

Substantial regional differences in human herpesvirus 8 seroprevalence in sub-Saharan Africa: insights on the origin of the “Kaposi’s sarcoma Belt”

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Equatorial Africa has among the highest incidences of Kaposi’s sarcoma (KS) in the world, thus earning the name “KS Belt.” This was the case even before the HIV epidemic. To date, there is no clear evidence that HHV-8 seroprevalence is higher in this region but interpretation of the available literature is tempered by differences in serologic assays used across studies. We examined representatively sampled ambulatory adults in Uganda, which is in the “KS Belt,” and in Zimbabwe and South Africa which are outside the Belt, for HHV-8 antibodies. All serologic assays were uniformly performed in the same reference laboratory by the same personnel. In the base-case serologic algorithm, seropositivity was defined by reactivity in an immunofluorescence assay or in 2 enzyme immunoassays. A total of 2,375 participants were examined. In Uganda, HHV-8 seroprevalence was high early in adulthood (35.5% by age 21) without significant change thereafter. In contrast, HHV-8 seroprevalence early in adulthood was lower in Zimbabwe and South Africa (13.7 and 10.8%, respectively) but increased with age. After age adjustment, Ugandans had 3.24-fold greater odds of being HHV-8 infected than South Africans ($p < 0.001$) and 2.22-fold greater odds than Zimbabweans ($p < 0.001$). Inferences were unchanged using all other serologic algorithms evaluated. In conclusion, HHV-8 infection is substantially more common in Uganda than in Zimbabwe and South Africa. These findings help to explain the high KS incidence in the “KS Belt” and underscore the importance of a uniform approach to HHV-8 antibody testing.

Key words: KSHV, HHV-8, Africa, adults

Potential conflicts of interest. All authors: no conflicts.

Presented In Part: Oral Presentation at the Seventh International Workshop on Kaposi’s Sarcoma Associated Herpesvirus and Related Agents, Santa Cruz, CA, 2004.

Grant sponsor: National Institutes of Health; **Grant numbers:** U01 CA078124, R01 CA119903, P30 MH062246, D43 TW000003, R03 TW006054, P01 DE007946, K01 HD052020, and P30 AI027763;

Grant sponsor: AIDS Malignancy Consortium; **Grant number:** U01 CA071375; **Grant sponsor:** University of California Universitywide AIDS Research Program; **Grant number:** CC99-SF-001; **Grant sponsor:** South Africa National Research Foundation (Thuthuka Programme); **Grant number:** 2054349; **Grant sponsor:** AIDS Care Research in Africa Program

DOI: 10.1002/ijc.25235

History: Received 29 Sep 2009; Accepted 8 Jan 2010; Online 8 Feb 2010

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Even prior to the HIV epidemic, the incidence of Kaposi’s sarcoma (KS) in equatorial Africa was among the highest in the world. In portions of Uganda, Tanzania, and what is now known as the Democratic Republic of Congo, the lifetime incidence of KS approached 16 per 1,000,¹ thus earning the region the name “KS Belt” (Fig. 1). Although KS case reporting is imperfect in Africa, it is clear that non-HIV-related KS incidence was about 3–10 times higher in this region as compared with countries further north and south on the continent.¹ Although several theories have been offered (e.g., regional volcanic salts²), none has been proven to explain this substantial difference in KS incidence over such a relatively small geographic area.

With the discovery that human herpesvirus 8 (HHV-8 also known as KS-associated herpesvirus) is the viral etiologic agent of KS,^{3–8} it was naturally hypothesized that regional differences in HHV-8 seroprevalence in Africa would explain differences in KS incidence, similar to how high HHV-8 seroprevalence in homosexual men in the U.S.^{9–11} and

