008 and 2017 (Table 1).

180

181 **Meta-analysis**

182 There was marked variation identified in the percentage of PWID infected with HBsAg within estimates,
183 countries, and regions. The overall I^2 statistic value for HBV prevalence among PWID across all estimates was
184 96.3% ($p < 0.001$): indicating considerable to substantial heterogeneity(11) (for individual country-level I^2 values,
185 see supplementary table 4). The global mean chronic HBV prevalence in PWID was 4.5% (95% confidence interval
186 [CI] 3.9-5.1) (Table 1; Figure 1). The region with the highest reported percentage of PWID infected with HBsAg
187 was East and Southeast Asia (17.6% (95%CI 13.3-22.3)), and the lowest was Western Europe (1.7% (95%CI 1.4-
188 2.1) (for regional forest plots see supplementary material pages 13-18). The country with the highest reported
189 mean percentage of PWID infected with HBsAg was Pakistan (43.0 (95%CI 37.3-48.8) – taken from one estimate),
190 and the lowest prevalence was found in Bosnia and Herzegovina (0.2 (95%CI 0.0-1.5)) and the Maldives (0.2
191 (95%CI 0.0-1.4)).

192

193 **Ecological Analysis**

194 For most country-level characteristics, the linear model demonstrated a superior fit. The model including the
195 quadratic term was deemed a superior fit for the HBV prevalence in the whole population and urbanisation
196 variables (Figure 2a-p). The results from the linear regression are displayed in Table 2.

197

198 There was some evidence of an association between HBV prevalence in the whole population and PWID in the
199 univariable model. HBV prevalence in PWID rises as HBV in the general population increases, then begins to fall
200 at the highest levels of HBV prevalence in the general population (Figure 2a). An association between these two
201 variables was also identified in most multivariable models containing other country-level characteristics when
202 HBV prevalence in the general population was adjusted for.

203

204 Some associations between the country-level characteristics were identified in the univariable model; however,
205 many were attenuated in the multivariable model when adjusting for HBV in the whole population. A weak
206 positive association between Gender Inequality Index and chronic HBV prevalence in PWID was identified (1.53
207 (95%CI -0.40, 3.10); $p=0.06$) but this was attenuated in the multivariable model (-0.49 (95%CI -2.97, 2.00);
208 $p=0.70$). There was a negative association between Human Development Index (-2.33 (95%CI -4.41, -0.25);
209 $p=0.03$), Gross National Income ($-2.52e^{-05}$ (95%CI $-5.11e^{-05}$, $7.70e^{-07}$); $p=0.06$), Sociodemographic Index (-1.69
210 (95%CI -3.27, -0.10); $p=0.04$), the percentage of women with secondary education (-0.01 (95%CI -0.02, $3.51e^{-04}$);
211 $p=0.06$), and male youth unemployment (-0.57 (95%CI -0.11, -0.01); $p=0.03$) and chronic HBV prevalence in
212 PWID. However, these associations were also attenuated (-0.18 (95%CI -3.56, 3.20); $p=0.92$), ($-4.43e^{-06}$ (95%CI -
213 $3.60e^{-05}$, $2.72e^{-05}$) $p=0.78$), (0.02 (95%CI -2.51, 2.56); $p=0.99$), ($4.81e^{-05}$ (95%CI 0.14, 0.01); $p=0.96$), and (-0.04
214 (95%CI -0.08, $3.85e^{-03}$); $p=0.07$), respectively) when adjusting for HBV prevalence in the whole population.

215

216 There was no evidence of an association between HCV and chronic HBV prevalence in PWID in the univariable
217 model (0.01 (95%CI -0.01, 0.02); $p=0.52$). However, once adjusting for the chronic HBV prevalence in the whole
218 population, there was some evidence of a positive association (0.01 (95%CI $2.82e^{-03}$, 0.03); $p=0.01$). There was
219 evidence of association between HIV and chronic HBV prevalence in the univariable model (0.03 (95%CI 0.02,
220 0.05); $p < 0.001$), which was maintained in the multivariable model (0.03 (95%CI 0.01, 0.04); $p < 0.001$).

221

222 There was no evidence of an association between the Gini coefficient, GDP, GNI, HepB3 (3-dose) immunisation
223 coverage among 1-year-olds, women's labour force participation rate, prison population (per 100,000 people of
224 the population), percentage of women with secondary education, urbanisation or female youth unemployment
225 of a nation and chronic HBV prevalence in PWID, in either the univariable or multivariable model when adjusting
226 for HBV prevalence in the whole population.

227 Discussion

228

229 We found that HCV and HIV prevalence in PWID were positively associated with chronic HBV prevalence among
230 PWID in multivariable analyses. There was some evidence also of associations between country-level measures
231 of social disadvantage and chronic HBV prevalence in PWID; including negative associations with Human
232 Development Index score, Gross National Income, Sociodemographic Index score, the percentage of women
233 with secondary education, levels of male youth unemployment, and a positive association with Gender
234 Inequality Index score. The general rule of these univariable results was that higher development or less
235 inequality was associated with lower HBV prevalence. The exception to this was the negative association
236 between male youth unemployment and HBV, which was perhaps acting as a marker of country-income as, for
237 example, male youth unemployment was 22.37% on average in high-income Western European countries that
238 have low HBV prevalence, whilst these figures were 10.95% in Sub-Saharan Africa and 11.80% in South Asia,
239 which contained high HBV prevalence low- and middle-income countries. However, these associations of
240 country-level measures of social disadvantage with HBV prevalence among PWID were all attenuated after
241 adjusting for prevalence of HBV in the whole population, which did not corroborate our hypothesis that
242 countries with higher levels of inequality and poverty would have higher HBV prevalence among PWID when
243 controlling through general population HBV prevalence. Thus, there is a clear rationale for investing in
244 comprehensive public health policies to reduce HBV in the whole population, not just in PWID(1), reinforced by
245 the inclusion of combatting HBV as one of the United Nation's Sustainable Development Goals(12). Our study
246 highlights that HBV is an ongoing and compared to other comorbidities, often neglected (due to the quantity of
247 countries missing data on HBV but not on HCV or HIV), issue among PWID, that is particularly problematic in East
248 and Southeast Asia.

249

250 *Other Evidence*

251 The positive association between HCV and HIV prevalence in PWID and chronic HBV prevalence in PWID was
252 expected due to the high co-infection rates of bloodborne viruses among PWID(13-15), highlighting the need for
253 interventions targeted at PWID to address multiple health harms. Evidence on drug related harm has highlighted
254 the need for improving harm reduction services(4), including introducing HBV vaccination for adults in high-risk
255 groups such as PWID(3). It has been suggested that less than one in three PWID have completed the vaccination
256 series as adults(4). As a result of exposure to increased high-risk environments and collective stigmatisation
257 towards drug use, PWID are one of the most socially and medically vulnerable populations(16). These disparities
258 in healthcare equality catalyse an environment where the help-seeking behaviours of PWID are impacted, often
259 making PWID reluctant to engage with clinical services, like vaccine uptake, increasing their susceptibility to
260 HBV(17).

