Mosquito electrocuting traps for directly measuring biting rates and host-preferences of *Anopheles arabiensis* and *Anopheles funestus* outdoors

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**Abstract**

**Background:** Mosquito biting rates and host preferences are crucial determinants of human exposure to vector-borne diseases and the impact of vector control measures. The human landing catch (HLC) is a gold standard method for measuring human exposure to bites, but presents risks to participants by requiring some exposure to mosquito vectors. Mosquito electrocuting traps (METs) represent an exposure-free alternative to HLCs for measuring human exposure to malaria vectors. However, original MET prototypes were too small for measuring whole-body biting rates on humans or large animals like cattle. Here a much larger MET capable of encompassing humans or cattle was designed, and its performance was evaluated relative to both the original small MET and HLC and for quantifying malaria vector host preferences.

**Methods:** Human landing catch, small human-baited METs (MET-SH), and large METs baited with either a human (MET-LH) or calves (MET-LC) were simultaneously used to capture wild malaria vectors outdoors in rural southern Tanzania. The four capture methods were compared in a Latin-square design over 20 nights. Malaria vector host preferences were estimated through comparison of the number of mosquitoes caught by large METs baited with either humans or cattle.

**Results:** The MET-LH caught more than twice as many *Anopheles arabiensis* than either the MET-SH or HLC. It also caught higher number of *Anopheles funestus* sensu lato (s.l.) compared to the MET-SH or HLC. Similar numbers of *An. funestus* sensu stricto (s.s.) were caught in MET-LH and MET-SH collections. Catches of *An. arabiensis* with human or cattle-baited large METs were similar, indicating no clear preference for either host. In contrast, *An. funestus* s.s. exhibited a strong, but incomplete preference for humans.

**Conclusions:** METs are a sensitive, practical tool for assessing mosquito biting rates and host preferences, and represent a safer alternative to the HLC. Additionally these findings suggest the HLC underestimate whole-body human exposure. MET collections indicated the *An. funestus* s.s. population in this setting had a higher than expected attack rate on cattle, potentially making eliminating of this species more difficult with human-targetted control measures. Supplementary vector control tools targetted at livestock may be required to effectively tackle this species.

**Keywords:** Mosquito electrocuting trap, Human landing catch, Mosquitoes, Malaria, Host preference, *Anopheles arabiensis*, *Anopheles funestus*, Sampling, Human biting densities
Background

Accurate estimation of mosquito biting rates and host preference are critical for assessing exposure risks of humans and animals to vector-borne diseases, and for optimizing the impact of vector control strategies [1–4]. Until relatively recently, measuring human exposure to mosquito bites outdoors necessitated laborious, potentially hazardous and ethically questionable human landing catches (HLCs) [5–7]. Despite these risks, the HLC remains the only technique considered reliable for direct estimation of human exposure to mosquito bites inside houses and outdoors, and over the course of the entire night [6, 8]. In addition to the limitations mentioned above, it is possible the HLC may also underestimate total human exposure to mosquito bites for a number of reasons. First, the HLC relies on the constant vigilance of the collectors throughout an entire night of sampling, and it is possible that capture efficiency drops as participants tire. Second, only mosquitoes attempting to feed on a person’s legs are collected with HLCs [6, 9]. While it is known that many malaria vectors are most attracted to the feet area [10–12], biting can occur on other parts of the body including the head and arms. While collectors may also attempt to capture mosquitoes landing on other parts of their body, these mosquitoes may go undetected given the focus on the lower leg area. Given the importance of measuring malaria transmission, there is a great need for surveillance tools that can accurately measure total human exposure to mosquito bites in both indoor and outdoor settings.

