A search for snail-related answers to explain differences in response of *Schistosoma mansoni*

to praziquantel treatment among responding and persistent hotspot villages along the Kenyan

shore of Lake Victoria

Running head: Snail-related answers for *S. mansoni* persistent hotspots

M. W. MUTUKU[1,2]*, M. R. LAIDEMITT[3], B. R. BEECHLER[4], I. N. MWANGI[1], F. O. OTIATO[5], E.

L. AGOLA[1], H. OCHANDA[2] , B. KAMEL[3], G. M. MKOJI[1] , M. L. STEINAUER[6], E. S. LOKER[3]

1. Centre for Biotechnology Research and Development, Kenya Medical Research Institute,
Nairobi, Kenya.

2. School of Biological Sciences, College of Biological and Physical Sciences, University of
Nairobi, Nairobi, Kenya

3. Center for Evolutionary and Theoretical Immunology, Parasitology Division, Museum of
Southwestern Biology, Department of Biology, University of New Mexico, Albuquerque,
New Mexico, United States of America

4. College of Veterinary Medicine, Department of Biomedical Sciences, Oregon State
University, Corvallis, Oregon, United States of America

5. Influenza Surveillance Program, Centers for Disease Control and Prevention, Nairobi,
Kenya

6. Department of Basic Medical Sciences, Western University of Health Sciences, Lebanon,
Oregon, United States of America

* Corresponding author

Centre for Biotechnology Research and Development, Kenya Medical Research Institute, P.O.

Box 54840-00200 Nairobi, Kenya. Email: mmutuku@kemri.org

Abstract

Following a four-year annual praziquantel treatment campaign the resulting prevalence

of *S. mansoni* was seen to differ among individual villages along the Kenyan shore of Lake

Victoria. We have investigated possible inherent differences in snail-related aspects of

transmission among such 10 villages, including six persistent hotspot (PHS) villages (≤30%

reduction in prevalence following repeated treatments) located along the west-facing shore of

the lake, and four PZQ-responding (RESP) villages (>30% prevalence reduction following

repeated treatment) along Winam Gulf. When taking into account all sampling sites and times

and water hyacinth presence/absence, shoreline-associated *B. sudanica* from PHS and RESP

villages did not differ in relative abundance or prevalence of *S. mansoni* infection. Water

hyacinth intrusions were associated with increased *B. sudanica* abundance. The deeper water

snail *Biomphalaria choanomphala* was significantly more abundant in the PHS villages and

prevalence of *S. mansoni* among villages both before and after control was positively correlated

with *B. choanomphala* abundance. Worm recoveries from sentinel mice did not differ between

PHS and RESP villages, and abundance of non-schistosome trematode species was not

associated with *S. mansoni* abundance. *Biomphalaria choanomphala* provides an alternative,

deepwater mode of transmission that may favor greater persistence of *S. mansoni* in PHS

villages. As we found evidence for ongoing *S. mansoni* transmission in all 10 villages, we

conclude conditions conducive for transmission and reinfection occur ubiquitously. This argues

for an integrated, basin-wide plan for schistosomiasis control to counteract rapid reinfections

facilitated by large snail populations and movements of infected people around the lake.

Key words: *Schistosoma mansoni*, *Biomphalaria sudanica*, *Biomphalaria choanomphala*, Lake

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INTRODUCTION

One of the most prevalent and persistent of the world's neglected tropical diseases is human schistosomiasis.[1] There is currently considerable momentum to bring schistosomiasis under control and to proceed to elimination efforts.[2-4] An essential weapon in the elimination of schistosomiasis is the drug praziquantel (PZQ) which has been used extensively in a variety of control programs and has considerably reduced prevalence and intensity of infection.[5] There is also a growing appreciation for the need for integrated control programs taking into account sanitation, provision of safe water, education, and acknowledging the fundamental role played by freshwater snails as vectors of the disease.[6,7] The persistent success of schistosome parasites is owed substantially to their molluscan hosts which support the prolific production of human-infective cercariae, and that often exist in huge populations across a variety of freshwater habitats.

Africa harbors 90+% of the global burden of *S. mansoni*.[8] In Kenya, there are three taxa of *Biomphalaria* snails that perpetuate transmission: 1) *B. pfeifferi*, whose distribution includes tributaries feeding Lake Victoria, and in small impoundments and both seasonal and perennial streams throughout the country, except in the tropical lowland belt along the coast; 2) *B. sudanica*, mainly found along the shores of Lake Victoria and Lake Jipe and their surrounding swamps; and 3) *B. choanomphala*, a deeper water inhabitant of Lake Victoria.[9,10] The latter two taxa are likely members of the same species, and are frequently referred to as ecophenotypes or ecomorphs.[11-13]

Prevalence of intestinal schistosomiasis in Kenya is highest (>50%) in the Mwea irrigation scheme in central Kenya and in the Lake Victoria basin in western Kenya.[14] Recent studies assessing the prevalence of *S. mansoni* infection in school children in the lake basin reported overall prevalences of 60.5% and 69%, with villages on lake islands having 2-fold higher prevalence rates than mainland villages.[15,16] Schools closer to the lake also had higher

prevalence rates than those further from the lake.[15] Both higher prevalences and likelihood of

reinfection following treatment would logically be associated with more frequent contact with

lake water containing infected snails.[17]

Several chemotherapy-based initiatives have been undertaken to control

schistosomiasis in the Lake Victoria region.[17-20] Among them, the Schistosomiasis Consortium

for Operational Research and Evaluation (SCORE) project implemented school-based and

village-wide mass drug administration (MDA) in varying treatment regimens over 4 consecutive

years (2011-2015) for 150 Kenyan villages situated on or near the eastern shore of Lake

Victoria.[17] Villages were randomized with respect to six different treatment strategies (study

arms) which involved combinations of school-based or community-wide treatments, with 3 study

arms receiving annual treatment and 3 receiving treatment twice over a four-year period.

Regardless of study arm or age group of treated individuals, using spatial scan statistics a

persistent hotspot area was identified in western Siaya County comprising west-facing villages

along the open waters of the lake.[17] Prevalence and intensity of infection did not decrease as

much as it did in areas outside of the persistent hotspot area, the latter including villages along

the more protected shores of Winam Gulf.[17] Although this study could not clearly identify factors

driving persistence of infection at hotspots, it reported an association between schistosome

persistence and vegetation density (as measured by remote sensing) indicating a potential role

for snail ecology.

