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Impact of Drought on the Spatial Pattern of Transmission of Schistosoma haematobium in Coastal Kenya

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Abstract. We analyzed temporal changes in spatial patterns of active Schistosoma haematobium infection in different age groups and associated them with ponds infested with Bulinus snails. A major drought between 2001 and 2009 resulted in drying of ponds that were known sources of infection, and we detected very few or no snails in ponds that were infested in the past. The household-level spatial pattern of infection for children of various age groups in 2009 was contrasted with historical data from 2000. The significant local clustering of high- and low-infection levels among school-aged children that occurred in 2000 was absent in 2009. We attribute the disappearance of significant clustering around historical transmission hotspots to a decade-long drought in our study area. The implications of extreme weather and climate conditions on risk and transmission of S. haematobium and their relevance to control strategies are discussed.

INTRODUCTION

Urinary schistosomiasis remains a significant public health problem, especially in sub-Saharan Africa.1–4 In Kenya, the coastal strip is hyperendemic for urinary schistosomiasis but with substantial spatial and temporal heterogeneities.5 Local water use behavior and proximity of snail intermediate host breeding sites typically drive the focal distribution of schistosomiasis.6–8 Schistosoma haematobium has been studied in the Msambweni area since 1984.9 Prevalence has remained high (> 50%) over the years, despite introduction of alternative water sources (boreholes) and chemotherapy programs targeting school children.7,9,10 The intermediate host in coastal Kenya is Bulinus nasutus snails that are commonly found breeding in rain-fed ponds.11,12 Because most of the important transmission foci are rain-fed, rainfall is the key abiotic factor. Prolonged droughts can drastically curtail the survival and reproduction of snail intermediate hosts and eliminate transmission sites.11,12 In the Msambweni study area, human infection clusters were detected primarily around one transmission site, Nimbozde pond (Figure 1) which is located within Milalani village.12,13 Based on data from 2000 parasitological survey, we previously reported the presence of spatial clustering of S. haematobium infection around Nimbozde pond and then studied the influence of multiple transmission sources on infection patterns and their changing effect over time in Msambweni District, Coast Province, Kenya.5 Now, a decade later, we revisited the same study area and applied similar spatial analyses after a decade-long drought to detect changes in spatial clustering of infections at the household level.

METHODS

Study area and population. This study was carried out in Milalani village, Msambweni division, Msambweni district (formerly Msambweni Division, Kwale District) of the Coast Province of Kenya. It is located (4.47°S and 39.45°E) approximately 50 km south of Mombasa on the coastal plain along the Indian Ocean, with an approximate area of 2.37 km² (Figure 1). In 2000 and 2003, parasitological surveys and treatment of schistosomiasis were done in Milalani, and this area was revisited in 2009. In 2009, there were 409 households in Milalani with a total of 1,645 people, of whom 47% participated in the parasitology survey (777 people in 240 households). The extent of the village differed slightly in 2009 compared with 2000; only houses within the 2009 extent were included in the analysis for the two study periods. With the new extent, we considered 269 of 279 houses that were analyzed in 2000 and a total of 1,074 people that were tested for S. haematobium.13 The area, population, and water sources have been described previously.1,12,13 Of the water sources mapped and studied in 2000, the spring-fed rivers and boreholes remained active, but the ponds were most often dry because of insufficient rainfall through 2009. Historically, despite the availability of clean water through boreholes and piped water supply, residents in the area prefer pond and stream/river water for laundry over the hard water from boreholes and open wells. They also use these surface sources for swimming and bathing.10,14

Of the water resources that have been monitored in the past from the Msambweni study area, we consider here the Nimbozde pond, which is located within Milalani village and was the subject of intensive study in 2000–2003; it was historically the most heavily infested with infected B. nasutus in 2000 data.

