Landscape and Residential Variables Associated with Plague-Endemic Villages in the West Nile Region of Uganda

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Landscape and Residential Variables Associated with Plague-Endemic Villages in the West Nile Region of Uganda

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Abstract. Plague, caused by the bacteria Yersinia pestis, is a severe, often fatal disease. This study focuses on the plague-endemic West Nile region of Uganda, where limited information is available regarding environmental and behavioral risk factors associated with plague infection. We conducted observational surveys of 10 randomly selected huts within historically classified case and control villages (four each) two times during the dry season of 2006 ($N = 78$ case huts and $N = 80$ control huts), which immediately preceded a large plague outbreak. By coupling a previously published landscape-level statistical model of plague risk with this observational survey, we were able to identify potential residence-based risk factors for plague associated with huts within historic case or control villages (e.g., distance to neighboring homestead and presence of pigs near the home) and huts within areas previously predicted as elevated risk or low risk (e.g., corn and other annual crops grown near the home, water storage in the home, and processed commercial foods stored in the home). The identified variables are consistent with current ecologic theories on plague transmission dynamics. This preliminary study serves as a foundation for future case control studies in the area.

INTRODUCTION

Plague is a severe, rodent-associated bacterial zoonosis caused by Yersinia pestis.1 Plague bacteria are maintained by various sylvatic and commensal rodent species and their fleas in enzootic and epizootic cycles.2,3 The majority of human infections are associated with epizootics or periods of rapid transmission that often cause local die-offs of susceptible rodent hosts, which forces potentially infectious fleas to infest alternative hosts, including humans. The severity of human illness depends, in part, on the clinical presentation. Bubonic plague, typically associated with flea-borne transmission, is the most common and usually the least severe form of the disease. Septicemic and primary pneumonic infections are commonly associated with cutaneous exposure or inhalation, respectively. These infections are rare but often more severe.4,5 Without treatment, human fatality rates range from 50% to 60% for bubonic plague to almost 100% for pneumonic infections.5 However, if plague is diagnosed and treated early with appropriate antibiotic therapy, outcome improves considerably.1,3,7

Although improvements in sanitation and availability of antibiotics have decreased the epidemiicity of plague, the disease remains a public health threat in many areas of the world.8 Over the last two decades, the majority of human plague cases has been reported from eastern and southern Africa and the neighboring island of Madagascar.9,10–12 Within plague-endemic regions, risk of exposure to Y. pestis is spatially heterogeneous.12 Several statistical models have been constructed using geographic information system (GIS)-based and remotely sensed (RS) data to evaluate landscape variables that are associated with plague risk and attempt to explain this spatial heterogeneity.10,12,17 For example, within the West Nile region of Uganda, which is the primary epidemiologic focus within that country,18 a fine resolution GIS-based model revealed that areas of higher elevation (above 1,300 m) that are wetter, with less vegetative growth and more bare soil during the dry month of January, pose an elevated risk of plague occurrence.12

Such models are useful for identifying areas of elevated risk based on landscape variables, which aids in targeting limited public health resources. However, even within areas that are considered ecologically conducive for plague activity, the likelihood of human exposure to the disease agent is influenced by environmental and behavioral factors that are not included in these landscape-level models. Factors influencing the critical contact between rodent reservoirs, flea vectors, and humans as well as human behaviors that may enhance or diminish this contact with infectious vectors or hosts are not well-understood, particularly within the West Nile plague focus in Uganda.19–21 This information is essential to establishing evidence-based recommendations on the prevention and control of plague in endemic regions, but few studies have sought to identify these potential risk factors. Within many of the world’s plague foci, humans are most often infected in the domestic and peri-domestic environment.8,13,14,21–23 Availability of harborage and food sources for rodents, often a result of human behaviors within the home environment, was shown to be associated with plague cases in North America.21 In addition, exposure to plague-infected companion animals has been previously identified as a risk factor.24–26 Other studies suggested that poor, unsanitary conditions are responsible for plague epidemics, because these conditions promote cohabitation of homes by humans and competent rodent hosts, which ultimately increases the potential of transmission.23,27,28