Particular locations where the prevalence or intensity of schistosome infection does not

fall to expected levels despite well-implemented, multiyear, MDA have been termed "persistent

hotspots" or PHS, in contrast to responding villages (RESP) in which intensity and prevalence

decline as expected. Why do some villages fail to respond to multi-annual treatment with PZQ?

Several explanations can be offered including variability in efficacy of praziquantel among worm

populations in different areas, levels of sanitation or availability of toilets, extent of drug

coverage or compliance, availability of safe water supplies, extent of water contact, defecation

behavior, or the extent to which they experience population movements such as resulting from

fishing activities or proximity to ferry landings. [19,22-25]

Conditions associated with snail vectors may also vary considerably including the nature

of the snail species present, their abundance and local compatibility with schistosomes,

longevity of infected snails and the rate at which they produce cercariae, and their patterns of

distribution relative to water contact sites and anthropogenic disturbance.[22,26-33] Shoreline and

bottom conditions can also play important roles in efficacy of transmission.[9,22,24,32,34]

In addition to the shoreline-associated vegetation effects noted by Wiegand et al. as a

possible factor associated with PHS, the dramatic presence and movements of large floating

mats of the introduced water hyacinth, *Eichhornia crassipes,* are other factors in lake ecology

that cannot be overlooked.[17] Hyacinths have been suggested to provide a suitable environment

for schistosome-susceptible snails and to facilitate their movements around the lake.[35] Plummer

noted that significantly more *B. sudanica* were recovered from experimental enclosures

containing hyacinths than control enclosures lacking them, and Ofulla et al. also concluded that

B. sudanica was significantly associated with hyacinths in Nyanza Gulf (now referred to as

Winam Gulf), Kenya.[36,37] Winam Gulf is one of the areas of the lake most affected by hyacinths,

and four of the ten villages we investigated are found on the Gulf.[38] Although hyacinth mats

seem to favor snails in the short term, it has been noted that longer-term residence of hyacinth

mats may be harmful to snails because of deficits in light penetration, water circulation, and

oxygen availability, the latter due to decomposing hyacinths.[36]

Here we focus on the relationship of *Biomphalaria* snails to PZQ-based control efforts for

S. mansoni in and around the Kenyan shoreline of Lake Victoria, one of the world's major

hyperendemic foci of human schistosomiasis. Upon the request of the SCORE program, during

the two years immediately following the cessation of their MDA project, on four separate

occasions, we surveyed *Biomphalaria* populations and their schistosome infection rates

adjacent to 10 villages along the Kenyan shore of Lake Victoria with different degrees of

responsiveness to treatment. In each of two collecting sites per village, we took note of the

general condition of the habitat and searched for snails in shoreline-associated and deeper

water habitats in the lake, as well as in smaller habitats adjacent to the lake. Building on the

results of Wiegand et al., we investigated the hypotheses that PHS and RESP villages differ

with respect to the species of vector snails present, relative snail abundance, likelihood that the

snails are infected with schistosomes or other trematodes, and numbers of cercariae in the

water as measured by infection rates of sentinel mice.[17] All of these factors could contribute to

persistently high prevalence in the face of control operations.

MATERIALS AND METHODS

Study sites

The study involved ten villages along the shores of Lake Victoria, western Kenya (Figure 1) that were part of the Wiegand et al., study already mentioned.[17] The villages had annual school-based mass drug administration (MDA) for 4 consecutive years (Table 1). Following one of the definitions of Kittur et al., six of the villages (Minya, Agok, Migiro, Miyandhe, Kanyibok, Usenge) were persistent hotspots (PHS), locations where the absolute change in *S. mansoni* prevalence from the beginning of the control program to the end was ≤30%.[21] By contrast, for the four remaining villages (Kotieno, Seka Dok, St. Douglas Weta, Mumbo) recorded a drop in prevalence >30% and are considered responding (RESP) villages. In each village, two shoreline habitats were identified where there was evidence of human-water contact activities, and these were established as our sampling sites. Each of the 20 sampling sites was visited four times (April 2016, August 2016, January 2017 and May 2017). .

Snail Sampling

Two methods were used to collect snails from each of the 20 designated sampling sites: scooping from the shore and dredging from a boat. From the shore, two experienced collectors from our team scooped snails for 30 minutes per sampling site using long-handled scoops (steel sieve with a mesh size of 2 × 2 mm, supported on an iron frame). Shoreline habitats were typical for Lake Victoria consisting of emergent and submerged vegetation that was swept by scoops for snails. Snails were also collected offshore by passing a dredge (0.75 m long and 0.4 m wide with attached sieve, 2 × 2 mm mesh size) along the bottom. At each sampling site dredge hauls covering a combined length of 150 meters was made, beginning at one meter depth and extending perpendicular to the shore to 10m depth. Live snails were picked from the dredged material. All snails were taken to the laboratory at KEMRI, Center for Global Health

162 Research, Kisumu where they were sorted into species based on shell morphology

163 characteristics, using standard taxonomic identification keys, counted and screened.[39]

Snail screening

Each snail was placed in an individual well of a 24-well plastic culture plate containing 1

ml of aged de-chlorinated tap water. The plate was placed in indirect sunlight for 2 hours

between 10:00-13:00hr. Individual wells were then examined with the aid of a dissecting

microscope for presence of cercariae. Using standard identification keys, cercariae were

identified to basic taxonomic groups.[40] Non-shedding snails were maintained in the laboratory

for four weeks and screened again to enable detection of snails that harbored pre-patent

infections at the time of initial collection.[31] Some cercariae were further identified with molecular

methods as parts of another study.[41] Additionally, samples of mammalian schistosome

cercariae recovered from *B. sudanica* (each sample consisting of 4 cercariae) were extracted

(QIAamp DNA Micro Kit, to elution volume of 45µl) and primers for the mitochondrial NADH

dehydrogenase subunit 5 gene (*nad5*) used for amplification, following which bands were

visualized on a gel (302 bp expected band size for *S. mansoni* and 800 bp for the related

schistosome *S. rodhaini*) as a check for the identity of the schistosome recovered.[28]

