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Data cannot be shared publicly because the study remains blinded. to outcome data. Data are available from the DeWorm3 Institutional Data Access Committee (contact via Barbra Richardson,

barbrar@uw.edu) for researchers who meet the criteria for access to these data.

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Introduction

 Soil-transmitted helminth (STH) infections are among the most common infections worldwide, affecting more than 1.5 billion of the poorest and most marginalized communities globally. (1) The most common STH species of humans include *Ascaris lumbricoides*, *Trichuris trichiura* and the hookworm species, *Necator americanus* and *Ancylostoma duodenale*. STH are transmitted by eggs present in human feces which in turn contaminate soil and water in areas with poor sanitation, conditions often found

 in low-resource countries. (2) STH are widely distributed in tropical and subtropical 38 areas, with the greatest numbers occurring in sub-Saharan Africa, the Americas, China and East Asia. (3)

 The World Health Organization (WHO) considers STH a public health problem in areas 42 where >1% of the at-risk population has moderate-to-heavy intensity infection – as 43 measured by number of eggs per gram of stool diagnosed by stool examination (4). These moderate to high intensity helminth infections are associated with poor cognitive and motor outcomes in infants, as well as with anemia. (5–9) Pre-school children (PSAC), school age children (SAC) and women of reproductive age (WRA), including adolescent girls, pregnant women, lactating women, and non-pregnant and non- lactating women living in endemic areas, are at highest risk of morbidity due to STH. Clear policy and guidance are essential to support country-level efforts to expand routine deworming of WRA, and recent WHO publications have provided the necessary policy framework. (2,3)

 The WHO Neglected Tropical Disease (NTD) Roadmap and London Declaration have 54 accelerated progress toward eliminating selected NTDs, including lymphatic filariasis and onchocerciasis, and formalized long-term disease-specific goals for other NTDs. $5 = (10)$ Global interest is shifting towards an elimination strategy for other NTDs, including the possibility of breaking the transmission of STH through community-wide mass drug administration (MDA). (11)

 In Benin, all major STH (hookworm sp., *A. lumbricoides* and *T. trichiura*) are a recognized public health problem, with more than 50% of districts requiring MDA based on the results of a recent national mapping exercise that sampled stool from SAC. (12,13) In Comé District, this recent national mapping showed a prevalence of STH in school-aged children of 20%, despite multiple rounds of school-based MDA with albendazole in 2015 (coverage 59%), 2016 (coverage 78%) and 2017 (coverage 83%). 66 (13) =

 In 2017, the DeWorm3 project (ClinicalTrials.gov Identifier NCT03014167) was initiated in Benin, and, in parallel, in India and Malawi. Using a cluster randomized controlled study design, the primary objective of the project is to determine whether

 the provision of an enhanced (twice yearly) level of high-coverage MDA, targeting all age groups in a whole community over a 3-year period, can interrupt transmission of STH (11). Here we report analyses of baseline data from a longitudinal monitoring cohort randomly selected from the whole population involved in the trial in order to determine the demographic and other parameters potentially associated with the STH infections detected by microscopy using a standard Kato-Katz procedure.

Materials and Methods

Study area and population

 The DeWorm3 trial in Benin is being conducted in the district of Comé. The study site 81 selection was based on criteria reported-previously. (14) Comé is located 70 km west of Cotonou in the Mono department, at latitude 6°24′N and longitude 1°53′E. The 83 district covers an area of 153 km² with a population estimated at 79,989 inhabitants in 84 the census of 2012, with an estimated yearly growth rate of 2.07% . (15) The district has five sub-districts (Central Comé, Akodéha, Oumako, Agatogbo and Ouèdèmè- Pedah) subdivided into 52 villages/areas or neighborhoods. The climate is sub- equatorial, tropical, alternating between two rainy seasons (April to July and September to November) and two dry seasons (December to March and August). Rainfall varies between 900 and 1,200 mm per year.

