

**Infection history and current co-infection with *Schistosoma mansoni*
decreases *Plasmodium* species intensities in pre-school children from
Uganda.**

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Summary: *Plasmodium-Schistosoma mansonii* co-infections are associated with a reduced *Plasmodium* intensity in pre-school aged children. This relationship, however, is modified by prior infections, host age and family wealth.

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Abstract

Malaria-schistosomiasis co-infections are common in sub-Saharan Africa but studies present equivocal results regarding the inter-specific relationships between these parasites. Through mixed model analyses of a dataset of Ugandan preschool children, we explore how current co-infection and prior infection with either *Schistosoma mansoni* or *Plasmodium* species, alter subsequent 1) *Plasmodium* intensity 2) *Plasmodium* risk and 3) *S. mansoni* risk. Co-infection and prior infections with *S. mansoni* were associated with reduced *Plasmodium* intensity, moderated by prior *Plasmodium* infections, wealth and host age. Future work should assess whether these interactions impact host health and parasite control efficacy in this vulnerable age group.

Key words: Co-infection, *Schistosoma mansoni*, *Plasmodium* species, Pre-school aged children, Prior infections.

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Introduction

Malaria and schistosomiasis are two of the most important tropical parasitic diseases, each infecting over 200 million people per year with substantial morbidity and mortality [1,2]. Preschool aged children (PSAC, aged <6 years old) can be at high risk for both infections [1,2]. Throughout sub-Saharan Africa, malaria and schistosomiasis are co-endemic and co-infections (i.e., both species infecting the host simultaneously) are common [3]. Several research studies suggest that *Plasmodium* species and schistosomes interact during co-infection, yet none have explicitly considered the consequences of co-infection in PSAC. This exposes a crucial knowledge gap in current options for effective control in younger children with no combined control strategy currently developed [1,2].

Prior studies of interactions between *Plasmodium* and schistosomes have provided equivocal outcomes, with some studies reporting increased *Plasmodium* intensity, prevalence or incidence when in co-infection with schistosomes [3-5], while others suggest a decrease [3,6]. All studies to date have been conducted on older cohorts. Older study cohorts are likely to have had multiple cycles of infection, treatment and developed acquired immunity for both *Plasmodium* and schistosomes, biasing the immune system in ways which may contribute to disparities observed between studies [7].

To shed light on the dynamics of malaria and schistosomiasis in young children, we examined *Plasmodium-Schistosoma mansoni* co-infections in PSAC, where we assessed whether prior infections and co-infections altered *S. mansoni* and *Plasmodium* prevalence, and *Plasmodium* intensity. We conducted an in-depth secondary analysis of the “Schistosomiasis In Mothers and Infants (SIMI)” dataset [8] which contains infection histories for each child, enabling us to tease

apart the relationship between prior infections and current co-infections for *Plasmodium* and *S. mansoni*. Previous work indicates that *P. falciparum* is the main species in our study regions [9]. The SIMI dataset does not, however, include species specific measures and, therefore, we will use *Plasmodium* throughout since we cannot be certain of the *Plasmodium* species. In line with studies in older children we hypothesise that *S. mansoni* will increase the risk and intensity of *Plasmodium* infections but that *S. mansoni* will be unaffected by *Plasmodium*.

Methods

The SIMI project was approved by the London School of Hygiene and Tropical Medicine, UK (LSHTM 5538.09) and the Ugandan National Council of Science and Technology. In brief, for primary data collection informed consent was provided in writing, or by a thumbprint in cases of illiteracy, by mothers for themselves and on behalf of their children before enrolment (further details in [8]).

