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Sexual behaviors and risk of extended-spectrum β -lactamase-producing Enterobacterales carriage: A cross-sectional analysis

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HAL Authorization

1 **Sexual behaviors and risk of extended-spectrum β -lactamase-producing**

2 **Enterobacterales carriage: a cross-sectional analysis**

3

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33 **ABSTRACT**

34

35 **Objectives:** Sexual transmission of extended-spectrum β -lactamase-producing
36 Enterobacterales (ESBL-E) is suspected. We aimed to identify clusters of sexual behavior
37 associated with ESBL-E carriage among individuals at risk of sexually transmitted infections
38 (STI).

39 **Methods:** In this cross-sectional study, patients attending an STI-screening center and HIV
40 outpatient clinic in Paris, France between 2018-2019 were asked questions on the following
41 sexual activities in the last 6 months: receptive/insertive anal intercourse, passive/active
42 rimming, receptive/insertive fellatio, receptive/insertive fisting, receptive/insertive fingering,
43 active/passive cunnilingus. ESBL-E carriage was determined from rectal swabs. Bayesian latent
44 class analysis was used to identify clusters of sexual activity, which were then associated with
45 ESBL-E carriage using logistic regression.

46 **Results:** Among 1211 men who have sex with men (MSM), those belonging to two latent
47 classes with higher prevalence of insertive fingering and active rimming (ESBL-E
48 prevalence=15.3%, $N=124$ and 16.0%, $N=100$) and one class with higher proportions of all
49 behaviors (24.3%, $N=70$) had a higher risk of ESBL-E carriage compared to those in a class
50 with few sexual behaviors (7.3%, $N=259$) after adjustment. Among 439 other men and 479
51 women, no clear associations between sexual clusters and ESBL-E carriage were observed.

52 **Conclusions:** Sexual behaviors are associated with varying degrees of ESBL-E carriage,
53 particularly among MSM.

54

55 **KEYWORDS:** antimicrobial resistance; Enterobacterales; sexually transmitted diseases;
56 transmission; epidemiology.

57 **INTRODUCTION**

58

59 Enterobacterales is a large order of Gram-negative bacillus, found as commensal organisms in
60 the digestive tract; and can lead to urinary, gastroenterological, and pulmonary infections. Of
61 the Enterobacterales, the production of extended-spectrum β -lactamase (ESBL) is the principal
62 mechanism of resistance to 3rd generation cephalosporin antibiotics. The therapeutic options for
63 individuals infected with these microorganisms are limited and almost always require the use of
64 carbapenems. *Escherichia coli* is the most common ESBL-producing Enterbacterales (ESBL-E)
65 in community [1]. The clinical progression of individuals with ESBL-E infection is more severe
66 compared to those without, and includes longer hospital stays and increased rates of morbidity
67 and mortality due to infection [2,3]. Carriage of ESBL-E is known to be a risk factor for
68 subsequent infection [4,5].

69

70 Of concern, the prevalence of ESBL-E has been increasing in community settings, with a recent
71 meta-analysis estimating an overall 10-fold increase in individuals colonized with ESBL-E
72 between 2001 and 2020 [6]. In France, one study in a general healthcare center has observed
73 an almost 10-fold increase in ESBL-E carriage, from 0.6% in 2006 to 6% in 2011 [7]. There are
74 a wide variety of risk factors for ESBL-E carriage described in the literature: recent antibiotic
75 therapy [8,9], hospitalization [10], travel to a region of high EBSL-E prevalence [11], and living
76 in communal environments [12]. Nevertheless, information on other potential risk factors for
77 developing ESBL-E carriage have been lacking and in 2010, the French High Counsel for
78 Public Health (*Haut Conseil pour la Santé Publique*) recommended conducting studies to
79 further understand other factors contributing to the dissemination of ESBL-E [13].

80

81 One putative mode of transmission of ESBL-E could be through sexual contact. Sexual
82 transmission of other pathogens has been observed via the fecal-oral route, such as *Shigella*

83 spp. [14,15] (and more recently *Shigella sonnei* carrying the blaCTX-M-27-producing plasmid
84 [16]), hepatitis A virus [17], and *Campylobacter coli* [18,19]. Indeed, a recent study conducted at
85 a sexual health center in Paris, France observed a high percentage of ESBL-E carriage, at
86 roughly 15%, among men who have sex with men (MSM), particularly those on pre-exposure
87 prophylaxis for human immunodeficiency virus (HIV) or living with HIV, when compared to other
88 men or women [20]. This difference was mostly explained by increased numbers of recent
89 sexual partners, which was associated with a higher probability of ESBL-E carriage.

90
91 These findings, coupled with others [21], illustrate that transmission of ESBL-E could be
92 possible though close contact during sex. However, it is unclear which sexual behaviors could
93 increase the risk of ESBL-E carriage. The aim of this study was then to examine the differences
94 in sexual behaviors between individuals with and without ESBL-E carriage at a larger sexual
95 healthcare clinic in Paris, France. Given the strong collinearity of sexual behaviors, we intended
96 to study the clusters of sexual behaviors that could be driving ESBL-E carriage.

97

98

99 **METHODS**

100

101 **Study design**

102

103 We analyzed cross-sectional data from the *Bactéries MultiRésistantes-Infections sexuellement*
104 *transmissibles* (BMR-IST) study [20]. In brief, this study is a two-part, mono-centric study
105 conducted at Sorbonne Université Saint-Antoine Hospital (Assistance Publique-Hôpitaux de
106 Paris, Paris, France) between 2018 and 2019. The first part was a cross-sectional assessment
107 of the prevalence and risk-factors for carriage of ESBL-E, Vancomycin Resistant *Enterococci*
108 (VRE) and Carbapenemase-Producing Enterobacterales (CPE) in various groups at risk of

109 sexually transmitted infections (STI). This analysis includes only data from the first part of this
110 cohort.

111
112 We included individuals who were ≥ 18 years, HIV-negative (seeking care at the Sexual Health
113 Clinic) or HIV-positive (seeking care at the Department of Infectious and Tropical Diseases),
114 and signed informed consent. We did not include HIV-positive men who exclusively have sex
115 with women (MSW) or HIV-positive women who have sex with men (WSM). We also did not
116 include individuals who were unable to proficiently read or speak French.

117 118 **Study procedures**

119
120 We asked individuals during regular outpatient consultations whether they would be interested
121 in participating in the study. After obtaining written informed consent, we asked participants to
122 fill out a questionnaire on demographic characteristics, ESBL-E risk factors and sexual
123 behaviour with the assistance of a study investigator. The choice of STI tests was at the
124 physician's discretion. We gave participants a self-swab (Copan ESwab™) and asked them to
125 collect a sample in the lavatory of the clinic by inserting the swab into the rectal canal. Only
126 swabs with visible fecal material were included in bacteriological analysis. After specimen
127 collection, we sent samples to the Microbiology Department of Saint-Antoine Hospital, where
128 they were tested for ESBL-E and stored at -80° C. All procedures relating to bacterial analysis
129 have been described elsewhere in detail [20,22]. Detection of ESBL-E is described in the
130 Supplementary materials.

131 132 **Study variables**

133
134 Demographic variables included age, geographic origin and whether participants were born in

135 France. Communal living was defined as living in a hotel, camp, or shelter, or being homeless.

136

137 Hospitalization within the past 12 months was recorded. We asked participants to list all

138 antibiotics used in the last 6 months, including initiation and discontinuation dates, and all

139 countries traveled to outside of France in the last 12 months, as well as entry and exit dates (or

140 duration of travel) for each country. We regrouped countries according to region and ESBL-E

141 country prevalence: low (<10%), medium (10-25%), and high (>25%) [11].

142

143 Gender was determined by the following categories: male, female, transgender male,

144 transgender female, other. Variables on sexual activity in the past 6 months included any stable

145 or known casual partners and any unknown casual partners, as well as questions on the gender

146 and number of partners. We categorized the number of partners as 0-7, 8-30, or >30. We asked

147 additional questions on whether a participant engaged in one of the following sexual behaviors

148 in the past 6 months: insertive and receptive vaginal sex, insertive and receptive anal sex,

149 insertive and receptive fellatio, active and passive cunnilingus, active and passive rimming,

150 receptive and insertive fingering, and active or passive fisting. For each behavior, we asked if a

151 condom was always, sometimes or never used. We also asked whether the participant was

152 taking preexposure prophylaxis (PrEP) for HIV.

153

154 **Statistical analysis**

155

156 From questions on sexual activity, we characterized three sexual groups: MSM, other men (*i.e.*,

157 men who reported having sex with exclusively women), and women. The limited numbers of

158 transgender persons and those with other genders precluded further analysis specific to these

159 individuals. Participant characteristics were summarized as count (%) for categorical and

160 median [interquartile range (IQR)] for continuous variables according to sexual groups. These

161 characteristics were compared between sexual groups using Pearson's χ^2 or Fisher's Exact test
162 for categorical variables and Kruskal-Wallis test for continuous variables.