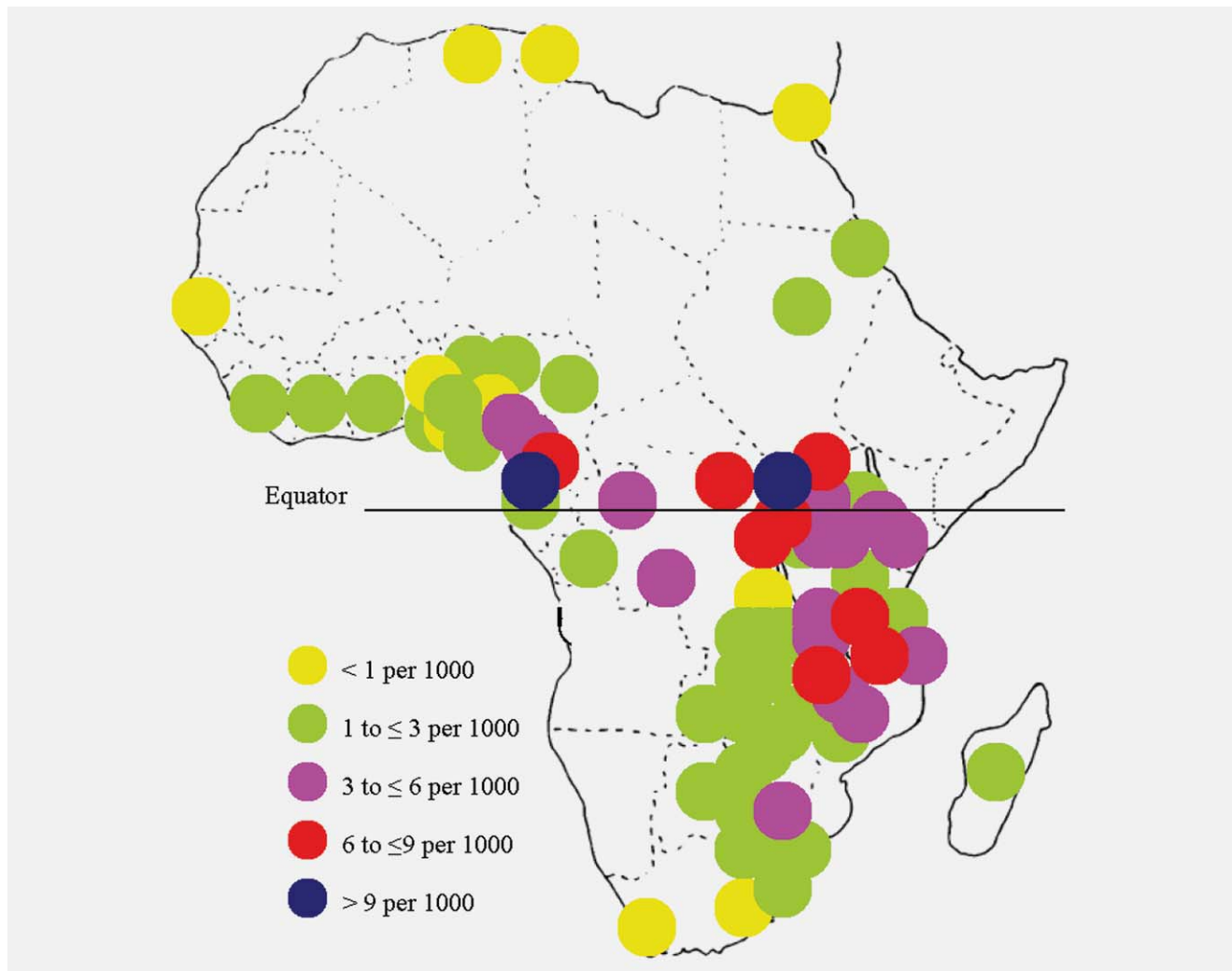


Figure 1. Cumulative incidence from birth to age 64 years of endemic (non-HIV related) Kaposi's sarcoma among men in Africa. Estimates are from Cook-Mozaffari *et al.*¹.

