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# Entomological impact of mass administration of ivermectin and dihydroartemisinin-piperaquine in The Gambia: a cluster-randomized controlled trial

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## Abstract

**Background:** Vector control interventions in sub-Saharan Africa rely on insecticide-treated nets and indoor residual spraying. Insecticide resistance, poor coverage of interventions, poor quality nets and changes in vector behavior threaten the effectiveness of these interventions and, consequently, alternative tools are needed. Mosquitoes die after feeding on humans or animals treated with ivermectin (IVM). Mass drug administration (MDA) with IVM could reduce vector survival and decrease malaria transmission. The entomological impact of MDA of combined IVM and dihydroartemisinin-piperaquine was assessed in a community-based, cluster-randomized trial.

**Methods:** A cluster-randomized trial was implemented in 2018 and 2019 in 32 villages in the Upper River Region, The Gambia. The with the inhabitants of 16 intervention villages eligible to receive three monthly rounds of MDA at the beginning of the malaria transmission season. Entomological surveillance with light traps and human landing catches (HLC) was carried out during a 7- to 14-day period after each round of MDA, and then monthly until the end of the year. The mosquitocidal effect of IVM was determined by direct membrane feeding assays.

**Results:** Of the 15,017 mosquitoes collected during the study period, 99.65% ( $n = 14,965$ ) were *Anopheles gambiae* sensu lato (*An. gambiae* s.l.), comprising *Anopheles arabiensis* (56.2%), *Anopheles coluzzii* (24.5%), *Anopheles gambiae* sensu stricto (*An. gambiae* s.s.; 16.0%) and *Anopheles funestus* sensu lato (*An. funestus* s.l.; 0.35%). No effect of the intervention on vector parity was observed. Vector density determined on light trap collections was significantly lower in the intervention villages in 2019 (adjusted incidence rate ratio: 0.39; 95% confidence interval [CI]: 0.20, 0.74;  $P = 0.005$ ) but not in 2018. However, vector density determined in HLC collections was similar in both the intervention and control villages. The entomological inoculation rate was significantly lower in the intervention villages than in the control villages (odds ratio: 0.36, 95% CI: 0.19, 0.70;  $P = 0.003$ ). Mosquito mortality was significantly higher when blood fed on IVM-treated individuals up to 21 days post-treatment, particularly in adults and individuals with a higher body mass index.

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**Conclusion:** Mass drug administration with IVM decreased vector density and the entomological inoculation rate while the effect on vector parity was less clear. Survival of mosquitoes fed on blood collected from IVM-treated individuals was significantly lower than that in mosquitoes which fed on controls. The influence of host characteristics on mosquito survivorship indicated that dose optimization could improve IVM efficacy. Future detailed entomological evaluation trials in which IVM is administered as stand-alone intervention may elucidate the contribution of this drug to the observed reduction in transmission.

**Keywords:** Ivermectin, Mass Drug Administration, Malaria, Vector

## Introduction

Vector control interventions such as long-lasting insecticidal nets (LLINs) and indoor residual spraying (IRS) are the main components of malaria vector control in sub-Saharan Africa [1, 2]. In The Gambia, prompt diagnosis and treatment with artemisinin-based combinations and the large-scale deployment of LLINs and IRS have resulted in a substantial decline of the malaria burden [3, 4]. Nevertheless, malaria transmission, which is highly seasonal, has not been stopped completely. Significant resistance to dichlorodiphenyltrichloroethane (DDT) and pyrethroids has been recently reported [5–7], which may partly explain, in addition to climate change [8], changes in the density distribution and biting and resting behaviors of *Anopheles gambiae* sensu lato (*An. gambiae* s.l.), the dominant malaria vector, and the heterogeneity of malaria transmission [9, 10]. Indeed, LLINs and IRS protect against those vectors which bite and rest indoors [11], but changes in vector behaviors, such as outdoor biting and/or biting earlier [10, 12], vector biodiversity and environmental change [13], may decrease the protection provided by these interventions. A recent study in The Gambia reported a significant preference of *Anopheles arabiensis* for outdoor resting [14], thereby decreasing the effect of standard vector control interventions such as IRS and LLINs. This behavior highlights the need for insecticides other than pyrethroids [1] and for the targeting of vectors currently able to escape standard control interventions [15].