163

164 To determine which individual sexual behaviors were associated ESBL-E carriage, we modeled
165 the probability of ESBL-E carriage using logistic regression. We included covariates in
166 univariable analysis, from which the odds ratio (OR) and 95% confidence interval (CI),
167 comparing the odds of ESBL-E carriage across levels of covariates, were calculated. We
168 constructed a multivariable model to obtain adjusted ORs, which were adjusted *a priori* on other
169 risk factors related to ESBL-E carriage (i.e., antibiotic use in the previous 6 months, highest
170 ESBL-E prevalence of region travelled in the previous 6 months, and number of sexual partners
171 in the previous 6 months). These models were carried out separately according to sexual group.

172

173 Given the highly correlated nature of sexual behaviors, we conducted a Bayesian latent class
174 analysis (LCA) to construct classes (or clusters) of behaviors. Briefly, a set of binomially-
175 distributed variables can be regrouped in to classes of participants with similar multivariate
176 distributions. This can be carried out using a finite mixture model framework with expectation-
177 maximization and uniform priors. We ran these models for each sexual group, separately, and
178 the variables used to construct latent classes differed between these groups. The model for
179 MSM included: receptive or insertive anal intercourse, passive or active rimming, receptive or
180 insertive fellatio, receptive or insertive fisting, and receptive or insertive fingering. The model for
181 other men included: insertive anal sex, passive or active rimming, active cunnilingus, receptive
182 fellatio, and receptive or insertive fingering. The model for women included: receptive anal sex,
183 receptive or insertive rimming, receptive cunnilingus, insertive fellatio, and receptive or insertive
184 fingering.

185

186 For each sexual group, we ran model with sequentially increasing numbers of classes, k . We

187 compared the fits between models using the Bayesian Information Criterion (BIC) and the
188 model with the BIC closest to 0 was selected as the final model. From the likelihood of this
189 model, we can calculate the posterior probability of belonging to each class k for individual i , π_{ik} .
190 Each individual can then be assigned to a class of sexual behavior for which their posterior
191 probability is the highest. The Bayesian LCA employed an expectation-maximization algorithm,
192 which was performed with the “BayesLCA” package in R [23]. We performed additional
193 diagnostic plots to assess the classification certainty of class membership across combinations
194 of behaviors.

195

196 To order classes in a more clinically meaningful way, we computed the proportion of total
197 sexual behaviors reported for each individual and calculated the mean of this proportion within
198 clusters (*i.e.*, “mean proportion of total behaviors”). The classes were then ordered from lowest
199 to highest mean proportion of total behaviors. We modeled the probability of ESBL-E carriage
200 using logistic regression and included class number as a covariate. We calculated the OR and
201 95%CI comparing the odds of carriage compared to the class with the lowest mean proportion
202 of total behaviors. OR were presented in both univariable (unadjusted) and multivariable
203 analysis (adjusted on antibiotic use in the previous 6 months and highest ESBL-E prevalence of
204 region travelled in the previous 6 months).

205

206 Data management was performed using STATA (v15.1, College Station, TX, USA) and analysis
207 carried out using R (version 4.0.3, Vienna, Austria). A p-value <0.05 was considered statistically
208 significant. Statistical code for these analyses can be found on a public repository
209 (https://github.com/boyd0094/BMRIST_sex_clusters).

210

211

212 **RESULTS**

213

214 **Description of the study population**

215

216 From 11 May 2018 to 30 June 2019, 2186 participants completed the questionnaire. 57
217 participants were excluded for the following reasons: no sample, $n=12$; did not consent for use
218 of data, $n=2$; no sexual partners in the last 6 months, $n=1$; transgender woman or man $n=14$. did
219 not answer any questions on specific sexual behavior, $n=28$. In total, 2129 were included in
220 analysis. Of them, 1211 were MSM, 439 were other men, and 479 were women.

221

222 Characteristics of the study population are described and compared between sexual groups in
223 Table 1. Compared to MSM, other men and women were younger ($p<0.001$), more likely to be
224 in communal living ($p=0.036$), and less likely to have used antibiotics in the past 6 months
225 ($p<0.001$). Other men and women also had a lower median number of sexual partners in the
226 past 6 months ($p<0.001$) compared to MSM. In the 6 months before the study visit, MSM used
227 PrEP ($p<0.001$), used drugs, and ($p<0.001$) had an STI ($p<0.001$) more often versus other men
228 or women. Among the 1211 MSM, 489 (40.4%) were HIV-positive and of the 722 MSM without
229 HIV, 246 (34.1%) were using PrEP.

230

231 The proportion of individuals engaging in the applicable sexual behaviors are described and
232 compared between sexual groups in Table 2. There were significant differences between
233 groups, where comparisons were possible, in all behaviors. Given the differences in study
234 population characteristics and sexual behaviors, we stratified analysis on sexual group.

235

236 **Association of individual sexual behaviors and ESBL-E carriage**

237

238 The prevalence of ESBL-E carriage was highest among MSM (12.3%, 95%CI=10.5-14.3),

239 followed by other men (10.0%, 95%CI=7.4-13.3) and women (6.9%, 95%CI=4.9-9.6)
240 ($p=0.0044$). Of those with ESBL-E carriage, the most frequent ST types were ST131 ($n=60$,
241 26.5%), followed by ST14 ($n=26$, 11.5%) and ST1193 ($n=22$, 9.7%), while almost all ST14 and
242 ST1193 strains were found among MSM. The majority of ESBL-E were CTX-M-15-producing
243 ($n=132$, 58.4%), followed by CTX-M-27 ($n=41$, 18.1%) and SHV-12 ($n=28$, 12.4%). All SHV-12-
244 producing ESBL-E strains were found in MSM.

245

246 The unadjusted and adjusted OR and 95%CI between each applicable sexual behavior and
247 ESBL-E carriage are given across sexual groups in Table 3. In MSM, the following sexual
248 behaviors were associated with a higher odds of ESBL-E carriage: receptive anal intercourse
249 ($p=0.034$), receptive fisting ($p=0.002$) and insertive fisting ($p=0.047$). In other men, only
250 receptive fellatio was associated with a lower odds of ESBL-E carriage ($p=0.030$). In women, no
251 sexual behaviors were associated with ESBL-E carriage. All behaviors associated with ESBL-E
252 carriage remained significant after adjustment, except for receptive anal intercourse and
253 insertive fisting for MSM.

254

255 **Association of latent classes of sexual behaviors and ESBL-E carriage**

256

257 In MSM, the LCA model with the lowest BIC (Supplementary Table 1) revealed 7 classes of
258 sexual behaviors, which are described in Figure 1A. The model produced high classification
259 certainty across combinations of behaviors (Supplementary Figure S1A). As shown in Table 4,
260 the mean proportion of total behaviors ranged from 29.9% in class 1 to 91.7% in class 7, while
261 at the same time the prevalence of ESBL-E carriage ranged from 7.3% in class 1 to 24.3% in
262 class 7. Characteristics of the study population are compared across these classes of MSM in
263 Supplementary Table S2.

264

265 When compared to class 1, class 3 (including higher proportions of individuals with insertive
266 fingering, insertive anal intercourse and active rimming), class 4 (including higher proportions of
267 individuals with insertive fingering and active rimming), and class 7 (including higher proportions
268 of individuals with all sexual behaviors) had significantly higher probability of ESBL-E carriage.
269 These differences remained after adjustment for other risk factors related to ESBL-E carriage.
270 Class 6 (including high proportions of individuals with all sexual behaviors except for insertive
271 and receptive fisting, and was the only class in which participants reported fisting) also had a
272 high prevalence of ESBL-E carriage compared to class 1 ($p=0.047$), but this was no longer
273 significant after adjustment.

274

275 In other men, the LCA model with the lowest BIC (Supplementary Table 1) revealed 3 classes of
276 sexual behaviors, which are described in Figure 1B. The model produced high classification
277 certainty across combinations of behaviors (Supplementary Figure S1B). As shown in Table 4,
278 the mean proportion of total behaviors ranged from 11.1% in class 1 to 79.0% in class 3, while
279 the prevalence of ESBL-E carriage was 16.7%, 8.1%, and 11.4% for classes 1, 2, and 3,
280 respectively. Characteristics of the study population are compared across these classes of other
281 men in Supplementary Table S3. Compared to class 1, class 2 (including high proportions of
282 individuals with insertive fingering, active cunnilingus and receptive fellatio) had a significantly
283 lower prevalence of ESBL-E carriage, even after adjustment.

284

285 In women, the LCA model with the lowest BIC (Supplementary Table 1) revealed 3 classes of
286 sexual behaviors, which are described in Figure 1C. The model produced high classification
287 certainty across combinations of behaviors (Supplementary Figure S1C). As shown in Table 4,
288 the mean proportion of total behaviors ranged from 7.6% in class 1 to 81.2% in class 3, while
289 the prevalence of ESBL-E carriage was 9.4%, 7.0%, and 4.2% for classes 1, 2, and 3,
290 respectively. Characteristics of the study population are compared across these classes of

291 women in Supplementary Table S4. Compared to class 1, there were no classes of sexual
292 behavior that were associated with ESBL-E carriage, also after adjustment.