Snail sampling. Five of six ponds in the Msambweni area (described in the work by Kariuki and others)15 were monitored for presence of S. haematobium intermediate host snails starting in July of 2008. These ponds (Bovo, Mwamagongo, Maridzani, Kiziamkala, and Nimbozde)12 had been studied in 2001–2003, and at that time, Nimbozde was the most heavily infested. In addition, we included a temporary pond, Chiziani, near Nganja village. The five known ponds were dry from July of 2008 to June of 2010, even during the long rains. However, B. nasutus snails were observed during both the long and short rains of 2009 in Chiziani pond. We initiated ongoing biweekly snail sampling in nine sites in the six ponds starting in June of 2010; snails were taken to the laboratory at Msambweni Hospital, separated by species, and tested for cercarial shedding.12,15

Ethical oversight. Before collecting urine samples for determining S. haematobium infection status, informed consent was obtained from area residents (or their parents). These studies were performed under human investigations protocols approved by the Ethical Review Board of Kenya Medical Research Institute (Nairobi, Kenya; Non-SSC protocol 087)
Infection prevalence and intensity. Residents of Milalani village submitted two midday urine specimens for examination for *S. haematobium* infection from 2000 to 2003; in 2009, they submitted one midday sample. Ten-milliliter aliquots from well-mixed urine samples were subjected to standard Nuclepore filtration,\textsuperscript{13, 16} and individual infectious burden was determined by microscopic counting of the parasite eggs recovered on the filters. Urine tests during both study periods were conducted in a parasitology laboratory situated within Msambweni District Hospital. Milalani residents had been surveyed before 2000 and were treated with praziquantel in 2000, 2003, and September of 2009.

Aggregated household infection levels are reported as geometric mean density (number of parasites per 10 mL urine per person in a household, regardless of infection status), which is a function of both prevalence (the proportion of people infected in a household) and geometric mean intensity (the geometric mean number of parasites for only infected people infected in a household).\textsuperscript{13, 17} We used mean density rather than mean intensity to allow for comparison with *S. haematobium* infection patterns in 2000.\textsuperscript{13} However, results were similar when only intensity was considered, because the household-level prevalence is high, resulting in similar patterns of mean intensity and mean density.

Statistical analysis. We analyzed the pattern of infections in children 6–17 years of age, the age group that exhibits the highest infection levels.\textsuperscript{18} Differences in prevalence between age groups were tested using the homogeneity $\chi^2$ test. The Mann–Whitney $U$ and Kruskal–Wallis tests were used for comparing distributions of infection intensity and density for two demographic groups and three or more demographic groups, respectively. These non-statistical analyses were performed using SPSS version 17.0 software (SPSS Inc., Chicago, IL; www.spss.com).

Geospatial processing. Households and water sources were mapped as described by Clennon and others\textsuperscript{5, 13} using an Ikonos high-resolution satellite image comprised of 1-m$^2$ panchromatic and 4-m$^2$ multispectral images of Msambweni complemented by global positioning system (GPS) readings. During the 2001 and 2009 demographic surveys, each household was assigned the same village-affiliated household identification (HID), and each person was assigned an individual identification (ID) number incorporating the HID.\textsuperscript{5, 13} Houses from 2000 that were still standing in 2009 and new houses were recorded, and information about movement of residents was collected.\textsuperscript{5} We were able to locate 269 homes from 2000 and 139 new homes in Milalani in 2009.

Household and water source locations were joined with demographic, parasitologic, malacologic, environmental (hydrological and topographical), and weather (rainfall) data. Location data were georectified to the Universal Transverse Mercator (UTM) zone 37S projection (1984 datum) in the GIS software package ArcGIS 9.3 (Environmental Systems Research Institute, Redlands, CA).