Europe^{12–14} was quickly determined to explain their high KS incidence.¹⁵ However, despite many studies of HHV-8 seroprevalence in Africa, clear regional differences have not been established resulting in the current view by many that the prevalence of HHV-8 is uniform throughout Africa.^{16–18} However, a closer inspection of work to date reveals that the estimates of HHV-8 seroprevalence from different African regions are difficult to compare. In addition to demographic differences in the subjects tested, differences in serologic testing importantly often preclude comparisons of studies across regions. This is because there is no commercially standardized HHV-8 antibody assay, a variety of different antigens are being targeted and serologic platforms being used across studies and there is proven assay discordance between assays.¹⁹

To address whether underlying regional differences in HHV-8 seroprevalence may in part explain differences in KS incidence, we compared HHV-8 seroprevalence in 3 countries with different pre-AIDS epidemic incidences of KS—

Uganda with high KS incidence, versus Zimbabwe and South Africa with lower KS incidence. Importantly, to preclude differences in serologic techniques, we tested all samples with assays for HHV-8 antibodies performed in the same laboratory.

Material and Methods

Participants

Participants were sampled in 3 cross-sectional studies of primarily young adults in Uganda, Zimbabwe, and South Africa. In Uganda, subjects were consecutive blood donors, age 17 years or older at the Nakasero Blood Bank in 2000–2001 who resided in Kampala and surrounding periurban and rural areas. Blood donation was voluntary and without monetary incentive. All blood donors in this analysis had been screened and found by the blood bank to be seronegative for HIV, hepatitis B virus, and *Treponema pallidum* infections. A portion of these subjects have been previously described.²⁰ In Zimbabwe, as part of a more intensive study of HHV-8

seroepidemiology and virologic shedding, consecutive women aged 17 years or older seeking family planning services at 4 clinics (Harare Hospital, Spilhaus Clinic, Chitungwiza Clinic, and Epworth Clinic) in Harare and surrounding periurban areas were recruited from 2001 to 2004. Participants were selected based on their HIV infection status in a 2:1 infected to uninfected ratio. In South Africa, as part of a larger study of horizontal transmission of HHV-8 to children, primary female caregivers of young children were selected from a community-based household sample conducted in 2003 in Cato Manor, an urban settlement in Durban, and, KwaXimba, a rural area outside Durban. All subjects provided their informed consent.

Measurements

Demographic characteristics. Age and gender were collected by questionnaire at all 3 sites.

Laboratory testing. All blood samples were tested for antibodies to KSHV with 2 enzyme immunoassays (EIAs) and 1 indirect immunofluorescence assay (IFA), each previously described and performed by the same personnel at the Centers for Disease Control and Prevention (CDC). The 2 EIAs target antibodies to open reading frame (ORF) K8.1 and ORF 65, respectively, using synthetic peptides as antigen substrates.^{21,22} The IFA uses KSHV-infected BCBL-1 cells as antigen substrate, in which KSHV is induced to its replicative phase²³; specimens were evaluated at a dilution of 1:80. In the base-case algorithm, specimens reactive in any 2 tests or the IFA alone were classified as KSHV-antibody positive. Specimens that were nonreactive in all tests or reactive in only 1 EIA were categorized as KSHV-antibody negative; all other patterns were considered equivocal and excluded from analysis. This serologic algorithm has an estimated specificity of 97.5%, as evidenced by testing blood donors,²⁴ and sensitivity of 96.3% based on testing patients with KS.²⁵ HIV antibody testing performed in Uganda used an EIA (MUREX HIV-1.2.O; Murex Biotech, Kent, United Kingdom), in Zimbabwe used 2 EIAs (Uni-Gold, Trinity Biotech, Bray, Ireland; and Determine HIV-1/2, Abbot Diagnostics, Abbott Park, IL) followed by a western blot for positive confirmation, and in South African samples used 2 EIAs (Vironostika HIV-1 Microelisa, bioMérieux, Durham, N.C.; and Genetic Systems rLAV EIA, Bio-Rad, Redmond, WA).

Statistical analysis

The primary analysis compared HHV-8 seroprevalence, determined by the base-case serologic algorithm, in Uganda, Zimbabwe, and South Africa. Multivariable logistic regression was performed to adjust for confounding and to assess for potential effect modification by age. As sensitivity analyses, we varied the serologic definition of HHV-8 seropositivity to include various permutations of single assays evaluated individually or in combination. All analyses were carried out using Stata version 9.1 (College Station, TX).