A new mosquito electrocuting trap (MET) has recently shown promise as a representative and safe alternative to the HLC for measuring human exposure to malaria vectors outdoors [13, 14]. This new tool produced similar estimates of relevant metrics of human exposure to malaria vectors (such as distribution of bites between indoors and outdoors and over the course of the night) as the HLC gold standard in urban Dar es Salaam [14]. Such measurements allow quantification of the proportion of human exposure occurring indoors and during hours when people are in bed, which are critical determinants of optimal deployment of bed nets, and other vector control tools [2, 4, 15]. Intervention choice should also be guided by the host preference of target mosquito vectors [2–4, 16]. For example, novel vector control approaches that target livestock with systemic insecticides [17, 18] will only be effective against vectors that commonly bite livestock as well as humans [2–4, 19].

Host preference is typically measured by comparing the number of mosquitoes attracted to different host types in a choice test [16]. However, such assays are often hard to implement under natural field settings due to the lack of standardized methods for sampling vectors attracted to humans and other animals. Consequently, the human blood index (HBI); defined as the proportion of blood meals that a vector species obtains from humans [20], is often used as a proxy for host choice instead of host preference. The HBI is estimated by sampling mosquitoes resting in and around houses [20–23], and identifying the source of their blood meal using molecular methods [24]. Although useful to confirm what mosquitoes actually feed on within a given environment, the HBI is dependent on the relative abundance of different host types and thus does not give an unbiased estimate of innate preference [16]. As it is difficult to measure preference across the range of potential hosts, HBI has been often used as a proxy of preference, even though it is only a measure of choice. Experimentally-controlled host preference assays, in which mosquitoes are given an equal opportunity to bite different hosts, provide the most reliable, direct and unambiguous measure of their innate behavioural preferences [25–28].

A range of field techniques have been proposed to estimate the host preference of mosquitoes [16, 29, 30]. These methods include a variety of net and stable-based traps that enclose either a human or animal host inside a physical structure, allowing mosquitoes to enter but impeding their exit [25]. Although useful, many of these methods have limitations that make them difficult to implement or interpret. For example, e-nets require a relatively large area for set up due to the requirement for a 10 m odour tube [30]. Other methods such as odour-baited entry traps (OBETs) involve luring mosquitoes to point source where they must enter a structure to be trapped. This requirement for entry behaviour may preferentially select endophilic (e.g. indoor biting) vector species; and thus not give a representative sample of outdoor biting mosquito taxa.

Electrocuting surfaces have been widely used to capture outdoor-biting tsetse flies [31], and have also been adapted for sampling host-seeking mosquitoes [13, 14, 30, 32–34]. A MET prototype previously applied to measure human exposure to malaria vectors across different times of the night performed consistently with the HLC gold standard method [14]. This original prototype was designed to operate similarly to HLC, with only mosquitoes approaching the lower leg area of a human bait being trapped [14]. Whilst this approach is appropriate when trying to replicate the estimates of human exposure gained from a HLC, it is not feasible for studies of mosquitoes host-seeking on other large animals as required to measure host preference. This study evaluated the performance of a new MET prototype designed to be capable of encompassing entire hosts (humans or livestock), and evaluated its performance relative to HLC and the original small sized MET for measuring human exposure.
to malaria vectors. Additionally large METs were used to measure the relative preference of African malaria vector species for biting human vs cattle hosts. In doing so, the versatility of MET-based sampling approaches to safely measure a range of epidemiologically-relevant mosquito vector ecological and behavioural traits were demonstrated.

**Methods**

**Study area and site**

The study was conducted in Sagamaganga village (S08°03.83′; E036°47.77′) [35–37], which is situated 15 km east of Ifakara town within the Kilombero Valley, south-eastern Tanzania. Most residents in this area live on subsistence farming, growing rice and maize as well as keeping livestock [38]. Despite the successful scale up of long lasting insecticide nets (LLINs), and undetectable level of the most historically-important vector *Anopheles gambiae* [39, 40], this area still experiences year-round malaria transmission [39, 40]. Most malaria transmission is mediated by *Anopheles funestus* sensu stricto (s.s.), a species that is thought to feed mainly inside house (endophilic) and on humans (anthrophilic) [41], and is highly physiologically resistant to pyrethroid insecticides [39]. *Anopheles arabiensis* is the most common and highly abundant anopheline species in the area but, has a much lower rate of malaria infection than *An. funestus* s.s. [39].

