

Seroprevalence and Risk Factors for Human Herpesvirus 8 Infection, Rural Egypt¹

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To determine whether human herpesvirus 8 (HHV-8) is associated with schistosomal and hepatitis C virus infections in Egypt, we surveyed 965 rural household participants who had been tested for HHV-8 and schistosomal infection (seroprevalence 14.2% and 68.6%, respectively, among those <15 years of age, and 24.2% and 72.8%, respectively, among those ≥15 years of age). Among adults, HHV-8 seropositivity was associated with higher age, lower education, dental treatment, tattoos, >10 lifetime injections, and hepatitis C virus seropositivity. In adjusted analyses, HHV-8 seropositivity was associated with dental treatment among men (odds ratio [OR] 2.4, 95% confidence interval [CI] 1.1–5.2) and hepatitis C virus seropositivity among women (OR 3.3, 95% CI 1.4–7.9). HHV-8 association with antischistosomal antibodies was not significant for men (OR 2.1, 95% CI 0.3–16.4), but marginal for women (OR 1.5, 95% CI 1.0–2.5). Our findings suggest salivary and possible nosocomial HHV-8 transmission in rural Egypt.

Human herpesvirus 8 (HHV-8, also called Kaposi sarcoma [KS]-associated herpesvirus) is the infectious cause of KS (1) and is prevalent in Africa (2). HHV-8 seroepidemiology parallels imperfectly KS epidemiology (3). Adult HHV-8 seropositivity is very high in eastern and central Africa (70%–90%), where KS is endemic, and lower in southern and northern Africa (10%–40%), including Egypt, where KS is more rare (4). This variation may be due, in part, to socioeconomic or environmental factors (5) influencing HHV-8 transmission or pathogenesis. HHV-8

is transmitted through contact with saliva (6–8), but sexual (9) and blood-borne (10) transmission also occur.

HHV-8 seroepidemiology in Egypt is incompletely described (3,11,12). Egypt offers the opportunity to investigate how HHV-8 correlates in the general population with well-characterized hepatitis C virus (HCV) (13) and schistosomal infections (14). Hundreds of thousands of Egyptians were exposed to multiple intravenous injections during treatment campaigns to control schistosomiasis from the 1950s until 1982, which resulted in an epidemic of HCV (15). Schistosomal infection has been reported to suppress the immune response to HCV, which could lead to more persistent infections in those who are co-infected (16–20). Whether a similar relationship exists with HHV-8 is not known. The rarity of KS in Egypt, despite its occurrence in organ transplant recipients (21), suggests that schistosomal-induced immunosuppression may not increase KS risk substantially. We sought to test the hypothesis that schistosomal seropositivity is associated with HHV-8 seropositivity, which would support the concept that schistosomal-induced immunosuppression modulates susceptibility to HHV-8 infection.

Methods

Patient Selection and Serologic Testing for HHV-8 and Schistosomal Infections

We randomly selected residual frozen serum samples from 965 of ≈6,000 persons who had participated in the HCV and schistosomiasis epidemiologic, population-based, household survey in Assiut Governorate, in rural southern Egypt in 1992 (13). Adults and parents of children

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≤15 years of age who had participated in the original survey gave informed consent and answered interviewer-administered questions about demographics, socioeconomic status, medical treatment, and parenteral exposures, including injections, transfusions, operations, dental treatment, and tattoos. Each participants gave a venous blood sample for serologic testing (13). The proportion of children included in our study was slightly lower than in the original survey because serum samples from some children had been exhausted by tests for HCV and other hepatitis viruses. However, the included children were otherwise representative of the original survey population.

Anti-HHV-8 antibodies were measured by using an enzyme immunoassay to K8.1 glycoprotein (a lytic-phase antigen) as previously reported (22,23). Antischistosomal antibodies were measured by using enzyme-linked immunoelectrotransfer blots (EITB) to detect species-specific antibodies against microsomal glycoprotein antigens from *Schistosoma hematobium* and *S. mansoni* (98% sensitivity and 99% specificity) (24). *S. hematobium* is the predominant species causing infection in Assiut Governorate; concurrent or single infection with *S. mansoni* among local inhabitants is rare (14).