Sentinel mice exposures

Mice were exposed to lake water at each near shore sampling site in floating cages that

ensured the mice were exposed to a depth of no more than 5 mm. Each cage was made from a

plastic container measuring 20 X 14 X 7 cm with the bottom of the container replaced with wire

mesh with openings measuring 5 x 5 mm. Styrofoam blocks were attached to the plastic

container to insure the cage floated at the desired position. Lids of the containers were dark to

provide shading and were perforated with 5 mm diameter holes to allow air circulation. The

abdomen of each mouse was shaved to facilitate cercariae penetration. For each of the

shoreline sampling sites, five cages each with 2 mice were tethered adjacent to locations where

snails were collected (total of 20 mice per village per sampling time). Mice were in contact with

the water for 3 hours (11:00 hr – 14:00 hr), an interval when shedding of cercariae by *S.*

mansoni-infected snails is expected to be at its peak.[42] Previous experience under lakeside

conditions indicated that a three-hour period of exposure did not jeopardize survival of the mice,

which was confirmed by the high survival rates of the mice retrieved. Mice from each sampling

site were then marked with an identifying ear tag and maintained together until perfused at eight

weeks post-exposure. Recovered worms were counted and tabulated with respect to village,

sampling site and sampling time.[43]

Ethics statement

All experiments involving mice were approved by institutional animal care and use

committees   (IACUCs)   at   the   Kenya   Medical   Research   Institute (Protocol

KEMRI/ACUC/03.10.15) and at the University of New Mexico (Animal Welfare Assurance #

D16-00565 (A4023-01, expiration 8/29/2021). All protocols and practices for the handling and

manipulation of mice were in accordance with the guidelines of the American Veterinary Medical

Association (AVMA) for humane treatment of laboratory animals.

Statistical analysis

To determine if PHS sites differed from RESP sites in snail-related factors, we compared

them with respect to relative abundance of each snail (*B. sudanica* or *B. choanomphala*), the

prevalence of infected snails, and the number of adult schistosome worms collected from the

sentinel mice (as a measure of the force of infection). Collection sites were nested into village

and were compared across the four time points. Because of the potential effects of large rafts of

floating water hyacinths that temporarily occupy some sites, we included the presence or

absence of hyacinths in the models.

Snail relative abundance: To determine whether relative abundance of either *B.*

sudanica or *B. choanomphala* differed between PHS and RESP villages, between sites with

hyacinths and no hyacinths, or over time, we performed generalized linear mixed models in R

using the package glmmTMB, with the random effect of village and using the family negative

binomial 1 (nbinom1).[44] Prior to settling upon this model structure we also explored a model that

included a nested random effect of sample site within village but the model fit was not

significantly better (anova, p>0.05) so we report the reduced model of only village as a random

effect.

Formula: glmmTMB(Abundance of either snail ~ Responder/Not + Hyacinth Y/N + timepoint +

(1 | School),family=nbinom1(link=log))

Prevalence of infected snails: To determine whether the likelihood of detecting an

infected snail was different in a RESP vs. PHS village, or in sites with and without hyacinths we

performed a generalized linear mixed model using R with the packages lmerTest and lme4 with

the random effect of village and the family of binomial (logit link).[45] We also included the offset

term of number of snails found, rescaled to be on the same scale as the binomial variables by

dividing by 2 standard deviations.[46] This is because there is a strong positive correlation

between the likelihood of finding an infected snail and the total number of snails found

(est=0.592+/-0.148, p=0.00005, statistics reported from the same model as described above but

with total number of snails found included as a main effect rather than an offset). We analyzed

only data for *B. sudanica* because the number of infected *B. choanomphala* was prohibitively

small.

Formula: glmer(smansoni.presence   Y/N~   Responder/Not+   Hyacinth   Y/N+   (1|

timepoint)+(offset(rescaledtotalsudanicacount))+(1| School),family=binomial(link="logit"))

Sentinel mice: To evaluate whether infection prevalence in the sentinel mice differed

between PHS and RESP villages, between sites with and without hyacinths and between time

points we performed a generalized linear mixed model in R using the package glmmTMB with a

random effect of village, and family of negative binomial 1. We also included the term of total

snails found at each site to account for snail relative abundance, however this variable was

rescaled by dividing by 2 standard deviations to place it on the same scale as the binomial

variables.

Formula: glmmTMB(total.number.of.schistosomes.in.mice ~ Responder/Not +   Time +

total.biomphalaria.snails + (1 | Village), family=nbinom1(link="log"))

We also evaluated whether the likelihood of finding a schistosome positive sentinel

mouse differed between PHS and RESP villages, between sites with and without hyacinths and

between time points. The only difference between this model and the one described above is

that the dependent variable is binomial (infected mouse found or not), rather than the burden

within the infected mouse. For this we used a model structure similar to the prevalence of

infected snails.   We performed a generalized linear mixed model with family of binomial

(link=logit) and the random effect of village. Once again we accounted for the total number of

snails collected, but rescaled it to be on the same scale as the binomial variables.[46]

Formula: glmer(smansoni.presenceinmice Y/N~ Responder/Not+ Hyacinth Y/N+ (1|

timepoint)+(offset(rescaledtotalbiomphalariacount))+(1| Village),family=binomial(link="logit"))

Correlation of human prevalence and snail density: To determine if the abundance

of snails at a particular site was associated with schistosome infection rates in humans, we

correlated the total number of snails collected at a village (summed between collecting sites and

all 4 time points) with both the initial and the final prevalence in humans at the corresponding

village using Pearson's correlation in Graphpad Prism 7.0. We performed separate analyses for

each snail species.

Biodiversity: To assess whether the likelihood of detecting a schistosome-infected *B.*

sudanica at a site correlated with the presence of other digenetic trematodes in the snails, we

included each of four commonly collected cercarial types (echinostome, strigeid,

xiphidiocercariae, and amphistome) as a main effect in the same model as described above in

the prevalence of infected snails.

Formula: glmer(smansoni.presence Y/N~ Responder/Not+ Hyacinth Y/N+amphistomes

Y/N+xiphido   Y/N,   echinostomes   Y/N,   strigieds   Y/N   +   (1| timepoint)   +

(offset(rescaledtotalsudanicacount)) + (1| village),family=binomial(link="logit"))

RESULTS

Descriptive overview: A total of 12,156 snails (10,249 *B. sudanica* and 1,907 *B. choanomphala*) were collected from the study sites during four collection periods (Table 2, Figure 2). No *B. pfeifferi* were found in the lake habitats sampled. We also searched small streams or impoundments near the villages and found them to be muddy, prone to drying and without *Biomphalaria* snails. At some times and collecting sites, hyacinth mats covered the water surface and were so dense as to preclude dredging (Figure 2). *Biomphalaria sudanica* was recovered along the shoreline from all 20 sampling locations and *B. choanomphala* was dredged from deeper water at all 12 sampling locations in PHS villages whereas Mumbo was the only RESP village from which *B. choanomphala* was found (Table 2). Of note is that Mumbo is located closer to the PHS villages than the remaining three RESP villages (Figure 1). *Biomphalaria choanomphala* was also sometimes found while scooping from shoreline collecting sites in three PHS locations, Kanyibok, Migoro and Usenge. At Usenge Beach, the shoreline *B. choanomphala* population persisted for at least five months on submerged vegetation along with another snail customarily recovered from deeper water, *Gabbiella humerosa*. Fifteen meters away, in a patch of emergent vegetation, *B. sudanica (*but not *B. choanomphala)* was found. *Biomphalaria choanomphala* was also occasionally recovered from the shore intermingled with *B. sudanica*, as at the Anyanga beach sampling site at Kanyibok.