293

294

295 **DISCUSSION**

296

297 In our study of a population of individuals at risk of acquiring STIs, we observed specific clusters
298 of sexual behaviors that were associated with a higher prevalence of ESBL-E carriage in MSM,
299 but not in other men or women. MSM belonging to the clusters that generally included insertive
300 fingering and active rimming had higher prevalence of ESBL-E carriage compared to the class
301 with few behaviors reported. In particular, MSM belonging to the group reporting almost all
302 sexual behaviors included in the questionnaire had a prevalence of ESBL-E carriage above
303 24%, which is almost four times higher than estimates in the general population in Paris, dating
304 from 2011 [7]. This study provides further insight into the role of sexual behaviors on ESBL-E
305 dissemination.

306

307 A handful of studies have highlighted the high prevalence of ESBL-E carriage among MSM,
308 particularly at risk of STIs [20,21]. For instance, one study conducted in Amsterdam, the
309 Netherlands, showed a prevalence of ESBL-E carriage at 16.3%, which corresponds to 2-times
310 higher than that of the general Dutch population [24]. Epidemics related to infections from
311 Enterobacteriales, such as *Shigella* and *E. coli*, have also been identified in MSM [25,26].
312 Coupled with the phylogenetic evidence that sexual behaviors are likely related to acquiring
313 ESBL-E [27,28], these findings suggest the potential for onward transmission in this key
314 population. However, very few of these studies have examined how sexual behaviors were
315 linked to either carriage or infection.

316

317 Emerging data have illustrated the association between increased numbers of recent sexual
318 partners and higher ESBL-E prevalence [20,21]. The LCA presented in this study allows to
319 understand specific groups of sexual behaviors related to ESBL-E carriage. In MSM, there is a
320 wide variability of concomitant sexual behaviors, as evidenced by the seven clusters of
321 behaviors, with the prevalence of ESBL-E carriage ranging from 7.3-24.3% across clusters. Of
322 note, the cluster with fisting, combined with other behaviors, had the highest prevalence. A
323 previous study in Amsterdam, which also used LCA, demonstrated that rimming and fisting,
324 combined with other sexual activities, was also associated with higher prevalence of ESBL-E
325 carriage [21]. Furthermore, phylogenetic data from these individuals would suggest putative
326 transmission clusters within sexual partners [28]. All these associations were independent of
327 risk factors commonly associated with ESBL-E carriage.

328

329 It is nevertheless difficult to pinpoint which transmission mechanism is responsible for the
330 increased risk of Enterobacterales [29], as sexual behaviors commonly co-occur. If we simply
331 model each behavior, only receptive fisting emerges as a determinant of ESBL-E carriage in
332 MSM. Simplistically, this result can only infer on the risk of ESBL-E carriage for that individual
333 behavior. However, LCA allows classes of co-occurring sexual behaviors to be constructed,
334 which can be used to model the probability of ESBL-E carriage. As such, the results from LCA
335 can infer on the risk of ESBL-E for the *ensemble* of behaviors within a class. This is not to say
336 the risk of ESBL-E carriage associated with receptive anal fisting was absent in LCA – rather
337 receptive anal fisting commonly occurred within many sexual behaviors, which together gave
338 rise to the highest probability of ESBL-E carriage. Given the composition of classes, it is
339 conceivable that ESBL-E was acquired through sexual behaviors more involved with the
340 gastrointestinal tract, *i.e.*, fecal-oral transmission.

341

342 It should be noted that MSM are not homogenous in their sexual behaviors and subsequently,

343 risk of STIs. One large study conducted in the Netherlands among approximately 13,000 MSM
344 demonstrated that the risk of STIs was highest in MSM who participate in group sex, chemsex
345 and increased numbers of sexual partners [30]. Accordingly, certain clusters could have a
346 higher prevalence of STIs, which then needed to be treated with antibiotics. The more frequent
347 exposure to antibiotic therapy would have likely induced more selective pressures for ESBL-E
348 to develop and thus higher prevalence of ESBL-E carriage. Nevertheless, we adjusted
349 comparisons on antibiotic exposure and differences remained significant.

350

351 In other men and women, LCA was unable to differentiate any cluster of sexual behavior
352 associated with higher ESBL-E prevalence, apart from one class bearing a lower risk of ESBL-E
353 prevalence in other men. These analyses do have several caveats. The variables on sexual
354 behaviors pertain to engaging in any one behavior during the past six months. As we did not
355 ask questions on the frequency of each behavior, it is not possible to determine how often these
356 behaviors needed to occur in order to attain a higher risk of ESBL-E carriage. From our
357 previous analysis of the same cohort, MSM have many more sexual partners compared to other
358 men or women [20]. The proportions of behaviors are likely to reflect many more acts in MSM,
359 thus a greater chance for exposure to ESBL-E. Second, based on phylogenetic data, there is
360 very little overlap between purported transmission networks of ESBL-E between MSM and other
361 men or women [28]. Coupled with the higher prevalence of ESBL-E in MSM, the
362 epidemiological context of ESBL-E transmission would be considered different to other men or
363 women, hence influencing the association on behavior clusters and ESBL-E prevalence. Finally,
364 it could be that other mechanisms of ESBL-E acquisition were more likely at play in other men
365 or women, namely travel to regions with high ESBL-E prevalence [11,31].

366

367 To the best of our knowledge, this study is one of the first to study the relationship between
368 sexual behaviors and ESBL-E carriage in sexual groups other than strictly MSM. The detailed

369 questionnaire allowed us to pinpoint sexual behaviors that are associated with ESBL-E carriage
370 and to add further evidence of how sexual contact could favor transmission of ESBL-E carriage.
371 The large numbers of included individuals also permitted a wide range of sexual behaviors,
372 from which we were able to differentiate different levels of risk of ESBL-E carriage. Taken these
373 advantages into consideration, these results could help guide which empirical antibiotics should
374 be given in groups with high risk of ESBL-E carriage.

375

376 Nevertheless, this study has certain limitations. First, the study pertains to only individuals who
377 are at risk of acquiring STIs and caution should be heeded when interpreting results to the
378 broader population. Second, we relied on self-reported sexual behaviors, which could be prone
379 to memory bias. The questionnaire was also administered with an interviewer and as questions
380 pertained to sensitive information about sexual behaviors, responses could have been biased
381 by non-reporting (e.g. from fear of stigmatization) or by over reporting (e.g., from social
382 desirability bias). Third, we did not include behaviors involving women (i.e., active cunnilingus,
383 insertive vaginal sex) in the LCA for MSM who also have sex with women. Different classes of
384 behaviors could have been observed because of adding these behaviors. However,
385 transmission of ESBL-E from these behaviors would be unlikely, given the analysis on other
386 men, and thus their inclusion would have only induced more statistical variation. Finally, we did
387 not ask questions related to specific activities during travel or the reasons for being hospitalized,
388 which could have been potential confounders [10,31].

389

390 In conclusion, our analysis was able to identify seven clusters of sexual behaviors among MSM
391 at risk of acquiring STIs, three of which were associated with a higher prevalence of ESBL-E
392 carriage compared to a cluster with few sexual behaviors. Clusters of sexual behaviors were
393 also observed in other men and women, but no clear association with higher ESBL-E
394 prevalence could be established. These data support the hypothesis that ESBL-E can be

395 sexually transmitted, but transmission is likely dependent on sexual behaviors associated with
396 fecal-oral routes and the epidemiological context of the key population. These data could be
397 useful in determining empirical antibiotic therapy and suggesting measures to prevent further
398 transmission of ESBL-E, especially to populations susceptible to these infections.

399

400

401 **CONFLICT OF INTEREST**

402

403 A.B. received speaker's fees from Gilead Sciences, Inc. T. C. received travel grants from Merck
404 Sharp & Dohme (MSD), Eumedica, Gilead Sciences, ViiV Healthcare, Pfizer PFE France, INC
405 Research, and Janssen-Cilag. P. L. W. received personal fees for participating on advisory
406 boards from MSD. K. L. received travel grants and personal fees for participating on advisory
407 boards and educational activities from MSD, ViiV healthcare, and Gilead, outside of this scope.
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409 potential conflicts.

410

411

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413

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416

417

418 **ETHICAL APPROVAL STATEMENT**

419

420 This study was approved by the "*Comité de protection des personnes Ouest II*" ethics

421 committee (2017-73) in accordance with the Helsinki Declaration. All participants were required
422 to provide consent for participation in the study and for the use of their data in analysis. This
423 study was conform with the European Union's General Data Protection Regulation.

424

425

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427

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431 responsible for coordinating the study.

432

433

434 **REFERENCES**

435

436 [1] Pitout JDD, Laupland KB. Extended-spectrum beta-lactamase-producing
437 Enterobacteriaceae: an emerging public-health concern. *Lancet Infect Dis* **2008**; 8:159–66.

438 [2] Schwaber MJ, Carmeli Y. Mortality and delay in effective therapy associated with
439 extended-spectrum beta-lactamase production in Enterobacteriaceae bacteraemia: a
440 systematic review and meta-analysis. *J Antimicrob Chemother* **2007**; 60:913–20.

441 [3] Schwaber MJ, Navon-Venezia S, Kaye KS, Ben-Ami R, Schwartz D, Carmeli Y. Clinical
442 and economic impact of bacteremia with extended- spectrum-beta-lactamase-producing
443 Enterobacteriaceae. *Antimicrob Agents Chemother* **2006**; 50:1257–62.