Our study focuses on the plague-endemic West Nile region of Uganda, where, from 1999 to 2007, a mean of 223 human plague cases was reported annually to the Ugandan Ministry of Health.29 During this time period, cases were not typically investigated to determine the location of exposure. Thus, in this study, we assume that cases acquired infection within the village of residence. The objectives of this study were (1) to identify landscape and residential risk factors associated

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with randomly selected huts within villages that historically reported human plague cases compared with randomly selected huts within villages that have not reported human plague cases from 1999 to 2004, (2) to identify residence-based variables associated with a previously published classification of elevated versus low risk based on RS variables, and (3) within areas that were classified as elevated risk based on RS variables, to identify residence-based predictors of huts within case versus control villages. This preliminary study, which couples a previously published fine-scale landscape model with an observational survey of environmental and behavioral variables, reveals risk factors for plague at the hut and village levels and serves as a foundation for future case control studies in this plague-endemic region of Uganda.

MATERIALS AND METHODS

Study area. This study was conducted in the plague-endemic West Nile region of northwestern Uganda (Figure 1). It focused primarily on the hut and village levels in Vurra and Okoro Counties, within Arua and Nebbi Districts, respectively. Previous plague-risk models identified elevation as a key predictor of plague risk in this region. Specifically, areas above 1,300 m represented greater risk for epizootic activity than areas below this elevation. In our study, all of the villages sampled were above 1,300 m (1,348–1,638 m).

Plague case definition and selection of case and control sites. In this area, plague was clinically diagnosed by symptomology alone: sudden onset of fever, chills, malaise, headache or prostration accompanied by painful regional lymphadenopathy (bubonic), hematemesis or hematochezia (septicemia), or cough with hemoptysis (pneumonic). The local health care delivery system did not include laboratory confirmation of provider diagnosis, and environmental investigations were not conducted to determine the exact location of exposure. We used a previously described database of suspected human plague cases reported from 1999 to 2004 to select villages with or without a history of human plague cases. Villages were categorized as case if a human plague case was reported in at least 3 of 6 years examined; these included Agore (N = 10 cases), Olii (N = 30 cases), Pomosi (N = 6 cases), and Upper Uyaru (N = 20 cases). Villages of similar estimated physical size and elevation that did not report any plague cases during the time period spanning 1999–2004 were selected as controls. These included Anyiku, Pembeleku, Kaza, and Gbalia (Figure 1).

Observational survey. Observational surveys were conducted at the hut level within two case villages (Olii and Pomosi) and two control villages (Kaza and Pembeleku) in Arua district and two case villages (Agore and Upper Uyaru) and two control villages (Anyiku and Gbalia) in Nebbi district (Figure 1).

The observational survey was used to collect information from huts within case and control villages on potential residential risk factors related to plague infection. Surveys were conducted, and latitude and longitude coordinates were collected for 10 randomly selected huts within the villages during
were conducted to identify (1) environmental variables associated with randomly selected huts within villages with (case) or without (control) a reported history of plague, (2) residence-based variables associated with a previously published RS classification of elevated versus low risk, and (3) residence-based predictors of huts that were situated in areas classified by the previously published model as elevated risk that were associated with case versus control villages. For each significant variable, odds ratios (ORs) and 95% confidence intervals (CIs) were calculated. Each variable that was significant in univariate tests was included for selection in a multivariable logistic regression model; predictive variables included in the model were chosen by forward stepwise procedure, with a probability to enter of 0.25. An adjusted OR (aOR) and 95% CI were calculated for each variable within the model. A goodness of fit test was used to determine if the variables included in the model were adequate. The receiver-operating characteristic curve (ROC) was used to assess the overall discrimination of the model. The ROC curve is a graphical representation of the relationship between false-positive (one-specificity) and true-positive (sensitivity) rates. Overall model accuracy was determined based on the area under the ROC curve, which provides a threshold-independent measure of overall accuracy. Data were analyzed using JMP v8 statistical software (SAS Institute, Cary, NC).
Comparisons were considered statistically significant when $P < 0.05$.