Results

A total of 2,375 participants were examined, 482 in South Africa, 539 in Zimbabwe, and 1,354 in Uganda. Overall, 42 (1.8%) of individuals were classified as equivocal in HHV-8 antibody testing and not considered in the primary analysis, leaving 471 participants in South Africa, 532 in Zimbabwe, and 1,330 in Uganda. Of those in the primary analysis, the median age was 28 years [interquartile range (IQR) 21–37] in South Africa, 29 years (IQR 25–33) in Zimbabwe, and 22 years (IQR 19–28) in Uganda. In Uganda, 21.4% of participants were women, and by design, all participants in South Africa and Zimbabwe were women. Prevalence of HIV infection among South African participants was 38.2%, while, by design, 63.7% of Zimbabwean participants and no Ugandan participants were HIV-infected.

Crude HHV-8 seroprevalence was 37.3% in Uganda, 20.3% in Zimbabwe, and 17.6% in South Africa. This equates to, unadjusted for age, Ugandans having 2.33-fold greater odds [95% confidence interval (CI): 1.83–2.95, $p < 0.001$] of being HHV-8 infected than Zimbabweans and a 2.77-fold greater odds (95% CI: 2.13–3.60, $p < 0.001$) than South Africans. The high seroprevalence of HHV-8 infection in Uganda was apparent early in adulthood with 35.5% of participants infected before age 21 years (Table 1). The higher seroprevalence in Uganda did not substantially change throughout adulthood, only rising to 43.4% in those 41 years or older ($p = 0.30$ for age trend). In contrast, HHV-8 seroprevalence was substantially lower among the youngest participants in Zimbabwe and South Africa (13.7% and 10.8%, respectively, in those 25 years and younger) but then increased significantly with age in both Zimbabwe ($p = 0.023$) and South Africa ($p < 0.001$). Gender differences were evaluable only in Uganda, where infection was more common among men (38.6%) than among women (32.0%, $p = 0.085$). In both South Africa and Zimbabwe, HHV-8 infection appeared to be unrelated to HIV infection status ($p = 0.44$ and 0.40, respectively).

To formally compare HHV-8 seroprevalence across countries, we restricted the analysis to only the 1,287 women—all of the participants from South Africa and Zimbabwe and the 284 women from Uganda—given the slightly higher seroprevalence among men in Uganda. In a multivariable logistic regression analysis adjusting for age, there was no strong evidence for a difference in HHV-8 seroprevalence in Zimbabwe versus South Africa [odds ratio (OR) = 1.32, 95% CI: 0.95–1.83; $p = 0.095$]. In contrast, Ugandans had a 3.24-fold greater odds (95% CI: 2.19–4.81; $p < 0.001$) of being HHV-8 infected than South Africans and a 2.22-fold greater odds (95% CI: 1.49–3.30; $p < 0.001$) than Zimbabweans. Because there was no significant difference between South Africa and Zimbabwe, we combined participants from these 2 countries and found that Ugandans had a 2.72-fold greater odds (95% CI: 1.96–3.79; $p < 0.001$) of being HHV-8 infected than the combination of South Africans and Zimbabweans (Table 2). When we further restricted the analysis to just those

Table 1. Factors associated with HHV-8 infection in South Africa, Zimbabwe, and Uganda