**Trapping methods**

**Mosquito electrocuting trap (MET)**

The METs evaluated here operate in similar fashion to previous prototypes where electrocuting surfaces are placed around a host so as to intercept and kill mosquitoes as they attempt to land and bite [13, 14]. The MET prototypes used in early trials were made of 30 cm × 30 cm wooden panel frames, which could be assembled into a square box to encompass the lower legs of a seated human. The prototypes used here were made with PVC panel frames and held together with hinges, so that they were lighter and could be more easily assembled into a box shape (Fig. 1a). Two MET prototypes were used: one with the original frame size (30 cm² panels) and a larger prototype with 125 cm × 122 cm panels, which could be assembled into a box to fully enclose the whole body of a sitting human volunteer or a standing/ sleeping bovine calf (Fig. 1b). Both MET designs were composed of parallel stainless steel wires with alternating polarity, spaced 5 mm apart, and held in place by passing them through evenly-spaced holes drilled into the PVC frame (Fig. 1c). The large MET was fitted with a dry bamboo protective fence on the inside, so as to prevent any accidental contact of the volunteer or calf with the electrified wires (Fig. 1d). The top of the large MET was also covered with an untreated net, secured with Velcro®, to prevent any possible host exposure to mosquitoes flying in from above (Fig. 1b). During experiments, each MET (whether baited with a human or calves) was placed on a square wooden platform measuring 2 × 2 m, which was covered with a white cotton sheet to increase the visibility of the electrocuted mosquitoes to collectors. The MET is powered by 2 12 V batteries in parallel to make 24 V which produce a DC output that creates electric potential between alternating wires. The current–voltage combination has been optimized to kill mosquitoes on contact, but without damaging the specimen so it can be morphologically identified [42] as confirmed in previous studies [13, 14].

**Human landing catch (HLC)**

The HLC was performed by an adult male sitting on a chair and exposing his lower legs to mosquitoes. The volunteer continually inspected his legs using a torch for 45 min of each sampling hour. Any mosquitoes observed to land upon their legs during this time were collected using a mouth aspirator as previously described [43, 44].

**Study design**

Experiments were conducted in an open field in Sagamaganga village. The field was bordered by two isolated compounds containing several houses, with one big cow shed on one side and a rice field on the other. Four sampling stations spaced approximately 20 m from each other along a straight line were set up in the field. Four different combinations of host and capture methods were evaluated: (1) a human volunteer conducting HLC, (2) a small MET with one adult male human (MET-SH), (3) a large MET containing one adult male human (MET-LH), and (4) a large MET baited with two female calves (~2 months old 60 to 80 kg average weight; MET-LC). Two calves were used as experience shows that calves are less stressed when kept together. The weight of male volunteers conducting HLC, MET-SH or MET-LH collections averaged ~ 65 kg. All four capture methods were used on each night of sampling, with one method allocated per sampling station. These capture techniques were then serially rotated through all four sampling stations in a 4 × 4 Latin square design (Fig. 2). Therefore, four nights were required to complete one replicate. The experiment was replicated five times, requiring a total of 20 sampling nights.

Each MET collection was made between 18:00 h and 6:00 h, with traps being run for 45 min of each hour. After the 45 min trapping period of each hour, MET traps were switched off and their outer surfaces and the white sheet below inspected for electrocuted mosquitoes.
Mosquitoes found dead on the white sheet were collected using forceps, and those electrocuted and stuck to the surface panels were first swept using a small brush and collected by forceps. A pair of volunteers participated in each of the three human-baited sampling techniques. One volunteer would catch mosquitoes between 18:00–00:00 h, and the second between 00:00–06:00 h.

Each pair of volunteers was randomly assigned to a particular station and remained associated with that station throughout the experiment, so that the systematic differences between individual station locations and volunteer pairs could be combined into a single source of variance. For the MET-LC, the same two female calves were systematically moved together with the large MET through
all four sampling stations. Mosquitoes collected by different trapping methods were stored separately for each hour of collection in labelled paper cups. The voltage for the electrocuting trap was checked regularly.

