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# **Evaluation of Respondent-Driven Sampling**

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## Abstract

**Background**—Respondent-driven sampling is a novel variant of link-tracing sampling for estimating the characteristics of hard-to-reach groups, such as HIV prevalence in sex-workers. Despite its use by leading health organizations, the performance of this method in realistic situations is still largely unknown. We evaluated respondent-driven sampling by comparing estimates from a respondent-driven sampling survey with total-population data.

**Methods**—Total-population data on age, tribe, religion, socioeconomic status, sexual activity and HIV status were available on a population of 2402 male household-heads from an open cohort in rural Uganda. A respondent-driven sampling (RDS) survey was carried out in this population, employing current methods of sampling (RDS sample) and statistical inference (RDS estimates). Analyses were carried out for the full RDS sample and then repeated for the first 250 recruits (small sample).

**Results**—We recruited 927 household-heads. Full and small RDS samples were largely representative of the total population, but both samples under-represented men who were younger, of higher socioeconomic status, and with unknown sexual activity and HIV status. Respondent-driven-sampling statistical-inference methods failed to reduce these biases. Only 31%-37% (depending on method and sample size) of RDS estimates were closer to the true population proportions than the RDS sample proportions. Only 50%-74% of respondent-driven-sampling bootstrap 95% confidence intervals included the population proportion.

**Conclusions**—Respondent-driven sampling produced a generally representative sample of this well-connected non-hidden population. However, current respondent-driven-sampling inference methods failed to reduce bias when it occurred. Whether the data required to remove bias and

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measure precision can be collected in a respondent-driven sampling survey is unresolved. Respondent-driven sampling should be regarded as a (potentially superior) form of conveniencesampling method, and caution is required when interpreting findings based on the sampling method.

Hidden or hard-to-reach population subgroups are often key to the maintenance of infectious diseases in human populations.<sup>1</sup> However, it is often difficult to investigate the factors that drive transmission in these groups by using representative samples, because there may not be an adequate sampling frame or the groups may be associated with illicit activity or subject to stigma. Researchers have therefore typically resorted to various types of convenience sampling to gather data on hidden populations.<sup>2</sup> While convenience sampling has its advantages, this approach is unable to generate unbiased population-based estimates of infection prevalence and risk factors.

In an attempt to address these limitations, respondent-driven sampling (a variant of a linktracing design), was proposed in 1997.<sup>3</sup> With this approach, a small number of "seed" respondents are selected by convenience sampling or other methods. Then, these initial recruits are given coupons (typically, 3 coupons) to recruit others from the target population, who in turn become recruiters. Recruits are given an incentive (usually money) for taking part in the survey and then for recruiting others. This process continues in recruitment "waves" until a pre-determined sample size is reached, or until the distribution of participant characteristics (such as the proportion infected) becomes similar between waves (called "reaching equilibrium" in respondent-driven-sampling terminology). Estimation methods are then applied to account for the non-random sample selection in an attempt to generate unbiased estimates for the target population. For this approach to be successful, the target population must be socially well-connected.

Two main estimation methods are generally used. The RDS-1 estimator, currently in wide use, can be implemented with the standard respondent-driven-sampling analysis software.<sup>3-6</sup> RDS-1 accounts for patterns of recruitment between subgroups and the average number of other members of the target group who the recruiters know (the "network size") in each subgroup.<sup>5,7</sup> RDS-2 is a more recently developed estimator that relates respondent-driven-sampling estimation to widely used survey estimation through the use of a generalized Horvitz-Thompson estimator.<sup>8</sup> RDS-2 accounts for network size only.<sup>8</sup> Initial theoretical analysis has asserted that the RDS-2 estimator is asymptotically unbiased as long as six key assumptions are met, including that respondents accurately report the size of their "network" (the number of other members of the target group they know), that respondents randomly recruit from their network, and that respondents have reciprocal relationships with members of the target population.<sup>8</sup>

A recent study simulating respondent-driven sampling and using empirical network data found that the variance of respondent-driven sampling estimates can be much higher than commonly assumed.<sup>9</sup> Nevertheless, respondent-driven sampling has rapidly become a popular and widely used survey method. Outside of the US, more than 120 respondent-driven-sampling studies, involving more than 30,000 participants, have been published,<sup>10</sup> and respondent-driven sampling is currently being employed to provide data for public-health decision-making by major funding bodies such as the US Centers for Disease Control and Prevention.