| | No. of tested | No. of HHV-8 antibody positive (%) | Adjusted odds ratio (95% CI) ¹ | <i>p</i> value |
|----------------------|---------------|------------------------------------|---|----------------|
| South Africa | | | | |
| Age (years) | | | 1.04 (1.02–1.06) ² | <0.001 |
| 14–20 | 106 | 13 (12.3) | | |
| 21–25 | 89 | 8 (9.0) | | |
| 26–30 | 81 | 15 (18.5) | | |
| 31–35 | 62 | 11 (17.7) | | |
| 36–40 | 43 | 9 (20.9) | | |
| >40 | 90 | 27 (30.0) | | |
| HIV infection status | | | | |
| Uninfected | 291 | 53 (18.2) | Reference | |
| Infected | 180 | 30 (16.7) | 1.23 (0.73–2.09) | 0.44 |
| Zimbabwe | | | | |
| Age (years) | | | 1.04 (1.004–1.08) ² | 0.026 |
| 14–20 | 20 | 1 (5.0) | | |
| 21–25 | 133 | 20 (15.0) | | |
| 26–30 | 172 | 37 (21.5) | | |
| 31–35 | 116 | 28 (24.1) | | |
| 36–40 | 52 | 12 (23.1) | | |
| >40 | 39 | 10 (25.6) | | |
| HIV infection status | | | | |
| Uninfected | 193 | 40 (20.7) | Reference | |
| Infected | 339 | 68 (20.1) | 0.82 (0.51–1.30) | 0.40 |
| Uganda | | | | |
| Age (years) | | | 1.01 (0.99–1.02) ² | 0.27 |
| 14–20 | 513 | 182 (35.5) | | |
| 21–25 | 367 | 132 (38.7) | | |
| 26–30 | 195 | 73 (37.4) | | |
| 31–35 | 118 | 41 (34.8) | | |
| 36–40 | 54 | 21 (38.9) | | |
| >40 | 83 | 36 (43.4) | | |
| Sex | | | | |
| Female | 284 | 91 (32.0) | Reference | |
| Male | 1046 | 404 (38.6) | 1.29 (0.97–1.72) | 0.081 |

¹Adjusted for age and other factor, either HIV infection (South Africa and Zimbabwe) or gender (Uganda). ²Assessed as a continuous variable per 1 year increase in age.

participants who were HIV-uninfected, the inferences were unchanged. Among HIV-uninfected individuals, Ugandans had a 3.47-fold greater odds (95% CI: 2.14–5.62; $p < 0.001$) of being HHV-8 infected than South Africans and a 2.06-fold greater odds (95% CI: 1.27–3.33; $p = 0.004$) than Zimbabweans. Finally, we used the multivariable model to determine whether the relationship between age and HHV-8 infection differed by country, as suggested by the within-in country analyses above. We found no evidence that the effect of age on HHV-8 seroprevalence differed in Zimbabwe *vs.* South Africa ($p = 0.89$ for interaction term). Although we did find

suggestive evidence that the effect of age differed in Uganda versus the combined population of South Africa and Zimbabwe, we cannot exclude that this is a chance occurrence ($p = 0.13$ for interaction term).

To determine if our finding of a significant difference in HHV-8 seroprevalence in Uganda versus either Zimbabwe or South Africa was dependent on the base-case algorithm, we used to determine seropositivity, we also evaluated a number of other algorithms by which to interpret our antibody assays. We found that the principal inferences—no difference between South Africa and Zimbabwe but substantially higher

Table 2. Comparison of HHV-8 seroprevalence in women in South Africa, Zimbabwe, and Uganda using different antibody interpretation algorithms