Study design

 The protocol and aims of the DeWorm3 study have been published elsewhere. (11) A 93 baseline census was conducted from January 8th to February 9^{th,} 2018 followed by 94 Cluster demarcation. The geospatial locations of all households were mapped using ArcGIS (Redlands, CA), and the study area was divided into 40 clusters with between 96 1,650 and 4,000 residents per cluster. From March 6th to April 5th, 2018, 6000 individuals (150 individuals by cluster) were randomly selected to constitute a longitudinal monitoring cohort (LMC) participating in annual follow-up STH infection surveys over 5 years. The LMC was selected from the censused population using stratified random sampling of PSAC aged 1-4 years old, SAC aged 5-14 years old and 101 adults aged 15 years old and above, at a ratio of 1:1:3. A sampling list of 150 individuals 102 (i.e. 30 PSAC, 30 SAC and 90 adults) was initially generated and backup lists of 75 individuals were issued to replace participants who could not be located or refused to 104 participate. LMC participants were *interviewed* and completed a more in-depth assessment of individual-level STH risk factors, including a survey of self-reported WASH access and use, history of deworming, and direct observation of WASH facilities and participants' use of footwear. Individuals participating in the LMC agreed to provide 108 Stool samples for immediate analysis using the Kato-Katz method (16) annually for the 109 duration of the study.

Data collection

Kato-Katz data

 Stool samples from LMC participants were collected by study staff and delivered to the laboratory within one hour. Samples were screened using the Kato-Katz technique, with results for each type of helminth (*A. lumbricoides*, *T. trichiura* and hookworm: *A. duodenale*/*N. americanus*) reported in eggs per gram (epg). Two slides were prepared from each sample, and each slide was examined by two experienced lab technicians. 118 A subset of 10% of slides was randomly selected for quality assurance by a laboratory **manager.** Prevalence was calculated both for individual STH types and cumulatively according to the following formulas:

- The prevalence per STH type:

 $p =$ Number of samples where at least one egg of STH species is found $t = \frac{t}{\text{total number of samples}}$ $\frac{124}{\text{total number of samples examined}}$ x 100

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- The cumulative STH prevalence:

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p = \frac{\text{Number of positive samples for one, two or three STH species}}{\text{total number of samples examined}} \times 100
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129 In cases of co-infection, the sample was counted to calculate the cumulative prevalence, and prevalence and intensity assessed separately for each species. The parasite intensity was calculated from a Kato-Katz smear made with 41.7 mg of stool, 132 by multiplying the egg count from the slide by a factor of 24 (24 x 41.7 mg \approx 1 g)-to get 133 the number of eggs per gram of stool $(EPG)_{\overline{n}}$

Outcomes

The primary outcomes were individual-level infection status for each STH type (positive

137 / negative) and intensity of infection in eggs per gram (epg).

Variables

 Individual factors (including age, gender, history of deworming during the past year and shoe wearing behavior), household factors (including highest educational level achieved, head of household occupation, household asset index, urbanization), water sanitation and hygiene (WASH) factors (household water service, household sanitation, household hand washing facility) and environmental factors (elevation, soil 144 sand fraction, soil acidity at average depth (0-5-15 cm), MODIS daytime land surface 145 temperature mean for 2018 (°Celsius), MODIS Enhanced Vegetation Index (EVI) mean for 2018, MODIS normalized difference vegetation index (NDVI) mean for 2018, aridity index) were collected or constructed based on existing data.

 Water, Sanitation and Hygiene variables: Water sources and sanitation facilities reported were grouped and categorized according to the 2017 WHO/UNICEF Joint Monitoring Program (JMP) methodology (none, improved, unimproved, limited or basic). (17) Improved drinking water sources are those that have the potential to deliver safe water by nature of their design and construction, while improved sanitation facilities are those designed to hygienically separate excreta from human contact. (18) **Distance** to the closest water source and sharing status for sanitation were also 156 collected. $\boxed{=}$

 Asset index: An asset index was compiled using principal components analysis. The procedure described by the Demographics and Health Survey *(Steps to constructing the new DHS Wealth Index)*(19) was followed, but factors associated with STH transmission (crowding [residents/room], WASH variables included in the risk factors analysis, and flooring materials) were excluded as they were evaluated separately in the model.