261

262 HBV elimination efforts have been described as 'successful but fragmented' by the World Health Organization.
263 Low-income countries traditionally lack the financial resources and health infrastructure to implement effective
264 prevention, such as universal vaccine programs or have only developed the propensity recently. These countries
265 often suffer the heaviest burden of HBV and score lower on multiple indices capturing socioeconomic and
266 development(18), as indicated by the associations identified in our univariable analyses between HBV
267 prevalence among PWID and various development indices. A comprehensive disease management strategy
268 requires both high level commitment, perhaps even legislation, and funding to increase health promotion,
269 screening, vaccination and treatment(19). A global investigation into the health governance of HBV and country-
270 level socioeconomic factors found that countries were more likely to have routine viral hepatitis surveillance
271 and a national strategy for preventing infection if they were in the higher binary categories for income level and
272 health expenditure, all traditionally associated with high-income countries(20).

273

274 Some countries are still struggling to implement universal vaccination programs in hyperendemic rural areas,
275 but, for the most part, coverage continues to increase(3, 18). However, the roll-out of HBV vaccine in the 1990s

276 means that a sizeable proportion of PWID may not have been inoculated as infants; the median age for PWID in
277 this analysis was 31.4 years. Whilst we did not find an association between immunisation coverage and HBV
278 prevalence in this analysis, repeating this analysis in five to ten years, when the effects of previous childhood
279 vaccination will have wider impact would be a further test of health inequalities. High vaccination rates will have
280 a powerful impact on eradicating HBV in the coming decades, however, there is a need to improve the
281 understanding of the associations between financial resources and the epidemiology of HBV infection in PWID,
282 to design and implement cost-effective public health interventions(1). Examples include catch-up vaccination
283 programs for key populations, harm reduction interventions like needle and syringe programs and opioid
284 substitution therapy and improving screening, testing, and antiviral therapy. The latter constitutes a substantial
285 financial burden and acts as a limiting factor in HBV infection management in LMICs(21).

286 *Strengths and Limitations*

287 To the best of our knowledge, this is the first ecological analysis exploring the association between country-level
288 chronic HBV prevalence in PWID and factors of development. One limitation of our study was that data were
289 incomplete for some country-level characteristics, so, some values were imputed. Furthermore, as the
290 exposures were measured at the national level rather than the individual level, the interpretation of these
291 results could be subject to an ecological fallacy and the association observed at a country/national level not hold
292 for an individual(22). For example, at a country-level, the number of prisoners per 100,000 of the whole
293 population was not associated with the country-level HBV prevalence among PWID, perhaps because this
294 variable may not adequately summarise differing prison-based exposure risks for HBV across countries.
295 However, at an individual-level, PWID who have been imprisoned may have higher odds of having HBV due to
296 exposures such as prison tattoos or a lack of clean needles and syringes being available in prisons(23). There also
297 was extensive missing data from many countries, due to the absence of studies on PWID and HBsAg, with only
298 55 of 179 countries included of those with reported evidence of IDU(5). Data were unavailable for Latin America,
299 the Caribbean and the Western Pacific, as well as large expanses of Sub-Saharan Africa (Figure 1), regions where
300 HBV is endemically high(24). Furthermore, estimates for 20 countries comprised of only one study, which limits
301 confidence in generalisability, and ultimately the external validity of the results. Finally, limited data on exposure
302 to IDU and risk of bloodborne viruses in PWID generates uncertainty in the estimates(8, 13). To emphasise this
303 uncertainty, we note that the estimate we used for Pakistan (43.0 (95%CI 37.3, 48.8)) was substantially higher
304 than the 22.4%(25) and 7.5%(26) found by other studies.

305 **Implications**

306
307
308
309 HBV is an ongoing issue among PWID, that is often neglected compared with other viral diseases. This is
310 evidenced by the substantial amount of missing data on HBV in countries with data on HIV and HCV, and well-
311 developed strategies in many countries to eliminate HCV and HIV, which do not always apply to HBV. The
312 association of HBV prevalence among PWID with the HBV prevalence among the general population shows
313 that there is also a need for investing in comprehensive public health policies to reduce HBV in the whole
314 population, as well as targeting vulnerable populations such as PWID. It is too early to assess whether
315 expanding childhood vaccination will also reduce chronic HBV in PWID populations. Our review also
316 emphasises the need for increased transparent epidemiological surveillance of HBV prevalence in PWID to
317 help establish effective strategies for harm reduction, particularly in countries where HBV is endemic(10, 27).

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Tables

Table 1. Characteristics of estimates included in the analysis, categorised by UNAIDS region.

Country of Report	Number of estimates	Publication Year(s)	Sample Size Range	% Infected HBsAg Range*	% Infected HBsAg (95%CI)
Australasia (1 country)	2	2009, 2012	382-402	3.0-4.0	3.6 (2.4, 5.0)
Australia	2	2009, 2012	382-402	3.0-4.0	3.6 (2.4, 5.0)
East and Southeast Asia (5 countries)	24	2009-2015	97-1049	2.4-51.6	17.6 (13.3, 22.3)
China	8	2009, 2010, 2011, 2014, 2017	97-1049	2.4-51.6	20.4 (11.4, 31.1)
Korea (Republic of)	1	2013	318	6.6	6.6 (4.1, 9.9)
Myanmar	2	2011	318-1029	12.3-43.1	15.3 (13.4, 17.3)
Thailand	1	2008	1535	30.5	30.5 (28.2, 32.9)
Vietnam	12	2011, 2012, 2015	272-1000	10.7-28.0	14.7 (12.0, 17.5)
Eastern Europe (12 countries)	59	2008-2016	42-9405	0.0-76.8	5.5 (4.0, 7.2)
Azerbaijan	14	2008, 2012	100-300	2.0-14.0	7.9 (5.9, 10.1)
Belarus	6	2015	160-400	0.4-31.9	9.6 (2.5, 20.4)
Bosnia and Herzegovina	2	2012, 2016	120-130	0.0-0.8	0.2 (0.0, 1.5)
Bulgaria	7	2016	661-1258	3.1-9.8	5.7 (3.9, 7.8)
Estonia	3	2010, 2012, 2016	326-351	4.0-76.8	23.1 (0.0, 75.9)
Hungary	6	2012, 2015, 2016	223-666	0.3-2.2	0.9 (0.3, 1.6)
Latvia	6	2016	81-1147	1.6-6.2	1.7 (1.1, 2.5)
Lithuania	1	2016	200	10.5	10.5 (6.6, 15.6)
Moldova (Republic of)	4	2013	115-365	0.0-12.4	4.6 (1.1, 10.3)
Romania	3	2011, 2016	45-522	5.0-10.6	7.3 (3.6, 12.2)
Slovakia	6	2016	42-67	1.7-5.1	2.6 (1.0, 4.8)
Ukraine	1	2016	9405	5.6	5.6 (5.1, 6.0)
Middle East and North Africa (7 countries)	19	2009-2016	40-4694	0.0-8.6	2.9 (2.0, 3.9)
Cyprus	6	2010, 2012, 2016	40-349	0.0-6.1	0.5 (0.0, 1.7)