**Mosquito processing**

All mosquito specimens collected were sorted, counted and morphologically identified in the field with the aid of a dissection microscope [45]. Mosquitoes were identified as members of the *An. gambiae* complex (*An. gambiae* sensu lato), the *An. funestus* complex (*An. funestus* s.l.), *Anopheles coustani*, *Anopheles zeimanni*, *Culex* species, or *Aedes* species, and classified in terms of sex and abdominal status. A subsample of 1839 out of 15,322 of *An. gambiae* s.l. and all *An. funestus* s.l. (n = 2067) collected were stored individually in 1.5 ml eppendorf tubes containing desiccated silica gel covered with a small ball of cotton wool. Subsample of 7 *An. gambiae* s.l. from each trapping hour of each collection (MET and HLC) were used for individual species identification using molecular analysis. These 7 were haphazardly selected from the total collected each hour. In cases where less than 7 were collected in an hour, all were subsampled. This generated a subsample of 3680 *An. gambiae* s.l. of which approximately 50% were analysed by PCR for species identification using molecular analysis. These 7 were haphazardly selected from the total collected each hour. In cases where less than 7 were collected in an hour, all were subsampled. This generated a subsample of 3680 *An. gambiae* s.l. of which approximately 50% were analysed by PCR for species identification (n = 1839 including representative samples from all trapping methods and dates [46, 47]. This subsample of individuals was analysed by enzyme-linked immune-absorbent assay (ELISA) for malaria sporozoite detection [48]. To avoid false positives, the ELISA lysates were heated in a boiling water bath for 10 min at 100 °C to inactivate heat-labile antigens other than *Plasmodium falciparum* circumsporozoite protein, which is not denaturable [49].

**Statistical analysis**

Statistical analyses were carried out using the R statistical software version 3.0.2, augmented with the *matrix*, *lattice* and *lme4* packages [50]. Generalized linear mixed models (GLMMs) [51, 52] were used to estimate and compare the mean abundance of malaria vector species in nightly catches with different trap and host types. The number of mosquitoes caught per night was treated as the dependent variable, with trapping method fit as an independent fixed effect and sampling night and station fit as random effects. Models were fitted using a Poisson distribution. Likehood ratio tests were performed to test the significance of the main effect of trap type. Separate statistical models were fit for each malaria vector species.

To estimate the relative host preference of malaria vectors, only data from MET-LH and MET-LC were considered. Here comparisons were made between the proportion of malaria vectors caught in the human vs cattle baited traps. The response variable was defined as the relative proportion feeding on human-baited traps (MET-LH/[MET-LH + MET-LC]), with sampling night and station treated as random effects. These models were fitted with a binomial distribution with a logit link function. These host preference models did not include any fixed effect variables, with the estimated intercept representing host preference in terms of difference from the null hypothesis of an equal distribution of bites on human and calf baits. Separate analyses were performed for all the distinct species identified within the *An. funestus* s.l. Here count data were obtained by aggregating the
total number of PCR-identified individuals from each species captured with each method in a single station on a single night. These aggregated count data were then analysed by GLMM as described above.

**Results**

A total of 23,820 female *Anopheles* mosquitoes were captured, of which most were *An. gambiae* s.l. (Table 1). The next most abundant mosquito genus was *Culex* (Table 1). A subsample of 1839 specimens from the *An. gambiae* complex were tested by PCR, out of which 1644 (89%) were successfully amplified. All of these *An. gambiæ* s.l. were confirmed to be *An. arabiensis*. Therefore, from this point onward, all results obtained for *An. gambiæ* s.l. are considered representative of *An. arabiensis* and referred to as such.