Ivermectin (IVM) is an endectocide, broad-spectrum systemic drug that is efficacious against nematodes and arthropods [16]. When IVM is ingested through a blood meal from an IVM-treated human or animal, it exerts a lethal effect by acting on the glutamate-gated chloride channels of arthropods [17, 18], disrupting their neuromuscular transmission and leading to paralysis and death [19]. IVM has the potential to target both insecticide-resistant and outdoor-biting *Anopheles* mosquitoes [2, 12]. Therefore, mass drug administration (MDA) of IVM may decrease the survival of human biting mosquitoes, regardless of their behavior [20–23] and insecticide resistance status [24, 25]. In Africa, MDA with IVM was found to temporarily alter the age structure of

mosquito populations and reduce malaria transmission by reducing vector survival and thus the capacity to complete the malaria parasite sporogonic cycle [26, 27]. The effect of MDA with IVM on malaria transmission can be assessed by implementing a community-based, cluster-randomized trial [28]. In Burkina Faso, repeated rounds of MDA with IVM reduced the incidence of clinical malaria, without concurrent reductions in entomological exposure indicators [29], highlighting the need for dedicated entomological evaluations on the impact of IVM on mosquito populations. Here, we report a detailed analysis of the entomological impact of MDA with IVM and dihydroartemisinin-piperaquine within a cluster-randomized trial that was carried out in eastern Gambia [30].

## Methods

### Study site and trial procedures

Thirty-two villages which were located at least 3 km apart were selected according to malaria prevalence determined by an earlier cross-sectional survey [31] and randomized to either the intervention (16 villages) or control group (16 villages). A buffer zone of 2-km radius was established around each intervention village to minimize contamination from neighboring untreated villages [30]. All villages within the buffer zone received the intervention although they were not included in the evaluation. In 2018 and 2019, monthly rounds of MDA with IVM (Laboratorio Elea, Los Polvorines, Argentina), administered at a dose of 300–400 µg/kg body weight per day for 3 consecutive days, and dihydroartemisinin-piperaquine (DP; Guilin Pharmaceuticals, Guilin, Guangxi, China), administered according to body weight following the manufacturer's guidelines, were performed for three consecutive months at the beginning of each malaria transmission season: in August, September and October of 2018 and in July, August and September of 2019. All drugs were administered orally. During each round of MDA, the research team covered all 16 intervention villages between 12 and 14 days, with each individual village covered within a period of about 3–5 days.

Malaria transmission in The Gambia is highly seasonal, with a peak in October–November [32]. Information on rainfall, temperature and humidity are presented in

Additional file 1: Table S1. The annual temperature and humidity were similar for both study years.

### Entomological collections

Mosquitoes were collected using standard US Center for Disease Control and Prevention (CDC) light traps (CDC-LT; CDC, Atlanta, GA, USA) hung from the ceiling at the foot end of a bed with the light 70–150 cm above the ground [33]. Intensive sampling to measure mosquito parity and density was carried out from 7 to 14 days after each MDA round, and then monthly until the end of the transmission season (December). Mosquitoes were collected over three consecutive nights in six randomly selected houses per village in all intervention villages and in six randomly selected houses in eight control villages. For the other control villages, collections were carried out at the same time for one night. CDC-LT were set up by trained field assistants and run for 12 h, from 19:00 h until 7:00 h. The CDC-LT were checked every 4 h.

Monthly human landing catches (HLC), both indoors and outdoors, were carried out in four intervention and four control villages that were randomly selected, both in 2018 and in 2019. In 2018, collections were done for three consecutive nights in six houses randomly selected using village census identification; in 2019, collections were done in three houses per village for two consecutive nights. HLC were done from 19:00 h to 07:00 h by four volunteers (2 indoors and 2 outdoors) who rotated every 2 h to avoid collector bias.

Each morning, all collected mosquitoes were transported to the laboratory where they were morphologically identified and stored in separate tubes with silica gel for further analysis, while other anophelines and culicine mosquitoes were counted and then discarded. A subset of *An. gambiae* s.l. ( $N = 12$  per night from HLC [6 outdoors and 6 indoors] and  $N = 10$  per room per day from CDC-LT) were used to estimate mosquito parity [34]. Head and thorax (500 per each collection round per arm if available) were used for the detection of *Plasmodium falciparum* circumsporozoite protein (CSP) by enzyme-linked immunosorbent assay (ELISA) [35]. Abdomens from a subset of samples processed for ELISA were used for mosquito identification by molecular methods [36].

### Membrane feeding experiment

A subset of the study population, comprising 80 randomly selected participants (50% aged 4–10 years old [children] and 50% aged  $\geq 18$  years [adults]) living in one intervention village ( $N = 40$ ) and one control village ( $N = 40$ ), were selected for participation in the direct membrane feeding assay (DMFA). These villages were chosen for their proximity to the insectary of the Medical Research Council Unit The Gambia (MRCG) field

station in Basse, The Gambia. Venous blood samples were collected in 4-ml tubes coated in lithium-heparin (BD, Franklin Lakes, NJ, USA) at 7, 14 and 21 days after the administration of the first dose of IVM from all participants in the intervention villages and in the control villages. For the intervention villages, in 2018, study participants were randomly selected, without confirming whether they had actually taken IVM; in 2019, only individuals who had taken all IVM doses under direct supervision were included in the analysis.