444 [4] Ruppé E, Lixandru B, Cojocaru R, et al. Relative Fecal Abundance of Extended-
445 Spectrum- β -Lactamase-Producing *Escherichia coli* Strains and Their Occurrence in Urinary
446 Tract Infections in Women. *Antimicrob Agents Chemother* **2013**; 57:4512–4517.

- 447 [5] Neulier C, Birgand G, Ruppé É, et al. Enterobacteriaceae bacteremia: Risk factors for
448 ESBLPE. *Med Mal Infect* **2014**; 44:32–38.
- 449 [6] Bezabih YM, Bezabih A, Dion M, et al. Comparison of the global prevalence and trend of
450 human intestinal carriage of ESBL-producing *Escherichia coli* between healthcare and
451 community settings: a systematic review and meta-analysis. *JAC Antimicrob Resist* **2022**;
452 4:dla048.
- 453 [7] Nicolas-chanoine MH, Gruson C, Bialek-Davenet S, et al. 10-fold increase (2006-11) in
454 the rate of healthy subjects with extended-spectrum β -lactamase-producing *Escherichia coli*
455 faecal carriage in a parisian check-up centre. *Journal of Antimicrobial Chemotherapy* **2013**;
456 68:562–568.
- 457 [8] Karanika S, Karantanos T, Arvanitis M, Grigoras C, Mylonakis E. Fecal Colonization
458 With Extended-spectrum Beta-lactamase-Producing Enterobacteriaceae and Risk Factors
459 Among Healthy Individuals: A Systematic Review and Metaanalysis. *Clin Infect Dis* **2016**;
460 63:310–8.
- 461 [9] Osthoff M, McGuinness SL, Wagen AZ, Eisen DP. Urinary tract infections due to
462 extended-spectrum beta-lactamase-producing Gram-negative bacteria: identification of risk
463 factors and outcome predictors in an Australian tertiary referral hospital. *Int J Infect Dis* **2015**;
464 34:79–83.
- 465 [10] Hagel S, Makarewicz O, Hartung A, et al. ESBL colonization and acquisition in a hospital
466 population: The molecular epidemiology and transmission of resistance genes. *PLoS One*
467 **2019**; 14:e0208505.
- 468 [11] Arcilla MS, van Hattem JM, Haverkate MR, et al. Import and spread of extended-
469 spectrum β -lactamase-producing Enterobacteriaceae by international travellers (COMBAT
470 study): a prospective, multicentre cohort study. *Lancet Infect Dis* **2017**; 17:78–85.
- 471 [12] Cochard H, Aubier B, Quentin R, van der Mee-Marquet N, Réseau des Hygiénistes du
472 Centre. Extended-spectrum β -lactamase-producing Enterobacteriaceae in French nursing

473 homes: an association between high carriage rate among residents, environmental
474 contamination, poor conformity with good hygiene practice, and putative resident-to-resident
475 transmission. *Infect Control Hosp Epidemiol* **2014**; 35:384–9.

476 [13] HCSP. Recommandations relatives aux mesures à mettre en oeuvre pour prévenir
477 l'émergence des entérobactéries BLSE et lutter contre leur dissémination. 2010.

478 [14] Aragón TJ, Vugia DJ, Shallow S, et al. Case-control study of shigellosis in San
479 Francisco: the role of sexual transmission and HIV infection. *Clin Infect Dis* **2007**; 44:327–34.

480 [15] Mook P, McCormick J, Bains M, et al. ESBL-Producing and Macrolide-Resistant *Shigella*
481 *sonnei* Infections among Men Who Have Sex with Men, England, 2015. *Emerg Infect Dis* **2016**;
482 22:1948–1952.

483 [16] Jacqueline C, Carrascoso GR, Gutiérrez-Fernández J, et al. Genetic Characterization of
484 Extensively Drug-Resistant *Shigella sonnei* Infections, Spain, 2021-2022. *Emerg Infect Dis*
485 **2023**; 29:2370–2373.

486 [17] Ndumbi P, Freidl GS, Williams CJ, et al. Hepatitis A outbreak disproportionately affecting
487 men who have sex with men (MSM) in the European Union and European Economic Area,
488 June 2016 to May 2017. *Euro Surveill* **2018**; 23.

489 [18] Gaudreau C, Rodrigues-Coutlée S, Pilon PA, Coutlée F, Bekal S. Long-Lasting Outbreak
490 of Erythromycin- and Ciprofloxacin-Resistant *Campylobacter jejuni* Subspecies *jejuni* from
491 2003 to 2013 in Men Who Have Sex with Men, Quebec, Canada. *Clinical Infectious Diseases*
492 **2015**; 61:1549–1552.

493 [19] Gaudreau C, Pilon PA, Sylvestre J-L, Boucher F, Bekal S. Multidrug-Resistant
494 *Campylobacter coli* in Men Who Have Sex with Men, Quebec, Canada, 2015. *Emerg Infect Dis*
495 **2016**; 22:1661–3.

496 [20] Surgers L, Chiarabini T, Royer G, et al. Evidence of Sexual Transmission of Extended-
497 Spectrum β -Lactamase-Producing Enterobacterales: A Cross-sectional and Prospective Study.
498 *Clin Infect Dis* **2022**; 75:1556–1564.

- 499 [21] van Bilsen WPH, van Dulm E, Matser A, et al. High carriage of ESBL-producing
500 Enterobacteriaceae associated with sexual activity among men who have sex with men. *Int J*
501 *Antimicrob Agents* **2021**; 57.
- 502 [22] Boyd A, El Dani M, Ajrouche R, et al. Gut microbiome diversity and composition in
503 individuals with and without extended-spectrum β -lactamase-producing Enterobacterales
504 carriage: a matched case-control study in infectious diseases department. *Clin Microbiol Infect*
505 **2024**;
- 506 [23] White A, Murphy TB. **BayesLCA** : An *R* Package for Bayesian Latent Class Analysis. *J*
507 *Stat Softw* **2014**; 61.
- 508 [24] Reuland EA, Al Naiemi N, Kaiser AM, et al. Prevalence and risk factors for carriage of
509 ESBL-producing Enterobacteriaceae in Amsterdam. *J Antimicrob Chemother* **2016**; 71:1076–
510 82.
- 511 [25] Simms I, Gilbert VL, Byrne L, et al. Identification of verocytotoxin-producing *Escherichia*
512 *coli* O117:H7 in men who have sex with men, England, November 2013 to August 2014. *Euro*
513 *Surveill* **2014**; 19.
- 514 [26] Tansarli GS, Long DR, Waalkes A, et al. Genomic reconstruction and directed
515 interventions in a multidrug-resistant Shigellosis outbreak in Seattle, WA, USA: a genomic
516 surveillance study. *Lancet Infect Dis* **2023**; 23:740–750.
- 517 [27] Vecilla DF, Urrutikoetxea Gutiérrez MJ, Nieto Toboso MC, et al. First report of *Shigella*
518 *sonnei* carrying a blaCTX-M-15 sexually transmitted among men who have sex with men.
519 *Infection* **2024**;
- 520 [28] de Korne-Elenbaas J, van der Putten BCL, Boek NDM, et al. Putative transmission of
521 extended-spectrum β -lactamase-producing *Escherichia coli* among men who have sex with
522 men in Amsterdam, the Netherlands. *Int J Antimicrob Agents* **2023**; 62:106810.
- 523 [29] Campos-Madueno EI, Moradi M, Eddoubaji Y, et al. Intestinal colonization with
524 multidrug-resistant Enterobacterales: screening, epidemiology, clinical impact, and strategies to

525 decolonize carriers. *Eur J Clin Microbiol Infect Dis* **2023**; 42:229–254.

526 [30] Slurink IAL, van Benthem BHB, van Rooijen MS, Achterbergh RCA, van Aar F. Latent
527 classes of sexual risk and corresponding STI and HIV positivity among MSM attending centres
528 for sexual health in the Netherlands. *Sex Transm Infect* **2020**; 96:33–39.

529 [31] Ruppé E, Armand-Lefèvre L, Estellat C, et al. High Rate of Acquisition but Short
530 Duration of Carriage of Multidrug-Resistant Enterobacteriaceae After Travel to the Tropics. *Clin*
531 *Infect Dis* **2015**; 61:593–600.