Israel	1	2010	199	6.3	6.0 (3.2, 10.3)
Lebanon	1	2010	81	2.5	2.5 (0.3, 8.6)
Saudi Arabia	1	2015	378	7.7	7.7 (5.2, 10.8)
Syrian Arab Republic	1	2014	394	0.5	0.5 (0.1, 1.8)
Tunisia	3	2009	62-712	0.0-3.5	2.7 (1.5, 4.1)
Turkey	6	2016	964-4694	3.6-8.6	4.7 (3.5, 6.2)
North America (1 country)	1	2015	462	4.8	4.5 (2.8, 6.9)
United States of America	1	2015	462	4.8	4.5 (2.8, 6.9)
South Asia (7 countries)	66	2008-2016	58-2292	0.0-43.0	6.7 (5.2, 8.2)
Afghanistan	12	2010, 2011, 2012, 2014	96-483	3.2-10.4	6.2 (5.1, 7.4)
Bangladesh	2	2008, 2015	400-561	7.0-2.9	8.4 (6.7, 10.2)
India	27	2008, 2009, 2010, 2011, 2012, 2013, 2014, 2015, 2016	58-2292	0.7-33.2	7.7 (6.3, 9.3)
Iran (Islamic Republic of Iran)	16	2009, 2010, 2011, 2012, 2013, 2014, 2016	60-1588	0.7-32.4	6.5 (2.7, 11.8)
Maldives	2	2008	129-147	0.0-0.8	0.20 (0.00, 1.40)
Nepal	6	2011, 2015	100-401	0.0-8.0	2.20 (0.60, 4.70)
Pakistan	1	2011	300	43.0	43.00 (37.30, 48.80)
Sub-Saharan Africa (7 countries)	10	2011-2016	57-620	0.3-10.5	4.5 (2.5, 7.0)
Côte d'Ivoire	1	2016	57	10.5	10.5 (4.0, 21.5)
Kenya	1	2015	371	5.4	5.4 (3.3, 8.2)
Madagascar	3	2012	176-211	3.1-8.2	5.3 (3.6, 7.3)
Mauritius	2	2011	500-511	7.0-9.0	6.3 (4.8, 7.9)
Nigeria	1	2013	328	6.7	6.7 (4.3, 10.0)
Seychelles	1	2011	346	0.3	0.3 (0.0, 1.6)
Tanzania (United Republic of)	1	2014	620	1.1	1.0 (0.4, 2.1)
Western Europe (15 countries)	123	2008-2016	11-2077	0.0-22.5	1.7 (1.4, 2.1)
Austria	5	2016	91-159	2.7-6.6	4.1 (2.7, 5.8)
Belgium	10	2016	32-405	0.0-4.3	1.6 (1.1, 2.3)

Croatia	3	2010	121-150	0.7-1.7	1.0 (0.1, 2.3)
Denmark	1	2010	241	1.3	1.2 (0.3, 3.6)
France	1	2016	908	0.8	0.8 (0.3, 3.6)
Germany	10	2009, 2016	130-2077	0.3-1.5	0.9 (0.6, 1.2)
Greece	70	2016	11-1911	0.0-11.8	1.9 (1.5, 2.2)
Luxembourg	1	2012	310	0.8	0.6 (0.1, 2.3)
Netherlands	6	2016	13-81	0.0-12.5	0.6 (0.0, 4.5)
Norway	4	2009, 2016	116-195	0.9-3.6	2.0 (1.0, 3.3)
Portugal	6	2016	503-1054	2.2-6.8	3.6 (2.1, 5.6)
Serbia	2	2014	199-300	2.8-5.0	3.9 (2.3, 5.8)
Spain	2	2008, 2010	516-1223	1.8-22.5	3.0 (2.2, 3.8)
Sweden	1	2014	277	1.9	0.7 (0.1, 2.6)
United Kingdom	1	2016	2344	1.0	0.9 (0.6, 1.4)
All included countries (55 countries)	304	2008-2016	11-9405	0.0-76.8	4.5 (3.9, 5.1)

95%CI = 95% Confidence Interval

*These values are unweighted so may differ slightly to the % Infected HBsAg

Table 2. Univariable and multivariable country-level linear regression analyses of country-level characteristics and the prevalence of chronic Hepatitis B virus (HBsAg positive) in people who inject drugs (logit transformed dependent variable). Notation = regression coefficient (95%CI); p-value.