PCR amplification was successful for only about half of *An. funestus* s.l. specimens (1098/2066). Within these samples, *An. funestus* s.s. was the most prevalent 69% (756/1098), followed by *Anopheles rivulorum* 25% (278/1098) and *Anopheles leesoni* 6% (46/1098) (Table 1). The proportion of *An. funestus* s.l. whose DNA could be successfully amplified varied between trapping methods as follows: HLC = 61% (n\textsubscript{total} = 297), MET-SH = 49% (n\textsubscript{total} = 546), MET-LH = 51% (n\textsubscript{total} = 39) and MET-LC = 56% (n\textsubscript{total} = 484). Amplification rates from specimens collected in HLC were significantly higher than from MET-SH ($\chi^2 = 11.16$, $P < 0.001$) and the MET-LH ($\chi^2 = 8.30$, df = 1, $P < 0.001$). Amplification rates were similar in *An. funestus* s.l. collected in HLC and MET-LC ($\chi^2 = 2.81$, $P = 0.09$). There was no significant differences in *An. funestus* s.l. amplification rates between any of the 3 MET types ($P > 0.05$ in all cases).

The proportion of *An. arabiensis* and PCR-confirmed *An. funestus* s.s. infected with sporozoites were 0.18% (3/1644) and 0.27% (2/756) respectively. None of the other sibling species from the *An. funestus* group or unidentified *An. funestus* s.l. were found to be infected with sporozoites.

The MET-SH sampled a similar number of *An. arabiensis* as the HLC (Table 2). The MET-LH caught more than twice as many *An. arabiensis* per night as either HLC (RR = 2.89, $P < 0.001$) or MET-SH (Tables 1, 2, Fig. 3). The MET-SH captured two times more *An. funestus* s.l. than HLC (Table 2), while the MET-LH consistently caught more *An. funestus* s.l. as either the HLC (RR = 3.63, $P < 0.001$) or MET-SH (Tables 1, 2, Fig. 3). However, MET-LH caught similar numbers of *An. funestus* s.s. and *An. rivulorum* as MET-SH (Table 2, Fig. 3).

On the basis of the numbers of mosquitoes caught in large METs baited with different host types, *An. arabiensis* was estimated to be attracted to human and cattle

| Species                          | Capture method | HLC | MET-SH | MET-LH | MET-LC | Total catch | %   |
|----------------------------------|----------------|-----|--------|--------|--------|-------------|-----|
| Anopheles spp.                   | Anopheles arabiensis\(^a\) | 1824 | 2331   | 5631   | 5536   | 15,322      | 39.8|
|                                  | Anopheles funestus s.s.     | 116  | 227    | 304    | 109    | 756         | 2.0 |
|                                  | Anopheles rivulorum         | 55   | 38     | 49     | 136    | 278         | 0.7 |
|                                  | Anopheles leesoni           | 11   | 4      | 27     | 22     | 64          | 0.2 |
|                                  | Anopheles funestus s.l.\(^b\) | 115  | 277    | 359    | 217    | 968         | 2.5 |
|                                  | Anopheles coustani          | 923  | 328    | 713    | 1330   | 3294        | 8.6 |
|                                  | Anopheles ziemanni          | 258  | 220    | 472    | 896    | 1846        | 4.8 |
|                                  | Anopheles pharoensis        | 118  | 86     | 182    | 193    | 579         | 1.5 |
|                                  | Anopheles squamosus         | 63   | 61     | 145    | 805    | 1075        | 2.8 |
|                                  | Anopheles maculipalpis      | 5    | 0      | 1      | 0      | 6           | 0.01|
|                                  | Anopheles wellcomei         | 87   | 66     | 153    | 300    | 606         | 1.6 |
| Other mosquito spp.              | Culex spp.                  | 1573 | 1012   | 4078   | 5673   | 12,336      | 32.0|
|                                  | Mansonia spp.               | 375  | 242    | 433    | 267    | 1317        | 3.4 |
|                                  | Coquillettidia              | 7    | 21     | 32     | 12     | 72          | 0.2 |
|                                  | Aedes spp.                  | 4    | 0      | 0      | 1      | 5           | 0.01|

\(^a\) Originally identified morphologically as *An. gambiæ* s.l. and then confirmed to be 100% *An. arabiensis* by PCR (All 1644 successfully amplified specimens)