532

Table 1. Characteristics of the study population stratified on sexual group

	Sexual group			<i>p</i> ^a
	MSM (<i>n</i> =1211)	Other men (<i>n</i> =439)	Women (<i>n</i> =479)	
Age, median (IQR), y	38 (30-48)	27 (23-33)	25 (22-29)	<0.001
Geographic origin ^b				<0.001
Europe	980 (80.9%)	348 (79.3%)	392 (82.0%)	
North America, DROM-COM or Australia	113 (9.3%)	23 (5.2%)	33 (6.9%)	
Africa or Middle East	92 (7.6%)	64 (14.6%)	46 (9.6%)	
Asia	26 (2.1%)	4 (0.9%)	7 (1.5%)	
Born in France	932 (77.0 %)	352 (80.2 %)	393 (82.0%)	0.051
Communal living ^c	16 (1.3%)	11 (2.5%)	15 (3.1%)	0.036
Antibiotic use in past 6 months	402 (33.2%)	47 (10.7%)	84 (17.5%)	<0.001
Type of antibiotic used				<0.001
None	809 (66.8%)	392 (89.3%)	395 (82.5%)	
β-Lactams	131 (10.8%)	26 (5.9%)	48 (10.0%)	
Doxycycline	64 (5.3%)	10 (2.3%)	9 (1.9%)	
Other	207 (17.1%)	11 (2.5%)	27 (5.6%)	
Antibiotic use in past 3 months	273 (22.5%)	30 (6.8%)	62 (12.9%)	<0.001
Travel outside of France	887 (73.2%)	313 (71.3%)	357 (74.5%)	0.54
Hospitalization in the past 12 months	69 (5.7%)	11 (2.5%)	19 (4.0%)	0.018
Number of sexual partners in the past 6 months				<0.001
0-7	610 (50.4%)	402 (91.6 %)	449 (93.7%)	
8-30	436 (36.0%)	34 (7.7 %)	30 (6.3%)	
>30	165 (13.6%)	3 (0.7 %)	0 (0.0%)	
HIV positive	489 (40.4%)	0 (0.0%)	0 (0.0%)	<i>ntp</i>
PrEP use in the past 6 months ^d	246 (34.1%)	1 (0.2%)	1 (0.2%)	<0.001
Any condomless sex in the past 6 months	1023 (84.5%)	371 (84.5%)	396 (82.7%)	0.63
Drug use ^e in the past 6 months	535 (44.2%)	161 (36.7%)	170 (35.5%)	<0.001
Any STI in the past 6 months	315 (26.0%)	26 (5.9%)	28 (5.8%)	<0.001
Sex work in the past 6 months	29 (2.4%)	7 (1.6%)	5 (1.0%)	0.16

Data were obtained from individuals in the BMR-IST cohort, Paris, France, 2018-2019. Data represent no. (%) of participants unless otherwise identified as median (IQR).

^aStatistical comparisons were performed using Pearson χ^2 or Fisher exact test for categorical and Kruskal-Wallis test for continuous variables.

^bMissing value for 1 participant in the women group.

^cCommunal living defined as living in a hotel, camp, or shelter or experiencing homelessness.

^dAmong participants without HIV.

^eDrug use excludes alcohol and tobacco use.

Abbreviations: DROM-COM, *départements et régions d'outre-mer et collectivités d'outre-mer* (Overseas France ou French overseas territories); ESBL-E, extended-spectrum β-lactamase–

producing Enterobacterales; HIV, human immunodeficiency virus; IQR, interquartile range; *ntp*, no test performed; PrEP, pre-exposure prophylaxis for HIV; STI, sexually transmitted infection.

Table 2. Description of sexual behaviors stratified on sexual group

Sexual behaviors ^a	Sexual group			<i>p</i> ^b
	MSM (<i>n</i> =1211)	Other men (<i>n</i> =439)	Women (<i>n</i> =479)	
Receptive anal intercourse	969 (80.0%)	-	160 (33.4%)	<0.001
Insertive anal intercourse	957 (79.0%)	154 (35.1%)	-	<0.001
Passive rimming	727 (60.0%)	42 (9.6%)	86 (18.0%)	<0.001
Active rimming	697 (57.6%)	94 (21.0%)	47 (9.8%)	<0.001
Passive cunnilingus	-	-	345 (72.0%)	<i>ntp</i>
Active cunnilingus	-	322 (73.3%)	-	<i>ntp</i>
Receptive fellatio	1071 (88.4%)	362 (82.5%)	-	<0.001
Insertive fellatio	1077 (88.9%)	-	408 (85.2%)	<0.001
Receptive fisting	104 (8.6%)	-	8 (1.7%)	<0.001
Insertive fisting	171 (14.1%)	5 (1.1%)	-	<0.001
Receptive fingering	611 (50.5%)	42 (9.6%)	388 (81.0%)	<0.001
Insertive fingering	704 (58.1%)	365 (83.1%)	83 (17.3%)	<0.001

Data were obtained from individuals in the BMR-IST cohort, Paris, France, 2018-2019. Data represent no. (%) of participants.

^aInsertive and receptive vaginal intercourse was not included, as these behaviors were not considered in the Bayesian latent class analysis.

^bStatistical comparisons were performed using Pearson χ^2 or Fisher exact test for categorical and Kruskal-Wallis test for continuous variables.

Abbreviations: MSM, men who have sex with men; *ntp*, no test performed.

Table 3. Association between individual sexual behaviors and ESBL-E carriage, stratified by sexual group

Sexual behavior ^a	ESBL-E carriage, <i>n</i> (%)		Non-adjusted		Adjusted ^b	
	No	Yes	OR (95%CI)	<i>P</i>	aOR (95%CI)	<i>P</i>
MSM (N=1211)	(<i>n</i> =1062)	(<i>n</i> =149)				
Receptive anal intercourse	840 (79.1%)	129 (86.6%)	1.70 (1.06-2.87)	0.034	1.60 (0.99-2.71)	0.063
Insertive anal intercourse	842 (79.3%)	115 (77.2%)	0.88 (0.59-1.35)	0.56	0.79 (0.52-1.21)	0.26
Passive rimming	630 (59.3%)	97 (65.1%)	1.28 (0.90-1.84)	0.18	1.04 (0.72-1.52)	0.83
Active rimming	608 (57.3%)	89 (59.7%)	1.11 (0.78-1.58)	0.57	0.92 (0.64-1.33)	0.65
Receptive fellatio	934 (87.9%)	137 (91.9%)	1.56 (0.88-3.05)	0.16	1.34 (0.74-2.64)	0.36
Insertive fellatio	944 (88.9%)	133 (89.3%)	1.04 (0.61-1.87)	0.89	0.89 (0.52-1.62)	0.69
Receptive fisting	81 (7.6%)	23 (15.4%)	2.21 (1.32-3.59)	0.002	1.99 (1.17-3.26)	0.008
Insertive fisting	142 (13.4%)	29 (19.5%)	1.57 (0.99-2.41)	0.047	1.41 (0.88-2.19)	0.14
Receptive fingering	83 (55.7%)	528 (49.7%)	1.27 (0.90-1.80)	0.17	1.14 (0.80-1.62)	0.47
Insertive fingering	611 (57.5%)	93 (62.4%)	1.23 (0.86-1.75)	0.26	1.06 (0.74-1.53)	0.76
Other men (N=439)	(<i>n</i> =395)	(<i>n</i> =44)				
Insertive anal intercourse	142 (35.9%)	12 (27.3%)	0.67 (0.32-1.31)	0.26	0.69 (0.33-1.36)	0.30
Passive rimming	37 (9.4%)	5 (11.4%)	1.24 (0.41-3.09)	0.67	1.29 (0.42-3.23)	0.61
Active rimming	82 (20.8%)	10 (22.7%)	1.12 (0.51-2.29)	0.76	1.16 (0.52-2.39)	0.70
Active cunnilingus	291 (73.7%)	31 (70.5%)	0.85 (0.44-1.74)	0.65	0.86 (0.44-1.76)	0.66
Receptive fellatio	331 (83.8%)	31 (70.5%)	0.46 (0.23-0.96)	0.030	0.46 (0.23-0.96)	0.030
Insertive fisting	5 (1.3%)	0	^c		^c	
Receptive fingering	41 (10.4%)	1 (2.3%)	0.20 (0.01-0.96)	0.12	0.20 (0.01-0.94)	0.11
Insertive fingering	333 (84.3%)	32 (72.7%)	0.50 (0.25-1.05)	0.056	0.50 (0.25-1.05)	0.056
Women (N=479)	(<i>n</i> =446)	(<i>n</i> =33)				
Receptive anal sex	148 (33.2%)	12 (36.4%)	1.15 (0.54-2.37)	0.71	1.25 (0.57-2.61)	0.56
Active rimming	83 (18.6%)	3 (9.1%)	0.44 (0.10-1.27)	0.18	0.48 (0.11-1.43)	0.25
Passive rimming	47 (10.5%)	0	^c		^c	
Receptive cunnilingus	322 (72.2%)	23 (69.7%)	0.89 (0.42-2.00)	0.76	0.93 (0.43-2.12)	0.85
Insertive fellatio	379 (85.0%)	29 (87.9%)	1.28 (0.48-4.42)	0.65	1.21 (0.45-4.23)	0.73
Receptive fisting	7 (1.6%)	1 (3.0%)	1.96 (0.10-11.50)	0.54	2.84 (0.14-18.90)	0.36
Receptive fingering	363 (81.4%)	25 (75.8%)	0.71 (0.32-1.74)	0.43	0.73 (0.33-1.80)	0.46
Insertive fingering	77 (17.3%)	6 (18.2%)	1.06 (0.39-2.51)	0.89	1.23 (0.44-2.99)	0.66

Data were obtained from individuals in the BMR-IST cohort, Paris, France, 2018-2019.

^aInsertive and receptive vaginal intercourse was not included, as these behaviors were not considered in the Bayesian latent class analysis.