| Algorithm | Age-adjusted odds ratios (95% CI) | | | |
|-------------------------------------|-----------------------------------|-------------------------|---------------------|---|
| | Zimbabwe vs. South Africa | Uganda vs. South Africa | Uganda vs. Zimbabwe | Uganda vs. pooling of South Africa and Zimbabwe |
| Base case ¹ | 1.32 (0.95–1.83) | 3.24 (2.19–4.81) | 2.22 (1.49–3.30) | 2.72 (1.96–3.79) |
| Es defined reactive ² | 1.25 (0.91–1.71) | 2.98 (2.04–4.36) | 2.15 (1.46–3.17) | 2.60 (1.88–3.59) |
| Es defined nonreactive ³ | 1.34 (0.97–1.85) | 3.25 (2.19–4.81) | 2.20 (1.48–3.25) | 2.71 (1.95–3.76) |
| IFA alone ⁴ | 1.22 (0.88–1.68) | 2.93 (1.98–4.33) | 2.13 (1.43–3.18) | 2.60 (1.87–3.63) |
| K8.1 EIA alone ⁴ | 0.92 (0.65–1.31) | 2.43 (1.61–3.68) | 2.58 (1.65–4.03) | 2.55 (1.78–3.67) |
| ORF65 EIA alone ⁴ | 1.23 (0.76–1.99) | 5.44 (3.16–9.37) | 3.43 (2.02–5.80) | 4.76 (3.07–7.37) |
| Both EIA reactive ⁵ | 1.07 (0.62–1.85) | 3.44 (1.82–6.50) | 2.42 (1.27–4.60) | 3.30 (1.94–5.61) |
| One EIA reactive ⁶ | 0.99 (0.71–1.38) | 3.24 (2.20–4.79) | 3.20 (2.11–4.84) | 3.26 (2.33–4.55) |
| All 3 assays reactive ⁷ | 1.09 (0.63–1.89) | 2.79 (1.46–5.33) | 1.92 (1.00–3.70) | 2.64 (1.53–4.55) |
| Any assay reactive ⁸ | 1.19 (0.88–1.60) | 3.13 (2.19–4.48) | 2.63 (1.81–3.82) | 2.82 (2.08–3.84) |

¹Seropositivity defined as either reactive on IFA or on both EIA assays; equivocal results discarded. ²As for base case, with equivocal results coded as reactive. ³As for base case, with equivocal results coded as nonreactive. ⁴Seropositivity defined as reactive on the assay listed, without regard to other assays; equivocal results discarded in case of IFA. ⁵Seropositivity defined as reactive on both the K8.1 EIA and the ORF65 EIA. ⁶Seropositivity defined as reactive on either the K8.1 EIA or the ORF65 EIA. ⁷Seropositivity defined as reactive on IFA, K8.1 EIA, and ORF65 EIA; equivocal results discarded. ⁸Seropositivity defined as reactive on either IFA, K8.1 EIA or ORF65 EIA; equivocal results discarded.

seroprevalence in Uganda than in either Zimbabwe or South Africa—were unchanged regardless of the algorithm used (Table 2). In the most strict algorithm, where reactivity in all 3 antibody assays is required to determine overall seropositivity, the unadjusted country-specific estimates of HHV-8 seroprevalence were predictably the lowest (South Africa: 6.8%; Zimbabwe: 5.6%; Uganda: 8.2%), but nonetheless after age adjustment, Ugandans again had a 2.64-fold greater odds of being HHV-8 infected than the South Africans and Zimbabweans. The least strict algorithm, where reactivity in any 1 of the 3 assays was deemed seropositive, produced the highest country-specific prevalence estimates (South Africa: 23.1%; Zimbabwe: 24.7%; Uganda: 47.1%) with Ugandans again having significantly greater seroprevalence. We also evaluated algorithms where we considered all the equivocal results as either reactive or nonreactive and again the principal findings were unchanged.

Discussion

The “KS Belt” in equatorial Africa has long been investigated to determine the reasons why KS is so common in the region. Prior work addressing whether underlying HHV-8 infection is more common in this region as compared with other parts of Africa has produced inconsistent results and has often been interpreted as the entire continent being monolithically HHV-8 infected.¹⁸ We believe that the inability of prior studies to establish regional differences in HHV-8 seroprevalence in Africa is partially explained by differences in the antibody assays used by these studies. Using the same assay, performed in the same laboratory by the same personnel, we here show the odds of HHV-8 seropositivity to be as much as 3-fold higher in the “KS Belt” (Uganda) than in

areas of Africa outside the Belt (Zimbabwe and South Africa).