\(^b\) *An. funestus* s.l. which could not be identified to species because did not amplify
Table 2: Comparisons of the mean number of females of mosquito species caught per night by HLC, MET-LH, MET-LC relative to reference mosquito electrocuting trap (MET-SH)

| Capture Method | Mean catch | RR [95% CI] | P value |
|----------------|------------|-------------|---------|
| Anopheles arabiensis | | | |
| HLC | 5.94 | 0.93 [0.88, 0.99] | 0.049 |
| MET-SH | 6.33 | 1a | NA |
| MET-LH | 17.17 | 2.71 [2.57, 2.85] | <0.001 |
| MET-LC | 16.80 | 2.65 [2.52, 2.79] | <0.001 |
| Anopheles funestus s.s. | | | |
| HLC | 4.42 | 0.53 [0.42, 0.68] | <0.001 |
| MET-SH | 8.26 | 1a | NA |
| MET-LH | 9.84 | 1.19 [0.99, 1.42] | 0.053 |
| MET-LC | 3.19 | 0.38 [0.31, 0.48] | <0.001 |
| Anopheles rivulorum | | | |
| HLC | 2.03 | 1.48 [0.97, 2.25] | 0.069 |
| MET-SH | 1.38 | 1a | NA |
| MET-LH | 1.38 | 1.00 [0.65, 1.53] | 0.996 |
| MET-LC | 3.58 | 2.61 [1.81, 3.75] | <0.001 |
| Anopheles funestus s.l. | | | |
| HLC | 1.23 | 0.49 [0.43, 0.57] | <0.001 |
| MET-SH | 9.37 | 1a | NA |
| MET-LH | 16.97 | 1.81 [1.64, 2.01] | <0.001 |
| MET-LC | 11.11 | 1.19 [1.06, 1.32] | 0.003 |

Results are based on 20 nights of collection with each trap type.
RR relative rate, CI confidence interval, NA not applicable because it is a reference capture method.

Discussion

Two sizes of electrocuting traps were evaluated for assessment of the biting densities and host preference of afrotropical malaria vectors in rural Tanzania. The large prototype baited with humans captured greater numbers of both An. arabiensis and An. funestus s.l. than the smaller prototype and the existing HLC gold standard. The large METs also proved effective for sampling mosquitoes attracted to cattle and human hosts. Estimates of host preference from MET collections indicated An. arabiensis is attracted to human and cattle hosts at a similar rate with no clear preference, whereas An. funestus s.s. clearly preferred humans over cattle. In contrast, An. rivulorum strongly preferred cattle. Notably, estimates of human biting rates in An. arabiensis and An. funestus s.l. were considerably greater when derived from the large MET than the small MET or the HLC. This raises the possibility that total human exposure to malaria vectors may be underestimated by current HLC gold standard method. The enhanced performance of large MET, is presumably due to the fact that, it samples a greater surface area around the host by encompassing their entire body and not just their lower legs. Such differences in sampling performance between small vs large METs were not detected for less abundant malaria vector species (An. funestus s.s. and An. rivulorum) [39]. However, this may be due to reduced statistical power to detect differences in these groups rather than a difference in their response to traps of different size.

Although the HLC is considered a gold standard approach for measuring human biting rates by mosquitoes, these results suggest it may underestimate total human exposure to Anopheles vector bites [43, 53, 54] and thus malaria transmission [55–57]. However, this hypothesis must be further investigated because it remains unclear whether all mosquitoes trapped with the large MET were actively host seeking, or whether some proportion was trapped during random or otherwise non-host directed flying. Ideally this could have been evaluated here by comparing the number of An. gambiae s.l. collected in baited and unbaited MET collections. Although this was not possible in this study due to limitations in the numbers of METs available and time period for study, it’s encouraged to be investigated in the future.

In summary, the results indicate that METs are efficacious for estimating both malaria vector biting rates and their host preference in outdoor environments.