Despite its popularity, it is not known whether respondent-driven sampling can generate unbiased estimates. This is primarily because the robust evaluation of respondent-driven sampling is methodologically challenging. By definition, gold-standard representative or total-population data are generally unavailable or are of poor quality for hidden/stigmatized

groups. Other methods of evaluation that have been attempted include in-silico studies,<sup>4,8,11,12</sup> comparisons of respondent-driven-sampling data with data from other convenience samples,<sup>13-19</sup> comparisons of serial cross-sectional respondent-driven sampling estimates on the same population over time,<sup>20</sup> and comparisons of Internet-collected respondent-driven sampling data with a population that has known characteristics.<sup>21,22</sup> Although all these studies have provided valuable information on respondent-driven sampling, none provides a robust assessment of whether respondent-driven-sampling could produce unbiased estimates — either because the required gold-standard comparison population was unavailable or because an Internet-based respondent-driven-sampling data collection method was used, whereas the vast majority of respondent-driven-sampling studies employ face-to-face data collection.<sup>10</sup>

We evaluate respondent-driven sampling by comparing field-collected respondent-drivensampling data with total-population data on the same population. Although the representative or total-population data required for such a comparison are generally unavailable, we dealt with this problem by evaluating respondent-driven sampling in a nonhidden/non-stigmatized population for which high-quality total-population data were available. This also allowed us to perform a range of analyses that are not possible in typical respondent-driven-sampling studies.

#### Methods

In order to evaluate whether respondent-driven sampling can generate representative data, we compared estimates from a respondent-driven-sampling survey of a rural Ugandan population with total-population data. The data used to define the target population were available from an ongoing general population cohort of 25 villages in rural Uganda covering an area of approximately 38km<sup>23,24</sup> (Figure 1). Each year, households in the study villages are mapped and, after obtaining consent, a total-population household census and an individual questionnaire and HIV-1 serosurvey are administered. The target population consisted of 2402 men who were recorded as a male head of a household within these villages between February 2009 and January 2010 (Figure 1). The characteristics of the target population are shown in the Table (population proportion).

To maximize the generalizability of our results, we employed where possible currently used respondent-driven-sampling data-collection methods.<sup>10</sup> Ten "seeds" (of varying village, age and tribe) were selected by convenience from the target population. Figure 1 shows their locations and eTable 1 (http://links.lww.com) summarizes their characteristics. Seeds and subsequent recruits were given three coupons to recruit other men into the study. The rate of early recruitment was high, and the number of people arriving each day for interviews became too large to be manageable. Because of this, between day nine and day 32 the probability of each recruit being offered three coupons was halved from 100% to 50%; other recruits received none. As incentives for participation and recruitment, seeds and recruits were offered soap, salt, or school books to the value of approximately \$1US. One incentive was offered for completing the first interview and another for each person successfully recruited.

Respondent-driven-sampling estimation requires information on how many other household heads each participant could potentially recruit. The primary network-size definition was created to be comparable with other respondent-driven-sampling studies<sup>25,26</sup> and was used here unless otherwise stated. Recruits were first asked the core question "How many men do you know who (i) were head of a household in the last 12 months in any of the Medical Research Council villages, and (ii) you know them and they know you, and (iii) you have

seen them in the past week.". More detailed network data were also collected (eAppendix [methods], http://links.lww.com).

Pre-processing of data was performed using Stata v11 (StataCorp, Texas).<sup>27</sup> Networks and "trees" were generated using scripts written in Stata and R v2.12.0 (R Foundation, Vienna)<sup>28</sup> and visualized using GraphViz (AT&T Research, New Jersey).<sup>29</sup> To maximize the comparability of our methods with those used in a typical respondent-driven-sampling study, we analyzed the dataset following current respondent-driven-sampling definitions and the statistical inference methods employed in RDSAT v6.0.1, the custom-written software package for the analysis of respondent-driven-sampling studies.<sup>6</sup> (ie the RDS-1 point estimator<sup>3-5</sup> and the bootstrap 95% confidence interval [CI] estimator<sup>11</sup>). We also analyzed the dataset using the more recently developed point estimator RDS-2 and the same bootstrap 95% CI estimator,<sup>11</sup> employing R. Simple respondent-driven-sampling sample proportions and respondent-driven-sampling estimates were calculated for two sample sizes. The first was the "Full" sample (n=927 including the 10 seeds). The second was a "Small" sample consisting of the first 250 recruits (including the 10 seeds); this was chosen to be more typical of the sample sizes used in respondent-driven sampling studies.<sup>10</sup>

Root mean squared errors were calculated for the differences between the population proportions and the full and small sample proportions, and for the differences between the population proportions and the RDS-1 and RDS-2 estimates, for each variable and in total. For comparison with the RDS-1 and RDS-2 estimates, we used the true population proportions to calculate recruitment probabilities for the target population using predictions from a logistic regression model<sup>30</sup> as weights. The variables shown in the Table were included in the model if they were significant at the 95% CI level.

Sensitivity analyses were used to assess the robustness of our results to various network size definitions, potential network-size bias and respondent-driven sampling sample size.