*Anopheles coluzzii* mosquitoes were reared in the insectary at 27 °C and approximately 70–80% relative humidity (RH) under a 12/12-h day/night cycle and fed on 5–10% glucose. Immediately after phlebotomy, two aliquots of 400–500 µl of whole blood were dispensed into two glass feeders, and 50 female *An. coluzzii* mosquitoes (aged 2–6 days) per feeder (total  $N = 100$ ) were allowed to feed through a Parafilm membrane for 20 min. After feeding, partially fed mosquitoes were removed and fully fed mosquitoes were kept at 27 °C in a specific container. Mortality was estimated daily, up to 14 days post-feeding.

### Entomological parameters and statistical analysis

Vector density, parity rate, species composition and sporozoite rate were determined from the CDC-LT collections and HLC. The biting rate and the entomological inoculation rate (EIR) were determined only from HLC collections.

Mosquito density was calculated as the number of collected mosquitoes divided by the number of trapping nights. For CDC-LT, mixed-effects generalized models with negative binomial distribution were used to determine the impact of MDA on mosquito density, controlling for MDA round as a categorical fixed effect. The village was included in the model as a random effect. Since the number of villages for HLC collections was small, the effect of the intervention was estimated based on village-level summaries [37]. The analysis was conducted in two stages: first, we used a Poisson regression model to compute a residual for each village after adjusting for the MDA round; second, we calculated the ratio of the observed to the predicted events for each village. The unpaired t-test to determine the mean difference between the two groups.

The parity rate was estimated for each collection method by dividing the number of parous mosquitoes by the total number of parous and nulliparous mosquitoes collected. For CDC-LT, mixed-effects logistic regression was used to model the impact of MDA on parity, with village included as a random effect. The models were adjusted for MDA round. For HLC collections, the effect of the intervention was based on village-level summaries.

**Table 1** Vector species composition by study arm and combined study years (2018 and 2019)

Vector species identification	Intervention, % (N)	Control, % (N)	Total, % (N)
<i>Morphological identification</i>			
<i>Anopheles gambiae</i> sensu lato	99.4 (5407)	99.8 (9557)	14,964 (99.65)
<i>Anopheles funestus</i> sensu lato	0.6 (30)	0.24 (23)	0.4 (53)
Total N	5437	9580	15,017
<i>Molecular identification<sup>a</sup></i>			
<i>Anopheles arabiensis</i>	59.5 (1192)	53.2 (1166)	56.2 (2,358)
<i>Anopheles coluzzii</i>	26.1 (524)	23.0 (504)	24.5 (1,028)
<i>Anopheles gambiae</i> sensu stricto	11.9 (238)	19.7 (432)	16.0 (670)
<i>Anopheles gambiae</i> /coluzzii	50 (2.5)	4.1 (91)	3.3 (141)
Total N	2004	2,193	4197

N Number of mosquitoes

<sup>a</sup> Only *Anopheles gambiae* sensu lato

The first step was to fit a logistic regression model that adjusted for the MDA round on individual-level data, ignoring the intervention effect and clustering. Next, we estimated the expected number of parous mosquitoes for each village and calculated residuals as a ratio of the number of expected parous mosquitoes to the number of observed parous mosquitoes. The intervention effect was calculated as the ratio of mean residuals between treatment arms. To determine the significance of the difference between the treatment arms, the unpaired t-test was applied to the village-level residuals.

Sporozoite rate was estimated by dividing the number of CSP-positive mosquitoes by the total number of mosquitoes analyzed. We used mixed-effects logistic regression to model the impact of MDA on sporozoite rate with village included as a random effect.

The biting rate was estimated by dividing the number of mosquitoes that were collected by the number of those capturing the mosquitoes. The effect of the intervention on the biting rate was estimated based on village-level summaries. The analysis was conducted in two stages: first, we used a Poisson regression model to compute a residual for each village after adjusting for the MDA round; second, we calculated the ratio of the observed to the predicted events for each village. The unpaired t-test was used to determine the significance of the mean difference between the two groups. Permutation tests were used to validate t-test P-values.

A summary EIR for the two transmission seasons (2018 and 2019) was estimated by multiplying the sporozoite rate for the HLC by the biting rate, multiplied by 360 days. A mixed-effects generalized model with negative binomial distribution was fitted to determine the impact of IVM on mosquito survival following DFMA. Study subject was included in the model as a random effect. The model was adjusted for age, gender and body

mass index (BMI) of the study subject. The analysis was further stratified by year of MDA administration and by age and gender of the study participant.