Statistical Methods

Because no standard for testing for HHV-8 infection exists, we previously based cutoff values for defining seropositivity on visual inspection of the distribution of the K8.1 assay optical density (OD) values (10), assuming that there are seronegative and seropositive subpopulations. However, this approach is subjective. To more objectively define seropositivity, we applied mixture models to the OD data (25). Briefly, the mixture model was based on the assumption that the OD value for each participant arises from either a seronegative or a seropositive subpopulation. The formulas and details on parameters for the statistical model are described elsewhere ([25]; online Appendix, available from www.cdc.gov/EID/content/14/4/586-app.htm). We assumed that when the calculated posterior probability of seropositivity was ≥0.5, then the person was seropositive. In sensitivity analyses (online Appendix), we used alternative parameters of the model and also excluded persons with an intermediate posterior probability (range: 0.4–0.6).

After defining seropositivity, we used logistic regression models (PROC GENMOD, SAS 8.0; SAS, Cary, NC, USA) to calculate odds ratios (ORs) for HHV-8 associations with demographic, behavioral, and clinical risk factors. We used generalized estimation equations that accounted for correlations between persons living in the same household to calculate 95% confidence intervals (CIs) (26). Because HHV-8 seropositivity is age dependent and modes of transmission may differ between children and adults, we performed univariate and multivariable analyses separately

for children (<15 years of age) and adults (≥15 years of age). Because we postulated a priori that antischistosomal antibodies were associated with HHV-8, we included schistosomal status in all multivariable models. To adjust for potential confounding, we included in our multivariable models those variables that were associated with HHV-8 seropositivity at $p \leq 0.1$ in univariate analyses. Age was fitted with a trend whenever this resulted in a statistically significantly improved model fit; otherwise, it was fitted as a categorical variable with dummy values. Two-tailed p values ($p < 0.05$) were considered statistically significant, while p values between 0.1 and 0.05 were suggestive of a trend.

Results

None of the original household survey participants had a history of KS (13). HHV-8 seroprevalence was lower among children compared with adults (14.3% vs. 24.2%, $p < 0.001$). Among children, in unadjusted analyses, HHV-8 seroprevalence was higher in girls than boys (20% vs. 9%; OR 2.4, 95% CI 1.1–5.3). HHV-8 seroprevalence was not significantly elevated in children with a history of ≥10 lifetime injections compared with those with <10 lifetime injections (18% vs. 11%, OR 1.8, 95% CI 0.8–3.8) nor among those with schistosomal antibodies (16% vs. 10%; OR 1.7, 95% CI 0.7–4.3). Age, education, dental treatment, tattoos, and HCV antibodies were unrelated to HHV-8 seropositivity among children.

Among adult men and women combined, unadjusted analyses showed that HHV-8 seropositivity was higher among older participants (≥45 years of age) compared with younger participants (15–24 years of age; OR 4.1, 95% CI 2.6–6.6); among those currently married (OR 1.9, 95% CI 1.2–3.0) or divorced, separated, or widowed (OR 3.3, 95% CI 1.7–6.4) versus never married; and among those with a history of dental treatment (OR 2.1, 95% CI 1.5–2.9), ≥10 lifetime injections (OR 1.5, 95% CI 1.0–2.3), tattoos (OR 1.7, 95% CI 1.1–2.7), or HCV seropositivity (OR 1.8, 95% CI 1.0–3.3) compared with participants without these characteristics. Conversely, HHV-8 seropositivity was lower among adults who reported primary (OR 0.6, 95% CI 0.4–1.0) or higher level of formal education (OR 0.3, 95% CI 0.2–0.6) compared with participants without formal education.

Antischistosomal antibodies, which indicate past as well current infection, were detected among 72.8% of participants. Current infection, as indicated by ova in stool or urine samples, was noted for 4% of participants, all of whom had *S. hematobium*. *S. mansoni* was only recently introduced in Assiut Governorate and remains rare and focal in distribution (14). Almost all participants who had *S. mansoni* antibodies were also positive for *S. hematobium*. Patterns of schistosomal seropositivity differed between women and men. Among women, antischistosomal antibody prevalence

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was inversely related with age (68.8% in those 15–24 years of age vs. 34.1% in those ≥ 45 years of age; $p < 0.001$); was lower in those who were widowed, divorced, or separated compared with those who were married or who had never married (38.8% vs. 63.4%; $p = 0.003$); but was unrelated to education ($p = 0.36$) or HCV seropositivity ($p = 0.12$). Among men, antischistosomal antibody prevalence was unrelated to age ($p = 0.61$), marital status ($p = 0.73$), education ($p = 0.64$), or HCV seropositivity ($p = 0.12$).