Spatial statistics. For spatial analyses, children were grouped by age (6–9, 10–13, and 14–17 years). Global, local, and focal spatial analyses were used to examine the spatial structure of *S. haematobium* infection patterns and identify significant clustering of elevated infection levels in the study area. To allow for comparison with the 2000 infection pattern, we applied the spatial extent of Milalani village in 2009 to the more extensive coverage of Milalani in 2000 (we also considered houses in neighboring villages that were not studied in 2009).\textsuperscript{5, 13}

We initially tested for global clustering (i.e., to determine whether there is clustering anywhere in the study area) of households and intensity of infection by household using global second-order spatial analysis, global Ripley’s $K$ function,\textsuperscript{19} and global weighted $K$ function.\textsuperscript{20} Details about these global spatial analysis tools were detailed in the work by Clennon and others.\textsuperscript{5, 13} After determination of global clustering patterns, we applied local and focal clustering measures to detect specific
locations where households with high-infection intensity were clustered and whether these high-intensity households were clustered around a suspected source of infection (pond containing infected snails).

We applied statistics developed by Getis and others,\textsuperscript{21} Getis and Ord,\textsuperscript{22,23} and Ord and Getis,\textsuperscript{24} \(G^i(d)/Gi(d)\), to test for local and focal clustering of the human infection patterns using Point Pattern Analysis software.\textsuperscript{25} The \(G^i(d)/Gi(d)\) statistic was applied both as local and focal measures of clustering of infection around each household and around the centroid of Nimbodze Pond, which was considered the source point for potential focal clustering analyses. Clustering was considered up to a realistic walking distance of 1,500 m surrounding the potential transmission focus (Nimbodze Pond). The 2009 data were compared with data from the 2000–2003 survey, where more details about these statistics are provided.

**Rainfall data.** Daily rainfall data were retrieved from the Ministry of Agriculture rain gauge located near Msambweni Hospital, which is \(\approx\) 2.5 km from Milalani village.

**RESULTS**

**Snail collections.** In 2010, \(B.\ nasutus\) snails were only recovered from three of five large ponds studied in 2000–2003. In two of these ponds, Mwamongo and Maridzani, 20 and 4 \(B.\ nasutus\), respectively, were recovered before drying. Six \(B.\ nasutus\) were recovered for the first time from the third pond, Nimbozde, which has water to date (March of 2011), 10 months after the sampling was started. One snail from Maridzani shed \(S.\ haematobium\) cercariae, but no cercariae shedding was observed in any snail from Mwamongo and Nimbozde. \(B.\ nasutus\) were consistently found in the Chiziani temporary pond every fortnight between June of 2010 and September of 2010. In total, 156 snails were recovered from Chiziani, and 6 of 156 snails shed \(S.\ haematobium\) cercariae. Among other snails, \(Lanistes\) purpureus was collected from all ponds except Kiziamkala, whereas \(B.\ forskali\) was only found in Kiziamkala and Mwamongo. \(B.\ forskali\) were not shed in this study, because they have not been historically incriminated in urinary schistosomiasis transmission in the study area\textsuperscript{13} or elsewhere in east Africa.\textsuperscript{26} These findings are in sharp contrast to 2001–2003, when \(B.\ nasutus\) were found in all the large ponds (often in large numbers) as well as rice fields, and cercariae shedding \(B.\ nasutus\) snails were collected from Kiziamkala, Maridzani, Boivo, Mwamongo, and in the highest proportion (2.5%), Nimbozde pond, where the highest number of \(Bulinus\) snails was also collected.\textsuperscript{12} In 2000, other snails (\(L.\ purpureus, B.\ forskali, and Melanoides\) tuberculata) were collected from several ponds.\textsuperscript{12}

**Infection prevalence and intensity.** In 2009, we were able to determine the locations of 409 households with 1,645 residents (Figure 1). From those households, 777 (47.2\%) residents submitted urine samples that were tested for \(S.\ haematobium\) eggs, and 336 (43.2\%) of these samples were positive. Among the tested individuals, the age distribution was \(\leq\) 5 years (31), 6–9 years (135), 10–13 years (152), 14–17 years (101), 18–21 years (48), and \(\geq\) 22 years (310). The overall geometric mean infection intensity was 28.9 eggs/10 mL urine. The mean ± standard deviation (SD) number of school-aged children tested per household was 1.61 ± 0.67. When all age groups were considered, infection prevalence and mean intensity varied significantly by age group (homogeneity \(\chi^2 = 201, P < 0.001;\) Kruskal–Wallis \(\chi^2 = 134, P < 0.001\) (Figure 2). Infection levels were highest in individuals 6–21 years, with infection prevalence peaking from 10 to 17 years and intensity peaking from 10 to 13 years.