To compare network size of the whole target population with the respondent-drivensampling recruits, 300 men in the target population who had not been recruited in the respondent-driven-sampling study were selected using simple random sampling to be interviewed using the first respondent-driven-sampling questionnaire. Mean network size of the whole target population was estimated as the weighted average of the mean network size of respondent-driven-sampling recruits and the mean network size of a simple random sample of eligible non-recruits. T-tests were used to test for differences between means. To help understand the quantitative study findings, 54 members of the population in the study villages and Medical Research Council staff were selected using random or purposive sampling for qualitative interview. Full details are shown in the eAppendix (Methods, http://links.lww.com).

#### Results

#### Recruitment

The dynamics of the respondent-driven-sampling survey recruitment are shown in Figure 2, and the recruitment networks from each seed are shown in Figure 3. A total of 1141 people (including the 10 seeds) were assessed for eligibility over a period of 54 days (8 March – 30 April 2010). No new coupons were distributed after day 47. 196 men attended but were ineligible, 16 were eligible but had already been recruited, 2 were eligible but did not give consent, and 927 were eligible, consented and were recruited. A video illustrating recruitment in space and time is provided online (http://links.lww.com). A roughly linear recruitment rate was achieved in the respondent-driven sampling survey (Figure 2A), due, in part, to changes in the probability of each recruit being offered coupons during the survey.

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All 10 seeds recruited people into the study, with one seed recruiting one person, four recruiting two people, and five recruiting three people. The total number of recruits originating from each seed ranged from 8 to 241 (1% to 26% of the full sample) (Figure 2B). 77% of the total recruitment was from four seeds. Full details of the seeds and recruitment by seed are given in eTable 1 (http://links.lww.com). The number of waves ranged from 3 to 16 for the full sample and from 2 to 6 for the small sample The highest recruitment occurred in wave five (12% of all recruits, excluding seeds) and 57% of recruits of seeds) were interviewed within 7 days of their recruiter's interview (Figure 2D).

Overall, 75% of recruits (including seeds) (n=684) were offered coupons to recruit others, and of these 90% (n=612) accepted (called "recruiters"). 66% of recruiters (n=401) returned to take part in a second interview and to collect their secondary incentives. A similar proportion of recruiters (including seeds) recruited zero, one, two or three recruits (Figure 2E, left bar). Recruits who returned to collect secondary incentives were more likely to have recruited (Figure 2E, middle and right bar). The proportion of the recruit's network already recruited at the time of their interview increased rapidly during the survey (Figure 2F; includes seeds). The average number of recruits per recruiter (including seeds) decreased from 2.6 in the first week of the study to 0.6 in the last week that coupons were given out. Only 30% of recruits had been named as a contact by their recruiter (and identified) at their recruiter's first interview,

In the simple random sample survey, 55% (164/300) of men selected were interviewed (4 - 28 May 2010; eAppendix [Simple random sample survey], http://links.lww.com). In the qualitative survey, 98% (53/54) of people selected were interviewed (16 June - 19 Oct 2010; eAppendix [Qualitative survey], http://links.lww.com).

The target population was well-connected. Data from the respondent-driven sampling and simple random sampling surveys showed that at least 73% were linked in a single network (eAppendix [Methods], http://links.lww.com). The distribution of the reported network size of respondent-driven sampling recruits, based on the primary definition of network size (NS-1), was approximately normal but with a slight positive skew, and shows likely overreporting of multiples of 5 (eFigure 1, http://links.lww.com; excluding seeds). The distributions of the other network size measures (as defined in the eAppendix [Methods], http://links.lww.com) were very similar, with the exception of definition NS-5, which showed a smaller proportion of larger network sizes because it was a subset of NS-4 (eFigure 2, http://links.lww.com; including seeds). Pearson correlations between different network size definitions reported by respondent-driven sampling recruits varied between 0.96 (NS-1 vs NS-2) and 0.75 (NS-1 vs. NS-5) (eTable 2, http://links.lww.com; including seeds). The mean network size (NS-1) of respondent-driven sampling recruits (including seeds) was higher than that of the whole target population (12.1 vs 9.2, p<0.001) (eFigure 3, http://links.lww.com). The number of times members of the target population were reported to be in the network of recruits ranged between 0 and 42 (eFigure 4, http://links.lww.com).

There was high within-group recruitment (homophily) by religion, tribe and village and in the highest socioeconomic status groups, but not by age, sexual activity or HIV status, or within the other socioeconomic-status groups, (eTable 3 and Table 4, http://links.lww.com). There was no evidence of low within-group recruitment for any characteristic, i.e. preferentially recruiting men who differed from themselves. Comparing actual recruitment proportions with expected recruitment proportions calculated from individual-level network data, there was evidence of non-random recruitment by age, tribe, socioeconomic status, village and sexual activity (eAppendix [Supporting results 'Recruitment pattern' section and eTables 5 and 6], http://links.lww.com).