#### Ethical considerations

The study protocol received ethical approval from The Gambia Government/MRC Joint Ethics Committee and the London School of Hygiene and Tropical Medicine Ethics Committee. Before any procedure was initiated, written informed consent was provided by adult participants and by parents/guardians of children. Children aged 12 to < 18 years provided their assent.

#### Results

Between August 2018 and January 2019 and between July and December 2019, a total of 4116 trapping nights involving CDC light-traps in 192 households and 924 trapping nights involving HLC in 72 households were completed. When the results of both trapping methods were combined, most vectors (99.6%, 14,964/15,017 mosquitoes) were morphologically identified as *Anopheles gambiae* s.l., with no difference between study arms (Table 1). The remaining mosquitoes were identified as *Anopheles funestus* sensu lato (*An. funestus* s.l.).

During the 2 years of the study, the mean number of *An. gambiae* s.l. caught per night by CDC-LT was 1.90 and 0.88 in the control and intervention arms, respectively, and by HLC, 5.46 and 3.44 in the control and intervention arms, respectively (Table 2).

Species identification by PCR (molecular method) was performed on 4197 *An. gambiae* s.l. samples. Of these, approximately one half (56.2%,  $N = 2358$ ) were identified as *An. arabiensis*, 24.5% (1028) as *An. coluzzii* and 16.0% (670) as *Anopheles gambiae* sensu stricto (*An. gambiae* s.s.), with no difference between intervention and control villages (Table 1).

**Table 2** Mean density of *Anopheles gambiae* sensu lato per trapping night by year and collection method

Year	Collection method	Arm	Number of mosquitoes collected	Trapping nights (N)	Mean density per trapping night
2018	CDC-LT	Control	1571	1002	0.81
2018	CDC-LT	Intervention	564	858	0.26
2018	HLC	Control	4583	360	12.49
2018	HLC	Intervention	2358	324	3.14
2019	CDC-LT	Control	3088	912	2.45
2019	CDC-LT	Intervention	1914	1344	1.04
2019	HLC	Control	315	120	2.35
2019	HLC	Intervention	571	120	4.56
2018 + 2019	LTC	Control	4659	1914	1.90
2018 + 2019	LTC	Intervention	2478	2202	0.88
2018 + 2019	HLC	Control	4898	480	5.46
2018 + 2019	HLC	Intervention	2929	444	3.44

CDC-LTC US Center for Disease Control and Prevention light trap, HLC human landing catch

**Table 3** Vector parity and density by year and study group

Parameter	Methods	2018				2019			
		Control, % (n/N)	Intervention, % (n/N)	Adj. RR (95% CI)	P-value	Control, % (n/N)	Intervention, % (n/N)	Adj. RR (95% CI)	P-value
Parity	CDC-LT	53.2 (91/171)	54.9 (50/91)	1.17 (0.93, 1.47)	0.181	81.7 (441/540)	84.2 (309/367)	1.01 (0.95, 1.52)	0.708
	HLC	68.4 (807/1180)	61.5 (475/772)	0.87 (0.75, 1.01)	0.055	89.5 (111/124)	76.3 (132/173)	0.93 (0.65, 1.34)	0.649
Density <sup>a</sup>	CDC-LT	282.2 (1571/1002)	118.3 (564/858)	0.47 (0.14, 1.57)	0.213	609.5 (3088/912)	256.3 (1914/1344)	0.39 (0.20, 0.74)	0.005
	HLC	2291.5 (4583/360)	1310.0 (2358/324)	0.45 (0.15, 1.35)	0.125	472.5 (315/120)	856.5 (571/120)	1.94 (0.88, 4.28)	0.088

Adj. IRR adjusted incidence rate ratio, CI confidence interval, n number of parous mosquitoes, N number of dissected mosquitoes, OR odds ratio, RR relative risk

<sup>a</sup> Number of mosquitoes per trapping nights multiplied by 180 days (30 days per month multiplied by 6 months per season per year)

Parity could be successfully assessed in 1169 mosquitoes captured by CDC-LT, representing about 17% (1169/6921) of all trapped mosquitoes, with the others being often desiccated prior to emptying CDC light-traps. The proportion of mosquitoes on which parity assessment could be performed was similar between the intervention (22.6%; 458/2029) and control (14.5%; 711/4892) villages, without any indication of bias. There was no difference in parity rates as estimated from CDC-LT collections between the intervention and control arms in both 2018 (adjusted relative risk [RR]: 1.17; 95% CI: 0.93, 1.47;  $P = 0.18$ ) and 2019 (adjusted RR: 1.01; 95% confidence interval [CI]: 0.95, 1.07;  $P = 0.71$ ) (Table 3). Parity as determined by HLC tended to be lower in the intervention group than in the control group, both in 2018 (adjusted RR: 0.87; 95% CI: 0.75, 1.01;  $P = 0.055$ ) and 2019 (adjusted RR: 0.93; 95% CI: 0.65, 1.34;  $P = 0.649$ ).