Hyacinth mats were noted during 3 of the collective total of 48 total visits (6.25%) we paid to PHS sampling sites, and two of these times precluded dredging at such sites (Figure 1). Hyacinth mats were found at 11 of 32 collective visits (34.3%) at RESP sites, and at five of these times precluded dredging.

Snail relative abundance: Overall, we collected far more *B. sudanica* than *B. choanomphala* (Table 2, Figure 2). However, it should be noted that because our collecting methods differed for each snail taxon, these numbers are not directly comparable (e.g. there is

not necessarily a larger population of *B. sudanica* than *B. choanomphala* in the lake). The

relative abundance of snails was highly variable among sites and collection times. For instance,

at Miyandhe, one of the PHS villages, with its rocky, wave-swept lakeshore habitat, few snails

were recovered relative to all the other villages. It was not unusual for snails to increase or

decrease several fold between collection time points at the same habitats (Figure 2).

In Lake Victoria*, B. sudanica* is considered a more important intermediate host of *S.*

mansoni than is B. choanomphala because *B. sudanica* with its shallow water habitat use is

more likely to contact miracidia from human feces deposited along the shore. Thus, we

predicted that snail driven differences between PHS and RESP sites would be reflected in this

species. However, we found that PHS sites did not contain more *B. sudanica* snails than RESP

sites (β = -0.0664, SE = 0.4603, P = 0.8854, Figure 3A). We note however that the three PHS

villages (Minya, Kanyibok and Usenge) with the highest residual prevalence of *S. mansoni* are

also the three villages with the highest overall counts of *Biomphalaria* snails. *B. sudanica*

density was positively associated with the presence of hyacinths (β = 0.7704, SE = 0.2980, P =

0.0097, Figure 4) and collecting time point (β = 0.37912, SE = 0.09532, P < 0.0001).

In contrast to the *B. sudanica* results, *B. choanomphala* was more abundant at PHS

sites (Figure 3B) compared to RESP sites (β = -2.1644, SE = 0.6975, P = 0.0091) and the

presence of hyacinths and time point were not significantly associated with *B. choanomphala*

density. Thus, results of the snail analysis indicated that only the deepwater taxon, *B.*

choanomphala, differed between PHS and RESP sites. The other novel finding was that *B.*

sudanica abundance was positively associated with the hyacinths.

Prevalence of infected snails: The overall number of positive snails and prevalence for

S. mansoni recovered among all 10 villages examined were higher for *B. sudanica* (83/10,249,

0.81%) than for *B. choanomphala* (3/1,907, 0.16%). Although few *B. choanomphala* were found

positive for *S. mansoni*, all came from PHS villages. For *B. sudanica*, the total number of *S.*

mansoni-infected snails and overall prevalence was higher in PHS villages (60/5,346, 1.12%)

than from RESP villages (23/4,903, 0.47%). As a check on the identity of schistosome cercariae

recovered, we amplified the *nad5* gene from cercariae from 20 different positive snails, five from

each of the time points, and representing several beach sites where schistosome cercariae

were found.[28] All 20 samples were verified as *S. mansoni* based on *nad5* band size, with one

sample (Maguti Beach) exhibiting a double infection with both *S. mansoni* and *S. rodhaini*.

The distribution of *S. mansoni* infections among snails from the different villages is

difficult to succinctly characterize. Two of the PHS villages with the highest residual post-

treatment prevalences of *S. mansoni* in children, Kanyibok and Usenge, had the highest overall

prevalence of positive snails (1.57% and 1.42% respectively (Table 2, Figure 2). From each

locality though, most of these infections were found in a single collection raising the possibility

they might have originated from one or a few human contamination events. Also, we found few

shedders in Minya and Migoro, and none in Miyandhe, all PHS villages, whereas Kotieno, a

RESP village, yielded the third most shedding snails, many of which were also found at a single

time point. Ten of 24 sampling times (41.7%) from PHS villages yielded positive snails whereas

7 of 16 sampling times (43.8%) from RESP villages yielded snails positive for *S. mansoni*.

Consequently, the number of *B. sudanica* infected with *S. mansoni* did not differ between PHS

and RESP sites when time point, hyacinths, and village were all taken into account. There was

a negative relationship that approached statistical significance between the number of infected

snails collected and the presence of hyacinths at a site (β = -0.12362, SE = 0.07186, P =

0.0854), a point considered further in the discussion.

Sentinel mice: Worm recoveries from sentinel mice (Figure 2) were low (53 worms

recovered, or 0.066 worms per mouse, or 0.022 worms/mouse/hour of exposure). The greatest

number of worms acquired by a single mouse was 7. Of the total number of worms recovered,

40 (75.5%) were from PHS villages (0.083 worms/mouse) while 13 (24.5%) were from RESP

villages (0.041 worms/mouse). Although the total number of adult schistosomes collected from

mice caged in PHS sites was greater than those caged in RESP sites, there was no statistical

difference between them when village, hyacinths, and time point were considered (Figure 5, β =

-0.36479   SE = 0.5720, P = 0.5237). Examination of the viscera of positive sentinel mice

revealed only the presence of *S. mansoni* eggs; no *S. rodhaini* eggs were found.

Correlation between snail abundance and human prevalence: Total number of *B.*

sudanica collected at a village was not significantly correlated with *S. mansoni* prevalence in

humans (Figure 6) either before or after the drug treatment intervention (before: Pearson r = -

0.0369, p = 0.460; after: Pearson r = -0.009, p = 0.49). However there was a strong correlation

between the number of *B. choanomphala* collected at a village with *S. mansoni* prevalence in

humans at the village (Figure 7) both before and after mass drug administration (before:

Pearson r = 0.587, r^2 = 0.345, p = 0.037; after: Pearson r = 0.9034, r^2=0.816, p = 0.0002).