The SIMI study took place in Uganda at six villages around Lake Albert (Piida, Bugoigo, Walukuba) and Lake Victoria (Bugoto, Bukoba, Lwanika), where infections in each child and mother were assessed at six-monthly intervals. We use data from the initial baseline assessment and the six-month follow up due to participant drop-out at later surveys. Extracted data details *S. mansoni* and *Plasmodium* infections of PSAC, presence/absence of any soil transmitted helminths (STH), information on infection risk behaviours, age, sex, tribe and mother's education level and occupation. Variables indicative of a family's socio-economic status (supplementary Table 1) were used to generate a wealth index using a principal component analysis (PCA, Gwatkin et al, 2000). Variables were included in the PCA unless greater than 95% of the participating families, or less than 5%, owned the asset to ensure a reasonable level of variation was included [10]. The first principal

component of the PCA was extracted to form the wealth index, then divided into equal wealth quintiles.

The presence/absence of *S. mansoni* was determined by a urine schistosome Circulating Cathodic Antigen (CCA) rapid diagnostic test. The intensity of a *S. mansoni* infection was inferred from Kato-Katz faecal egg counts, as eggs per gram of faeces (EPG) and split into, low (<100), moderate (100-399) and high intensity (>399) according to WHO guidelines [11]. Parasitaemia/ μ l blood, assessed from Giemsa-stained blood films, was used both to determine presence/absence and intensity of *Plasmodium* infection. Presence of STH was determined using the Kato-Katz method. Praziquantel and Albendazole were provided to all children at the baseline survey, regardless of infection status in line with mass drug administration protocols. Antimalarials (Lonart[®]) were provided if *Plasmodium* was detected by microscopy or by a fingerpick rapid diagnostic test [9].

Of the 1211 children in the original dataset, 520 were excluded due to an incomplete sample profile (supplementary Figure 1). Data from missing participants was representative of those individuals included for all variables in the analyses (supplementary Table 2). The remaining 706 children provided a blood sample for *Plasmodium* diagnosis, a urine sample for schistosome CCA, and a stool sample to detect schistosome and STH egg patent infections, at both baseline and the six-month survey. The participants summary statistics are included in Supplementary Table 3.

All analyses were completed in R version 3.5.2 statistical software (R core team, 2018). The effect of infection history (infection at baseline) and co-infection (co-infection at the six-month survey) on 1) *Plasmodium* intensity, log transformed to normalise the distribution ($\ln(x+1)$), 2) *Plasmodium* risk and 3) *S. mansoni* risk were assessed in a series of general linear (*Plasmodium* intensity) and

generalised linear (risk models) mixed models using the ASReml-R v4 mixed modelling package (VSN International Ltd., Hemel Hempstead, UK). In each assessment two models were created, with one set incorporating all infections as presence/absence data and the other including *S. mansoni* and *Plasmodium* (Ln(x+1)) data as inferred intensities. These latter models allowed the investigation of potential non-linear relationships between the parasites. *Plasmodium* intensity models had a Gaussian error distribution and an identity link function. Risk models used a binomial error distribution with a logit link function.

Baseline prevalence of STH was included in all models, however, the six-month follow up STH prevalence was only 3.9% and therefore excluded. All starting models included a child's age, sex, family's wealth quintile, their tribe and their mother's occupation and education level, behavioural variables associated with *S. mansoni* infection (the number of times a child bathed and how long they spent in water) and *Plasmodium* infection (the use of bed nets, the presence of mosquitoes in the home and whether a person sleeps outside) and the random terms of family identifier, the child's village, nested within lake, and age fitted with a cubic smoothing spline, as parasite-age profiles are often non-linear [12]. Detailed model structures can be found in supplementary Table 3. For all models, the random terms were refined using maximum likelihood ratio tests, then, stepwise deletion of insignificant fixed terms was conducted using the Wald test and assessment of the F-statistic (significance limit taken as $p=0.05$).

Results

Baseline *Plasmodium* infection (presence/absence), in interaction with an *S. mansoni* co-infection, was associated with a significant reduction in intensity of *Plasmodium* ($p=0.001$; Figure 1A). For children with no prior history of *Plasmodium*, current co-infection with *S. mansoni* was associated

with a lower intensity of *Plasmodium* infection ($p < 0.001$, Table 1). In children that had a baseline *Plasmodium* infection, there was no significant association with a current *S. mansoni* co-infection ($p = 0.126$; Table 1). In *S. mansoni* co-infected children *Plasmodium* infection intensity had a U-shaped relationship with wealth, with significant differences at moderate wealth quintiles (3-4; Table 1).