In 2000, 269 households were located within the same village extent as the 2009 data. From those households, 1,053 residents submitted urine samples that were tested for \(S.\ haematobium\) eggs, and 587 (55.7\%) of these samples were positive, with geometric mean infection intensity of 25.9 eggs/10 mL urine. Among the tested individuals, the age distribution was 5 years (35), 6–9 years (153), 10–13 years (158), 14–17 years (128), 18–21 years (87), and \(\geq\) 22 years (492). Within age groups, infection intensity levels were comparable between the two study periods, with most age groups recording higher infection intensity levels in 2000 (Figure 2). However, the overall infection intensity was lower in 2000 compared with 2009 because of the lower proportion of children among those people tested in 2009. Infection prevalence was lower across all age groups in 2009, with an overall decline in prevalence of 12.5\% (\(\chi^2 = 28, P < 0.0001\)).

**Spatial infection patterns.** Global patterns. In 2009, the significantly high spatial aggregation (\(K\) function) of households with high-intensity (> 100 eggs/10 mL urine) infection observed in 2000\textsuperscript{3} was not repeated when all ages were pooled, indicating that heavy household-level infections were distributed more randomly throughout Milalani village. When the effect of age was considered and different ages were considered separately, a small significant peak in global clustering at 300–400 m was detected for 10- to 13-year-old children. In contrast, when applied to the same spatial extent for the 2000 Milalani infection data, weighted \(K\) function revealed global clustering of household infection density, with peak clustering at around distances of 150 m for the pooled data and each age group, indicating tight clustering of households with high intensity of infection.

Local patterns. When we tested for local clusters of infection using the \(Gi(d)\) statistic, we detected no significant local clusters in 2009, either for the pooled data or when age groups were considered separately. This lack of local clustering stands in sharp contrast to the 2000 Milalani data (again using the 2009 spatial extent), when significant local clusters of households with high-infection levels were detected for all age groups at distances of 500 m from Nimbodze pond.

**Focal clustering of infection.** Focal clustering of infection density around Nimbodze pond varied by age and between the

![Figure 2. Prevalence of infection in Milalani village in 2000 (—Δ—) and 2009 (—□—) and geometric mean intensity of eggs/10 mL urine in 2000 (open bars) and 2009 (shaded bars).](image-url)
two study periods. In 2009, there were no noticeable infection clusters in children 6–9 and 10–13 years, and the only significant clustering of infection was found in adolescents (14–17 years) at a distance of 300 m from Nimbodze (Figure 3A). When patterns of infection density are compared with data from 2000, differences in the patterns of clustering are apparent. In 2000, significant clustering of high-infection density for all age groups of children was detected starting at different distances from Nimbodze (Figure 3B). Clustering of infection levels in 6- to 9-year-old children was significant starting at 350 m, and for children 10–13 and 14–17 years, significance was from 600 to 700 m. In all age groups in 2000, clustering was significant up to a distance of 1,100 m. High-infection levels for all age groups were significantly clustered at a range of 400–800 m, but the degree and extent of clustering were higher for 6–9 and 10–13 years than for the older children (14–17 years). Infections were not clustered around any other permanent or temporary pond in either 2000 or 2009.

Rainfall patterns. In general, rainfall followed the typical bimodal pattern of the Kenyan coast. The long rainy season usually occurs from April to June, and the short rainy season occurs between October and December. There is also a short dry season between July and September and a long dry season between January and March (Table 1). The classification of seasons on the coast is not fixed; sometimes, overlaps of about 1 month may occur. Major peaks in the amount of rainfall were usually reported in April through June from 2001 to 2010, with the exception of 2006, when the major peaks were seen in October/November (Table 1).