This study also demonstrates that the MET can be practically applied to quantify vector host preferences by increasing its size so it can accommodate large non-human hosts. Many existing methods for assessing host preference rely on luring mosquitoes towards the odour component of host stimuli [28, 32, 59]. Although useful, these approaches may fail to capture the full range of visual, odour, heat and other stimuli arising from a host individual. The large MET evaluated here overcomes these limitations by presenting hosts to mosquitoes in a relatively natural way, with mosquitoes attempting to feed being intercepted just before they land. By increasing the size of the MET so it can encompass an entire host, this tool has potential to be applied to assess mosquito biting rates on a range of hosts including wildlife and domestic animals. This could be particularly useful to study the transmission of mosquito-borne zoonotic diseases [60]. The MET prototypes assessed here were also found to be reasonably practical for field use. These METs are made of durable but lightweight materials that are stackable, compact and easy to assemble.
Despite their advantages, METs also have practical limitations. The most notable limitation is the MET’s reliance on electrical components which may break or short circuit if exposed to excessive moisture. However, this specific limitation can be overcome by placement of traps under a tarpaulin cover and on platform which would enable their use even during the rainy seasons as exemplified in Fig. 1b of the study by Maliti and colleagues [13]. Furthermore, the METs used here are currently research prototypes that are produced individually by the Bioelectronics Department at the University of Glasgow, UK. The Ifakara Health Institute and the University of Glasgow have submitted a joint UK patent application for the MET (application number 1708369.2) which is currently under review. This technology is available for licensing through an Easy Access IP agreement, with design details available on request from the University of Glasgow. The costs of bespoke production per unit on this basis are currently quite high, ~£1100 for a large MET, which is too high for most routine vector density surveillance applications. However the costs of these devices are anticipated to be significantly lower if produced in volume and could become much more economically viable in the future.

Use of METs has helped to confirm key aspects of the feeding behaviour of the main malaria vectors in the Kilombero Valley, and shed light on their ecology and potential response to control. For example, An. funestus s.s. was confirmed as having a preference for humans over cattle, in contrast to An. arabiensis which was attracted to humans and cattle at a similar rate. A previous study in Zimbabwe using e-nets baited with host odour [30] and a modelling analysis of host demography and choice data from northern Tanzania [61] indicate that An. arabiensis had a modest preference for cattle over humans. The lack of preference for cattle in this study may be due to the use of relatively small calves, rather than full-grown adults [30] or entire herds of all ages [61]. Variation in biomass and attractiveness to mosquitoes with age and pregnancy have such strong influence upon human exposure that these demographic factors play a defining role in shaping malaria burden distribution across at-risk populations [62–65]. Evaluation of potential changes in mosquito host preference over time or between sites should ideally use consistent trapping methods with standardized host density and biomass or standardized synthetic odour lures.

Use of METs also helped update and further elucidate the ecology of An. funestus s.l. vectors in Kilombero. Historically, this species group has received relatively little attention in southern Tanzania, but may now be the most important vector of persisting transmission [39]. The species composition of the An. funestus s.l. reported here is consistent with other studies in the Kilombero valley [39, 40], with An. funestus s.s. being the most prevalent, followed by An. rivulorum and An. leesoni. However, only about half of An. funestus s.l. could be successfully identified to species level. The much lower amplification rates for An. funestus s.l. than for An. gambiae s.l. is most probably due to restricted availability of primers (only available for 4 of the 9 species in the r group). The lack of amplification of some An. funestus s.l. could indicate the presence of other species within the An. funestus group.