Infection of *B. sudanica* with other digenetic trematodes: In all 10 villages, the

number of trematode infections other than *S. mansoni* observed from *B. sudanica*

(Supplementary Table 1) exceeded those of *S. mansoni* (totals of 241 and 83, respectively). In

PHS villages 60 of 184 trematode infections were of *S. mansoni* (32%), whereas in RESP

villages, 23 of 150 (15%) were of *S. mansoni*. Overall prevalence of trematode infections

including *S. mansoni* was similar between PHS and RESP villages (3.5 and 3.1%, respectively).

Echinostomes were the most abundant non-*S. mansoni* infections with 112 snails infected from

the ten study villages. We did not observe any snails to have double infections (e.g. to be

simultaneously shedding two different kinds of cercariae). Although *S. mansoni* infections were

consistently outnumbered by the aggregate totals of all other digenetic trematode infections,

there were no statistical associations between the number of snails infected with *S. mansoni*

and any of the other trematode species measured with the model employed.

With respect to having high overall total trematode prevalence, three villages stood out

from the rest, the two PHS villages Kanyibok and Usenge and the one RESP village Kotieno

(Supplementary Table 1). All three villages had relatively high *Biomphalaria* populations, and

yielded the most *S. mansoni* infections found in snails relative to other villages (Table 2).

DISCUSSION

The SCORE-supported PZQ-based treatment program in the Lake Victoria region in

western Kenya noted that PHS villages comprised approximately a third of the 150 study village.

The village prevalence was determined by testing 9-12 year old students each year. These

students had a relative risk of infection that was nearly 4 times higher than for students outside

the PHS area.[17] The SCORE program is by no means unique in identifying difficulties in

reducing prevalence of infection in some villages adjacent to Lake Victoria.[19] The PHS for *S.*

mansoni transmission identified by Wiegand et al., was in Siaya County, Bondo region, of

western Kenya, associated with west-facing villages adjacent to the open, unprotected waters of

the lake. In contrast, the PZQ RESP villages faced the more protected waters of Winam Gulf.[17]

Among several factors postulated to be associated with the PHS we considered in our

investigation were larger snail populations, or presence of different species of vector snails

factors.[17]

Of the three taxa of *Biomphalaria* commonly implicated in transmission of *S. mansoni* in

Kenya, two were collected in this study, the shoreline-associated taxon *B. sudanica* and the

deeper water taxon *B. choanomphala*. There was no obvious indication that *B. pfeifferi* played a

role in *S. mansoni* transmission in any of our study villages, as no snails of this species were

found in the lake, or in streams or small impoundments within the study localities. We found *B.*

sudanica at all 20 collecting sites and it was typically abundant as it is along much of the Lake

Victoria shoreline.[9,11,22,34] We recovered approximately five times as many *B. sudanica* as *B.*

choanomphala.

When taking into account all sampling sites, sampling times and presence of hyacinths,

mean abundance of *B. sudanica* did not differ statistically between PHS and RESP villages. We

did note a significant trend for more *B. sudanica* to be present from our collecting sites if mats of

hyacinths were present at the time we took our samples. Hyacinth mats were found in collecting

sites from both PHS and RESP villages but were more common in the RESP villages located

along Winam Gulf, an area known to be prone to hyacinth intrusions.[38] The recovery of *B.*

sudanica from hyacinths and the association we noted for them with hyacinths supports the

assertion that floating hyacinth mats favor the dispersal of *B. sudanica* along the lakeshore.[35]

Hyacinth movements are likely to have major effects on the population structure of *B. sudanica*

in the lake.

By contrast, *B. choanomphala* was significantly more likely to be recovered from PHS

than RESP villages. We found *B. choanomphala* from all 12 of the collecting sites in PHS

villages and from only 2 of the 8 collecting sites from RESP villages, both from the village of

Mumbo. Mumbo is located closer to the mouth of Winam Gulf than the remaining RESP villages

which are located in more protected waters of the Gulf. Previous studies have also shown *B.*

choanomphala to be rare or absent within Winam Gulf, possibly because the bottom substrate is

too muddy, whereas, the substrate found off the shore of the PHS villages was a combination of

sand and mud, known to be favorable to *B. choanomphala*.[11,22] Gouvras et al., found *B.*

choanomphala to be focally distributed in the Tanzanian waters of the lake, including in some

more protected inlets.[34] They noted, as did we, a general trend for this species to be more

common in western than eastern locations, and considered a possible gradual change in

substratum conditions from west to east might be responsible. Many other factors such as levels

of dissolved oxygen, pollution and seasonal effects like rainfall may also influence *B.*

choanomphala populations.[33,34] We did not observe any association between the abundance of

B. choanomphala and water hyacinths. Although hyacinth mats can cover areas of the lake

where deeper water and hence *B. choanomphala* populations are present and could have

adverse shading or other effects, our impression was that hyacinth mats along the more

exposed shores away from the Gulf were more prone to be dispersed by winds.

420   With respect to the schistosome infections among the snails recovered, a couple of

caveats should be kept in mind. One is that some infections are prepatent at the time the snails

are collected such that these infections would be missed. Our approach here was simply to

retain all non-shedding snails in lab aquaria and to re-shed them after four weeks. This is

because we have excellent snail-rearing facilities near our field sites at our disposal in which

mortality of snails during the four-week interval can be minimized. This approach has the

advantage of detecting those infections that successfully transition to patency (cercariae are

actually produced). An alternative approach to detect prepatent infections is to employ

molecular xenomonitoring techniques to detect nascent infections at the time of

collection.[24,28,34,47] Such techniques can be very sensitive and specific, but are prone to error if

the infection is very young at the time of assay, and some detected infections may have been

doomed to fail, so transmission potential could potentially be overestimated.[28] Also, extractions

of individual snails for PCR-based or other types of molecular detection of prepatent infections

becomes prohibitively expensive and time-consuming when large numbers of snails need to be

screened. Pooling of samples or use of eDNA-based techniques may help but are likely to be

accompanied by a loss of precision in estimates of snail infection rates.[48] The second caveat is

that the rodent-transmitted *Schistosoma rodhaini*, the sister species of *S. mansoni*, is known to

be present in the Lake Victoria basin though it is rare by comparison with *S. mansoni*.[42] We

extracted mammalian schistosome cercariae (four cercariae per infected snail) from 20 different

positive snails and all yielded *nad5* bands consistent with *S. mansoni*, with one of these

samples indicating a co-infection with *S. rodhaini*.[28] Molecularly-based techniques

unquestionably increase the accuracy of species identifications, but again pose some

challenges insofar as an infected snail can produce thousands of cercariae over a period of

several months and it is conceivable that the species composition of the schistosome cercariae

could change during the interval. Again, with a large survey with many positive snails recovered,

resources limit the ability to repeatedly sample individual cercariae produced by the same snail.