 Environmental variables: We examined the association of the following environmental and sociodemographic factors with STH infection: mean enhanced vegetation index and land surface temperature during the study period; elevation; aridity; soil acidity and sand content; and population density. These environmental, topographical, and sociodemographic measures were extracted for each household using point-based extraction using ArcGIS 10.3 (Environmental Systems Research Institute Inc., Redlands, CA, USA). Data sources and methods have been described previously (20). Estimates of population density were obtained by calculating the number of individuals 173 living within 1km² buffer around each household, which was used to classify areas as high, medium or low population density. Continuous variables were categorized by tertiles for analysis.

Descriptive statistics

 Categorical variables were described using proportions and 95% confidence intervals 179 and the continuous variables were described by the median and interquartile ranges. To compare proportions, we used the Chi-square, and Cuzick trend tests. Continuous 181 variables were compared using the Student _T-test and analysis of variance (ANOVA). For each STH species we determined the cluster level prevalence (proportion of individuals infected in the cluster) and cluster level arithmetic mean of individual's egg density per gram of feces. We plotted the cluster level mean egg density against the cluster level prevalence and assessed the strength of the linear relationship using Pearson's correlation coefficient test. Descriptive statistics were generated using Stata® 14.0 (Stata Corp, College Station, Texas).

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Factors associated with STH infection

 Factors associated with presence and intensity of baseline infection with each STH species were identified using mixed effects models with random effects at the household and cluster levels and exchangeable correlation matrix. For binary infection status, mixed effects logistic regression was used, while for intensity of infection negative binomial mixed effects regression was used.

 For the negative binomial regression, the output was the infection intensity ratio (IIR): $\mathsf{IIR} = e^{\beta} = e^{\left[\log(\mu_{X0+1}) - \log(\mu_{X0})\right]} = e^{\left[\log(\mu_{X0+1}/\mu_{X0})\right]}$

 where β is the regression coefficient, μ is the expected intensity of infection (epg) and 198 the subscripts represent where the predictor variable, say x, is evaluated at x_0 and x_{0+1}

199 (implying a one unit change in the predictor variable x). The IIR are interpreted as the ratio of expected intensity of infection for a one unit increase in the predictor variable given the other variables are held constant in the model.

 All models were adjusted for age and sex. Groups of socio-economic status indicators, environmental factors and WASH factors hypothesized to be associated with infection were proposed *a priori* in the multivariable analysis. For groups of indicators with similar variables, the factor from each group with the lowest Akaike Information Criterion (AIC) in univariate analyses was selected for inclusion in the multivariable model. Models were further simplified by backward stepwise elimination until AIC was no longer further reduced in the adjusted model.

 Random effects predicted by the fully adjusted model were compared to those predicted by a model containing only age and sex and the proportion of clustering explained by the explanatory variables was quantified.

Ethics statement

 Ethical approval of the DeWorm3 trial protocol was obtained both from the Human Subjects Division at the University of Washington and the National Ethics Committee for Health Research of Benin. (CNERS ethical clearance reference No: 002- 2017/MS/DC/SGM/DFR/CNERS-Ministry of Health, Benin). The trial was registered at Clinical Trials.gov NCT03014167. Written consent was obtained from each participant $21 =$ (or participants' parents, when participants were under 18 years of age). For children aged 1-6 years old, verbal assent was obtained and for adolescents aged 7-17 years written assent was obtained. Data were collected electronically using password protected smartphones and was stored in datasets. Although WHO guidelines do not 223 recommend MDA for adults, following the stool analysis any **adults (** \geq **15 years of age)** 224 in control clusters presenting **moderate to heavy** intensity STH infection according to WHO definitions (21) or requiring treatment according to local guidelines, were treated 226 with albendazole by study staff. $\boxed{=}$