**RESULTS**

Summary of landscape- and residence-based variables associated with case and control villages based on historic reports of human plague case occurrence. During the period of March to June 2006, a total of 158 huts were surveyed. Among these huts, 78 were located in villages classified as case based on records of human plague case occurrence from 1999 to 2004, and 80 huts were in control villages that did not report any human plague cases during this time. Based on univariate logistic regression analyses, three residential variables (pigs observed in the peridomestic setting, bedding material, and distance to neighbor) and four landscape variables (elevation, wetness, brightness, and Landsat ETM+ band 6) were significantly associated with village classification (Table 1). Compared with huts within control villages, huts within case villages were more likely to have pigs observed in the peridomestic environment, less likely to have beds made of material other than reed, and more likely to be farther from neighboring homesteads than huts within control villages (case villages median [range] = 10 m [3–50 m], control villages = 8 m [2–45 m]). In addition, huts within case villages were more likely to be situated in pixels that were wetter with more bare soil (higher wetness values, and lower brightness and band 6 values) compared with huts in control villages. Although elevation was included in the model and had a statistically significant effect, the OR revealed little or no association with hut classification (OR = 1, 95% CI = 1–1.01). However, this may be because elevation is quantified in meters and changes in risk are not expected at such small increments of measurement. Furthermore, there was limited variation in elevation between huts in case and control villages, and all were situated above the previously described 1,300-m threshold.

Using the variables described above that were significantly different in univariate logistic regression analyses, we used a forward stepwise selection process to create multivariable logistic regression models predicting the odds of a hut being classified as belonging to a case village. The model included positive associations with presence of pigs in the peridomestic setting and distance to neighbor but negative associations with bedding material other than papyrus mats and Landsat ETM+ band 6 (Table 2). Based on the area under the ROC curve, this model had an overall accuracy of 78%.

**Summary of residence-based variables associated with huts situated in areas classified as elevated or low risk.**

As described in the subsection above, landscape-level RS variables that were used in a GIS-based risk assessment of plague in this region remained as significant indicators of risk. Although the previous study indicated that these variables were suggestive of certain agricultural practices posing a risk, the study did not include data to verify this assertion. Here, we sought to identify residence-based correlates of the previously published plague-risk model based on RS variables. Of the 158 huts surveyed and described above, 100 huts were located within pixels classified by the model as elevated risk, and 58 huts were located in pixels classified as low risk. When comparing huts within pixels classified as elevated risk versus low risk based on RS data, nine residence-based variables differed significantly between these categories (Table 3). Huts within pixels classified as elevated risk were less likely to have water available to rodents inside the hut than low-risk areas, and they were also less likely to have coffee growing within 30 m of the residence and processed commercial foods stored within the residence. Compared with huts located in pixels classified as low risk, those in areas classified as elevated risk were more likely to have sheep observed in the peridomestic environment, to have corn grown or dried within 30 m of the residence, and to grow melons on both vines and trees than low-risk areas. Melons growing on trees were defined as a

**Table 1**

<table>
<thead>
<tr>
<th>Risk factor</th>
<th>Case huts ($N = 78$)</th>
<th>Control huts ($N = 80$)</th>
<th>OR</th>
<th>95% CI</th>
<th>$P$ value</th>
</tr>
</thead>
<tbody>
<tr>
<td>Peridomestic pig</td>
<td>41 (72)</td>
<td>28 (73)</td>
<td>2.31</td>
<td>1.09–4.13</td>
<td>0.025</td>
</tr>
<tr>
<td>Bedding material (other)</td>
<td>1 (64)</td>
<td>9 (61)</td>
<td>0.09</td>
<td>0.01–0.75</td>
<td>0.007</td>
</tr>
<tr>
<td>Distance to neighboring homestead</td>
<td>76 (76)</td>
<td>77 (77)</td>
<td>1.07</td>
<td>1.03–1.21</td>
<td>0.002</td>
</tr>
<tr>
<td>Elevation</td>
<td>78 (78)</td>
<td>80 (80)</td>
<td>1</td>
<td>1–1.01</td>
<td>0.014</td>
</tr>
<tr>
<td>Wetness</td>
<td>78 (78)</td>
<td>80 (80)</td>
<td>1.1</td>
<td>1.02–1.08</td>
<td>0.002</td>
</tr>
<tr>
<td>Brightness</td>
<td>78 (78)</td>
<td>80 (80)</td>
<td>0.98</td>
<td>0.96–1.0</td>
<td>0.031</td>
</tr>
<tr>
<td>Landsat ETM+ band 6</td>
<td>78 (78)</td>
<td>80 (80)</td>
<td>0.95</td>
<td>0.92–0.98</td>
<td>0.0004</td>
</tr>
</tbody>
</table>

*Percentages were calculated only for nominal variables.