In unadjusted sex-specific analyses (Tables 1 and 2), HHV-8 seropositivity was not associated with antischisto-

somal antibodies in women (OR 1.0, 95% CI 0.6–1.5); it was not significantly associated in men (OR 2.3, 95% CI 0.3–19.0), because only 9 men had no antischistosomal antibodies. The HHV-8 associations with age, formal education, marital status, and history of dental treatment among men and women combined were largely recapitulated in the sex-specific analyses, except for associations with having tattoos, ≥ 10 lifetime injections, and HCV seropositivity, which were evident in women but not men (Tables 1 and 2). HHV-8 was significantly associated with cigarette smoking; cigarette smoking was only recorded for men.

Table 1. Prevalence and crude OR of association of HHV-8 seropositivity with demographic, behavioral, and clinical characteristics among male patients, Egypt*

Characteristic	n/N	%	OR	95% CI	p value†
Total	52/235	22.1	–	–	–
Age group, y					<0.001
15–24	8/96	8.3	Ref	–	–
25–34	16/54	29.6	4.6	2.0–11.0	–
35–44	10/30	33.3	5.5	1.9–15.7	–
≥ 45	18/55	32.7	5.3	2.2–13.2	–
Education					0.001
None	19/50	38.0	Ref	–	–
Primary	22/100	22.0	0.5	0.2–1.0	–
Secondary/past secondary	11/85	12.9	0.2	0.1–0.6	–
Job					0.001
Student	2/38	5.3	Ref	–	–
Not working	6/31	19.3	4.3	0.8–23.3	–
Farmer	25/87	28.7	7.3	1.6–32.4	–
Trade, service, production	13/63	20.6	4.7	1.1–20.0	–
Technician/secretary	6/16	37.5	10.8	1.9–61.3	–
Marital status					0.004
Not married	11/91	12.1	Ref	–	–
Married	39/136	28.7	3.1	1.4–6.5	–
Separated/divorced/widowed	2/8	25.0	7.6	1.0–60.8	–
Dental treatments					<0.001
No	16/128	12.5	Ref	–	–
Yes	36/107	33.6	3.5	1.8–7.1	–
Tattoos					0.53
No	50/220	22.7	Ref	–	–
Present	2/15	13.3	0.5	0.1–2.5	–
HCV serostatus					0.72
Negative	48/214	22.4	Ref	–	–
Positive	4/21	19.0	0.8	0.3–2.5	–
Injections (lifetime)					0.74
<10	20/72	20.8	Ref	–	–
≥ 10	37/163	22.7	1.1	0.6–2.1	–
Smoking					0.02
Never	23/138	16.7	Ref	–	–
Ever	29/97	29.9	2.1	1.1–4.0	–
Goza‡					0.77
Never	42/193	21.8	Ref	–	–
Ever	10/42	23.8	1.1	0.5–2.4	–
Schistosomiasis					0.43
Negative	1/9	11.1	Ref	–	–
Positive	51/226	31.0	2.3	0.3–19.0	–

*OR, odds ratio; HHV-8, human herpesvirus 8; CI, confidence interval; Ref, referent; HCV, hepatitis C virus.

†p value for age group and education is test for trend; otherwise, p value is test for heterogeneity

‡Goza is a method of tobacco smoking in which tobacco smoke passes through a water pipe.