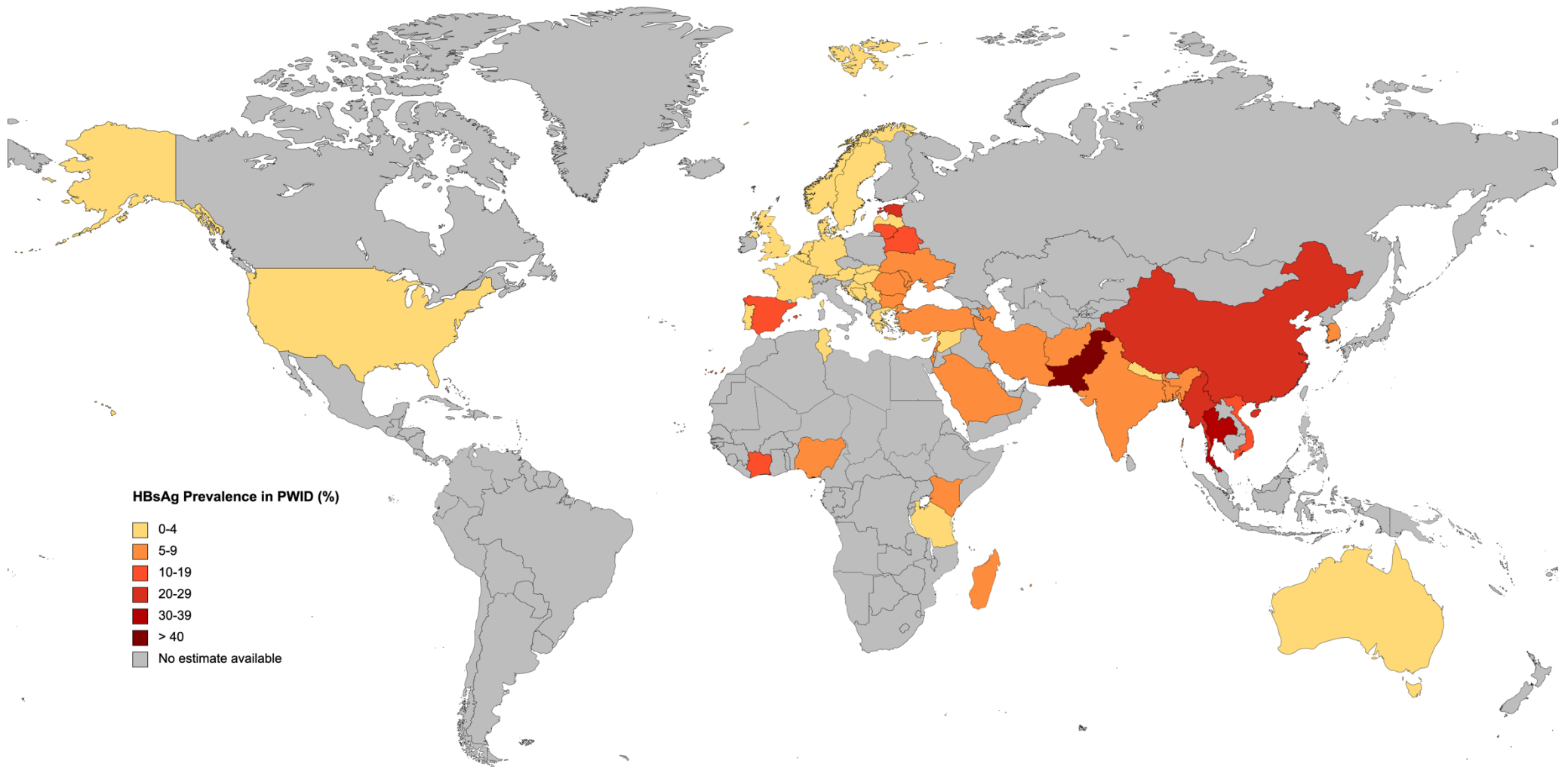
Country-level characteristic (Independent variable)	Univariable Model*		Multivariable Model**			
	Country-level independent variable		HBV Prevalence Among General Population		Country-level independent variable	
	Linear term	Quadratic term	Linear term	Quadratic term	Linear term	Quadratic term
HBV Prevalence in the whole population (%)	0.10 (0.02, 0.18); p=0.02	-0.02 (-0.05, 4.20e ⁻⁰³); p=0.10				
Gender Inequality Index	1.53 (-0.40, 3.10); p=0.06	NA	0.50 (0.02, 0.98); p=0.04	-0.03 (-0.06, 3.79e ⁻⁰³); p=0.09	-0.49 (-2.97, 2.00); p=0.70	NA
Gini Coefficient	0.02 (-0.04, 0.08); p=0.60	NA	0.51 (0.08, 0.94); p=0.02	-0.03 (-0.05, 2.84e ⁻⁰³); p=0.08	-0.03 (-0.09, 0.03); p=0.36	NA
Gross Domestic Product (US\$ Billion)	2.61e ⁻⁰⁵ (-6.10e ⁻⁰⁵ , 1.13e ⁻⁰⁴); p=0.56	NA	0.48 (0.07, 0.89); p=0.02	-0.03 (-0.06, 4.67e ⁻⁰³); p=0.10	5.65e ⁻⁰⁵ (-1.64e ⁻⁰⁵ , 1.29e ⁻⁰⁴); p=0.13	NA
Gross National Income (US\$)	-2.52e ⁻⁰⁵ (-5.11e ⁻⁰⁵ , 7.70e ⁻⁰⁷); p=0.06	NA	0.41 (-0.09, 0.91); p=0.10	-0.02 (-0.05, 0.01); p=0.17	-4.43e ⁻⁰⁶ (-3.60e ⁻⁰⁵ , 2.72e ⁻⁰⁵); p=0.78	NA
Human Development Index	-2.33 (-4.41, -0.25); p=0.03	NA	0.44 (-0.04, 0.92); p=0.07	-0.02 (-0.05, 6.00e ⁻⁰³); p=0.12	-0.18 (-3.56, 3.20); p=0.92	NA
Immunization coverage among 1-year-olds (%)	1.27e ⁻⁰³ (-0.02, 0.02); p=0.90	NA	0.45 (0.03, 0.87); p=0.04	-0.02 (-0.05, 0.01); p=0.14	0.01 (-0.02, 0.04); p=0.54	NA
Labour Force Participation Rate (women) (%)	-1.21e ⁻⁰³ (-0.03, 0.02); p=0.92	NA	0.45 (0.04, 0.87); p=0.03	-0.02 (-0.05, 4.43e ⁻⁰³); p=0.10	-4.03e ⁻⁰⁴ (-0.02, 0.02); p=0.96	NA
Prisoners per 100,000 of the whole population	2.28e ⁻⁰⁴ (-1.82e ⁻⁰³ , 2.29e ⁻⁰³); p=0.83	NA	0.48 (0.05, 0.90); p=0.03	-0.03 (-0.06, 4.17e ⁻⁰³); p=0.09	-4.11e ⁻⁰⁴ (-2.43e ⁻⁰³ , 1.61e ⁻⁰³); p=0.69	NA
Sociodemographic Index	-1.69 (-3.27, -0.10); p=0.04	NA	0.45 (-0.02, 0.93); p=0.06	-0.02 (-0.05, 0.01); p=0.11	0.02 (-2.51, 2.56); p=0.99	NA
Urbanisation (% Growth)	0.13 (-0.05, 0.31); p=0.16	-0.66 (-1.60, 0.28); p=0.17	0.48 (0.07, 0.89); p=0.02	-0.02 (-0.05, 4.13e ⁻⁰³); p=0.09	0.55 (-0.19, 1.28); p=0.15	-0.13 (-0.27, 0.01); p=0.07
Women with Secondary Education (%)	-0.01 (-0.02, 3.51e ⁻⁰⁴); p=0.06	NA	0.45 (-0.03, 0.93); p=0.07	-0.02 (-0.05, 0.01); p=0.12	4.81e ⁻⁰⁵ (-0.14, 0.01); p=0.96	NA
Female Youth Unemployment (%)	-0.03 (-0.07, 0.01); p=0.16	NA	0.42 (0.04, 0.80); p=0.03	-0.02 (-0.05, 4.22e ⁻⁰³); p=0.10	-0.02 (-0.06, 0.01); p=0.15	NA
Male Youth Unemployment (%)	-0.57 (-0.11, -0.01); p=0.03	NA	0.36 (-0.02, 0.74); p=0.06	-0.02 (-0.05, 0.01); p=0.13	-0.04 (-0.08, 3.85e ⁻⁰³); p=0.07	NA
HCV Prevalence in PWID (%)	0.01 (-0.01, 0.02); p=0.52	NA	0.49 (0.11, 0.87); p=0.01	-0.02 (-0.05, 4.42e ⁻⁰³); p=0.10	0.01 (2.82e ⁻⁰³ , 0.03); p=0.01	NA
HIV Prevalence in PWID (%)	0.03 (0.02, 0.05); p<0.001	NA	0.27 (-0.12, 0.65); p=0.17	-0.01 (-0.04, 0.02); p=0.39	0.03 (0.01, 0.04); p<0.001	NA

HBsAg = Hepatitis B Surface Antigen; 95%CI = 95% Confidence Interval; NA = Not applicable; HCV = Hepatitis C virus.