**Table 2**

<table>
<thead>
<tr>
<th>Parameter estimate</th>
<th>Likelihood ratio test</th>
<th>Odds ratios</th>
</tr>
</thead>
<tbody>
<tr>
<td>Model covariates</td>
<td>Estimate</td>
<td>SE</td>
</tr>
<tr>
<td>Intercept</td>
<td>0.63</td>
<td>1.07</td>
</tr>
<tr>
<td>Peridomestic pig</td>
<td>0.34</td>
<td>0.22</td>
</tr>
<tr>
<td>Bedding material (other)</td>
<td>-1.36</td>
<td>0.59</td>
</tr>
<tr>
<td>Distance to neighboring homestead</td>
<td>0.07</td>
<td>0.03</td>
</tr>
<tr>
<td>Landsat ETM+ band 6</td>
<td>-0.06</td>
<td>0.02</td>
</tr>
</tbody>
</table>

SE = standard error; aOR = adjusted odds ratio; df = degrees of freedom.
model presented above that did not constrain the analysis to areas of elevated risk, pigs observed in the peridomestic environment and distance to neighbor were positively associated with classification of case villages (Table 6). Although the ORs were similar between the models for distance to water, when restricting the model to include only huts within elevated risk areas, the OR associated with pigs in the peridomestic environment notably increased (Tables 2 and 6). The overall accuracy of this multivariable logistic regression model was 74%.

DISCUSSION

Few studies have focused on environmental and behavioral variables associated with plague infection, especially in the plague-endemic West Nile region of Uganda. Information gleaned from such studies may aid in tailoring evidence-based recommendations for the prevention and control of plague in this region.

Using these significant variables, a multivariable regression model was created based on a forward stepwise selection process. The final model that predicted the odds of classification as elevated risk, which had an overall accuracy of 75%, was based on a positive association with corn and other crops grown or dried within 30 m of the residence and negative associations with water availability to rodents and processed commercial foods stored within the residence (Table 4).

Summary of risk factors for plague in case and control villages within areas classified as posing an elevated risk for plague.

A total of 100 huts located within pixels classified as elevated risk for plague were included in this analysis, with 53 huts being in historic case villages and 47 huts in control villages. Of the variables considered in univariate logistic regression analyses, four residential variables were significantly associated with village classification. Compared with huts in control villages, huts within case villages were more likely to have pigs in the peridomestic environment, were more likely to have a greater number of males under the age of 12 years sleeping in the hut on a regular basis, and were more likely to be farther from their neighbors (case village median [range] = 15 m [3–40 m], control village median [range] = 10 m [6–32 m]). Huts within case villages were less likely to have bedding material other than reed mats compared with huts within control villages (Table 5). Bedding material within the category other included foam pads and stuffed mattresses.

Of the variables that were significant in univariate tests, only two were chosen by forward stepwise regression to be included in the multivariable regression model. As in the

<table>
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<th>Odds ratios</th>
</tr>
</thead>
<tbody>
<tr>
<td>Model covariates</td>
<td>Estimate</td>
<td>SE</td>
</tr>
<tr>
<td>Intercept</td>
<td>–1.23</td>
<td>0.54</td>
</tr>
<tr>
<td>Water available to rodents</td>
<td>–0.26</td>
<td>0.44</td>
</tr>
<tr>
<td>Corn grown or dried within 30 m</td>
<td>1.13</td>
<td>0.46</td>
</tr>
<tr>
<td>Other crops grown or dried within 30 m</td>
<td>0.97</td>
<td>0.51</td>
</tr>
<tr>
<td>Processed commercial foods stored within residence</td>
<td>–1.06</td>
<td>0.46</td>
</tr>
</tbody>
</table>

SE = standard error; aOR = adjusted odds ratio; df = degrees of freedom.
practices that increase the availability of food or harborage for rodents. Rodent movement is often driven by food availability and appropriate habitat. Therefore, behaviors that attract flea-infested rodents into peridomestic settings may increase the probability of human exposure to plague-infested fleas or rodent hosts.