Results

Descriptive

 Based on the census data, 11,979 individuals were selected for participation in three consecutive stages (Stage 1: n=5,979; Stage 2: n=3,000; Stage 3: n=3,000), with the goal to reach 150 individuals in each cluster: 30 PSAC, 30 SAC and 90 adults. Characteristics of the longitudinal monitoring cohort (LMC) population in comparison 234 to censused population of the DeWorm3 site are presented in Table 1, and Fig 1 presents the study flow chart. Individuals selected were listed as living in 9,265 households from which 8,741 were located and visited. In those households 7,045 individuals were present, among whom 6,814 consented to participate in the LMC cohort. Stool samples were collected from 6,153 individuals. The most common reasons for stool samples not being collected were (i) no sample visit documented (319), (ii) sample could not be collected after 3 visits (n=111), (ii) refusal to provide sample (n=231). As no documented survey could be verified for 14 individuals, Kato- Katz tests performed were confirmed for 6,139 samples comprising 1,184 PSAC (98.7% of 1,200 expected), 1,335 SAC (>100% of 1,200 expected), and 3,620 adults 244 (>100% of 3,600 expected). In total 6,139 tests $\overline{5}$ d two slides read by laboratory $_{\text{max}}$ 245 technicians. A random subset of Kato-Katz tests was selected for reading by the 246 supervisor and compared against the original readings for quality assurance. $\frac{1}{1}$

Prevalence of STH

 Among the 6139 individuals tested by Kato-Katz, STH infections of any type were found in 324 (5.3%), 199 (3.2%) due to hookworm spp. and 126 (2.0%) due to *Ascaris. Trichuris* was found in just 5 (0.1%) individuals (Table 2). Six individuals were co- infected with hookworm and *Ascaris*. Due to the small number of *Trichuris* infections, only analyses focused on hookworm spp. and *Ascaris* are presented. Among all infections, 258 (79.7%) were light-intensity, 54 (16.7%) moderate-intensity and 12 255 (3.7%) heavy intensity infections. Hookworm spp. were more prevalent in adults than 256 in SAC or PSAC (4.4% *versus* 2% *versus* 1% *respectively*, Chi², p<0.001). SAC were more frequently infected with *Ascaris* compared to PSAC or adults (3% *versus* 2% *versus* 1.7% respectively, Chi², p=0.02) (Fig 2). A higher proportion of males than females was infected with hookworm spp. (4% *versus* 2.6%; p=0.002) and *Ascaris* (2.6% *versus* 1.6%; p=0.004).

Intensity of STH infection

262 The median egg density for hookworm spp. was 108 eggs per gram (epg) (IQR: 48- 312; range: 12-12,960), 3,840 epg for *Ascaris* (IQR: 312-15,180; range: 12-135,084) and 120 epg for *Trichuris* (IQR: 60-468; range: 36-20,124). The intensity of infection was similar in all age groups for hookworm spp. (ANOVA, p=0.22), with a median egg density of 264 epg (IQR: 36-384; range: 12-3,048) in PSAC, 96 epg (IQR: 24-312; range: 12-11,100) in SAC and 108 epg (IQR: 48-288; range: 12-12,960) in adults. We found a difference in intensity of infection with *Ascaris* between age-groups (ANOVA, p=0.005), this difference was between SAC and adults (Bonferroni, p=0.004). Median egg densities were 6,972 epg for PSAC (IQR: 264-26292; range: 12-60000), 7,848 epg for SAC (IQR: 3,714-25,314; range: 84-56,412) and 780 epg for adults (IQR: 36-8,772; range: 12-135084).