In the models where infection intensities were considered, the baseline *S. mansoni* intensity, in interaction with age, had a non-linear relationship with *Plasmodium* infection intensity ($p = 0.011$; Figure 1B). *Plasmodium* intensity fell as age increased except for children with a moderate ($n = 34$) *S. mansoni* infection, where *Plasmodium* intensity increased with age. The age-related decline in *Plasmodium* intensity was steeper for those children with prior low ($n = 101$) and high ($n = 13$) level *S. mansoni* infections. In children with moderate baseline infections, although *Plasmodium* intensity increased with age it remained lower than in either infection category for the equivalent age group. Cumulative *Plasmodium* intensity is therefore lowest in children with moderate prior *S. mansoni* infections (supplementary Figure 3).

Co-infection was not associated with a change in infection risk of either parasite, however, each species was associated with an increase in its own subsequent risk of infection (Supplementary Results). Age and sex were also significantly associated with changes in *Plasmodium* and *S. mansoni* risk (Supplementary Results).

Discussion

In our analysis we detected, for the first time, an association between *Plasmodium* and *S. mansoni* in PSAC. In contrast to our hypothesis, we see that co-infection with *S. mansoni* and *Plasmodium* are

associated with a reduced *Plasmodium* infection intensity. This relationship between *Plasmodium* and *S. mansoni* is, however, complex and moderated by host age, family wealth and prior infections. There was no apparent effect of *S. mansoni* on the risk of *Plasmodium* infection nor *Plasmodium* on the risk of *S. mansoni*.

The body's response to *S. mansoni* infection is generally considered to be biased toward a T-helper 2 (Th2) response which is largely directed against schistosome eggs [13]. As the Th2 response is co-downregulatory to the T-helper 1 (Th1) response, the primary response required for *Plasmodium* clearance [13,14], it might be expected that co-infection would lead to increased *Plasmodium* intensity whereas we observed a reduction. Early-stage schistosome infections (i.e. before egg production) however, stimulate a Th1 response [13]. Of the PSAC in our study, only 29% had egg patent infections, suggesting that many *S. mansoni* infections may have been in the Th1 phase of immune stimulation. An early *S. mansoni* induced Th1 response could negatively impact *Plasmodium* and explain the reduced intensity of *Plasmodium* we observe. Nevertheless, there is the possibility that the relationship between these parasite species is driven by resource competition. *Plasmodium* species and hookworms have been observed to compete for the same resource, blood, with hookworm co-infection reducing *Plasmodium* intensity [15]. Blood is also a resource for schistosomes, however, resource competition would not explain the increased *Plasmodium* intensity seen in schistosome co-infections in older children [4].

When we explored the relationship between *Plasmodium* and prior *S. mansoni* infection in respect to different schistosome intensities, a child's age was an important moderator of this relationship. Age was associated with lower *Plasmodium* intensity at low and high prior *S. mansoni* intensities,

however, *Plasmodium* intensity increased with age for moderate infections. Non-linear effects of schistosomes on *Plasmodium* have been observed elsewhere, with moderate schistosome infections reducing the incidence of *Plasmodium* infection while high and low intensity infections are associated with increased incidence [5]. It is feasible that the interactions between these parasites is both immune and resource mediated, and it may be that the interplay between these two mechanisms varies, with the effect of one becoming more dominant than the other at different stages of infection. The low number of high intensity *S. mansoni* infections is a limitation of our data. In older children most studies have observed positive associations between schistosome infection and *Plasmodium* intensity [4,6]. This shift to a positive association could be observed if more children of this age have moderate infection intensities, as the child age-*Plasmodium* intensity trajectory we observe is positive.