Although the available literature on HHV-8 seroprevalence in Africa nominally gives the impression of no substantial differences between regions, a close dissection of studies that, in part, use identical or very similar antibody assays substantiates our conclusions. Specifically, 2 studies of adults from South Africa^{26,27} and 1 from Uganda²⁸ each include an IFA targeting antibodies to the latency-associated nuclear antigen (LANA) of HHV-8. When comparing only serologic results from the LANA-specific IFA, HHV-8 seroprevalence in Uganda was found to be 48%²⁸ compared with 14%²⁶ and 16%²⁷ in South Africa. These estimates are very similar to our findings. Furthermore, separate reports from Tanzania (within the “KS Belt”) and Nigeria (outside the Belt), both using the same K8.1-based EIA found HHV-8 seroprevalence to be substantially higher in Tanzania.^{29,30} In the only other report we are aware of that used the same antibody assay in 2 different populations representing within and outside the “KS Belt”, there was higher HHV-8 seroprevalence in the Democratic Republic of Congo compared with Botswana when using an ORF65 EIA but not when using the LANA IFA or K8.1 EIA.³¹ The very high seroprevalence of HHV-8 found in this report in Botswana (>75% by K8.1 EIA or LANA IFA) has not been replicated in other parts of the country and may possibly reflect the isolated nature of the population studied.

Not only is the overall level of HHV-8 seropositivity is higher in Uganda than in Zimbabwe and South Africa but the patterns of HHV-8 infection for these 3 countries also appear different. In Uganda, high levels of infection are already apparent by early adulthood with only minimal

increases thereafter, suggesting scant primary infection during adulthood. In South Africa and Zimbabwe, in contrast, prevalence was low in the youngest adults and increased steadily throughout adulthood, suggesting ongoing transmission. This finding that most of the HHV-8 transmission in Uganda has occurred prior to adulthood, whereas most transmission in South Africa occurs after adulthood is supported by recent direct work in children, which found that HHV-8 seroprevalence among Ugandan children was much higher than among South African children.³²

In addition to geographic differences in HHV-8 seroprevalence, we found evidence of differences by gender in the 1 region where this was evaluable. In Uganda, men had a trend toward higher seroprevalence than women, which is consistent with some,²⁹ but not all,²⁸ prior reports from this region. This is notable because endemic KS, the form of KS that existed in Africa prior to the spread of HIV, was about 10 times more common among males than females.¹ However, the magnitude of the difference we found in HHV-8 seroprevalence between men and women, if true, would explain only a small portion of the difference observed for KS. We did not find any difference in HHV-8 seroprevalence by HIV infection status in Zimbabwe or South Africa, which is compatible with some reports,^{26,28,33–37} but not others.^{30,38–41} This inconsistency in the relationship between HIV and HHV-8 in Africa across studies has defied explanation to date.

A potential limitation of this study is that the participants were not sampled in identical ways in the 3 countries. In particular, the Ugandan sample is from blood donors who were prescreened for 3 sexually transmitted diseases. If HHV-8 is

sexually transmitted in Africa among adults, as been suggested in some reports,^{30,42,43} then our analysis underestimates the true prevalence of HHV-8 infection among Ugandans and underestimates the difference between Uganda and Zimbabwe and South Africa. Similarly, it could be argued that the inclusion in South Africa of only women who were caregivers to a young child might have selected for those who were naturally exposed to more infections. However, if this were the case, then our sample overestimates the true prevalence of HHV-8 in South Africa and our analysis would again underestimate the difference between Uganda and South Africa. A more significant limitation is that we could only formally compare HHV-8 seroprevalence across countries in women and it is not known if these same results will hold true for men.

In conclusion, the use of a standardized approach to antibody testing, some of the strongest evidence that HHV-8 seroprevalence is substantially higher in the “KS Belt” than regions south of this Belt. We hypothesize that this difference in underlying HHV-8 seroprevalence in part explains the higher rate of endemic KS within the Belt. The question remains as to why these regional differences in HHV-8 seroprevalence exist. Possible explanations include differences in viral strain infectiousness, host susceptibility to infection, environmental factors, and behavioral practices. That HHV-8 infection, unlike many other herpesvirus infections, does not appear to be ubiquitous in any of the African regions studied should provide opportunities to study HHV-8 transmission dynamics that may ultimately be exploitable for the development of effective prevention strategies.

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