 λ ₂ Moderate to heavy intensity (MHI) infections were found in $66/6,139$ individuals overall (1.1%) amongst whom 10 (0.2%) MHI with hookworm spp., 55 (0.9%) MHI with *Ascaris* and 1 (<0.1%) MHI with *Trichuris* (Table 2). The burden of MHI was greatest in SAC with 2.1% (25/1,184) prevalence of MHI of *Ascaris* (Table S1). 68.2% (45/66) of MHI were found in males (Table S2). MHI were distributed in 15/40 clusters. MHI with hookworm spp. were present in 7/40 clusters, MHI with *Ascaris* in 7/40 clusters and MHI with *Trichuris* in 1 cluster. There were two clusters showing a particularly high burden of *Ascaris*, with respectively 19 (12.7%) and 30 (20%) individuals with MHI with *Ascaris*.

Age- and sex-related prevalence and intensity of STH infection (hookworm and *Ascaris***)**

 Figs 3 and 4 show the age-infection profile for hookworm spp. and *Ascaris,* respectively. The prevalence of hookworm spp. increased with age in both sexes. The 287 prevalence was similar in males and females among PSAC and SAC_n but in adults, the prevalence in males was higher than in females except for 50-60 year olds, in whom females were more frequently infected. The intensity of hookworm infection was similar in males and females regardless of age, and was higher in adults than in children. The prevalence of *Ascaris* infection was similar in males and females across all ages, with the period of adolescence and early adulthood (between 10 and 18 years old) corresponding to the period with highest prevalence of *Ascaris* infection in males and the lowest in females (6% for males *versus* 1% for females). Intensity of *Ascaris* infection followed the same profile as prevalence in both sexes.

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Community-level correlation between intensity and prevalence of

STH infection

 We found a positive linear relationship between STH infection prevalence and the intensity of infection at cluster level in our study population (Fig 5). This correlation was strong for both hookworm spp. (ρ=0.73, p<0.0001) and *Ascaris* (ρ=0.98, p<0.0001).

Factors associated with hookworm infection

 The results of univariate analyses of factors associated with hookworm infection 306 prevalence are presented in Table S2. Here, the results of multivariable analyses-are presented (Table 3).

- 308 At the individual level, PSAC and SAC were significantly less likely to be infected with hookworm spp. than adults (aOR=0.21, 95%CI 0.10-0.44, p< 0.001 and aOR=0.49, 95%CI 0.29-0.83, p=0.008, respectively). Females were also significantly less likely to be infected than males (aOR=0.56, 95%CI 0.38-0.83, p=0.004). Individuals who reported a history of deworming during the past year were significantly less likely to be infected (aOR= 0.45, 95%CI 0.27-0.75, p< 0.002).
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 Among household factors, the household asset index, a proxy measure for family wealth, showed that individuals in the poorest households had a significantly higher 317 odds of infection than the richest $(5th$ quintile) with a significant dose-response effect (Cuzick test of trend, p<0.001), (First quintile: aOR= 5.03, 95%CI 2.10-12.01, p<0.001, $2nd$ quintile aOR= 3.62, 95%CI 1.51-8.66, p=0.001 and 3rd quintile aOR= 2.51, 95%CI 1.05-6.00, p=0.02). With respect to occupational exposure, farmers were more likely to be infected with hookworm spp. than others (aOR= 1.79, 95%CI 1.11-2.90, p=0.02). Individuals living in medium population density settings were more likely to be infected than those living in high density settings, (aOR= 2.59, 95%CI 1.25-5.40, p=0.01). 324 Among WASH factors, household sanitation, and especially open defecation, was

found to be strongly associated with hookworm infection. Individuals using improved

 unshared sanitation facilities had half the odds of hookworm infection compared to those defecating outdoors (aOR=0.48, 0.24-0.98, p=0.04).