^bAdjusted for antibiotic use in the past 6 months, highest ESBL-E prevalence of country travelled (i.e., none, low, medium, high), and number of sexual partners (i.e., 0-7, 8-30, ≥ 30 for MSM and 0-7 and ≥ 8 for other men and women).

^cParameter estimates were not obtained due to 0 prevalence in certain categories.

Abbreviations: aOR, adjusted odds ratio; CI, confidence interval; ESBL-E, Extended-Spectrum β -Lactamase-Producing Enterobacterales; MSM, men who have sex with men; OR, odds ratio.

Table 4. Association between latent classes and ESBL-E carriage, stratified by sexual group

Latent class ^a	n	Mean proportion of behaviors	ESBL-E prevalence	Non-adjusted OR (95%CI)	<i>P</i>	Adjusted ^b aOR (95%CI)	<i>P</i>
MSM (N=1211)							
Class 1	259	29.9%	7.3%	Ref.		Ref.	
Class 2	102	42.3%	12.7%	1.85 (0.86-3.86)	0.11	1.74 (0.81-3.66)	0.15
Class 3	124	44.8%	15.3%	2.29 (1.16-4.52)	0.017	2.19 (1.11-4.34)	0.023
Class 4	100	60.0%	16.0%	2.41 (1.17-4.89)	0.015	2.31 (1.12-4.72)	0.021
Class 5	198	62.2%	10.6%	1.50 (0.78-2.89)	0.22	1.41 (0.73-2.73)	0.31
Class 6	358	79.7%	12.3%	1.77 (1.02-3.18)	0.047	1.65 (0.95-2.97)	0.084
Class 7	70	91.7%	24.3%	4.05 (1.96-8.34)	<0.001	3.78 (1.81-7.83)	<0.001
Other men (N=439)							
Class 1	72	11.1%	16.7%	Ref		Ref	
Class 2	297	45.0%	8.1%	0.44 (0.21-0.95)	0.031	0.44 (0.21-0.96)	0.031
Class 3	70	79.0%	11.4%	0.65 (0.24-1.67)	0.37	0.64 (0.24-1.67)	0.37
Women (N=479)							
Class 1	64	7.6%	9.4%	Ref		Ref	
Class 2	343	44.7%	7.0%	0.73 (0.30-2.03)	0.51	0.72 (0.29-2.04)	0.50
Class 3	72	81.2%	4.2%	0.42 (0.09-1.67)	0.23	0.45 (0.09-1.82)	0.28

Data were obtained from individuals in the BMR-IST cohort, Paris, France, 2018-2019.

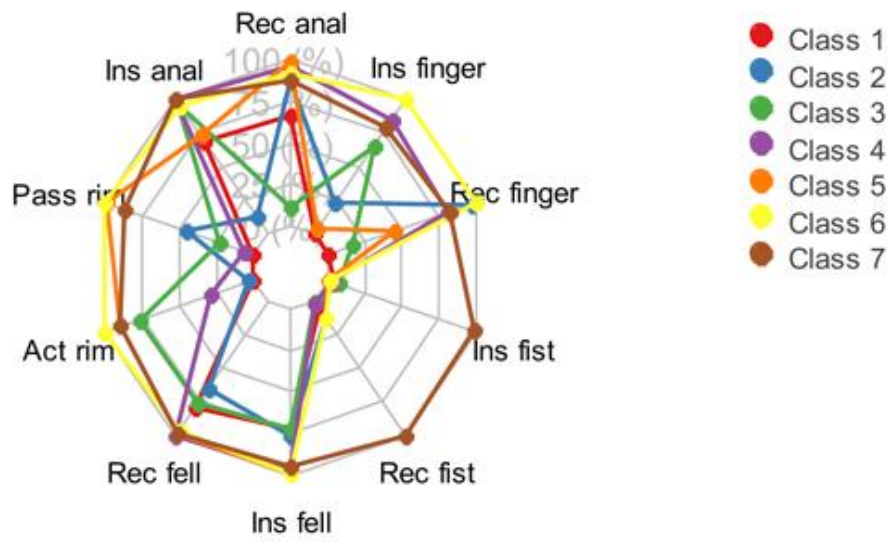
^aComposition of latent classes for MSM, other men and women are described in Figures 1A, 1B, and 1C, respectively.

^bAdjusted for antibiotic use in the past 6 months and highest ESBL-E prevalence of country travelled (i.e., none, low, medium, high).

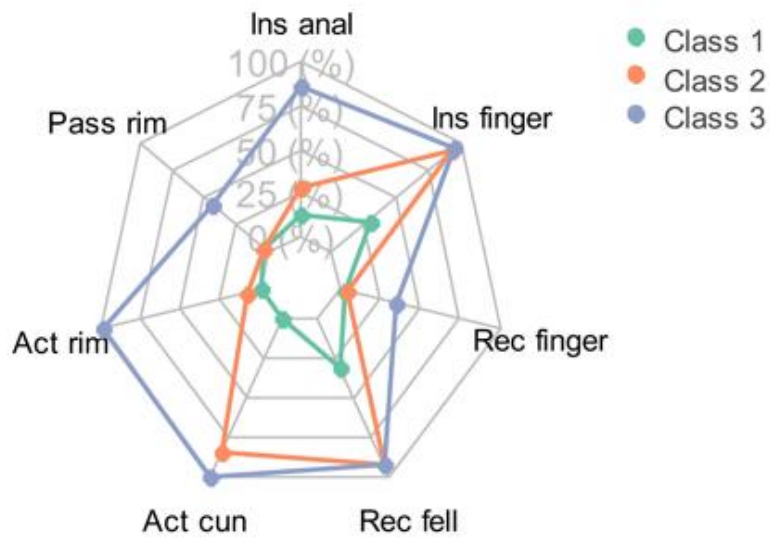
Abbreviations: aOR, adjusted odds ratio; CI, confidence interval; ESBL-E, Extended-Spectrum β -Lactamase–Producing Enterobacterales;

OR, odds ratio; ref, reference group.

A



B



C

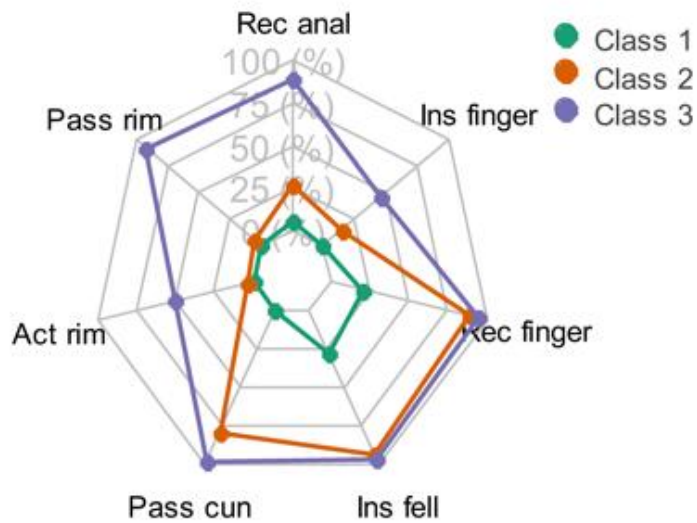


FIGURE LEGENDS

Figure 1. Description of sexual behavior clusters

Plots depict the proportion (in %) of individuals engaging in a given sexual behavior in the last 6 months, according to latent class. Analysis is stratified by men who have sex with men (**A**), other men (**B**), and women (**C**). Data were obtained from individuals in the BMR-IST cohort, Paris, France, 2018-2019.

Abbreviations: Act, active; cun, cunnilingus; fell, fellatio; fist, fisting; finger, fingering; Ins, insertive; Pass, passive; Rec, receptive; rim, rimming

Supplement to: Boyd A, Mathieu P, Françoise U, et al. Sexual behaviors and risk of extended-spectrum β -lactamase-producing Enterobacterales carriage: a cross-sectional analysis.

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SUPPLEMENTARY METHODS

Determining presence of ESBL-E in rectal swab samples

To ensure detection of small bacterial loads, we grew three separate precultures in Brain Heart Infusion (BHI) broth with 10µg/ml cefotaxime (to select for ESBL-E) after inoculation from the same e-rectal swab. Precultures were grown for 18h at 37°C. We then plated each culture on bioMerieux ESBL medium for selective isolation of ESBL-E. After incubating plates for another 18h at 37°C, identification of bacterial colonies was performed using mass spectrometry (MALDI BiotyperR, Bruker, Billerica, MA, USA). We chose colonies of ESBL-E strains picked from selective media and stored them at -80°C for further analysis. To ensure that Enterobacterales exhibiting antibiotic resistance via indirect pathways (e.g., overproduction of AmpC) were not selected, all enterobacterial or enterococcal species isolates were subjected to a standard antibiogram using Mueller-Hinton agar (Biorad), according to recommendations from the European Committee on Antimicrobial Susceptibility Testing (<https://www.eucast.org/>).