**Table 3** Proportion of attack of *Anopheles* species on human showing the 95% confidence interval around the preference estimates as were observed from host seeking MET-LC and MET-LH as estimated by binary logistic GLMM regression

| Anopheles species | $P_h$ | 95% CI    | Z value | P value |
|-------------------|-------|-----------|---------|---------|
| An. arabiensis    | 0.56  | [0.43–0.67] | 0.937   | 0.349   |
| An. funestus s.s. | 0.76  | [0.68–0.82] | 5.85    | 0.001   |
| An. rivulorum     | 0.27  | [0.18–0.37] | 0.48    | 0.001   |

In the $P_h$ column, numbers in bracket represent the denominator (e.g. total number of caught host seeking on humans and cattle combined). $P_h$ is the proportion of attack on human.
On the basis of samples that could be identified, these results indicate An. funestus s.s. has a strong preference for humans over cattle as consistent with previous studies [22, 41, 66]. However, the degree of anthropophily in An. funestus s.s. found here is somewhat lower compared than previously reported in northern Tanzania where this species was estimated to feed almost entirely on humans [22]. The results of the present study are consistent with more recent reports from Zambia where the human attack rate of An. funestus was only 41.2% [27], and another in western Kenya where the human blood index of An. funestus was only 60% [21]. These results indicate that An. funestus s.s. in the Kilombero Valley and elsewhere in Africa can exploit cattle as a source of blood. Therefore, this species may be more difficult to eliminate with LLINs and indoor residual spray (IRS). Thus, complementary interventions that target livestock as alternative blood sources may also be required to tackle this species.

**Conclusions**

This study adds to a growing body of evidence that, METs are a sensitive, practical, exposure-free alternative to the HLC gold standard tool for assessing human biting rates and measuring host preferences. Estimates of malaria vector biting rates were considerably higher in large METs, suggesting that total human exposure to bites may be underestimated by conventional methods. METs showed a higher than expected preference upon cattle for An. funestus s.s. suggesting that supplementary interventions may be needed to tackle this important vector.

**Abbreviations**

HLC: human landing catch; MET-SH: small mosquito electrocuting trap baited with human volunteer; MET-LH: large mosquito electrocuting trap baited with human volunteer; MET-UC: large cattle-baited mosquito electrocuting trap; HBI: human blood index; LLINs: long-lasting insecticide nets; IRS: indoor residual spray; PCR: polymerase chain reaction; ELISA: enzyme-linked immunosorbent assay; GLMMs: generalized linear mixed models.

**Authors’ contributions**

FCM, ATM, NJG conducted the experiment. NJG and HMF designed the study and mosquito sampling protocol. GFK, contributed to the study design, analysis and interpretation. FCM, NJG performed data analysis, interpreted the results and drafted the manuscript. KK, GFK, HMF, NM and DFM provided comments upon the manuscript. All authors read and approved the final manuscript.

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**Competing interests**

University of Glasgow and Ifakara Health Institute have filed a patent application for the MET with NJG, DFM, KK, NM, and HMF named in it as investigator. All other authors declare that they have no competing interest.

**Availability of data and materials**

Access and use of data supporting this article will have to comply with the Ifakara Health Institute data sharing policy. If data are requested and no competing interest is apparent, the requested data will be made available under defined conditions expressed in writing through an exchange of letters between parties stipulating those conditions and any agreed limits to use thereof.

**Consent for publication**

Written informed consent was obtained from livestock owners and volunteers for participation in the study, and for publication of this report and any accompanying images. Consent and approval for publication was also obtained from the National Institute of Medical Research in Tanzania.

**Ethics approval and consent to participate**

Before trapping, village leaders were consulted, informed about the purpose of our study and asked for permission to proceed. On gaining their approval, a suitable field was identified, and members of nearby households consulted to seek permission for collection of mosquitoes. Participants in MET and HLC mosquito collections were IHI technical staff who were trained in methods for catching mosquitoes. They agreed to participate by giving informed consent.
after the nature of their participation was explained. As these volunteers participated in HLC procedures, they were all provided with prophylactic atovaquone-proguanil (Malarone®), GlaxoSmithKline, taken daily before mosquito collection. All mosquito catchers were screened weekly for malaria parasites by a rapid diagnostic test. No participant tested positive for malaria at any stage of the study, but if they had they would have been treated with the standard first-line treatment in Tanzania, Artemisinin-Lumefantrine (Co-artem®). No participant tested positive for malaria by a rapid diagnostic test. No participant tested positive for malaria vector populations occurring indoors in rural Africa. Int J Epidemiol. 2013;4:235–47.

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