The SIMI data does not contain complete information on all *Plasmodium* species present. Three species of *Plasmodium* can be found within the studied communities, but *P. falciparum* dominates (75%) [11]. One study has reported differential outcomes of co-infection between hookworms and different *Plasmodium* species [15] and further work should assess whether our findings are robust to all *Plasmodium* species.

Assessing infections in PSAC with known infection histories, has enabled us to elucidate the complex relationship between *Plasmodium* and *S. mansoni*. We show that prior infections can have long lasting effects and suggest that *Plasmodium*-schistosome interactions are likely to be mediated by the host immune response since any infections have been removed or reduced below detection limits by chemotherapeutic interventions. Determining the order of infection was essential to understanding co-infection outcomes in this system. Our results also suggest that stage of infection

(early or later-stage) may have an important influence on the relationship between *Plasmodium* and schistosomes, again, likely mediated by the host immune response. Our work has focussed on the effect parasites have on one another, but it will now be important to identify whether, and how, these interactions affect host health and the efficacy of the individual parasite control strategies, highlighting the potential need for an integrated *Plasmodium*-schistosome control strategy for this more vulnerable age group.

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Conflict of interest

I confirm that the authors do not have any conflicts of interest

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Footnotes Page

Conflict of interest

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Previous meetings where the work has been presented

This work has been presented at the British Ecological Society (10th-13th December 2019) in Belfast, Northern Ireland, UK and at the British Society for Parasitology (21st-25th June 2021) virtually.

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Table 1: GLMM analyses of the relationship between *Plasmodium* intensity (Ln(X+1)) and a) *S. mansoni*, prior *Plasmodium* infection (presence/absence) and wealth, b) Prior *S. mansoni* EPG and Age of 520 preschool-aged children in Uganda 2009-2011. Significant explanatory variables and groups are denoted by asterisks *. Std. Error = Standard Error and CI₉₅ = 95% Confidence Intervals. Median = median age of PSAC (3).

Variables	Group	Estimate	Std. Error	CI	F-statistic	P value
a)						
<i>S. mansoni</i> : prior <i>Plasmodium</i>	<i>S. mansoni</i> -: <i>Plasmodium</i> -	7.60	0.32	6.97-8.24	$F_{1,506}=9.2$	0.001*
	<i>S. mansoni</i> +: <i>Plasmodium</i> -	-1.77	0.45	5.18-6.49	-	<0.001*
	<i>S. mansoni</i> -: <i>Plasmodium</i> +	-0.41	0.27	6.71-7.68	-	0.128
	<i>S. mansoni</i> +: <i>Plasmodium</i> +	-0.95	0.42	6.10-7.20	-	0.022*
<i>S. mansoni</i> : Wealth Quintile	<i>S. mansoni</i> -:1	7.34	0.31	6.72-7.96	$F_{4,506}=3$	0.018*
	<i>S. mansoni</i> -:2	0.37	0.39	7.00-8.41	-	0.351
	<i>S. mansoni</i> -:3	0.26	0.35	6.97-8.24	-	0.455
	<i>S. mansoni</i> -:4	0.66	0.35	7.37-8.62	-	0.064
	<i>S. mansoni</i> -:5	-0.26	0.36	6.44-7.72	-	0.474
	<i>S. mansoni</i> + :1	-0.36	0.42	6.38-7.58	-	0.398
	<i>S. mansoni</i> + :2	-0.97	0.42	5.79-6.95	-	0.022*
	<i>S. mansoni</i> + :3	-1.41	0.44	5.17-6.49	-	0.001*
	<i>S. mansoni</i> + :4	-1.19	0.45	5.48-6.83	-	0.009*
	<i>S. mansoni</i> + :5	-1.14	0.50	5.40-7.00	-	0.025*
Family ID		0.27	0.29			
Age (Fitted with a spline curve)		0.0003	0.006			
Residual Variation		3.01	0.33			
b)						
Prior <i>S. mansoni</i> EPG:Age	None : 1	7.48	0.15	7.18-7.78	$F_{3,511}=3.8$	0.011*
	None : Median	-0.32	0.10	6.96-7.36	-	0.023*
	None : 5	-0.67	0.17	6.47-7.15	-	0.010*
	Low : 1	0.27	0.41	6.94-8.56	-	0.541
	Low : Median	-0.33	0.21	6.75-7.55	-	0.203
	Low : 5	-0.96	0.26	6.02-7.02	-	0.001*
	Moderate : 1	-2.34	0.92	3.34-6.93	-	0.012*
	Moderate : Median	-1.20	0.43	5.43-7.13	-	0.009*
	Moderate : 5	-0.10	0.40	6.60-8.17	-	0.827
	High : 1	1.86	1.51	6.36-12.32	-	0.223
High : Median		-0.49	0.65	5.72-8.25	-	0.458
	High : 5	-2.88	0.81	3.02-6.18	-	<0.001*
Family ID		0.09	0.29			
Age (Fitted with a spline curve)		<0.0001	0.006			
Residual Variation		3.11	0.35			