SUPPLEMENTARY TABLES

Supplementary Table S1. Fit statistics for Bayesian latent class models according to sexual group

Number of classes (<i>k</i>)	MSM		Other men		Women	
	AIC	BIC	AIC	BIC	AIC	BIC
2	-11004	-11111	-2685	-2746	-3027	-3090
3	-10692	-10855	-2579	-2673	-2920	-3016
4	-10530	-10750	-2573	-2700	-2900	-3029
5	-10365	-10640	-2570	-2729	-2891	-3054
6	-10221	-10552	-2569	-2761	-2888	-3085
7	-10145	-10532	-2571	-2796	-2887	-3137
8	-10089	-10533	-2577	-2835	-2892	-3154

Latent class models were fit, as described in the Statistical analysis section, on data obtained from individuals in the BMR-IST cohort, Paris, France, 2018-2019. Models were fit to increasing numbers of classes (*k*). The Akaike Information Criteria (AIC) and Bayesian Information Criteria (BIC) were calculated and the model with the BIC closest to 0 was selected in further analysis.

Supplementary Table S2. Characteristics of the study population stratified on membership of sexual behavior cluster (in men who have sex with men)

	Sexual clusters							<i>p</i> ^a
	Cluster 1	Cluster 2	Cluster 3	Cluster 4	Cluster 5	Cluster 6	Cluster 7	
	(<i>n</i> =259)	(<i>n</i> =102)	(<i>n</i> =124)	(<i>n</i> =100)	(<i>n</i> =198)	(<i>n</i> =358)	(<i>n</i> =70)	
Age, median (IQR), y	46 (34-53)	39 (28-47)	34 (28-46)	38 (30-45)	37 (30-45)	35 (28-45)	42 (35-49)	<0.001
Geographic origin								<0.001
Europe	185 (71.4%)	75 (73.5%)	100 (80.6%)	82 (82.0%)	164 (82.8%)	310 (86.6%)	64 (91.4%)	
North America, DOM, TOM or Australia	32 (12.4%)	8 (7.8%)	13 (10.5%)	10 (10.0%)	20 (10.1%)	28 (7.8%)	2 (2.9%)	
Africa or Middle East	32 (12.4%)	16 (15.7%)	9 (7.3%)	6 (6.0%)	11 (5.6%)	15 (4.2%)	3 (4.3%)	
Asia	10 (3.9%)	3 (2.9%)	2 (1.6%)	2 (2.0%)	3 (1.5%)	5 (1.4%)	1 (1.4%)	
Born in France	179 (69.1 %)	74 (72.5 %)	96 (77.4%)	76 (76.0%)	152 (76.8%)	296 (82.7%)	59 (84.3%)	0.004
Communal living ^b	9 (3.5%)	2 (2.0%)	2 (1.6%)	1 (1.0%)	1 (0.5%)	1 (0.3%)	0 (0)	0.023
Antibiotic use in past 6 months	58 (22.4%)	33 (32.4%)	42 (33.9%)	31 (31.0%)	71 (35.9%)	133 (37.2%)	34 (48.6%)	<0.001
Type of antibiotic used								0.0020
None	201 (77.6%)	69 (67.6%)	82 (66.1%)	69 (60.0%)	127 (64.1%)	225 (62.8%)	36 (51.4%)	
β-Lactams	29 (11.2%)	11 (10.8%)	13 (10.5%)	9 (9.0%)	22 (11.1%)	38 (10.6%)	9 (12.9%)	
Doxycycline	10 (3.9%)	7 (3.9%)	3 (6.9%)	2 (2.0%)	14 (7.1%)	22 (6.1%)	6 (8.6%)	
Other	19 (7.3%)	15 (7.3%)	26 (14.7%)	20 (20.0%)	35 (17.7%)	73 (20.4%)	19 (27.1%)	
Antibiotic use in past 3 months	40 (15.4%)	21 (20.6%)	21 (16.9%)	24 (24.0%)	50 (25.3%)	91 (25.4%)	26 (37.1%)	0.0017
Travel outside of France	171 (66.0%)	77 (75.5%)	90 (72.6%)	73 (73.0%)	147 (74.2%)	276 (77.1%)	53 (75.7%)	0.17
Hospitalization in the past 12 months	14 (5.4%)	1 (1.0%)	8 (6.5%)	8 (8.0%)	12 (6.1%)	22 (6.1%)	4 (5.7%)	0.47
Number of sexual partners in the past 6 months								<0.001
0-7	187 (72.2%)	60 (58.8%)	63 (50.8%)	52 (52.0%)	81 (40.9%)	151 (42.2%)	16 (22.9%)	
8-30	61 (23.6%)	31 (30.4%)	45 (36.3%)	32 (32.0%)	83 (41.9%)	145 (40.5%)	39 (55.7%)	
>30	11 (4.2%)	11 (10.8%)	16 (10.8%)	16 (16.0%)	34 (17.2%)	62 (17.3%)	15 (21.4%)	
HIV positive	167 (64.4%)	27 (26.4%)	52 (41.9%)	33 (33.0%)	64 (32.3%)	109 (30.4%)	37 (52.9%)	<0.001
PrEP use in the past 6 months ^c	32 (34.8%)	18 (24.0%)	24 (33.3%)	30 (44.8%)	46 (34.3%)	79 (31.7%)	17 (51.5%)	<0.001
Condomless sex in the past 6 months	199 (76.8%)	79 (77.5%)	109 (87.9%)	89 (89.0%)	171 (86.4%)	321 (89.7%)	55 (78.6%)	<0.001
Drug use ^d in the past 6 months	69 (26.6%)	40 (39.2%)	56 (45.2%)	40 (40.0%)	97 (49.0%)	189 (52.8%)	44 (62.9%)	<0.001
Any STI in the past 6 months	55 (21.2%)	28 (27.5%)	36 (29.0%)	26 (26.0%)	52 (26.3%)	93 (26.0%)	25 (35.7%)	0.30
Sex work in the past 6 months	3 (1.2%)	0 (0%)	6 (4.8%)	3 (3.0%)	7 (3.5%)	8 (2.2%)	2 (2.9%)	0.19

Data were obtained from individuals in the BMR-IST cohort, Paris, France, 2018-2019. Data represent no. (%) of participants unless

otherwise identified as median (IQR). Composition of latent classes for MSM are described in Figure 1A.

^aStatistical comparisons were performed using Pearson χ^2 or Fisher exact test for categorical and Kruskal-Wallis test for continuous variables.

^bCommunal living defined as living in a hotel, camp, or shelter or experiencing homelessness.

^cAmong participants without HIV.

^dDrug use excludes alcohol and tobacco use.

Supplementary Table S3. Characteristics of the study population stratified on membership of sexual behavior cluster (in other men)

	Sexual clusters			p^a
	Cluster 1	Cluster 2	Cluster 3	
	(n=72)	(n=297)	(n=70)	
Age, median (IQR), y	28 (24-35)	26 (23-32)	29 (25-38)	0.0040
Geographic origin				0.0045
Europe	47 (65.3%)	245 (82.5%)	56 (80.0%)	
North America, DOM, TOM or Australia	3 (4.2%)	15 (5.1%)	5 (7.1%)	
Africa or Middle East	22 (30.6%)	34 (11.4%)	8 (11.4%)	
Asia	0 (0%)	3 (1.0%)	1 (1.4%)	
Born in France	47 (65.3%)	249 (83.8%)	56 (80.0%)	0.0019
Communal living ^b	7 (9.7%)	3 (1.0%)	1 (1.4%)	0.0001
Antibiotic use in past 6 months	9 (12.5%)	29 (9.8%)	9 (12.9%)	0.65
Type of antibiotic used				0.41
None	63 (87.5%)	268 (90.2%)	61 (87.1%)	
β -Lactams	3 (4.2%)	18 (6.1%)	5 (7.1%)	
Doxycycline	2 (2.8%)	7 (2.4%)	4 (1.4%)	
Other	4 (5.6%)	4 (1.3%)	3 (4.3%)	
Antibiotic use in past 3 months	6 (8.3%)	17 (5.7%)	7 (10.0%)	0.38
Travel outside of France	48 (66.7%)	215 (72.4%)	50 (71.4%)	0.63
Hospitalization in the past 12 months	1 (1.4%)	8 (2.7%)	2 (2.9%)	0.80
Number of sexual partners in the past 6 months				<0.0001
0-7	68 (94.4%)	278 (93.6%)	56 (80.0%)	
8-30	4 (5.6%)	19 (6.4%)	11 (15.7%)	
>30	0 (0%)	0 (0%)	3 (4.3%)	
HIV positive	0 (0%)	0 (0%)	0 (0%)	<i>ntp</i>
PrEP use in the past 6 months ^c	1 (1.4%)	0 (0%)	0 (0%)	<i>ntp</i>
Condomless sex in the past 6 months	54 (75.0%)	257 (86.5%)	60 (85.7%)	0.050
Drug use ^d in the past 6 months	15 (20.8%)	108 (36.4%)	38 (54.3%)	0.0002
Any STI in the past 6 months	0 (0%)	21 (7.1%)	5 (7.1%)	0.066
Sex work in the past 6 months	0 (0%)	5 (1.7%)	2 (2.9%)	0.39

Data were obtained from individuals in the BMR-IST cohort, Paris, France, 2018-2019. Data represent no. (%) of participants unless otherwise identified as median (IQR). Composition of latent classes for other men are described in Figure 1B.

^aStatistical comparisons were performed using Pearson χ^2 or Fisher exact test for categorical and Kruskal-Wallis test for continuous variables. *ntp*, no test performed.