Table 2. Prevalence and crude OR of association of HHV-8 seropositivity with demographic, behavioral, and clinical characteristics among female patients, Egypt*

Characteristic	n/N	%	OR	95% CI	p value†
Total	125/495	25.3	–	–	–
Age group, y					<0.001
15–24	23/148	15.5	Ref	–	–
25–34	21/106	19.8	1.4	0.7–2.7	–
35–44	31/115	27.0	2.0	1.1–3.6	–
≥45	50/126	39.7	3.6	2.1–6.3	–
Education					<0.001
None	88/307	28.7	Ref	–	–
Primary	34/154	32.7	0.7	0.4–1.1	–
Secondary/past secondary	4/38	22.5	0.3	0.1–0.8	–
Job					0.11
Other‡	2/21	9.5	Ref	–	–
Housewife	121/470	25.7	3.3	0.8–14.3	–
Dental treatments					0.02
No	49/239	20.5	Ref	–	–
Yes	77/260	29.6	1.6	1.1–2.5	–
Tattoos					0.006
No	93/410	22.7	Ref	–	–
Present	33/89	37.1	2.0	1.2–3.3	–
HCV serostatus					0.006
Negative	112/470	23.8	Ref	–	–
Positive	14/29	48.3	3.0	1.4–6.5	–
Injections (lifetime)					0.03
<10	23/128	18.0	Ref	–	–
≥10	103/371	27.8	1.7	1.1–2.9	–
Schistosomiasis					0.94
Negative	48/189	25.4	Ref	–	–
Positive	76/303	25.1	1.0	0.6–1.5	–

*OR, odds ratio; HHV-8, human herpesvirus 8; CI, confidence interval; Ref, referent; HCV, hepatitis C virus.

†p value for age group and education is test for trend; otherwise, p value is test for heterogeneity.

‡Job category for women: "Other" includes women who are students, are not working, or have technical or scientific jobs.

In a multivariable analysis, HHV-8 seropositivity was higher among girls than boys (OR 2.6, 95% CI 1.2–5.8) and marginally associated with antischistosomal antibodies (OR 2.2, 95% CI 0.8–5.6). Among adult men, HHV-8 seropositivity was independently associated with older age and history of dental treatment but not with schistosomal antibodies (OR 2.3, 95% CI 0.3–16.1; Table 3). Among women, HHV-8 seropositivity was associated with older age, HCV seropositivity, and marginally with antischistosomal antibodies (OR 1.5, 95% CI 1.0–2.5; $p = 0.07$).

Discussion

We report HHV-8 seroepidemiology in a rural population in Egypt in which correlates of schistosomal and HCV were previously characterized (13,14). As in other populations, HHV-8 seropositivity in Egypt rose with increasing age (3,5). In subgroups, we found associations of HHV-8 seropositivity with history of dental therapy, lifetime injections, tattoos, and HCV seropositivity.

Previous studies of HHV-8 in Egypt (11,12) reported a seroprevalence of ≈40%, which is ≈2× the prevalence we observed. Those studies were hospital based, were conducted in urban areas, and detected anti-HHV-8 antibod-

ies with lytic immunofluorescence assays; all of these factors may have contributed to higher prevalence estimates. Despite these differences, the patterns in our associations support their validity. First, we detected HHV-8 antibodies in children, consistent with earlier reports and nonsexual HHV-8 spread in Egypt (11). Second, HHV-8 seroprevalence increased with age, in accord with the general pattern of HHV-8 observed in other populations (5). Among men, HHV-8 seropositivity was significantly associated with dental treatment, which may be a marker for transmission through saliva- or blood-contaminated dental instruments. Our HHV-8 associations with a history of ≥10 lifetime injections, tattoos, and HCV seropositivity also point to possible blood-borne transmission and agree with other studies (10,27,28). Sexual transmission might be suggested by our HHV-8 association with marital status, but this association did not persist after adjustment for age. Rather than ongoing transmission among adults, higher HHV-8 seropositivity in older persons may be due to a birth-cohort effect, i.e., reflecting periods of elevated HHV-8 transmission risk in the past. In this regard, the widespread use of intravenous injections for schistosomiasis treatment and control programs from 1950 to 1982 would be consistent with a

Table 3. Adjusted OR of association of HHV-8 seropositivity with demographic and clinical variables among adults, Egypt*

Characteristic	Men			Women		
	OR	95% CI	p value	OR	95% CI	p value
Age group, y†			0.002			
15–24	Ref	–	–	Ref.	–	–
25–34	1.6	1.2–2.2	–	0.8	0.4–1.6	0.53
35–44	2.6	1.4–4.9	–	1.5	0.8–2.9	0.15
≥45	4.3	1.7–11.0	–	3.1	1.5–6.4	<0.001
Dental treatments‡			0.04			
No	Ref	–	–	–	–	–
Yes	2.3	1.1–4.9	–	–	–	–
HCV serostatus‡			–			0.007
Negative	–	–	–	Ref	–	–
Positive	–	–	–	3.3	1.4–7.9	–
Schistosomiasis§			0.47			0.07
Negative	Ref	–	–	Ref	–	–
Positive	2.3	0.3–16.1	–	1.5	1.0–2.5	–

*OR, odds ratio; HHV-8, human herpesvirus 8; CI, confidence interval; Ref, referent; HCV, hepatitis C virus.