Acknowledging this constraint, given the ubiquity of human infections in our study area, and that

20 out of 20 positive snails cercariae samples we tested were identified as *S. mansoni*, we are

confident the conclusions we reach here pertain to *S. mansoni* infections.

With these caveats in mind, for snail infections of *S. mansoni*, both the number of

positive snails found and prevalence were higher for *B. sudanica* than *B. choanomphala*. The

overall prevalence of *S. mansoni* infections in all the *Biomphalaria* snails collected was almost

2-fold higher for the PHS villages than for the RESP villages, and the highest recovery of

positive snails was found in Kanyibok (1.57%) and Usenge (1.42%), two of the villages with the

highest residual prevalence of infections in schoolchildren. However, one RESP village Kotieno

also showed a high overall prevalence of infected snails (1.36%). When all sampling sites,

sampling times and presence of hyacinths were taken into account, PHS and RESP villages did

not differ with respect to numbers or prevalence of infected *B. sudanica*. The few *B.*

choanomphala found to be positive were recovered only from PHS villages.

We also noted a suggestive negative relationship between hyacinth presence and

infection rates of *B. sudanica* with *S. mansoni*. If a true association, this finding may reflect a

decreased likelihood of collecting infected snails when large hyacinth mats are present. Several

explanations could account for this including higher mortality of infected snails due to oxygen

depletion under hyacinth mats, reluctance of people to enter water for either defecation or

washing purposes if hyacinths are present, or increased difficulty of miracidia in finding snails in

spatially complex hyacinth mats. In general, the role of hyacinths in understanding the dynamics

of schistosomiasis transmission in the lake requires further investigation.

Although fewer *B. choanomphala* than *B. sudanica* were collected, and their prevalence

of infection with *S. mansoni* was relatively low, it would be incorrect to assume the former

species has limited significance with respect to schistosome epidemiology in Lake Victoria. Our

dredge hauls across the extensive bottom surface of the lake can only be expected to retrieve a

small proportion of the snails present and likely grossly underestimate the role of this snail

species in perpetuating transmission. At least two factors might facilitate infections in *B.*

choanomphala: currents moving from offshore to deeper water, and fishermen who may

defecate directly into deeper water from their boats. In this regard, failure to include fishermen in

control programs may favor *B. choanomphala*-mediated transmission. Another factor potentially

favoring involvement of *B. choanomphala* in transmission is that experimental infection studies

of Kenya-derived specimens suggest this species, like *B. pfeifferi*, is inherently more susceptible

to *S. mansoni* than *B. sudanica*.[29,31] Gouvras et al., working in southern Lake Victoria found

more *B. choanomphala* positive for *S. mansoni* than we did: 12.2% of all *S. mansoni* shedders

were *B. choanomphala* in their study vs. only 3.5% in our study.[34] In both studies though, *B.*

sudanica was identified as the major vector for *S. mansoni* in the lake.

Considering the second measure of the force of transmission we examined, namely

sentinel mice infections, we did not observe a significant difference in mean number of worms

recovered between PHS and RESP villages, though worm recoveries were approximately twice

as high in the former and a positive correlation was noted across villages between the total

number of snails shedding *S. mansoni* and the number of worms recovered from mice. Results

of snail collections and worm recoveries were concordant in 60% of our 40 sampling times, but

presence of *S. mansoni*-shedding snails by no means guaranteed the recovery of worms from

sentinel mice placed nearby, and vice versa: recovery of worms from sentinel mice was

frequently noted at sampling locations and times we found no shedding snails. This suggests

that neither method perfectly reflects the possibility of transmission at a particular locality at a

certain time.

Not unexpectedly, and as reported from previous studies, *Biomphalaria* snails shedding

other trematode cercariae were also frequently collected.[49-52] Though non-schistosome

infections exceeded *S. mansoni* infections by a factor of 2.9, we noted no significant positive or

negative associations with any of the groups of trematodes with *S. mansoni* infections. The

three villages that yielded the most *S. mansoni* snail infections (PHS villages Kanyibok and

Usenge and RESP Kotieno) also yielded by far the most infections with other trematodes,

suggesting they are general "trematode-transmitting hotspots." One possibility to explain this is

that the overall susceptibility to trematode infection is higher in snails from collecting sites in

some villages than others. This possibility is currently being examined with respect to

susceptibility to *S. mansoni*. Another explanation is that certain water contact points along the

lakeshore attract intense activity by a variety of potential definitive hosts including people,

domestic animals like cattle or goats and wild birds, the latter often thronging to locations where

fishermen bring their catches. People often contact water extensively at such sites, making

them potentially dangerous potential transmission sites.

In comparison to *B. sudanica*, *B. choanomphala* had fewer *S. mansoni* infections and

fewer infections with trematodes of other species (only 4 other species noted). Deep water has

been considered a refugium, or in the parlance of King et al., a "coevolutionary cold spot" that

limits exposure of snails to digenetic trematodes.[52] In comparison, shoreline habitats are visited

by numerous definitive hosts and can be considered as "coevolutionary hot spots" for digenetic

trematode infections, supported by the many trematode species noted to be transmitted by *B.*

sudanica.[52]

This study poses several questions requiring clarification. For instance, it might be

argued that PHS villages have higher prevalence of human infections because the snails

adjacent to such villages are more susceptible to *S. mansoni* infection. Experiments are

underway to test the relative compatibility of *B. sudanica* from different villages to sympatric and

allopatric isolates of *S. mansoni*. Our null hypothesis is that strongly differentiated patterns of

schistosome-snail compatibility will not be seen, because both human migration (and

consequently *S. mansoni* migration) and snail movements mediated by hyacinths, winds and

currents will overcome local differentiation. Also in need of additional clarification is the extent

to which the gene pools of *B. sudanica* and *B. choanomphala* co-mingle. Studies of

mitochondrial and nuclear markers indicate the two taxa are not highly divergent genetically and

should probably be considered as a single species with the species name with precedence yet

to be decided.[12,13] It is clear though that pockets of snails with the characteristic *B.*

choanomphala phenotype can be recovered from the shoreline over a period of several months.