^bCommunal living defined as living in a hotel, camp, or shelter or experiencing homelessness.

^cAmong participants without HIV.

^dDrug use excludes alcohol and tobacco use.

Supplementary Table S4. Characteristics of the study population stratified on membership of sexual behavior cluster (in women)

	Sexual clusters			p^a
	Cluster 1	Cluster 2	Cluster 3	
	(<i>n</i> =64)	(<i>n</i> =343)	(<i>n</i> =72)	
Age, median (IQR), y	26 (23-32)	24 (22-28)	25 (22-29)	0.016
Geographic origin				0.0012
Europe	42 (66.7%)	286 (83.4%)	64 (88.9%)	
North America, DOM, TOM or Australia	4 (6.3%)	23 (6.7%)	6 (8.3%)	
Africa or Middle East	15 (23.8%)	29 (8.5%)	2 (2.8%)	
Asia	2 (3.2%)	5 (1.5%)	0 (0%)	
Born in France	38 (59.4%)	295 (86.0%)	60 (83.3%)	<0.0001
Communal living ^b	7 (10.9%)	6 (1.7%)	2 (2.8%)	0.0005
Antibiotic use in past 6 months	14 (21.9%)	58 (16.9%)	12 (16.7%)	0.62
Type of antibiotic used				0.61
None	50 (78.1%)	285 (83.1%)	60 (83.3%)	
β -Lactams	8 (12.5%)	36 (10.5%)	4 (5.6%)	
Doxycycline	2 (3.1%)	5 (1.5%)	2 (2.8%)	
Other	4 (6.3%)	17 (5.0%)	6 (8.3%)	
Antibiotic use in past 3 months	11 (17.2%)	41 (12.0%)	10 (13.9%)	0.50
Travel outside of France	43 (67.2%)	260 (75.8%)	54 (75.0%)	0.35
Hospitalization in the past 12 months	0 (0%)	15 (4.4%)	4 (5.6%)	0.20
Number of sexual partners in the past 6 months				<i>ntp</i>
0-7	62 (96.9%)	325 (94.8%)	62 (86.1%)	
8-30	2 (3.1%)	18 (5.2%)	10 (13.9%)	
>30	0 (0%)	0 (0%)	0 (0%)	
HIV positive	0 (0%)	0 (0%)	0 (0%)	<i>ntp</i>
PrEP use in the past 6 months ^c	0 (0%)	0 (0%)	1 (1.4%)	<i>ntp</i>
Condomless sex in the past 6 months	48 (75.0%)	285 (83.1%)	63 (87.5%)	0.15
Drug use ^d in the past 6 months	7 (10.9%)	127 (37.0%)	36 (50.0%)	<0.0001
Any STI in the past 6 months	2 (3.1%)	2 (0.6%)	1 (1.4%)	0.58
Sex work in the past 6 months	2 (3.1%)	2 (0.6%)	1 (1.4%)	0.18

Data were obtained from individuals in the BMR-IST cohort, Paris, France, 2018-2019. Data represent no. (%) of participants unless otherwise identified as median (IQR). Composition of latent classes for women are described in Figure 1C.

^aStatistical comparisons were performed using Pearson χ^2 or Fisher exact test for categorical and Kruskal-Wallis test for continuous variables. *ntp*, no test performed.

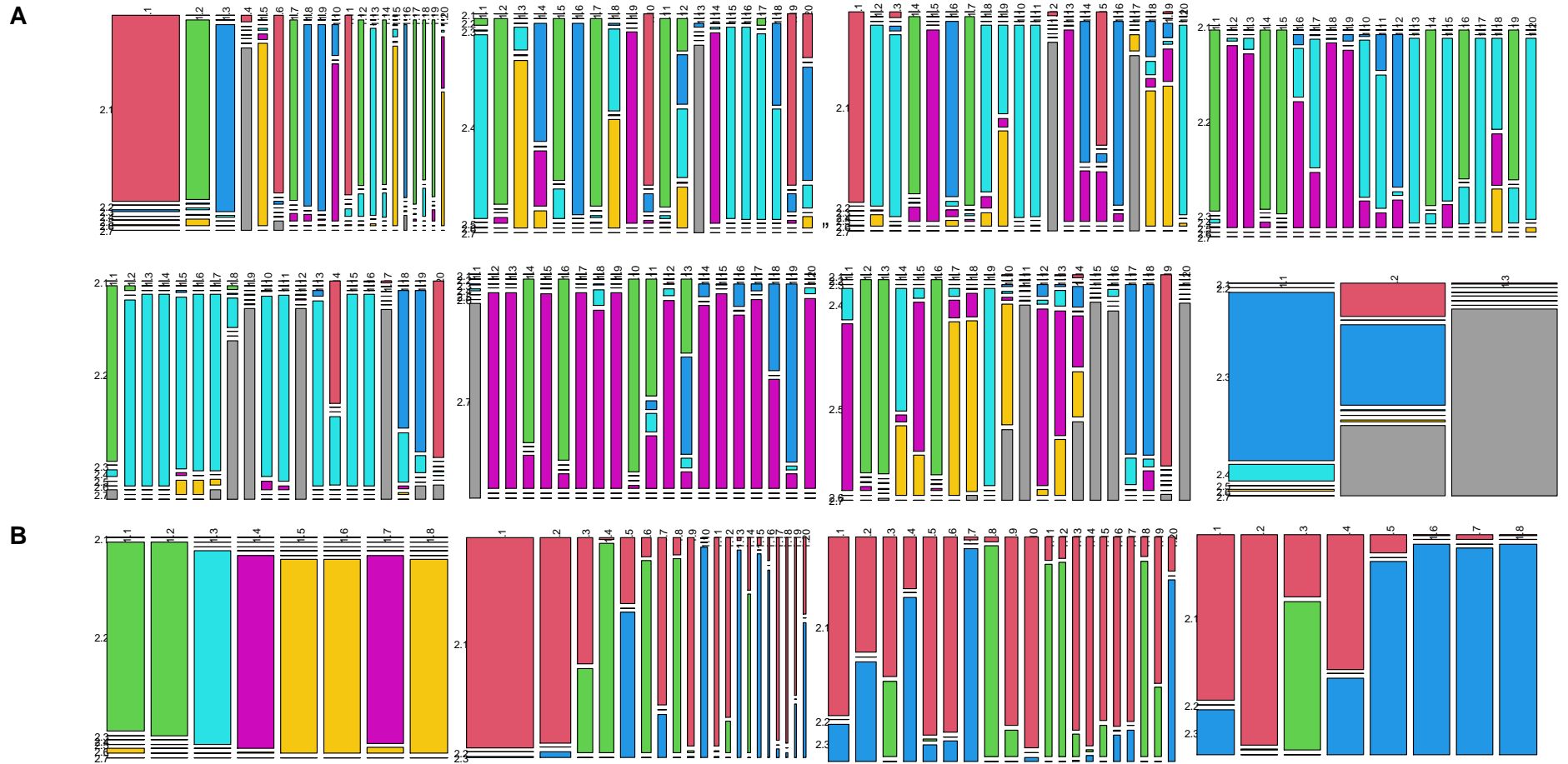
^bCommunal living defined as living in a hotel, camp, or shelter or experiencing homelessness.

^cAmong participants without HIV.

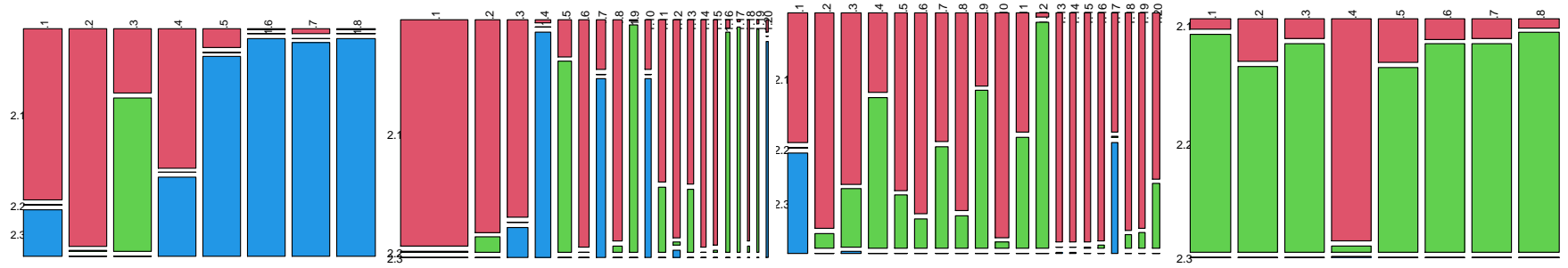
^dDrug use excludes alcohol and tobacco use.

SUPPLEMENTARY FIGURES

Supplementary Figure S1. Certainty plots for Bayesian latent class models according to sexual group



C



Plots are constructed in which combinations of sexual behaviors are categorized on the x-axis and the probability of belonging to each class is depicted on the y-axis. High degree of certainty is present when most categories have bars with one color. Given the high numbers of combinations, these diagnostic plots had to be presented in several panels. Plots are given for men who have sex with men (A), other men (B), and women (C).