The other RDS-2 estimator assumptions<sup>8</sup> were not met. In common with current practice for all respondent-driven-sampling studies, respondents were not limited to recruiting only one other person, and recruited persons were ineligible for re-recruitment. It is likely that only a low proportion of the relationships between members of the target population were reciprocated and/or the population may not have accurately reported their network size, as only 30% of recruits were mentioned by their recruiter during the recruiter's first interview.

#### Comparison with target population data

The Table shows the comparison between the population proportions, sample proportions, and RDS-1 and RDS-2 estimates, for the full and small sample. The sample proportions were often similar to population proportions, with the following exceptions. In both samples, younger men (<30 years) were underrepresented and older men ( $\geq$ 40 years) were overrepresented. In the small sample, Catholics were overrepresented. In both samples men in the highest socioeconomic group were underrepresented and men in the lowest socioeconomic group were overrepresented. The proportions of men with unknown numbers of sexual partners or unknown HIV status were underrepresented in both samples. It is unlikely that the differences between the population and sample proportions occurred by chance (p $\leq$ 0.0001 for all except p=0.04 for the highest socioeconomic status group using the small sample).

Respondent-driven-sampling inference methods generally failed to reduce bias where it occurred. Adjustment resulted in an improved estimate of the population proportion in only 37% (19/52) of comparisons using RDS-1 and 33% (15/52) using RDS-2 for the full sample, and 31% (8/26) using RDS-1 and 37% (18/49) using RDS-2 for the small sample. Based on these estimates, the 95% bootstrap confidence intervals included the target population proportion in 69% (36/52) of comparisons using RDS-1 and 50% (13/26) using RDS-2 for the full sample, and 69% (18/26) using RDS-1 and 74% using RDS-2 for the small sample.

The root mean squared error for the difference between the population proportions and the sample proportions was 6% for the full sample. The root mean squared error for the difference between the population proportions and the respondent-driven sampling estimates for the full sample were 7% for both RDS-1 and RDS-2 (eTable 7, http://links.lww.com). Root mean squared errors were slightly larger for the small sample.

In general, if the respondent-driven sampling adjustments did not improve the estimates, the adjustments were small and did not add substantial bias. The exception to this was the variable village. Due to the large number of subgroups for "village," however, the sample size was not sufficiently large to reliably estimate the parameters used to make RDS-1 adjustments.

In comparison, using the predictions from the logistic regression models as recruitment probability weights, adjustment improved the estimate of the target population proportion for 88% (46/52) of the full-sample estimates, and for 57% (28/49) of the small-sample estimates (eTable 6, http://links.lww.com), showing that recruitment was associated with characteristics other than network size.

For specific cases in which the sample estimates of population proportions were biased, current respondent-driven-sampling inference methods generally failed to reduce bias. For age group, using either the RDS-1 or the RDS-2 estimator, only 2 out of 5 estimates were closer to the population proportion when applied to the full sample, and only 1 out of 4 when applied to the small sample. Neither RDS-1 nor RDS-2 improved the over-representation of Catholics in the small sample, the over-representation of the lowest socioeconomic group in the full sample, the under-representation of the highest

socioeconomic group in either sample, or the underrepresentation of men with unknown number of sexual partners in either sample. Applying RDS-2 to the full sample very slightly reduced the under-representation of men with unknown HIV status. Applying RDS-2 to the small sample or RDS-1 to either sample slightly increased the under-representation of men with unknown HIV status.

Respondent-driven-sampling inference methods failed to reduce bias because groups tended to be under- or over-recruited by all groups, rather than being under-recruited by some groups and over-recruited by other groups (limiting the usefulness of RDS-1 to improve estimates), and because under-represented groups tended not to have markedly smaller network sizes (limiting the ability of RDS-1 and RDS-2). For example, men aged 50+ years were over-recruited by all age groups, and network sizes in all age groups were relatively similar (eTable 3, http://links.lww.com). Therefore neither RDS-1 nor RDS-2 improved the estimates.

Qualitative data suggested possible explanations for these findings. Recruiters did not consider younger unmarried men to be household heads, in contrast with the definition used in the ongoing general population cohort ("...they were being left out because some of the older men didn't take them as household heads because they didn't have any wives" [45-year-old respondent-driven-sampling recruit]). The respondent-driven-sampling incentives were likely to be a greater incentive to men in lower socioeconomic groups ("...the token might look small to some people and big to others." [42-year-old female community member]). The under-recruitment of men with unknown number of sexual partners or unknown HIV status was likely, at least in part, to be because men who had refused to participate in the ongoing general population cohort in the past were also less likely to participate in the respondent-driven-sampling study.