In 2001 and 2003, after sufficient rains during the long rainy season, all the ponds were filled but were not sustained throughout the year, because the short rains failed (Table 1). The rainfall pattern experienced in 2002 (small amounts of rain over a long period of time) did not allow accumulation of sufficient amounts of water to fill most ponds. There was insufficient rainfall in both of the rainy seasons to fill most ponds in 2004, 2005, 2008, and 2009 (Table 1). Thus, our study area has experienced two cycles of drought over the 2001–2010 decade: from 2002 to 2006 and from 2008 to 2010 (Figure 4).

Although we characterize the decade of 2001–2010 as a drought period, rainfall patterns for the period varied, and heavy rains did fill up the transmission sites in April–June of 2001, 2003, 2006, 2007, and 2010. However, lack of sufficient rains after these rainfall events resulted in the transmission sites drying up within a few months, even in the years with average amounts of total rainfall, resulting in no long-term establishment of infected *B. nasutus* snail populations.

**DISCUSSION**

In our previous studies in Milalani village, we detected highly significant spatial clusters of infected school children on the household level in 1984 and 2000. No mass drug administration for schistosomiasis has taken place in the village during the last decade, and routinely, only a small proportion of those people infected seek treatment (data not shown). Thus, the disappearance of schistosomiasis infection clusters and the relatively modest (although significant) 12.5% decline in prevalence from 55.7% to 43.2% during the current study period are noteworthy. We attribute these changes to the

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**Table 1**

<table>
<thead>
<tr>
<th>Year</th>
<th>January to March</th>
<th>April to June</th>
<th>July to September</th>
<th>October to December</th>
<th>Total</th>
<th>Pond*</th>
</tr>
</thead>
<tbody>
<tr>
<td>2001</td>
<td>150.5</td>
<td>1,145</td>
<td>118</td>
<td>57</td>
<td>1,470.5</td>
<td>Water/dry</td>
</tr>
<tr>
<td>2002</td>
<td>186.8</td>
<td>336.2</td>
<td>401.9</td>
<td>353.2</td>
<td>1,278.1</td>
<td>Dry</td>
</tr>
<tr>
<td>2003</td>
<td>63.9</td>
<td>865.1</td>
<td>183.9</td>
<td>212.3</td>
<td>1,325.2</td>
<td>Water/dry</td>
</tr>
<tr>
<td>2004</td>
<td>128.4</td>
<td>279.4</td>
<td>113.4</td>
<td>256.7</td>
<td>1,172.5</td>
<td>Dry</td>
</tr>
<tr>
<td>2005</td>
<td>109.3</td>
<td>533</td>
<td>279.8</td>
<td>250.4</td>
<td>727.3</td>
<td>Water/dry</td>
</tr>
<tr>
<td>2006</td>
<td>94.9</td>
<td>625.6</td>
<td>340.8</td>
<td>727.3</td>
<td>1,788.6</td>
<td>Water/dry</td>
</tr>
<tr>
<td>2007</td>
<td>132</td>
<td>902.6</td>
<td>236.7</td>
<td>147</td>
<td>1,418.3</td>
<td>Water</td>
</tr>
<tr>
<td>2008</td>
<td>59.8</td>
<td>413.9</td>
<td>143.5</td>
<td>264.2</td>
<td>881.4</td>
<td>Dry</td>
</tr>
<tr>
<td>2009</td>
<td>56.8</td>
<td>417.8</td>
<td>210.9</td>
<td>335.9</td>
<td>1,021.4</td>
<td>Dry</td>
</tr>
<tr>
<td>2010</td>
<td>135.6</td>
<td>1,514.7</td>
<td>218.7</td>
<td>265.1</td>
<td>2,134.1</td>
<td>Water/dry</td>
</tr>
<tr>
<td>Average rainfall</td>
<td>111.8</td>
<td>703.3</td>
<td>224.8</td>
<td>286.9</td>
<td>1,526.8</td>
<td></td>
</tr>
</tbody>
</table>