*With country-level HBV prevalence among PWID as the dependent variable and the country-level characteristics as independent variables (including a linear and sometimes a quadratic term depending on fit).

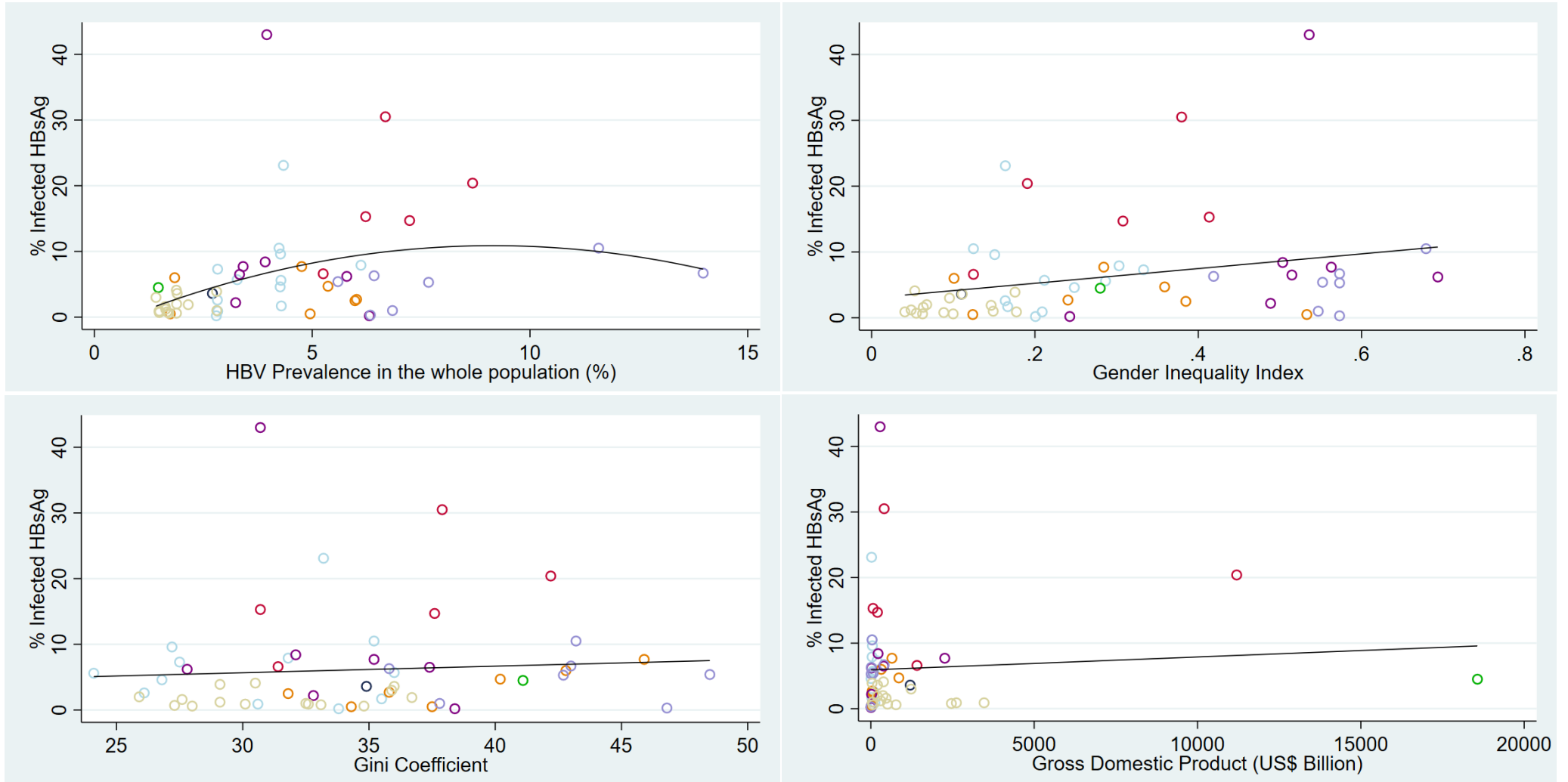
**As with the univariable model, but additionally adjusting for country-level HBV prevalence among the general population.

Figure 1. Estimated chronic Hepatitis B virus (HBsAg positive) prevalence in people who inject drugs (PWID).