There was very little difference in the performance of the respondent-driven-sampling estimators when different network size definitions were used (eAppendix [Results], http://links.lww.com). There was no evidence that collecting detailed network size data reduced the performance of the respondent-driven sampling estimators (eAppendix [Results], http://links.lww.com).

### Discussion

In our study, recruitment by respondent-driven sampling produced a largely representative sample of the target population for most variables. The exceptions were an underrepresentation of men who were younger, men of higher socioeconomic status, men of unknown HIV status, and men with unknown number of sexual partners in both samples, and an overrepresentation of Catholics in the small sample. The most plausible reason for sample bias by age is that younger men were not considered to be heads of household. The most plausible reason for sample bias by socioeconomic status is that men of higher status were less attracted by the incentives. Men who refused to participate in the ongoing general population cohort were probably more likely to also have refused to participate in respondent-driven sampling and that was probably at least partially responsible for the under-recruitment of men of unknown HIV status or with an unknown number of sexual partners. These biases may increase the design effect of respondent-driven sampling. Neither of the respondent-driven-sampling inference methods was designed to correct for these sources of bias.

The bias in recruitment by socioeconomic status is likely to be generalizable to most, if not all, respondent-driven-sampling studies because different sub-groups of the target population are likely to be differentially motivated by whatever incentives are offered. An

"unknown" category for HIV status and other variables will not exist in most other respondent-driven-sampling studies. The differential recruitment of persons in the population by willingness to participate in surveys is nevertheless likely to be a generalizable finding, but it is not limited to respondent-driven sampling. However, it is especially difficult to estimate the size of this bias using respondent-driven-sampling data, as information on people who refuse to participate can be obtained only indirectly from the subset of recruiters who return to collect their secondary incentives. The bias in recruitment by age may not exist in other respondent-driven-sampling studies, but this finding does highlight the challenge created when the community understands a definition of target-group membership differently from the researcher. As in this case, the bias may be quite subtle and difficult to detect. Quantification of the size of the bias would require triangulation with other sources of quantitative data, and the explanation for the bias may become clear only with qualitative data.

Overall, the sample proportions were closer to the population proportions than were the respondent-driven sampling estimates more than 60% of the time, for both sample sizes. Both RDS-1 and RDS-2 adjustments slightly increased the total root mean squared error compared with the sample proportions. The overall failure of the respondent-driven-sampling inference methods to reduce bias is probably because the assumptions behind the respondent-driven-sampling method were not met, and so the methods imperfectly accounted for the patterns of recruitment between subgroups (RDS-1) and differences in network size (RDS-1 and RDS-2). Recruitment was associated with characteristics other than network size. It is surprising that respondent-driven sampling inference methods increased bias more often than not. This occurred because when the respondent-driven sampling adjustments were in the right direction, they often greatly over-compensated. That is, the magnitude of the adjustment was often more than twice the size of the bias, so that after adjustment the respondent-driven sampling estimate was even further away from the population proportion, in the other direction.

The reason that the 95% confidence intervals included the population proportions substantially less than 95% of the time may be due either to the fact that the CIs are too narrow (as has been suggested in another study<sup>9</sup>) or because the respondent-driven sampling estimates were biased, or both.

There are at least four potential limitations to our study. First, empirical evaluation of respondent-driven sampling is problematic. The representative or total-population data that are required for robust evaluation are generally unavailable on the hidden and stigmatized groups that respondent-driven sampling is most commonly used to survey. We evaluated respondent-driven sampling in a non-hidden/non-stigmatized population of male household heads, because of the availability of high-quality total-population data. This may limit the generalizability of our results. However it may also be a best-case scenario for an empirical evaluation of respondent-driven sampling. Respondent-driven sampling data on hidden and stigmatised populations may suffer from higher levels of bias than our sample. If respondent-driven sampling estimators are as unsuccessful at reducing this bias as our findings suggest, then estimates on hidden populations may be even less representative than ours.

Second, the findings of this study are based on only one respondent-driven-sampling sample, and the biases that we observed in the sample proportions could have arisen by chance. The differences between the population and sample proportions were highly unlikely to have occurred by chance, however ( $p \le 0.0001$  for all differences except the under-representation of men in the highest socioeconomic group, where p=0.04). In

addition, in each case where we identified a likely bias, the qualitative data suggested a plausible reason why the bias occurred.

Third, although we ordered the network-size questions so that the first to be asked was similar to the question asked in most respondent-driven-sampling studies,<sup>25,26</sup> statements made by respondent-driven-sampling interviewers during the qualitative study suggested that the more detailed network questions may have caused later recuits to under-report network size so that the interview could be conducted in less time. However, sensitivity analysis showed there was no evidence that collecting detailed network data reduced the performance of the respondent-driven sampling estimators. Therefore we believe that our results and conclusions are robust to this potential limitation.

