On September 20, 2022, Uganda’s Ministry of Health declared an Ebola disease outbreak after a case caused by Sudan virus (SUDV) was confirmed in a village in Mubende District (1). Suspicious deaths in the same district had occurred earlier in the month. Investigations conducted by the National Rapid Response Team allowed the identification of probable SUDV cases dating back to mid-August 2022 (2,3). As of November 2, 2022, a total of 149 cases (131 PCR-confirmed and 18 probable) were reported in the country; most cases occurred in the districts of Mubende (63 confirmed, 17 probable), Kassanda (42 confirmed, 1 probable), and Kampala (18 confirmed) (2) (Figure 1, panels A–C). On January 11, 2023, the outbreak was declared over with a total of 164 cases (142 confirmed, 22 probable) (4).

The Study

We provide 2 estimates of the serial interval distribution (the time elapsed between the symptom onset in an index case-patient and in their secondary case-patients) by using observed serial intervals in infector–infectee pairs as identified during contact-tracing operations conducted in 2 SUDV outbreaks in Uganda, during 2000–2001 (24 pairs) (5) and the 2022 outbreak (12 pairs) (6). We fitted 3 families of distributions (Weibull, Gamma, and log-normal) with a possible offset (5,6). We obtained the best fit for the serial interval distribution for both datasets with a Weibull distribution. We estimated the mean serial interval to be 12.0 days (95% CI 10.0–14.2 days) by using the 2000–2001 outbreak data and 11.7 days (95% CI 8.2–15.8 days) by using the 2022 outbreak data.

We then used estimates of the serial interval as a proxy of the generation time to compute the basic ($R_0$) and net ($R_t$) reproduction numbers. We defined $R_0$ as the average number of secondary infections generated by an infectious person in a fully susceptible population. If $R_0 < 1$, transmission is expected to fade out, whereas if $R_0 > 1$, the epidemic has the potential to continue; the larger $R_0$, the more difficult it is to control the epidemic. We defined $R_t$ as the average number of secondary cases per infectious person at time $t$; $R_t$ is key to monitor the effectiveness of interventions throughout the epidemic. In the main analysis, we computed $R_t$ and $R_0$ by using a method based on the renewal equation in the formulation by Cori et al. (7). In additional analyses, we used the assumption of exponential (8) or subexponential (9) growth of the cumulative case incidence curve to compute $R_0$. We also provided an alternative estimate of $R_t$ obtained by applying a recently proposed approach (10) that has been suggested to perform better with low case counts (Appendix, https://wwwnc.cdc.gov/EID/article/29/7/22-1718-App1.pdf).

$R_0$ of the 2022 outbreak, as estimated from the epidemic curve of Mubende District, was 2.7 (95% CI 1.9–3.5) based on the serial interval distribution from the 2000–2001 outbreak. $R_0$ was 2.4 (95% CI 1.7–3.3) based on the serial interval distribution from the 2022 outbreak.
We estimated $R_t$ in the 3 districts and according to the 2 estimated serial interval distributions (Figure 1, panels D–F). For convenience and given their similar values, numbers reported hereafter refer to the 2000–2001 serial interval; corresponding numbers for the 2022 serial interval are reported separately (Appendix). In Mubende District, $R_t$ reached a peak during September 21–23, 2022, with an estimated value for $R_t$ that was close to $R_0$ (mean 2.4 [95% CI 1.5–3.5]). $R_t$ fell rapidly below the epidemic threshold during September 28–October 15 (mean 0.71 [95% CI 0.50–0.91]), possibly because of control interventions and population behavior changes after awareness of the outbreak had increased. In the second half of October, $R_t$ increased again, reaching a peak of 1.34 (95% CI 0.78–2.13) in the week October 18–24. In the districts of Kassanda and Kampala, $R_t$ increased rapidly in the second half of October. In Kassanda, $R_t$ reached a peak of 3.5 (95% CI 2.5–4.9) during October 20–24. In Kampala, the peak $R_t$ value was 2.0 (95% CI 1.3–3.2) during October 18–22.

Estimates of $R_t$ at the national level (Figure 2) are characterized by 2 peaks in September, which were 2000–2001 outbreak (5) (red) and from the 2022 outbreak (6) (blue). Shaded areas represent 95% CIs of estimates. We assumed that the first case of the epidemic curve in each district was imported and that all the others were locally transmitted. SUDV, Sudan virus.
driven by SUDV transmission in Mubende. A third marked peak that occurred in the second half of October was sustained mainly by increasing transmission in the districts of Kassanda and Kampala, as well as by a resurgence in Mubende.

Conclusions
We estimated the distribution of the serial intervals for SUDV by using 2 different datasets from the 2000–2001 outbreak and from the ongoing outbreak in Uganda, finding similar distributions and an average serial interval of ≈12 days. On the basis of those estimates and publicly available data on the epidemic curve made available by the Ugandan Ministry of Health (2), we found the \( R_0 \) in Mubende District, the district first and most affected by the current outbreak, was ≈2.4–2.7, although with broad uncertainty (95% CI 1.7–3.5). Those estimates are in line with previous estimates for SUDV, which ranged from 1.3 to 4.1 (11), and with estimates obtained using alternative methods (8,9) (Appendix). After a temporary containment of the outbreak from the end of September until mid-October 2022, with \( R_t \) hovering around 0.7, the third week of October marked a resurgence of transmissibility in Mubende (\( R_t \approx 1.3 \)) and the emergence of new outbreaks in the Kassanda (\( R_t \approx 3.5 \)) and Kampala (\( R_t \approx 2.0 \)) districts. The \( R_0 \) associated with the national aggregation shows the same temporal features but suggests even higher numbers for the \( R_t \) in the fourth week of October (mean \( R_t =4 \)), demonstrating the important role played by spatial heterogeneity of transmission in the 2022 outbreak.

Our estimates should be interpreted with caution, considering the following limitations. Estimates of the serial interval distributions are based on small numbers of infected–infectee pairs. Reproduction numbers have a broad uncertainty because of limited case numbers and may be substantially affected by superspreading events, biasing estimate upward with respect to the average transmissibility in the general population. However, the proposed estimates are in line with those obtained using an alternative method that was suggested to be more robust for low case counts (10) (Appendix). Moreover, a potential increase in reporting rates of confirmed cases after the discovery of the first cases in each district may inflate the estimate of the reproduction numbers.

Given the geographic expansion of the outbreak, which included urban settings, and the absence of therapeutics and licensed vaccines to treat and prevent SUDV, by the end of October 2022, the World Health Organization assessed the risk for infection at the national level to be very high (12). However, the rapid deployment of interventions (including contact tracing, isolation of case-patients, and informational activities to promote community engagement) was sufficient to contain the outbreak, which was declared over on January 11, 2023. Our analysis provides quantitative information on the evolution of SUDV transmissibility in the different districts of Uganda during the 2022 outbreak. Estimates provided for the serial interval may be instrumental in planning control interventions in possible future outbreaks of SUDV.

This research was supported by European Union funding within the NextGeneration EU-MUR PNRR Extended Partnership initiative on Emerging Infectious Diseases (project no. PE00000007, INF-ACT). The funders had no role in study design, data collection and analysis, decision to publish, or preparation of the manuscript.

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