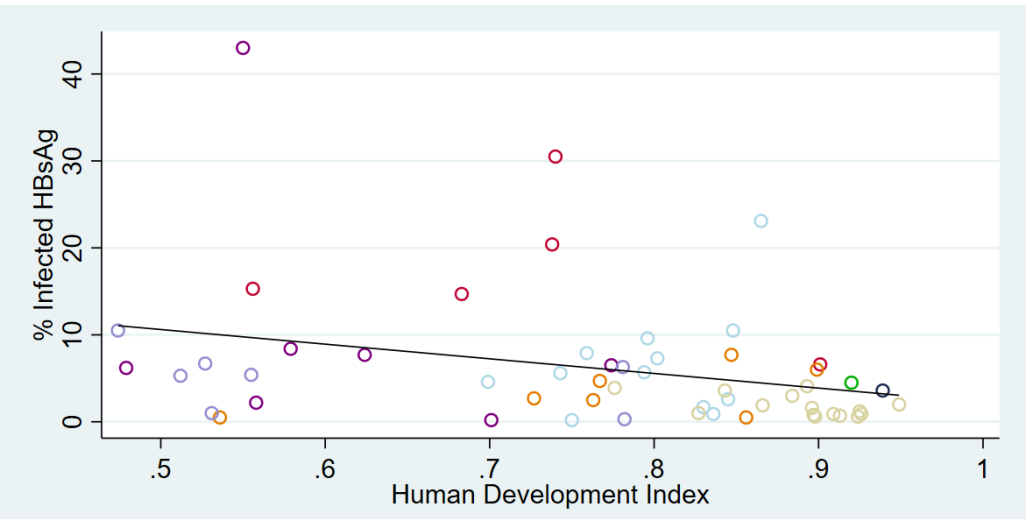
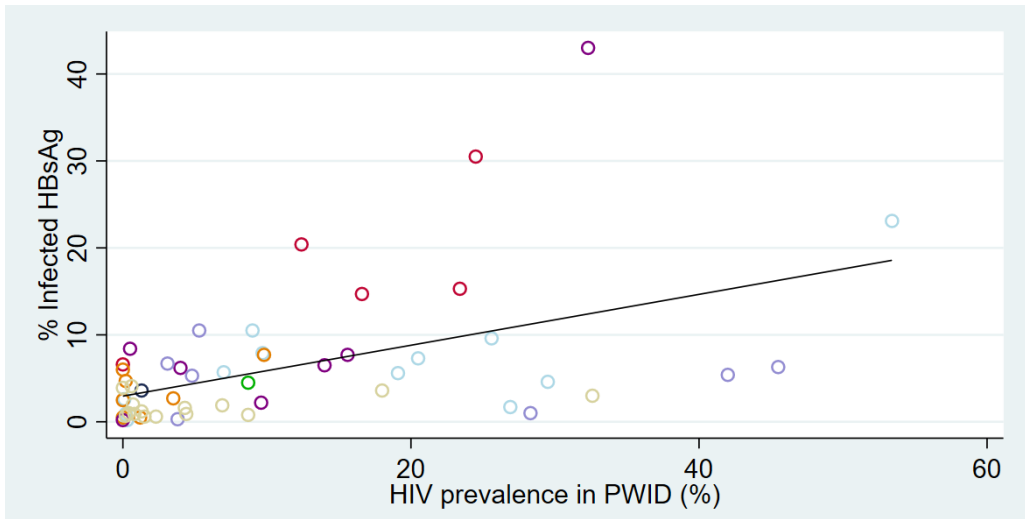
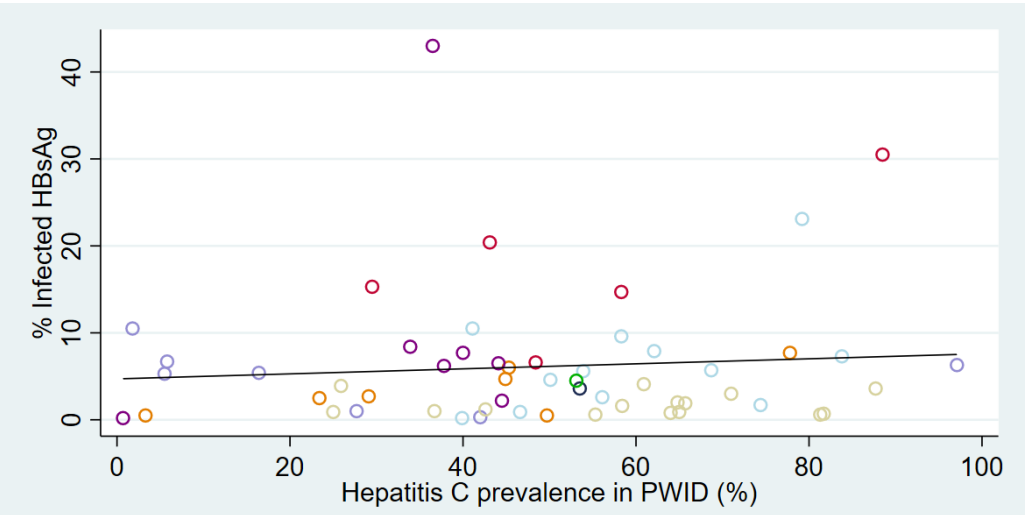
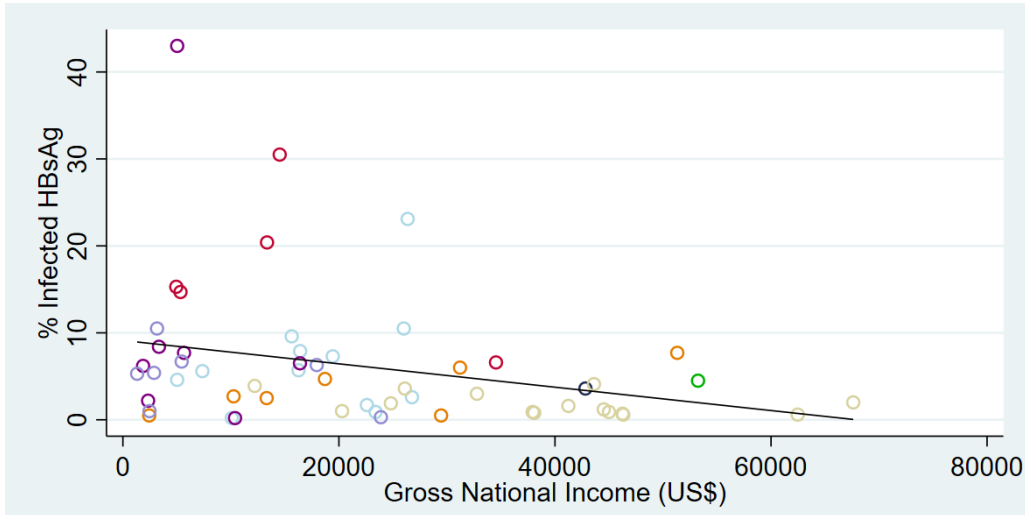