†p value is for age group fitted with trend among men; p values for heterogeneity for categories given for women (see Statistical Methods).

‡Missing values in sex-specific analyses mean the variable was not significant and was excluded from final multivariable model.

§Schistosomiasis seropositivity was included in models even when not significant because we hypothesized a priori that it was associated with HHV-8 seropositivity (see online Appendix, available from www.cdc.gov/EID/content/14/4/586-app.htm).

birth-cohort effect for high HHV-8 seroprevalence among our older participants. However, most populations, including those with no similar historical programs, have higher HHV-8 seroprevalence among older persons, which suggests that ongoing HHV-8 transmission among adults is a more likely explanation.

We found a 2-fold higher HHV-8 seroprevalence in persons who also had schistosomal antibodies. In multivariable analyses, the CI for this association was 1.0–2.5 among women, but it was wide among children (0.8–5.6) and especially among men (0.3–16.1). Chance association and serologic cross-reactivity between HHV-8 and schistosomal antibodies are possible explanations, but schistosomal antibodies may be a valid marker for exposure to injections in the historical anti-schistosomal program; these findings are consistent with parenteral transmission of HHV-8 as discussed above. Another possibility is that anti-schistosomal antibodies may be a marker for contact with surface water sources or walking barefoot, which are also risk factors for HHV-8 seropositivity and KS (5,7,29). No biologic explanation has been advanced for these environmental correlations with HHV-8 and KS. Our study suggests that perhaps contact with surface water or walking barefoot is a marker of exposure to and potential infection with *Schistosoma* or other water-related parasites. Infection with such parasites could influence the natural history of HHV-8 by shifting the immune response from a T helper 1 (Th1)-type response, which is central to controlling viral infections, to a Th2-dominant response (30), which is less effective against viral infections. If this model is correct, schistosomal infection could increase susceptibility to HHV-8 infection at relatively low exposure to the virus. In parallel, Th2-dominant hosts

may fail to effectively control HHV-8 infection and thus shed infectious virions in saliva more frequently and at higher levels, resulting in higher HHV-8 transmission. If our findings are confirmed, they could drive investigations of environmental characteristics, including exposures to volcanic soil or plants (31), to explain variation in HHV-8 infection and possibly KS.

Our study has several limitations. First, current HHV-8 serologic assays have imperfect specificity and sensitivity (32), which could have contributed to the lower HHV-8 seroprevalence observed. Except for the possible cross-reactivity mentioned above, serologic misclassification is likely to be random, which would attenuate associations toward the null. Second, our HCV and schistosomal antibody assays cannot distinguish current from resolved infections, diminishing the strength of observed associations as well. Third, with our cross-sectional design, we cannot determine the temporality of associations. This limitation may be particularly relevant to our findings of HHV-8 with antischistosomal antibodies. The antischistosomal programs surely reduced the prevalence and load of schistosome eggs, but they may also have contributed to HHV-8 transmission through injections. Finally, we studied only ≈15% of the participants in the original survey, which limited our statistical power to estimate some associations.

The strengths of our study include our state-of-the-art serologic methods, our model-based approach to estimating infection risk, and our well-characterized general population with detailed socioeconomic and clinical data. HHV-8 seropositivity was associated with older age, dental therapy, lifetime injections, and HCV and schistosomiasis seropositivity. These findings suggest salivary and possible nosocomial HHV-8 transmission in rural Egypt and

a potential biologic explanation for geographic variation of HHV-8 seropositivity and KS.