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Factors associated with hookworm infection intensity

 Children had a significantly lower intensity of hookworm infection as compared to adults (PSAC: adjusted IIR=0.10, 95%CI 0.03-0.31, p<0.001; SAC: adjusted IIR=0.29, 95%CI 0.12 – 0.72, p=0.01. Females had significantly lower intensity infections than males (adjusted IIR=0.32, 95%CI 0.16–0.64, p=0.001), as did individuals dewormed the year before (adjusted IIR=0.20, 95%CI 0.08–0.48, p<0.001). (Table 3).

 \blacksquare

 At the household level, less educated people (those with no education, primary school or secondary school) had higher intensity infections with hookworm spp. compared to those with university level education (adjusted IIR=40.13, 95%CI 2.47–652.77, p=0.01; adjusted IIR=30.92, 95%CI 1.86–513.88, p=0.02; adjusted IIR=19.34, 95%CI 1.21– 308.81, p=0.04, respectively). Being a farmer, living in a house with natural floor material versus man-made floor material, and living in a peri-urban setting were also all factors found to be associated with a significantly increased intensity of hookworm infections (farmer: adjusted IIR=3.94, 95%CI 1.67–9.27, p=0.002; natural floor material: adjusted IIR=0.23, 95%CI 0.05-1.03, p=0.06; peri-urban settings: adjusted IIR=6.18, 95%CI 1.82 - 20.90, p=0.003).

 Access to unimproved water available more than 30 minutes away from the house was associated with significantly higher intensity hookworm infection (adjusted IIR= 13.47, 95%CI 1.62–111.55; p=0.02) compared to improved water available less than 30 minutes from the house. Compared to open defecation behavior, using an improved and unshared toilet was associated with significantly lower intensity hookworm infections (adjusted IIR=0.23, 95%CI 0.07 – 0.70, p=0.01). No environmental factor was found to be associated with intensity of hookworm infections in multivariable analyses.

Factors associated with *Ascaris* **infection prevalence.**

 Among the individual factors assessed, SAC (5-14 years) were significantly more likely to be infected with *Ascaris* than adults (aOR= 2.0, 95%CI 1.1-3.6, p=0.01). However, no difference in odds of infection was found between PSAC and adults. Female individuals were less likely to be infected with *Ascaris* than males (aOR= 0.5, 95%CI 0.3-0.9, p= 0.02).

 Amongst environmental factors, low soil acidity was significantly associated with increased odds of *Ascaris* infection compared to the highest soil acidity (aOR=4.8, 95%CI 1.8-13.1, p=0.002). Moderate [29.6-31.9°C] and high [31.9; 32.8°C] daytime land surface temperatures were associated with lower odds of infection with *Ascaris* compared to lower temperatures [26.2-29.6°C[(aOR=0.12, 95%CI 0.03-0.44, p=0.001; and aOR=0.17, 95%CI 0.03-0.91, p= 0.04 respectively). The summary of the multivariable analysis with *Ascaris* is presented in Table 4.

Intra-Class Correlation statistics for hookworm and *Ascaris*

infection prevalence.

 Comparison of the Intra-Class Correlation values between models containing only age and sex (model 1) and the fully adjusted multivariable final model with all the fixed effect covariables (model 2) showed decreased ICC values in the fully adjusted model, considering either level-3 ICC at the cluster level or level-2 ICC at the household-within-cluster level (Table S4).

 When only adjusting for age and sex, the prevalence of hookworm infection was correlated between individuals within the same cluster (ICC=0.16, 95%CI 0.10-0.26), and this correlation increased significantly between individuals within the same household and cluster level (ICC=0.58, 95%CI 0.40-0.74). Prevalence of *Ascaris* infection was moderately correlated within the same cluster (ICC=0.54, 95%CI 0.34- 0.73), with a small increase within the same household and cluster level (ICC=0.60, 95%CI 0.26-0.76). In this model, household and cluster random effects compose approximately 58% and 60% of the total residual variance for hookworm spp. and *Ascaris* infection prevalence respectively.