HBsAg = Hepatitis B Surface Antigen

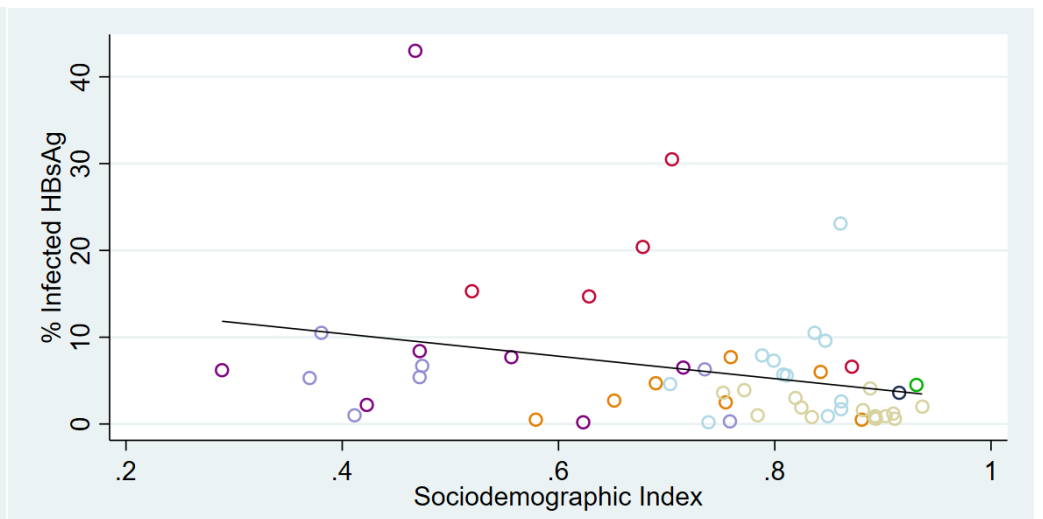
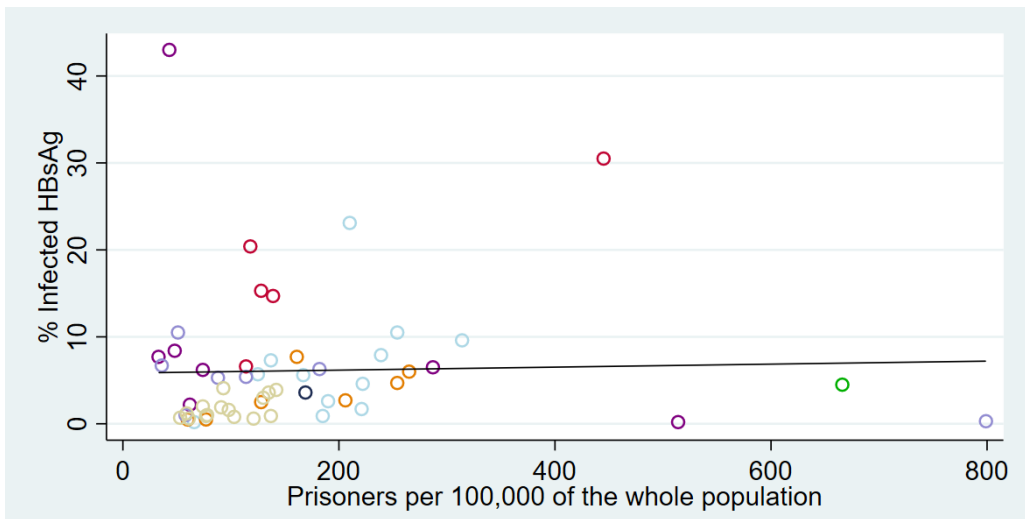
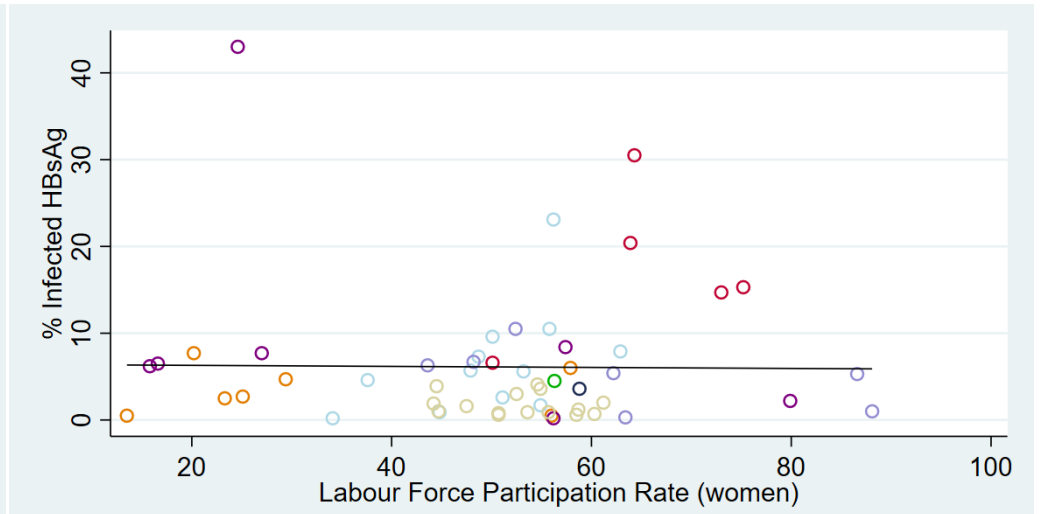
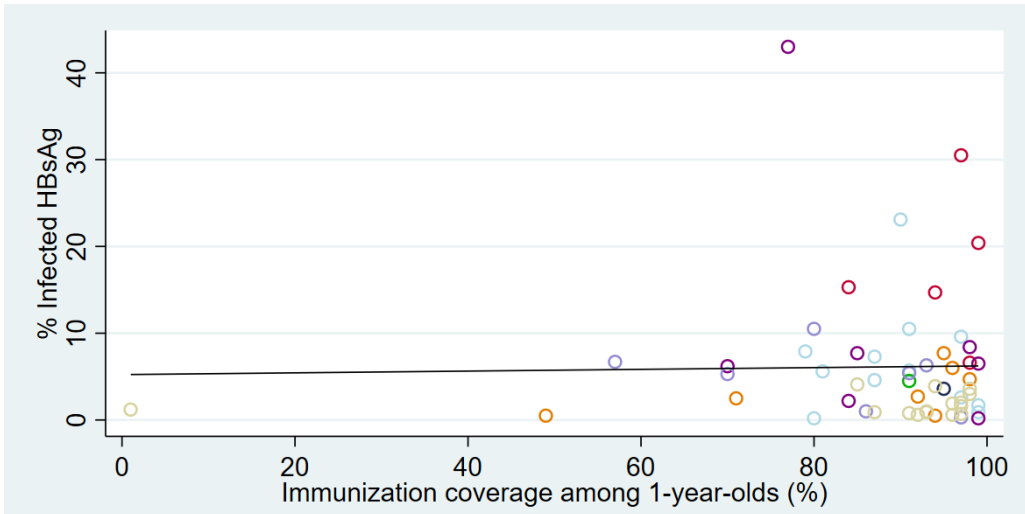
Figure 2a-p. Scatter plot for country-level characteristics and the percentage infected with HBsAg per country, grouped by UNAIDS region.