The extent to which snails of the two phenotypes interbreed, the appearance of the progeny,

and the nature of genes and alleles dictating compatibility to schistosomes and other

trematodes all remain to be investigated.

CONCLUSIONS

Results of this study did not find significant differences between PHS and RESP villages

for *B. sudanica* with respect to relative abundance or numbers of snails shedding *S. mansoni*

cercariae, or in the number of worms recovered from sentinel mice. Although *B. sudanica* was

present in all the villages we studied and based on the number of positive snails of this species

recovered is also the major vector for *S. mansoni* in all villages, *B. choanomphala* was found in

all PHS villages and in only one RESP village. The relative abundance of *B. choanomphala* was

significantly higher in PHS villages. In our view, *B. choanomphala* may provide an under-

appreciated and more diffuse and difficult to measure alternative route of transmission along

much of the shore of the lake. Many other trematode species infect *Biomphalaria* snails in Lake

Victoria but their overall prevalence is low and suggests they do not have a dramatic primary

effect on *S. mansoni* prevalence rates in snails. Evidence of active *S. mansoni* transmission

was found in all 10 villages investigated, not surprising from the standpoint that the lake

supports massive populations of *Biomphalaria* snails and sanitary facilities in rural areas around

the lake are scarce. However, distinct differential responsiveness to annual MDA with PZQ was

somewhat surprising. Nevertheless, opportunities for transmission and reinfections are very

likely to continue to occur, even in RESP villages. Control of *S. mansoni* in and around the lake

will remain a daunting challenge and its persistence argues more persuasively than ever for

implementation of improved sanitation and provision of safe water supplies as important parts of

an integrated and sustained three-country basin-wide control approach, and as fundamental

rights for the people living in the area.

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Authors' addresses

Martin W. Mutuku, Centre for Biotechnology Research and Development, Kenya Medical Research Institute, Nairobi, Kenya and School of Biological Sciences, College of Biological and Physical Sciences, University of Nairobi, Nairobi, Kenya, E-mails: mmutuku@kemri.org and mwmutuku81@gmail.com. **Martina R. Laidemitt**, Center for Evolutionary and Theoretical Immunology, Parasitology Division, Museum of Southwestern Biology, Department of Biology, University of New Mexico, Albuquerque, New Mexico, United States of America, E-mail: mlaidemitt@unm.edu. **Brianna R. Beechler,** College of Veterinary Medicine, Department of Biomedical Sciences, Oregon State University, Corvallis, Oregon, United States of America, E-mail: breebeechler@gmail.com. **Ibrahim N.Mwangi,** Centre for Biotechnology Research and Development, Kenya Medical Research Institute, Nairobi, Kenya, E-mail: indungu@kemri.org. **Fredrick. O. Otiato**, Influenza Surveillance Program, Centers for Disease Control and Prevention, Nairobi, Kenya, E-mail: frede516@gmail.com**. Eric L. Agola**, Centre for Biotechnology Research and Development, Kenya Medical Research Institute, Nairobi, Kenya, E-mail: elelo@kemri.org. **Horace Ochanda**, School of Biological Sciences, College of Biological and Physical Sciences, University of Nairobi, Nairobi, Kenya, E-mail: hochanda@uonbi.ac.ke. **Bishoy Kamel**, Center for Evolutionary and Theoretical Immunology, Parasitology Division, Museum of Southwestern Biology, Department of Biology, University of New Mexico, Albuquerque, New Mexico, United States of America, E-mail: bishoyh@unm.edu.

Gerald M. Mkoji, Centre for Biotechnology Research and Development, Kenya Medical

Research Institute, Nairobi, Kenya, E-mail: gmkoji5@gmail.com. **Michelle. L. Steinauer**,

Department of Basic Medical Sciences, Western University of Health Sciences, Lebanon,

Oregon, United States of America, E-mail: msteinauer@westernu.edu. **Eric S. Loker**, Center for

Evolutionary and Theoretical Immunology, Parasitology Division, Museum of Southwestern

Biology, Department of Biology, University of New Mexico, Albuquerque, New Mexico, United

States of America, E-mail: esloker@unm.edu.

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761   **Table 1: *Schistosoma mansoni* prevalence in schools within the study villages showing**
762   **response to praziquantel treatment following four years of Mass Drug Administration.**
763   **Data acquired from and shown with permission of the SCORE project. Sample sizes**
764   **shown in parentheses.**

SCHOOL	Prevalence (%) Year 1 (2011)	Prevalence (%) Year 2 (2012)	Prevalence (%) Year 3 (2013)	Prevalence (%) Year 4 (2014)	Prevalence (%) Year 5 (2015)	Absolute Prevalence Change Year 1 Minus

						Year 5 as %
			Persistent Hotspot Villages			
MINYA	68.6 (n=97)	77(n=100)	75(n-97)	74(n-92)	82(n-97)	+13.4
AGOK	60.5(n-99)	59.8(n-93)	60.2(n-97)	67(n-86)	39(n-91)	-21.5
MIGIRO	69.4(100)	76.7(n-94)	49.2(n-87)	72(n-87)	48(n-88)	-21.4
MIYANDHE	78.6(86)	70.5(n-91)	71.4(n-87)	73.5(n-92)	54(n-94)	-24.6
KANYIBOK	91.8(100)	93(n-96)	83.7(n-98)	84(n-94)	83(n-100)	-8.8
USENGE	69.4(100)	68(n-100)	65(n-100)	46.5(n-100)	65(n-100)	-4.4
			Responding Villages			
KOTIENO	52.9(87)	40.9(n-92)	19(n-93)	16(n-84)	12(n-88)	-40.9
SEKA DOK	81.6(100)	73.2(n-97)	55.8(n-91)	48(n-90)	19(n-96)	-62.6
ST.DOUGLAS WETA	55.2(100)	27.4(n-94)	24(n-93)	19(n-87)	19(n-91)	-36.2
MUMBO	57.7(77)	30(n-74)	11.5(n-78)	15.6(n-79)	5.5(n-77)	-52.5

766 **Table 2: The 20 sampling sites and their GPS coordinates, total number of *Biomphalaria***

767 ***sudanica* and *B. choanomphala* collected from each, and the number of each species**