*Water/dry denotes that most ponds did not have water throughout the year; dry denotes that most ponds were dry throughout the year, whereas water means that most ponds had water throughout the year.*
disappearance of the most influential transmission sites and resulting changes in water contact. The transmission sites in Msambweni area, including Nimbodze pond, which is described extensively elsewhere, have largely remained dry during the course of the study period and short-lived when available because of unstable rains. Marked declines in infection prevalence after prolonged droughts have been reported from Ethiopia, Nigeria, and Zimbabwe.

Rainfall patterns have changed considerably in the study area, with two drought cycles reported over the last 10 years (Figure 4 and Table 1). Not only did the transmission sites remain mostly dry during the drought years, but also, even in the few years with average rainfall (except for 2006), the short rains (October and November) did not fill the ponds in preparation for the long dry season (December to March) failed. Snails survive the seasonal drying out of habitats by aestivation, but they are likely to die when drought conditions persist for more than 8 months and when these conditions are more frequent. This condition is precisely the situation that has been ubiquitous in our study area over the last 10 years, and it is a likely explanation for the continuing absence of intermediate host snails from the relatively large ponds that were historically the important transmission foci (i.e., Nimbodze, Kiziamkala, and Maridzani). Indeed, this condition is still the situation in March of 2010, 10 months after being refilled by the heavy 2010 April to May rains.

_Bulinus_ snails usually aestivate around the margins of the transmission sites and only emerge when the sites are flooded. As a result of this strategy, it seems that, although in the relatively large ponds, most of the aestivating snails and especially those snails that were infected died, because the ponds remained dry for 12 months or more, some snails did survive in temporary ponds, such as Chiziani, that are more frequently filled, even when the rains are insufficient to fill the larger ponds. In aestivation, uninfected snails usually survive longer than the infected snails, and therefore, snails that do survive the drought are likely not infected. These findings also suggest that transmission is currently sustained by temporary snail habitats rather than the more permanent ones that were in the past foci of transmission, although spatial analysis did not detect clustering of infection around the temporary pond, which is > 1 km from Milalani. Whether this relatively high importance of ephemeral habitats will hold in the future is a subject for additional investigations.

Overall, the loss of local/focal clustering of infection in the community that continues to maintain high-infection prevalence and intensity indicates that the process of schistosomes’ transmission is more uniformly distributed across the community. This shift in spatial infection pattern from clustering around well-defined transmission hot spots means that, in this setting, there is a need to consider the value of community-wide interventions rather than just those measures concentrated on focalized efforts aimed at only a few hot spots that may not drive transmission at all times.

The absence of snails in Nimbodze pond for 9 months after being refilled may be explained in part by the destruction of the well-defined dispersal corridors among irrigation scheme and stream/river described by us elsewhere. Since 2008, there is a shift from rice cultivation (2000–2007) to sugar cane cultivation in the western parts of Nimbodze, especially near the Koromonjo reservoir, the main source of reflooding of Nimbodze and other ponds after sufficient rains. Sugar cane cultivation may have impeded snail dispersal through the two routes, which was identified by Clennon and others. Unlike rice cultivation, sugar cane cultivation reduces water pooling, subsequently restricting habitat connectivity.

In conclusion, inadequate and unpredictable precipitation in Msambweni area during the study period has diminished the historical importance of Nimbodze as a focus for urinary schistosomiasis transmission. The long-term drought and hydrological changes have resulted in the absence of _Bulinus_ intermediate host snails in the pond for at least 9 months after the refilling of the pond. Our current findings point to the potential impact of climatic variability and anthropogenic changes on transmission patterns of _S. haematobium_. Such changes need to be considered when implementing urinary schistosomiasis surveillance and control programs.

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REFERENCES


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REFERENCES