Finally, our decision not to offer all recruits the chance to recruit others, in order to slow the rate of recruitment, could have biased the results. However, in general, the respondentdriven sampling sample estimate was representative of the population proportions, and where they were not, plausible explanations were identified for these biases. Our results and conclusions are therefore likely to be robust to this limitation as well.

In line with other studies, our study showed that respondent-driven sampling was an effective data-collection method.<sup>10,31</sup> However, our data suggest that the current respondent-driven-sampling statistical-inference methods can fail, and the confidence intervals may be too narrow. Whether the data required to reliably remove bias and measure precision can be collected in a respondent-driven-sampling survey is unresolved. Respondent-driven sampling should be regarded as a (potentially superior) form of convenience sampling method, and caution is required when interpreting respondent-driven sampling study findings.

Further empirical studies should investigate the size of biases in respondent-driven-sampling studies in other populations, particularly in those rare examples of hidden/stigmatized populations on which representative data might be available. In addition, the effect of these biases on both simple and adjusted estimates should be investigated using simulations of respondent-driven-sampling recruitment, and theoretical work should attempt to develop improved point and interval estimators.

#### Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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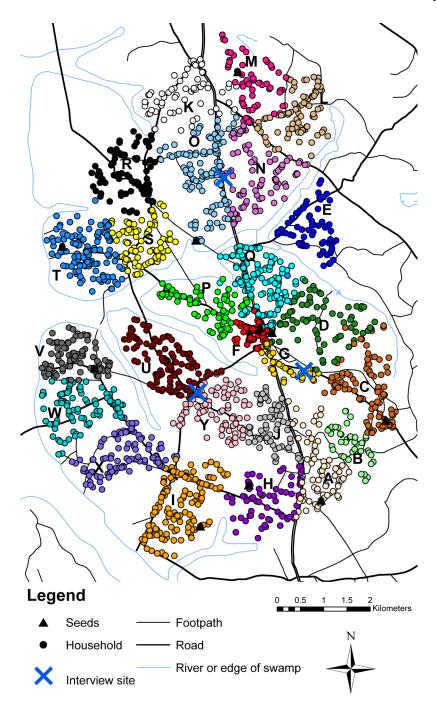
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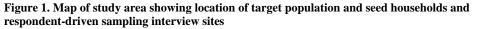
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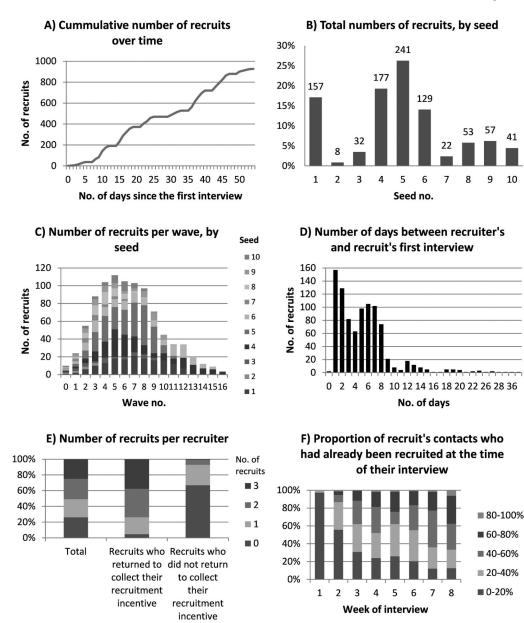
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Colors are used to represent households in different villages. Each village has been labelled with a letter for confidentiality.

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**Figure 2. Summary of the dynamics of respondent-driven sampling survey recruitment** (A)The cumulative number of recruits over time (including seeds). (B) The total number of recruits per seed (excluding seeds). (C)The number of recruits by wave and seed (including seeds).(D) The number of days between recruiters' interview and their recruits' first interview. (E) The number of recruits per recruiter, overall and by whether the recruiters returned for incentive collection (including seeds). (F) The proportion of recruit's network who had already been recruited at the time of their interview (using network size definition NS-5, including seeds).

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#### Figure 3. Recruitment networks showing HIV infection status, by seed

Seeds are shown at the top of each recruitment network. Symbol area is proportional to network size. HIV serostatus is shown by shading: black indicates HIV positive; white, HIV negative; grey, HIV status unknown. HIV status omitted for seeds for confidentiality.

Table

Population proportions, sample proportions, and RDS-1 and RDS-2 estimates with 95% confidence intervals (CIs), for the full and small samples

Respondent-driven sampling estimates are shown in bold if they are closer to the population proportion than the sample proportion. C's are shown in bold if they include the population proportion.