388 In the fully adjusted multivariable final model, \bar{w} , found a correlation of hookworm infection prevalence within the same cluster (ICC=0.03, 95%CI 0.01-0.10), although this correlation increased within the same household and cluster level (ICC=0.39, 95%CI 0.17-0.65). *Ascaris* infection prevalence was moderately correlated between individuals within the same cluster (ICC=0.42, 95%CI 0.23-0.64), and this correlation increased slightly within the same household and cluster level (ICC=0.51, 95%CI 0.26- 0.76). We estimated that household and cluster random effects compose approximately 39% and 51% of the total residual variance of hookworm and *Ascaris* infection prevalence, respectively, in the fully adjusted model.

Discussion

 We observed a relatively low prevalence of STH in this region of Benin. Hookworm spp. were the most prevalent infections, and were more prevalent in adults, while *Ascaris* was more prevalent in children. Females were generally less infected than males across all ages. Females, children, those dewormed during the previous year and those using improved unshared sanitation facilities had lower odds of hookworm infections, while being a farmer, living in peri-urban settings versus urban and being poor was associated with a higher odds of hookworm infection. In addition to those factors, the intensity of hookworm infection was also decreased if an improved water source was available at less than 30 minutes distance.

 Since 2013, the Ministry of Health in Benin has focused its efforts on developing and implementing strategies for the control of five NTDs considered to be of highest priority, 410 namely trachoma, onchocerciasis, lymphatic filariasis, schistosomiasis and soil-411 transmitted helminths. Those efforts were bolstered markedly through the **ENVISION** program (22), a USAID-funded initiative that ran from 2013 through 2019 in Benin. A nationwide STH prevalence survey was completed in 2015, that reported 20% prevalence (13) in school-aged children in Comé district. Following that national 415 mapping effort, 3-rounds of school-based MDA with albendazole were undertaken according to the recommendations of WHO, i.e. primarily targeting school-age (SAC) 417 and pre-school age children (PSAC) for either once or twice yearly treatment as a function of the estimated prevalence of infection in any given district. (23) Coverage of

 SAC with school MDA between 2015 and 2017 was estimated between 59% and 83%. Albendazole and/or Mebendazole are also distributed in health facilities and to 421 pregnant women during routine antenatal care starting from the $2nd$ -trimester of 422 pregnancy. (24) In the context-described, we sought to better understand patterns of STH infection in order to move towards the elimination STH as a public health problem, 424 by reaching a prevalence of STH less than 1%, as prescribed by the WHO Neglected Tropical Disease (NTD) Roadmap and London Declaration on NTD. (10,23)

 When focusing on the at-risk population of SAC, the prevalence of STH infection found 427 in the current study is lower than that reported in the same district using the same diagnostic technique in 2015 during the national mapping exercise (5.2% versus 20.0% 429 respectively, p<0.001). \Box at survey was conducted with a total of 250 stool samples from SAC collected from schools located in 5 rural villages. (13) The prevalence of infections with *Ascaris* (3.0% *versus* 15.6% respectively, p<0.001) or *Trichuris* (0.15% in 2018 *versus* 4.8% in 2015, p<0.001) decreased while the decrease in prevalence of hookworm spp. in SAC was less marked (2.0% in 2018 versus 4.0% in 2015, p=0.054) compared to the findings of the national STH mapping 3 years earlier. (13) The decline in STH prevalence in the study area might be related to differences in sampling, as the 436 current study was conducted in the community instead of in-schools, with more than 6,000 stools randomly selected from three age groups (PSAC, SAC and adults). (11) STH prevalence estimates can vary depending on the sampling strategies used. (25) The reasons for the observed variations of prevalence between hookworm spp. and other STH species in SAC could also be that STH rate of reinfection post-treatment 441 varies across species, with a faster reinfection with *Ascaris* than hookworm spp. systematic review of helminth reinfection at 3, 6, and 12 months (95% CI), after drug 443 treatment shows that *Ascaris* prevalence reached 26% (16-43%), 68% (60-76%) and $=$ 94% (88-100%) of pretreatment levels, respectively and for hookworm spp., 30% (26- 34%), 55% (34-87%), and 57% (49-67%). (26) These results may also be partly explained by the fact that the current STH program does not include adults. The suggestion that hookworm spp. prevalence only decreased slightly between 2015 and 2018, may be due to the persistent untreated adult reservoir in which hookworm spp. are most common. Data from several worm expulsion studies show that the proportion of hookworms harbored by adults ranged from 70 to 85%, (27–32) and a reinfection– infection study in Indonesia show that adults have higher reinfection rates with