Figure 1

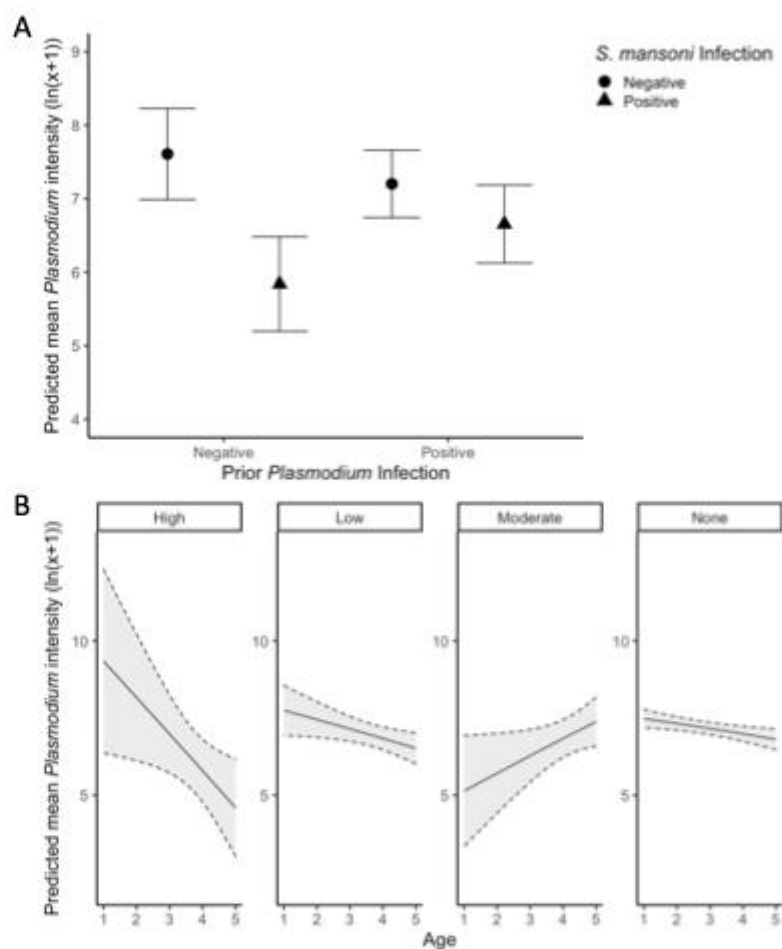


Figure 1: The change in the predicted mean *Plasmodium* intensity $\ln(x+1)$ at the six-month survey in relation to A) *Schistosoma mansoni* coinfection, with and without prior *Plasmodium* infection, and B) prior *S. mansoni* infection intensity with child age. For A), children were either uninfected with *S. mansoni* at the six-month survey (circles) or infected (triangles). For B), none, low, moderate and high represent the intensity of the prior *S. mansoni* infection. For A) predictions were made with age at the median (3 years), sex set to male and the wealth quintile set to 3. For B) Age and a prior *S. mansoni* infection were the only significant terms. Error bars or dotted lines represent 95% confidence intervals for the predictions.