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		Population proportion (n=2402)	F	II RDS s	Full RDS sample (n=927 including seeds)	ncluding	seeds)	Sn	all RDS	Small RDS sample (n=250 including seeds)	including	seeds)
			Sample		Estimate (95% CI)	95% CI		Sample		Estimate (95% CI)	(95% CI)	
					RDS-1		RDS-2			RDS-1		RDS-2
Age group (years)	0-19	0.020	0.004	0.005	(0.000-0.012)	0.005	(0.001 - 0.009)	0.000	ı	ı	ı	ı
	20-29	0.202	0.133	0.133	(0.106 - 0.156)	0.129	(0.109 - 0.159)	0.104	0.104	(0.052 - 0.150)	0.108	
	30-39	0.275	0.250	0.243	(0.208 - 0.275)	0.242	(0.215 - 0.279)	0.267	0.251	(0.175 - 0.313)	0.239	ı
	40-49	0.207	0.240	0.225	(0.193 - 0.263)	0.230	(0.194 - 0.257)	0.246	0.233	(0.169 - 0.313)	0.248	,
	50+	0.297	0.373	0.394	(0.356-0.437)	0.395	(0.351-0.429)	0.383	0.412	(0.334 - 0.505)	0.405	
Tribe	Ganda	0.697	0.667	0.663	(0.619-0.709)	0.661	(0.624 - 0.709)	0.654	0.714	(0.620 - 0.812)	0.662	(0.623 - 0.806)
	Rwanda/kole	0.179	0.210	0.222	(0.182-0.261)	0.212	(0.174-0.244)	0.167	0.152	(0.084 - 0.220)	0.165	(0.085 - 0.201)
	Kiga	0.018	0.021	0.016	(0.005 - 0.029)	0.022	(0.009-0.033)	0.038	0.011	(0.000-0.039)	0.036	(0.004-0.044)
	Rundi	0.047	0.061	0.065	(0.005-0.089)	0.069	(0.046 - 0.089)	0.092	0.086	(0.036 - 0.140)	0.100	(0.039-0.133)
	Other <sup>a</sup>	0.060	0.040	0.035	(0.021-0.051)	0.036	(0.025 - 0.054)	0.050	0.037	(0.010-0.073)	0.038	ı
Religion	Catholic	0.598	0.624	0.640	(0.595-0.684)	0.645	(0.597-0.685)	0.733	0.740	(0.659 - 0.812)	0.752	(0.669 - 0.816)
	Protestant	0.170	0.171	0.151	(0.121 - 0.182)	0.153	(0.131 - 0.188)	0.100	0.086	(0.046 - 0.133)	0.085	(0.051-0.126)
	Muslim	0.227	0.202	0.207	(0.166 - 0.249)	0.198	(0.159 - 0.233)	0.158	0.170	(0.099-0.250)	0.155	(0.098-0.228)
	$Other^b$	0.005	0.003	0.003	(0.000-0.008)	0.004		0.008	0.004	(0.000-0.014)	0.008	
Socioeconomic status	Highest	0.257	0.179	0.167	(0.136 - 0.200)	0.170	(0.141-0.200)	0.200	0.190	(0.120 - 0.271)	0.188	(0.135 - 0.269)
	Higher	0.249	0.242	0.231	(0.197 - 0.266)	0.238	(0.207 - 0.270)	0.258	0.242	(0.174 - 0.314)	0.247	(0.186-0.314)
	Lower	0.229	0.275	0.266	(0.233 - 0.300)	0.269	(0.238-0.302)	0.238	0.247	(0.181 - 0.319)	0.244	(0.179 - 0.302)
	Lowest	0.214	0.266	0.303	(0.263-0.346)	0.290	(0.252-0.326)	0.254	0.281	(0.206 - 0.358)	0.284	(0.206-0.348)
	Unknown	0.052	0.038	0.033	(0.019 - 0.047)	0.033	(0.022 - 0.046)	0.050	0.040	(0.011-0.069)	0.036	(0.015 - 0.064)
Village	A	0.033	0.021	0.032	(0.001-0.096)	0.025		0.042			0.043	
	в	0.017	0.017	0.017	(0.000-0.075)	0.017		0.021			0.022	ı
	С	0.042	0.070	0.028	(0.000-0.094)	090.0		0.104	,	ı	0.073	ı
	D	0.033	0.047	0.019	(0.000-0.052)	0.040		0.017			0.014	

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sample (n=250 including seeds)

Small RDS

Full RDS sample (n=927 including seeds)

Population proportion (n=2402)