768 **positive for *S. mansoni*.**

Village	Habitat	Latitude	Longitude	*Biomphalaria sudanica*	No. of snails (+) for *S. mansoni*	*Biomphalaria choanomphala*	No. of snails (+) for *S. mansoni*
				Persistent Hotspot Villages			
	Kochilo beach	$00^014'10.96"S$	$034^014'45.79"E$	522	0	96	0
MINYA	Luthi beach	$00^014'04.58"S$	$034^013'58.74"E$	758	1	282	0
	Gudwa beach	$00^021'26.52"S$	$034^019'48.37"E$	209	1	238	2
AGOK	Kogoye beach	$00^020'51.94"S$	$034^020'74.32"E$	220	6	85	0
	Uyawi beach	$00^012'18.58"S$	$034^009'18.36"E$	728	1	89	0
MIGIRO	Kadede beach	$00^012'12.09"S$	$034^009'11.74"E$	132	0	100	0
	Kamumbo beach	$00^011'49.04"S$	$034^010'28.17"E$	48	0	15	0
MIYANDHE	Daraja beach	$00^011'46.65"S$	$034^009'44.11"E$	103	0	71	0
	Anyanga beach	$00^005'22.49"S$	$034^005'09.34"E$	897	28	380	1
KANYIBOK	Ndiara beach	$00^005'27.33"S$	$034^005'06.24"E$	407	0	161	0
	Police beach	$00^004'34.80"S$	$034^003'24.74"E$	833	19	4	0
USENGE	Usenge beach	$00^004'21.49"S$	$034^003'35.84"E$	489	4	292	0
			B. sudanica 60/5,346 (1.12%), *B. choanomphala* 3/1,813 (0.17%)				
				Responding villages			
	Kotieno beach	$00^021'09.07"S$	$034^040'03.52"E$	674	2	0	0
KOTIENO	Maguti beach	$00^021'17.57"S$	$034^039'32.50"E$	649	16	0	0
SEKA DOK	Seka beach	$00^021'18.58"S$	$034^040'59.95"E$	462	2	0	0

ST. DOUGLAS WETA	Kagwa beach	00^{0}21'19"S	034^{0}40'52.12"E	834	1	0	0
	Kobala beach	00^{0}21'05"S	034^{0}45'31.21"E	735	0	0	0
	Alara beach	00^{0}21'08"S	034^{0}45'16.01"E	584	0	0	0
MUMBO	Kabuong beach	0^{0}20'20.06"S	034^{0}21'04.09"E	576	1	93	0
	Mumbo beach	0^{0}20'0.99"S	034^{0}21'33.03"E	389	1	1	0
B. sudanica 23/4,903 (0.47%), *B. choanomphala* 0/94 (0%)							

LEGEND

Figure 1: Map showing position of study villages along the shore of Lake Victoria, Kenya. The

map was constructed using QGIS version 2.18.22 and 3.2.1, the basemap layer was added

from the standard terrain tiles provided from http://maps.stamen.com through the OpenLayers

function. The tiles are © Stamen Design, under a creative commons attribution (CC BY 3.0)

license. The data for the locations were obtained manually using cellular phone GPS

functionality according to the WGS1984 mercator projection.

Figure 2: Number of *B. sudanica* (blue bars) and *B. choanomphala* (black bars) collected at

each of the four sampling times (April and August 2016, January and May 2017) from the 6

persistent hotspot villages on the top versus the 4 responding villages on the bottom. Snail

numbers are summed between collection sites at a village. Numbers over bars indicate number

of snails infected with *S. mansoni* shedders/total number of snails found. Numbers shown within

mouse figures represent the number of worms recovered from sentinel mice at that time point

(summed across mice and collection sites at that village). An H in a closed circle indicates

hyacinths were present and dredging was done. An H in an open circle indicates hyacinths were

present but dredging could not be done. An asterisk (*) indicates Kabuong Beach was dredged

but not Mumbo Beach.

Figure 3. Relative abundance of *Biomphalaria* snails at persistent hotspot and responding sites

of Lake Victoria in Kenya. **A.** Persistent hotspot sites did not contain more *B. sudanica* snails

than responding sites when time point, village, and hyacinths were taken into

account. **B.** *B. choanomphala,* was far more abundant at persistent hotspot sites compared to
responding sites (P = 0.0091). Error bars indicate standard error of the mean.

Figure 4. The number of *B. sudanica* collected was greater when hyacinths
were present than when hyacinths were not present (P=0.00973). The presence of
hyacinths did not influence *B. choanomphala* abundance (not shown). Error bars
indicate standard error of the mean.

Figure 5. The total number of adult schistosomes collected from sentinel mice was
greater in persistent hotspot than responding sites, but there was no statistical
difference between them when village, hyacinths, and time point were considered.

Figure 6. Total number of *B. sudanica* collected at a village was not significantly
correlated with prevalence of *S. mansoni* in humans either before or after the drug
treatment intervention (before: Pearson r = -0.0369, p = 0.460; after: Pearson r = -0.009,
p = 0.49). Yellow = persistent hotspot villages; Blue = responding villages.

Figure 7. Total number of *B. choanomphala* collected at a village was significantly
correlated with prevalence of *S. mansoni* in humans both before and after mass drug
administration (before: Pearson r = 0.587, r^2 = 0.345, p = 0.037; after: Pearson r =
0.9034, r^2=0.816, p = 0.0002). Yellow = persistent hotspot villages; Blue = responding
villages (* two villages with same prevalence).

Usenge beach

Police beach   KANYIBOK

USENGE   Anyanga beach

Ndiara beach

Daraja beach

MIYANDHE

MIGIRO   Kamumbo beach

Kadede beach

Kochilo beach

Uyawi beach   MINYA

Luthi beach

Mumbo beach   Kotieno beach

Kabuong beach   Maguti beach   Seka beach

MUMBO

Kogoye beach   KOTIENO GUMBA   SEKA DOK

Gudwa beach   Alara beach

AGOK

Kagwa beach   Kobala beach

ST. DOUGLAS WETA

🏠 Persistent hotspot villages

🏠 Responding villages

10   0   10   20   30   40 km

Top panel: scatter plot with x-axis "Total *Biomphalaria choanomphala*" (0 to 600) and y-axis "Prevalence in Humans Before MDA" (50 to 100).

Bottom panel: scatter plot with x-axis "Total *Biomphalaria choanomphala*" (0 to 600) and y-axis "Prevalence in Humans After MDA" (0 to 100). An asterisk (*) appears near a blue square point around y = 19.