Vector density estimated from CDC-LT collections was lower in the intervention villages than in the control

villages, particularly in 2019 (adjusted RR: 0.39; 95% CI: 0.20, 0.74;  $P = 0.005$ ), but in 2018 this difference did not reach statistical significance (Table 3). However, vector density as estimated by HLC tended to be lower in 2018 and higher in 2019 in the intervention villages compared to the control villages (Table 3).

Using HLC data, no statistically significant differences in biting and sporozoite rates were observed between the intervention and control groups (Table 4). However, the overall EIR was significantly lower in the intervention group than in the control group (odds ratio [OR]: 0.36; 95% CI: 0.19, 0.70;  $P = 0.003$ ).

The mortality of mosquitoes fed on blood collected from individuals in the intervention villages was significantly higher than that of those fed on blood collected from individuals in the control villages. The highest mortality was observed during the first days after blood-feeding (Fig. 1), particularly in mosquitoes fed on blood collected from individuals 7 and 14 days after the first

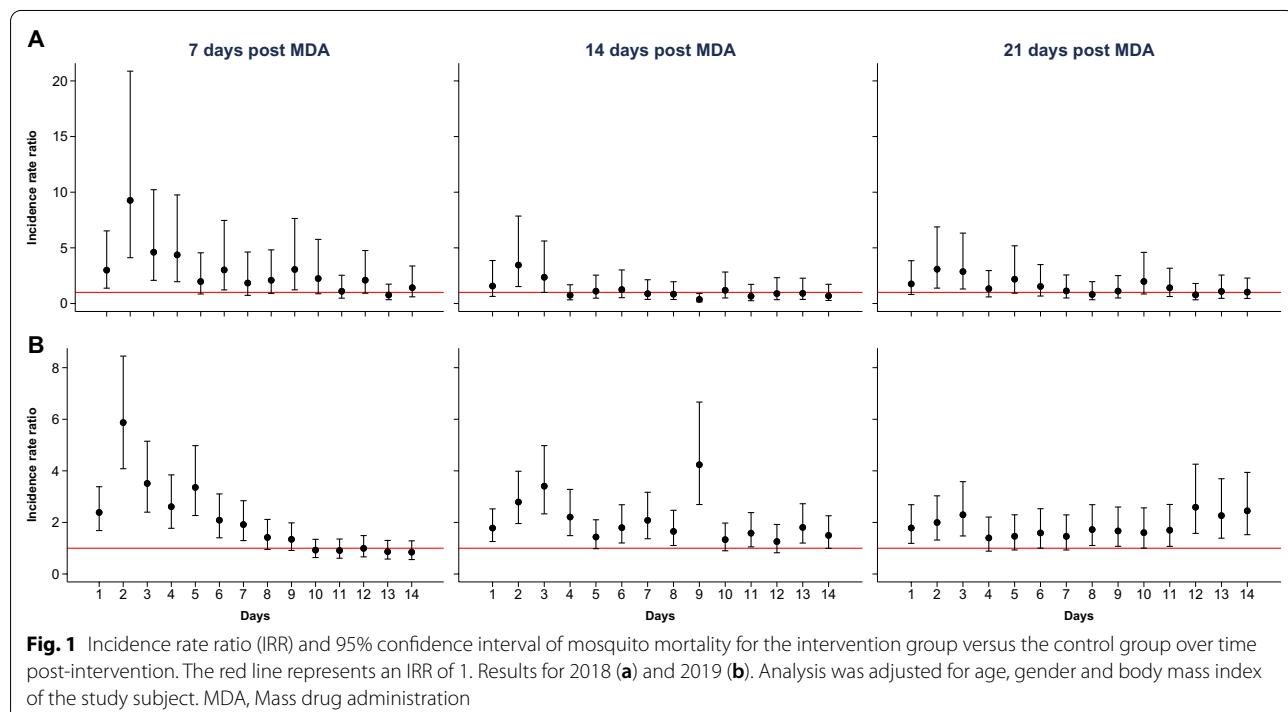
**Table 4** Vector sporozoite rate, biting rate and entomological inoculation rate for 2018 and 2019 combined