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								,				
			Sample		Estimate (95% CI)	<u>, 12 % cr</u>		Sample		Estimate (95% CI)	(I) %e()	
					RDS-1		RDS-2			RDS-1		RDS-2
	E	0.027	0.027	0.072	(0.011-0.259)	0.025		0.000	,	ı		
	F	0.067	0.016	0.013	(0.000-0.026)	0.013		0.013	,		0.007	
	G	0.025	0.028	0.012	(0.000-0.059)	0.030		0.050	,	·	0.046	
	Н	0.031	0.010	0.004	(0.000-0.052)	0.012		0.021	,	ı	0.024	
	Ι	0.060	0.045	0.047	(0.000-0.144)	0.042		0.008	,	ı	0.005	
	ſ	0.028	0.034	0.014	(0.000-0.111)	0.045		0.075	,	·	060.0	
	К	0.031	0.045	0.060	(0.007 - 0.232)	0.037		0.000	,	ı		
	L	0.040	0.047	0.026	(0.006-0.082)	0.056		0.071	,	ı	0.082	
	М	0.026	0.033	0.016	(0.004-0.052)	0.035		0.071	,	·	0.066	
	Z	0.033	0.038	0.030	(0.007 - 0.074)	0.041		0.038	,	ı	0.041	
	0	0.049	0.062	0.026	(0.004-0.073)	0.067		0.079	·	ı	0.081	,
	Р	0.034	0.023	0.024	(0.000-0.057)	0.020		0.021	,	ı	0.016	
	0	0.086	0.047	0.034	(0.001 - 0.067)	0.041		0.025	,	ı	0.019	
	R	0.038	0.055	0.061	(0.003 - 0.151)	0.045		0.013	·	ı	0.015	,
	S	0.038	0.038	0.107	(0.002 - 0.266)	0.040		0.071	,	ı	0.055	
	T	0.050	0.061	0.147	(0.002 - 0.367)	090.0		0.046	ï	I	0.048	·
	U	0.050	0.065	0.064	(0.000-0.161)	0.064		0.042	·	ı	0.051	,
	Λ	0.039	0.045	0.034	(0.000-0.318)	0.043		0.017	,	ı	0.022	
	W	0.040	0.033	0.054	(0.001 - 0.273)	0.028		0.004	,	ı	0.002	
	X	0.043	0.035	0.030	(0.000-0.126)	0.033		0.004	·	ı	0.005	,
	Υ	0.041	0.059	0.030	(0.000-0.124)	0.082		0.150	,	ı	0.175	
Number of sex	0	0.113	0.148	0.170	(0.136 - 0.206)	0.161	(0.133 - 0.190)	0.133	0.142	(0.087 - 0.203)	0.139	(0.095 - 0.192)
partners in last year	1	0.419	0.577	0.572	(0.534 - 0.609)	0.574	(0.537 - 0.611)	0.558	0.573	(0.498-0.652)	0.571	(0.502-0.648)
	2-3	0.114	0.140	0.125	(0.099-0.154)	0.128	(0.104 - 0.151)	0.163	0.147	(0.091 - 0.207)	0.141	(0.093 - 0.189)
	4+	0.037	0.035	0.039	(0.021 - 0.059)	0.040	(0.024 - 0.056)	0.033	0.029	(0.006-0.058)	0.036	(0.011-0.065)
	Unknown	0.316	0.100	0.094	(0.069 - 0.122)	0.098	(0.077-0.122)	0.113	0.108	(0.054 - 0.174)	0.113	
HIV status	Positive	0.063	0.079	0.075	(0.054 - 0.097)	0.074	(0.054 - 0.096)	0.075	0.075	(0.032 - 0.126)	0.078	(0.033 - 0.124)
	Negative	0.600	0.817	0.820	(0.794 - 0.848)	0.820	(0.790 - 0.846)	0.813	0.821	(0.763-0.872)	0.818	(0.758 - 0.874)
	Unknown	0.337	0.105	0.104	(0.082-0.128)	0.106	(0.084-0.132)	0.113	0.104	(0.065-0.153)	0.105	(0.064-0.156)

	<b>Closer to target</b>	Within CI	to target Within CI Closer to target Within CI	Within CI	<b>Closer to target</b>	Within CI	Closer to target Within CI Closer to target Within CI	Within CI
Number of comparisons	52	52	52	26	26	26	49	19
Number met criteria	19	36	17	13	8	18	18	14
% met criteria	37%	%69	33%	50%	31%	%69	37%	74%
- indicates estimate that could not be calculat	be calculated because	subgroups we	re recruited exclusi	vely from withir	themselves or becaus	se (excluding s	eeds) no one was re	ecruited exclusively from within themselves or because (excluding seeds) no one was recruited from certain subgroups.

aIncludes other known tribe and unknown tribe.

 $\boldsymbol{b}_{\textrm{Includes}}$  other known religion, no religion, and unknown religion.