In an attempt to identify areas of elevated risk of human exposure to plague bacteria in the West Nile region, a fine-resolution model was recently created based primarily on RS variables. As previously mentioned, plague risk was higher at elevations greater than 1,300 m than below, and RS variables included in the model implied that risk was elevated in wetter areas with less vegetative growth and more bare soil during the dry month of January when fields of seasonal crops are typically fallow. The reasons for why these variables were indicative of elevated risk were not evaluated in the previous study, but it was hypothesized that these spectral signatures were associated with areas of more intensive agriculture.

Presumably, agricultural crops would increase the food supply for rodents and may result in higher densities of rodents. Our residence-based observations showed that some seasonal crops, such as corn and other grains, were associated with elevated risk areas and may be more attractive to rodents than perennial plantation crops such as bananas or coffee beans, which were either similar between elevated- and low-risk areas or more frequently associated with low-risk areas. Indeed, previous studies rarely observed rodent damage to banana and coffee crops compared with other seasonal crops, suggesting that, perhaps, rodents are less attracted to banana and coffee crops. The differentiation between these crop types is consistent with the spectral signatures used to differentiate elevated- and low-risk areas. Specifically, corn and other grain fields are often fallow during the dry month of January and would be expected to yield high brightness values indicative of bare soil and little vegetative growth.

The RS model also identified a positive association between wetness and the probability of an area being classified as elevated risk. Our multivariable model indicated a negative association with availability of water for rodents (e.g., presence of water in containers) within huts situated in pixels classified as elevated risk. We believe that water storage in huts may be more common in drier areas that are more distant from water sources. Finally, storage of processed commercial foods within the hut was more frequently associated with areas of low risk. Again, this variable may relate back to agricultural practices, such that in areas of elevated risk, more subsistence crops such as corn are grown, and thus, there is less need for purchasing commercial foods. Alternatively, it is possible that commercially processed foods are more commonly purchased in areas that engage in growing cash crops, such as coffee, which is found in areas classified as low risk, as stated above. In addition, the packaging of this processed food could be a deterrent for the rodents if enough easily accessible food is available. Because our study did not quantify the abundance of such foods but rather surveyed simple presence or absence of these food types, it is difficult to draw conclusions about this variable.

Our observational survey was conducted before the identification of RS landscape variables associated with elevated plague risk in this region, and thus, we did not control for many of these variables in our village selection. Confirming and reemphasizing the importance of these RS variables in assessing the likelihood of human plague case occurrence, our univariate analyses (objective 1) (Table 1) showed that three of four variables included in the previous model (e.g., elevation, brightness, and Landsat ETM+ band 6) remained significant predictors of case or control designation. The multivariable analyses revealed that huts in case villages typically had lower Landsat ETM+ band 6 values compared with huts in control villages; this is indicative of wetter conditions in case areas compared with control areas and shows consistency with the model predictions.

In addition to these landscape variables, we identified three residence-based variables that were predictive of case or control village designation. Specifically, when huts in case villages were compared with huts in control villages, pigs were more commonly observed in the peridomestic setting, and huts were spaced farther from their neighbors. We speculate that peridomestic pigs were observed more frequently in historic case villages because of a similar dietary pattern between rodent

### Table 5

<table>
<thead>
<tr>
<th>Risk factor</th>
<th>Case (N = 53)</th>
<th>Control (N = 47)</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Present (N)</td>
<td>Percent*</td>
</tr>
<tr>
<td></td>
<td>OR 95% CI</td>
<td>P value</td>
</tr>
<tr>
<td>Peridomestic pig</td>
<td>28 (48)</td>
<td>58</td>
</tr>
<tr>
<td>Bedding material (other)</td>
<td>1 (46)</td>
<td>2</td>
</tr>
<tr>
<td>Number of males ≤ 12 years old sleeping in hut</td>
<td>49 (49)</td>
<td>42 (42)</td>
</tr>
<tr>
<td>Distance to neighboring homestead</td>
<td>52 (52)</td>
<td>47 (47)</td>
</tr>
</tbody>
</table>

*Percentages were calculated only for nominal variables.