Parameter	Methods	Control, % (n/N)	Intervention, % (n/N)	Adj, IRR (95% CI)	P-value
Biting rate, %	HLC	3.4 (4898/1440)	2.2 (2929/1344)	0.78 (0.23, 2.77)	0.701
Parameter	Methods	Control, % (n/N)	Intervention, % (n/N)	Crude OR (95% CI)	P-value
Sporozoite rate, %	CDC-LT and HLC	1.1 (68/6001)	0.8 (31/3939)	0.69 (0.46, 1.06)	0.091
EIR	HLC	56.9 (4898/360) × (29/2498) × 360	20.8 (2929/360) × (13/1835) × 360	0.36 (0.19, 0.70)	0.003

EIR Entomological inoculation rate

Biting rate: n number of mosquito collected; N number of capturers

Sporozoite rate: n number of mosquito positive for sporozoite; N number of mosquito processed for sporozoite



**Fig. 1** Incidence rate ratio (IRR) and 95% confidence interval of mosquito mortality for the intervention group versus the control group over time post-intervention. The red line represents an IRR of 1. Results for 2018 (a) and 2019 (b). Analysis was adjusted for age, gender and body mass index of the study subject. MDA, Mass drug administration

IVM dose. Mortality remained high among mosquitoes fed on blood collected from individuals 21 days after the first IVM dose (Fig. 1; Table 5).

The mosquitocidal effect of IVM at day 7 post-MDA did not vary by age, gender, year of MDA and BMI (Table 5). Nevertheless, at days 14 and 21 post-treatment, the effect was significantly higher in 2019 than in 2018. Mosquito mortality was also higher at days 14 (incidence rate ratio [IRR]: 1.57; 95% CI: 1.22, 2.02;  $P < 0.001$ ) and 21 (IRR: 1.78; 95% CI: 1.27, 2.49;  $P < 0.001$ ) post-treatment among individuals with a BMI of at least 25 than in those with a lower BMI. The mosquitocidal effect at day 21 post-treatment was also significantly higher in older children (IRR: 1.67; 95% CI: 1.11, 2.50;  $P = 0.013$ ) and adults

(IRR: 1.90; 95% CI: 1.27, 2.85;  $P = 0.002$ ) than in children aged < 5 years.

## Discussion

We report here the entomological effect of an intervention that aimed to reduce the human reservoir of infection using dihydroartemisinin-piperaquine and vector survival and density using IVM. While both the incidence of clinical malaria and the prevalence of infection were significantly lower in the intervention villages than in the control villages [30], the effect of IVM on the vector was less evident, with some apparently contradictory results.

Vector parity, the primary entomological endpoint, was similar between the intervention and control groups

**Table 5** Incidence rate ratios for the effects of mass drug administration on mosquito mortality based on combined 2018 and 2019 data

Variables	7 days post-MDA			14 days post-MDA			21 days post-MDA		
	IRR	95% CI	P-value	IRR	95% CI	P-value	IRR	95% CI	P-value
<i>Treatment day</i>									
1	2.38	1.72, 3.29	<0.001	1.64	1.16, 2.31	0.005	1.69	1.16, 2.45	0.006
2	6.20	4.42, 8.69	<0.001	2.87	2.03, 4.07	<0.001	2.04	1.39, 3.00	<0.001
3	3.81	2.69, 5.40	<0.001	3.14	2.18, 4.53	<0.001	2.34	1.57, 3.49	<0.001
4	3.02	2.12, 4.29	<0.001	1.64	1.13, 2.37	0.009	1.33	0.88, 2.01	0.171
5	2.91	2.03, 4.17	<0.001	1.35	0.93, 1.94	0.113	1.52	1.00, 2.29	0.048
6	2.13	1.48, 3.08	<0.001	1.64	1.12, 2.41	0.012	1.51	1.00, 2.30	0.051
7	1.82	1.26, 2.63	0.001	1.67	1.13, 2.48	0.010	1.33	0.88, 1.99	0.176
8	1.53	1.06, 2.20	0.023	1.42	0.97, 2.07	0.072	1.46	0.97, 2.19	0.069
9	1.44	1.01, 2.06	0.046	2.18	1.47, 3.24	<0.001	1.48	0.99, 2.21	0.055
10	0.99	0.70, 1.40	0.959	1.33	0.91, 1.93	0.143	1.64	1.07, 2.51	0.022
11	0.97	0.67, 1.39	0.848	1.33	0.90, 1.98	0.154	1.55	1.03, 2.35	0.037
12	1.23	0.86, 1.77	0.263	1.15	0.77, 1.72	0.504	1.82	1.18, 2.80	0.007
13	0.86	0.59, 1.23	0.403	1.58	1.07, 2.34	0.022	1.81	1.17, 2.79	0.007
14	0.96	0.66, 1.40	0.846	1.32	0.89, 1.95	0.170	1.88	1.24, 2.86	0.003
<i>Age group (years)</i>									
<5	1			1			1		
5–15	1.02	0.82, 1.26	0.891	1.07	0.80, 1.44	0.651	1.67	1.11, 2.50	0.013
>15	1.07	0.87, 1.33	0.515	1.25	0.93, 1.68	0.140	1.90	1.27, 2.85	0.002
<i>Gender</i>									
Female	1			1			1		
Male	1.05	0.93, 1.19	0.400	0.97	0.83, 1.14	0.716	0.99	0.80, 1.23	0.961
<i>Year</i>									
2018	1			1			1		
2019	1.06	0.91, 1.23	0.456	1.72	1.41, 2.10	<0.001	1.73	1.34, 2.24	<0.001
<i>Body mass index (kg/m<sup>2</sup>)</i>									
<25	1			1			1		
≥25	1.02	0.84, 1.24	0.838	1.57	1.22, 2.02	<0.001	1.78	1.27, 2.49	<0.001