Dr Mbulaiteye is a tenure-track investigator in the Viral Epidemiology Branch, Division of Cancer Epidemiology and Genetics, National Cancer Institute, Bethesda, Maryland, USA. He is interested in characterizing the relationship between immunity, human herpesvirus infection, and cancer, particularly in Africa.

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Appendix

Statistical Methods

Because there is no standard for testing for human herpesvirus 8 (HHV-8) infection, we defined *infection status* by using a statistical mixture, or latent class, model for the optical density (OD) readings of the K8.1 assay (I). Briefly, the model considered the OD assay readings for each participant to arise from either an uninfected ($I = 0$) or an infected ($I = 1$) population. The probability density function of $y = (x^\lambda - 1) / \lambda$, where x denoted the assay readings of the K8.1 assay, is $g(y) = p f(y; \alpha_0) + (1-p) f(y; \alpha_1)$. The probability density function $f(y; \alpha_1)$ reflected the infected population, whereas $f(y; \alpha_0)$ was the probability density function corresponding to uninfected. P is the proportion of uninfected persons in the population. Details on parameters of the component densities and the estimation of the parameters are described elsewhere (I). Patients were classified as infected, $I = 1$, if the posterior probability of infection, given x , i.e., $\text{pr}(I = 1 | x) = p f(y; \alpha_0) / \{ p f(y; \alpha_0) + (1-p) f(y; \alpha_1) \}$ was ≥ 0.5 and as uninfected otherwise. This classification rule minimized the overall misclassification probability based on the mixture model when discriminating infected from uninfected subjects.

The association of HHV-8 seropositivity with demographic, behavioral, and clinical risk factors was determined by fitting logistic regression models to computed odds ratios (OR) for infection status (PROC GENMOD, SAS 8.0, SAS, Cary, NC, USA). Because HHV-8 seropositivity is age dependent, we performed analyses separately for children and adults to minimize confounding from age. In addition, because we found significant differences by sex and age in the seroepidemiology of schistosomiasis (all p values < 0.05), which we had postulated a priori would be associated with HHV-8, we performed overall and sex-specific analyses for the HHV-8 associations among adults. Because many exposures are age dependent, we constructed multivariable models including variables that were significant at $p \leq 0.1$ in univariate analyses separately for children, men, and women to determine the independent contribution of variables to HHV-8 seropositivity. Age was fitted with a trend whenever this resulted in a statistically significant ($p < 0.05$) improved model fit; otherwise, it was fitted as a categorical variable with dummy values for the categories. Schistosomal seropositivity was included in all models because we postulated a priori that it was associated with HHV-8 seropositivity. We used generalized

estimation equations to calculate 95% confidence intervals to account for correlations between participants living in the same household (2). We assumed an equally correlated working correlation matrix when computing the variances of the OR estimates, but other working correlations yielded similar results. We assumed that a 2-tailed p value <0.05 was statistically significant and that p values between 0.1 and 0.05 represented a trend.

Sensitivity Analyses

Appendix Tables 1–3 show our sensitivity analyses to address the concern that different model parameterizations may give very different estimates of posterior probabilities of infection. Appendix Table 1 shows HHV-8 seropositivity classification based on the mixture models with the Box Cox transformation and a) normal component densities, b) polynomial degree 1 and 2, c) polynomial degree 2, which was used in the paper. The table shows that no large variations in HHV-8 prevalence are observed, and that the classification of infection based on the posterior probabilities of the respective models leads to different classification of at most 4/734 individuals.

Appendix Table 2 addresses the concern that the degree of uncertainty in the models may lead to widely varying estimates of HHV-8 prevalence and thus, widely varying odds ratios in multivariable association models. The adjusted logistic regression associations before and after exclusion of individuals whose posterior probability was between 0.4-0.6 (38/734 subjects) and using HHV-8 status based on classification from 2 other parameterizations are presented in Appendix Tables 2 and 3. These tables show that the results presented in the manuscript do not appear to be overly sensitive to inclusion of “indeterminate range” individuals (Appendix Table 2) nor to the specific parameterization of the mixture model (Appendix Table 3). The different models gave very similar associations. The stability in the results obtained suggests that the associations that we present are likely valid.