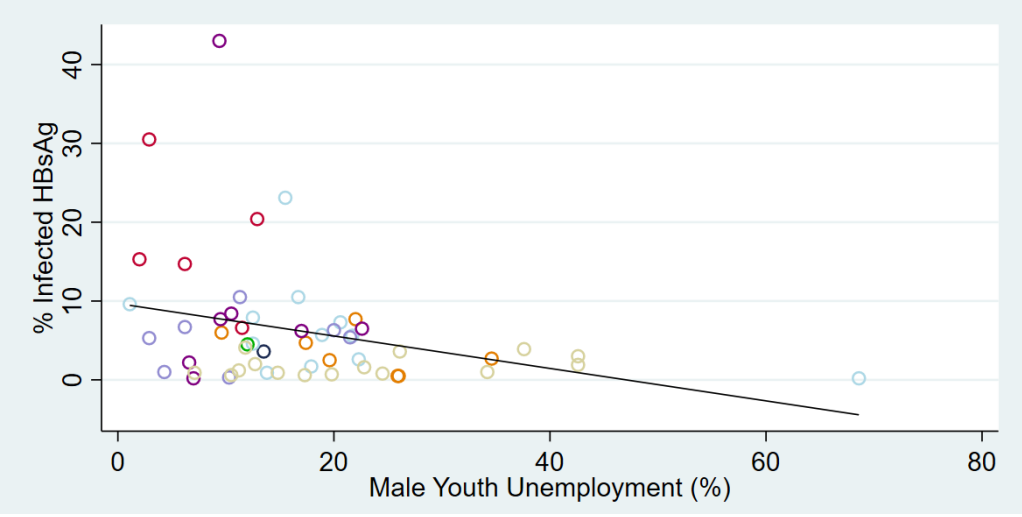
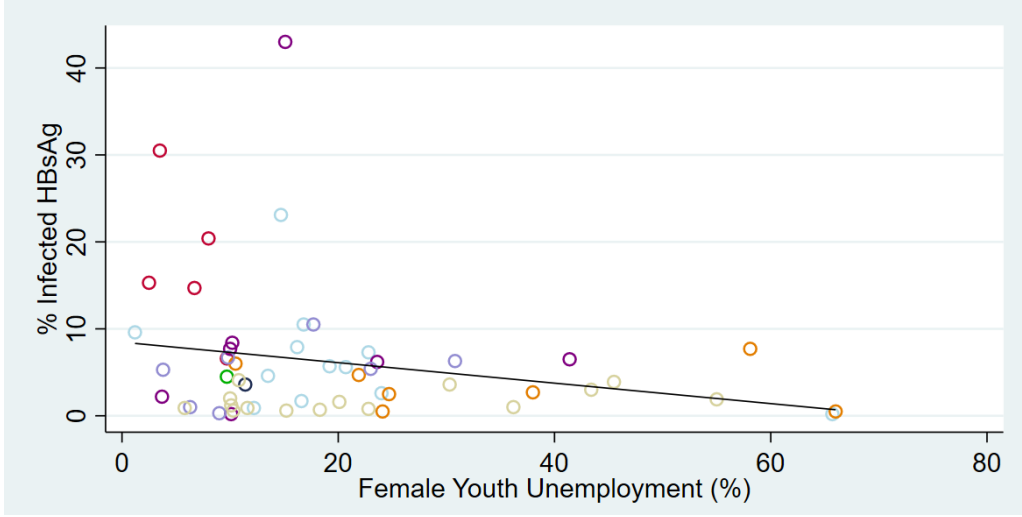
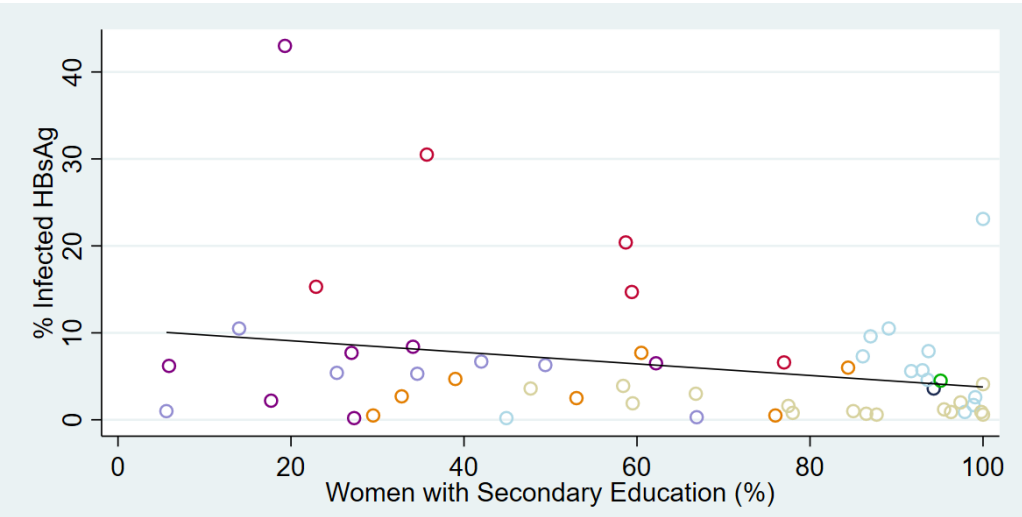
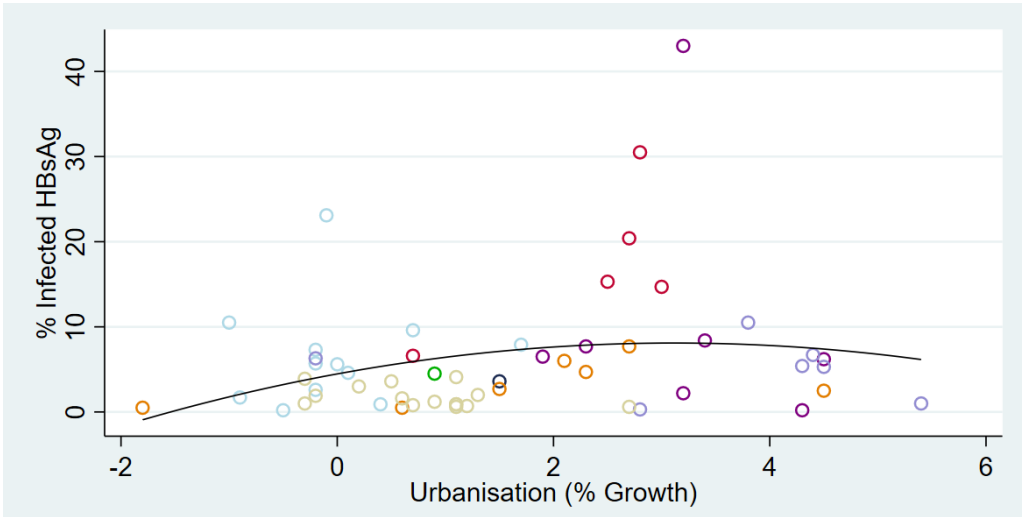
a. HBV Prevalence in the whole population (%) **b.** Gender Inequality Index **c.** Gini Coefficient **d.** Gross Domestic Product (US\$ Billion)



e. Gross National Income (US\$) f. HCV Prevalence in PWID (%) g. HIV prevalence in PWID (%) h. Human Development Index



i. Immunisation coverage among 1-year-olds (%) j. Labour Force Participation Rate (women) k. Prisoners per 100,00 of the whole population l. Sociodemographic Index



m. Urbanisation (% Growth) **n.** Women with Secondary Education (%) **o.** Female Youth Unemployment (%) **p.** Male Youth Unemployment (%)

- Australasia
- Eastern Europe
- North America
- Sub-Saharan Africa
- East and Southeast Asia
- Middle East and North Africa
- South Asia
- Western Europe

The scatterplots have been superimposed by either a linear or quadratic line of best fit depending on whether the linear or quadratic term had a superior fit in the model. These graphs exhibit a regular scale and have not been logit transformed.