MDA Mass drug administration

when estimated by both the CDC-LT and HLC collections, although when estimated used HLC collections parity tended to be lower in the intervention group, both in 2018 and in 2019. Parity could be determined in only 15–20% of mosquitoes captured in CDC-LT, and mostly in those captured in the early hours of the morning as those captured earlier may have died and dried up or been damaged, making dissection impossible. Conversely, mosquitoes captured in HLC can remain alive until the following morning when ovary dissection can be carried out. However, in the present study, HLC could be conducted in only a small number of villages. We therefore did not achieve the number of observations for parity we intended, reducing the study power to detect a statistically significant difference. A similar lack of effect of MDA with IVM on vector parity was observed in Burkina Faso, although serological reactivity to an anopheles

salivary gland protein was significantly lower in the intervention group than in the control group, suggesting a lower exposure of individuals to mosquito bites [38]. Another potential explanation for the lack of effect on vector parity may be spillover from neighboring villages, despite a 2-km buffer zone around intervention villages, as mosquitoes can fly for a distance of up to 3 km when unfed, 9 km when sugar fed and 10 km when blood fed [39], as has been observed previously in The Gambia [40].

In 2018, vector density determined on CDC-LT collections tended to be lower in the intervention than control villages, but the difference was not statistically different; in 2019 this difference became statistically significant, possibly reflecting the higher coverage achieved in 2019. However, vector density determined on HLC collections did not differ between the intervention and control groups, although there was a tendency in 2019 for it to

be higher in the intervention villages, but without reaching statistical significance. This apparent discrepancy between CDC-LT and HLC data may be due to the low number of villages sampled using HLC (8 from a total of 32 study villages), which limited the power to detect a possible difference. Considering the marked heterogeneity in entomological parameters between villages, which reflected both actual differences and the variability within the entomological collection methods [41], the tendency for a higher vector density in intervention villages observed in 2019 could be due to differences that were present prior to the intervention.

The effect of IVM on vector survival is shown by the DMFA results, obtained by feeding colony mosquitoes with blood samples from inhabitants of one intervention and one control village, respectively. Previous studies have reported a >90% mortality of different anophelines fed on human blood collected individuals immediately after IVM treatment, with a subsequent rapidly declining efficacy over time [42]. Nevertheless, mosquito survival was found to decrease significantly for at least 28 days after feeding on blood collected from individuals after treatment with IVM at either 300 or 600 µg/kg per day for 3 days [43]. It is likely that IVM metabolites contribute to the observed mortality [44], and this should be further investigated [45]. In our study, the mosquitocidal effect of IVM was predictably stronger at 7 days after the first dose, although the effect remained detectable at 14 and 21 days after treatment [43].

Our findings on the daily survivorship allowed a detailed examination of the kinetics of mosquito mortality. Mosquitoes fed on treated blood continued to experience increased mortality up to 10–14 days after feeding, particularly when feeding on blood collected 21 days post-treatment. Such a delayed mortality could partly explain the lower-than-expected effect on vector parity as the reproductive cycles in *An. gambiae* could be as short as 2 days [40], and thus the female could lay eggs and become parous before IVM has a mosquitocidal effect. It is unclear to which extent IVM would alter vector behavior and the reproductive cycle. In Tanzania, blood digestion in mosquitoes fed on IVM-treated cattle was much slower and egg production decreased up to 15 days post-feeding [46]. Moreover, *An. arabiensis* fed on blood from IVM-treated individuals (7 and 10 days post-treatment) produced significantly fewer eggs than those fed on untreated controls [47]. Therefore, IVM may significantly alter the mosquito reproductive cycle although a subset of the mosquitoes exposed to IVM can successfully complete the gonotrophic cycle, hence diluting any effect IVM may have on the age structure of the vector populations.