Appendix Table 1. Cross-tabulation of classification of patients by different models to estimate human herpesvirus 8 seropositivity*

Model	Negative	Positive	Total
Model II		Model I	
Negative	556	4	560
Positive	0	174	174
Total	556	178	734
Model III		Model II	
Negative	559	0	559
Positive	1	174	175

Total	560	174	734
Model I	Model III		
Negative	556	0	556
Positive	3	175	178
Total	559	175	734

*I: Classification based on posterior probability from model $\lambda = 1, K = 2$ (used in this study); II: Classification based on posterior probability from model $\lambda = 1, K = 1$; III: Classification based on posterior probability from model $\lambda = 1, K = 0$ (Box Cox transformation included, normal component densities).

Appendix Table 2. Adjusted human herpesvirus 8 associations with and without indeterminate participants

Characteristic	Model I*			Model II†		
	OR	95% CI	p value	OR	95% CI	p value
Men						
Age group, y			<0.002			0.002
15–24	Ref			Ref		
25–34	1.6	1.2–2.2		1.6	1.2–2.2	
35–44	2.6	1.4–4.9		2.6	1.4–5.1	
45+	4.3	1.7–11.0		4.2	1.6–11.4	
Dental treatments			<0.04			0.05
No	Ref			Ref		
Yes	2.3	1.1–4.9		2.3	1.0–5.1	
HCV serology			–			
Negative	–			Ref		
Positive	–	–				
Schistosomiasis			0.47			0.43
Negative	Ref			Ref		
Positive	2.3	0.3–16.1		2.1	0.3–15.5	
Women						
Age group, y						
15–24	Ref			Ref		
25–34	0.8	0.4–1.6	0.53	0.9	0.5–1.9	0.9
35–44	1.5	0.8–2.9	0.15	1.7	0.9–3.2	0.1
45+	3.1	1.5–6.4	<0.001	3.7	2.0–6.9	<0.001
Dental treatments			–			
No	–			–		
Yes	–	–		–		
HCV serology			0.007			0.008
Negative	Ref			Ref		
Positive	3.3	1.4–7.9		3.3	1.4–8.0	
Schistosomiasis			0.07			0.09
Negative	Ref			Ref		
Positive	1.5	1.0–2.5		1.5	0.9–2.5	

*Classification based on posterior probability from model $\lambda = 1, K = 2$ (used in the paper). OR, odds ratio; CI, confidence interval; Ref, referent; HCV, hepatitis C virus.

†Results based on statistical model excluding subjects in the model indeterminate range (posterior probability 0.4–0.6).

Appendix Table 3. Adjusted human herpesvirus 8 associations by using alternative models*

Characteristic	Model II			Model III		
	OR	95% CI	p value	OR	95% CI	p value
Men						
Age group, y			<0.002			0.002
15–24	Ref.			Ref.		
25–34	1.6	1.2–2.3		1.6	1.2–2.2	
35–44	2.7	1.4–5.1		2.6	1.4–4.9	
45+	4.5	1.7–11.7		4.3	1.7–11	
Dental treatments			<0.03			0.04
No	Ref			Ref		
Yes	2.4	1.1–5.3		2.3	1.1–4.9	
HCV serology			–			

Negative	–			Ref		
Positive	–	–				
Schistosomiasis			0.43			0.41
Negative	Ref			Ref		
Positive	2.2	0.3–15.7		2.3	0.3–16.1	
<hr/>						
Women						
Age group, y						
15–24	Ref.			Ref.		
25–34	0.8	0.4–1.5	0.43	0.8	0.4–1.5	0.43
35–44	1.5	0.8–2.7	0.23	1.5	0.8–2.7	0.23
45+	3.6	2.0–6.5	<0.001	3.6	2.0–6.5	<0.001
Dental treatments						
No	–				–	
Yes	–	–			–	
HCV serology						
Negative	Ref.		0.02	Ref		0.02
Positive	2.9	1.2–6.9		2.9	1.2–6.9	
Schistosomiasis						
Negative	Ref		0.07	Ref		0.07
Positive	1.5	1.0–2.5		1.5	0.1–2.5	

*II, classification based on posterior probability from model $\lambda = 1$, $K = 1$; III, classification based on posterior probability from model $\lambda = 1$, $K = 0$ (Box Cox transformation included, normal component densities). R, odds ratio; CI, confidence interval; Ref, referrent; HCV, hepatitis C virus.