### Table 6

<table>
<thead>
<tr>
<th>Parameter estimate</th>
<th>Likelihood ratio test</th>
<th>Odds ratios</th>
</tr>
</thead>
<tbody>
<tr>
<td></td>
<td>Estimate</td>
<td>SE</td>
</tr>
<tr>
<td>Intercept</td>
<td>-0.70</td>
<td>0.45</td>
</tr>
<tr>
<td>Peridomestic pig</td>
<td>0.79</td>
<td>0.24</td>
</tr>
<tr>
<td>Distance to neighboring homestead</td>
<td>0.07</td>
<td>0.03</td>
</tr>
</tbody>
</table>

SE = standard error; aOR = adjusted odds ratio; d.f = degrees of freedom.
species and pigs. Commensal and sylvatic rodents that are sus-
cceptible to plague infection could be attracted to food pro-
vided to the pigs. These food sources could serve to increase
the local carrying capacity for these rodents. Regarding hut
spacing, local observations suggest that the spaces between
neighboring homesteads are often occupied either by agricul-
tural plots that could serve as food sources for rodents (e.g.,
corn and other seasonally grown grains) or brush that could
serve as rodent harborage. Land converted to agriculture
increases the habitat and food availability for sylvatic rodents
and potential interactions with peridomestic rodents. 1, 2, 3, 8, 9
It is also believed that areas with increased primary production
of food crops, which result from elevated precipitation rates
also believed that areas with increased primary production
of food crops, which result from elevated precipitation rates
and wetter conditions, may increase the carrying capacity of
rodents. 1, 2, 13, 33, 38, 39 After repeating the analysis using only huts
that occurred within pixels recently classified as elevated risk
(objective 3) (Tables 5 and 6), the association with pigs in the
peridomestic setting and distance to neighbor remained as sig-
nificant predictors of case or control designation.

In addition to identifying variables that may increase the
risk for plague infection, other factors were more commonly
found within control villages, suggesting that these behaviors
or environmental conditions may be protective against ex-
posure to plague bacteria. Beds made of material other than reed
mats (e.g., foam pad or stuffed mattress) were observed more
frequently within huts in control villages compared with case
villages. We speculate that reed mats may be more likely to
be infested with fleas than other bedding, because the organic
content or dust media in these mats provides a suitable place
for fleas to breed. 41

Although our models yielded overall accuracies of 74–78%,
several additional studies are required to improve our under-
standing of plague risk factors. Because case investigations
were not conducted to determine the location of exposure, our
study was based on randomly selected huts within case and
control villages. Within villages, huts are not homogenous with
respect to many of the surveyed variables. Focusing future sur-
veys on the residence of laboratory-confirmed plague patients
and appropriate controls may improve model accuracy. As sug-
gested before, targeting areas classified as elevated risk that
continue to not report plague cases may provide insight into
subtle differences in housing, crops, or behaviors that prove
protective. 12 In addition to focusing future efforts within eco-
logically conducive areas and in conjunction with laboratory-
confirmed plague case investigations, it would be beneficial to
conduct the surveys during plague outbreaks or during the wet
season, when most plague cases have been reported. Many of
the variables that we surveyed related to crop production and
storage. These practices change seasonally and may be asso-
ciated with observed differences over time and space in the
distribution of zoonotic hosts and their vectors. 12, 19, 20, 32, 35
Future studies could also include making adjustments to the envi-
ronmental and behavioral questions on the observational survey.
Questions should include type and frequency of mud smear-
ning, quantity of seasonal crops growing around huts, quantity of
foods being stored in huts, and frequency and type of inter-
actions between peridomestic animals and humans. Lastly, fur-
ther investigation of the variables that presented as significant
in this study should be conducted to test our hypotheses for
why they may enhance or diminish plague risk. Understanding
the critical link between fleas, their reservoir hosts, and
humans is essential to prevention and control of plague within
this endemic region of Uganda. 23 We believe that this study
provides important preliminary information about environ-
mental and behavioral factors that might be contributing to
plague risk.

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Environmental Health Officer for Logiri Subcounty. We wish to es-
pecially recognize the lifetime contributions of author Asaph Ogen-Odoi,
who died in December 2006 while conducting field work to control the
large plague outbreak mentioned in this paper that immediately fol-
lowed this study. His contributions to this study and to the under-
standing and control of plague in Uganda can never be overestimated.

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