Nevertheless, the effect of IVM on vectors' reproductive cycle would, by slowing down blood digestion as well as egg laying and hatching, translate into lower vector density, as observed in this trial.

Interestingly, the mosquitocidal effect of IVM, as determined in the feeding assays, was associated with host characteristics. Although in a previous study the results on blood collected at day 7 post-treatment did not show any variation by age, year of intervention and BMI [29], blood samples collected at day 14 and 21 post-treatment, when the effect of IVM was waning, did show some important differences that may be related to the pharmacokinetics of IVM. In that study, the effect of IVM was still visible at day 21 post-treatment in older children and adults but not in children aged < 5 years. Considering that one of the inclusion criteria for treatment with IVM was a body weight ≥ 15 kg or height ≥ 90 cm, this young age group included mainly children aged 4 years [48]. Nevertheless, results suggest that IVM may be more rapidly eliminated in these children than in older children and adults. Recent pharmacokinetics analyses indicate that IVM-treated children aged < 12 years reach half the peak concentration and total exposure as adults [49]. In addition, a relative underdosing in children for other drugs used in malaria control has been reported [50, 51].

In individuals with a BMI ≥ 25, IVM had a much longer and significant effect at days 14 and 21 post-treatment than in "thinner" individuals, and this effect remained apparent after controlling for age, year of intervention and gender. This effect is possibly explained by the accumulation of IVM in fat tissue, which would be released slowly, increasing the concentration of IVM in the blood over time and thus resulting in a higher and prolonged mosquitocidal effect [43]. Interestingly, a previous study showed that despite a predicted higher concentration of IVM in capillary blood, which mosquitoes probe naturally rather than venous blood, vector mortality after direct skin feeding was similar to that after membrane feeding [52].

This study has a number of limitations. In addition to the relatively low coverage achieved in 2018, a major limitation is the lack of baseline entomological data from all study villages. Due to limited resources, HLC could be implemented in only a few study villages, limiting the capacity of estimating the vector parity rates, which was the primary entomological endpoint and an important parameter to disentangle the effect of IVM from that of dihydroartemisinin-piperaquine. It was also not possible to estimate the human blood index, which limited our capacity to determine the proportion of vectors not feeding on humans and thus not exposed to the intervention.

## Conclusions

Mass drug administration efforts with IVM decreased vector density and EIR while the effect on vector parity is less clear. The individual contribution of IVM to the observed reduction in transmission cannot be clearly defined as the intervention combined IVM with dihydroartemisinin-piperaquine. A more extensive entomological evaluation of the impact of MDA with IVM alone is needed, and ongoing studies will hopefully provide such information.

## Abbreviations

BMI: Body mass index; CDC-LT: US Center for Disease Control and Prevention's light trap; CI: Confidence interval; CSP: Circum sporozoite protein; DMFA: Direct membrane feeding assay; DP: Dihydroartemisinin-piperaquine; EIR: Entomological inoculation rate; HLC: Human landing catch; IRR: Incidence rate ratio; IVM: Ivermectin; MDA: Mass drug administration; MRC: Medical Research Council; OR: Odds ratio; RR: Relative risk.

## Supplementary Information

The online version contains supplementary material available at <https://doi.org/10.1186/s13071-022-05557-4>.

**Additional file 1: Table S1.** Weather information of the study area (2018 and 2019 rainy seasons).

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## Author contributions

HMS: Investigation, methodology, writing of original draft, formal analysis, review and editing. EDD and UDA: Investigation, methodology, review and editing. JB, SWL, CD, AE and TB: review and editing. MMC, LJ, PMG, SK, EAJ, AKN, FS, MON, BC, SC: Investigation, methodology. NIM and MO: Data analysis. UDA: Conceptualization, funding acquisition, investigation, methodology, project administration, supervision, validation, visualization, review and editing. All authors read and approved the final manuscript.

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## Availability of data and materials

After publication, trial data will be made available on reasonable request to the corresponding author. A proposal with a detailed description of study objectives and a statistical analysis plan is needed for assessment of requests. Additional materials may also be required during the process. Deidentified participant data will be provided after approval by the sponsor and trial management group.

## Declarations

### Ethics approval and consent to participate

The study protocol received ethical approval from The Gambia Government/MRC Joint Ethics Committee and the London School of Hygiene and Tropical Medicine Ethics Committee. Before any procedure was initiated, written informed consent was provided by adult participants and by parents or guardians of children. Children aged 12 to <18 years provided their own assent.

### Competing interests

We declare no competing interests.

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