 hookworm spp. than children. (33) Children cleared of hookworms through annual school de-worming could easily be re-infected at home through contact with adult members of their households.

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 Hookworm prevalence was higher in adults while *Ascaris* prevalence was higher in 456 children. One explanation of these findings might be helminth species transmission modes. (34) The three species of STH (*A. lumbricoides*, *T. trichiura*, hookworm spp.) 458 have relatively similar cycles involving the presence of adult worms in the intestine $_s$ </sub> however the main mode of transmission of *Ascaris* and *Trichuris* is through contaminated food and water (parasite egg ingestion) whereas hookworm spp. are mainly transmitted by skin penetration, although they can be transmitted by ingestion. (35,36) The eggs of *Ascaris* and *Trichuris* are found in soil contaminated by human feces or in uncooked food contaminated by soil containing eggs of the worm. A person becomes infected after accidentally swallowing the fertile eggs. Children may be more likely to be infected with *Ascaris* because they are more likely to put their contaminated fingers in their mouths after playing in contaminated soil. (37) Unlike *Ascaris* infection, which declines in prevalence with age, hookworm infects all ages throughout life with prevalence increasing in adults. (27)

 Community-level prevalence and the arithmetic mean of infection intensity were significantly correlated for all STH infections in our study, with a strong prevalence- intensity correlation for infection with hookworm and *Ascaris*. Similar trends were recently found in Kenya for hookworm spp. and *Trichuris.* (20) At the individual level, prevalence and intensity of hookworm infection followed the same trend. Markers of poverty and exposure to environmental sources of STH infection, including being a farmer, lack of improved or private sanitation facilities, low income, poor access to water, no or limited education, or living in a house with natural floor material were all associated with a higher prevalence or intensity of hookworm infection. These findings are linked with the mode of hookworm spp. transmission, which is direct either by ingestion (for *A. duodenale*) or by skin penetration (both *N. americanus* and *A. duodenale*) of infective larval stages living in the soil. (38,39) These findings are consistent with the results of a recent study in Kenya where there was a strong association between hookworm infection prevalence and intensity and socio-economic status, with those in the poorest households having the heaviest infections and highest prevalence, and wealthier individuals having the lightest intensity and reduced odds of infection. (20) Globally, a negative correlation between hookworm infections and income level is demonstrated in cross-country comparisons. (40–42) Moderate population density, corresponding to a peri-urban environment, was also associated with both high prevalence and heavy intensity of hookworm infection when compared to the higher population density observed in urban environments. (43)

 Although we found no association between hookworm prevalence and water source, quality of water seems to affect intensity of hookworm infection. Heavier intensity infections were found in participants with access only to unimproved water, such as unprotected wells, unprotected springs and surface water available at more than 30 minutes from the house. In a school survey in Togo, unimproved drinking water was associated with higher odds and intensity of hookworm. (44) Malaysian children with access to piped water were less infected with hookworm. (45) However, other researchers have found no statistically significant associations between piped water access and hookworm infection (46,47). We did not find any association between WASH variables and either prevalence or intensity of *Ascaris* infection. However, there is evidence that integrated water, sanitation and hand hygiene intervention, treatment of water with chlorine (48), drinking piped water, as well as hand washing before eating and after defecating reduce the odds of *Ascaris* infection. (49)

 This study had a number of strengths, including the large population size, completeness and quality of data and the high level of quality control for Kato-Katz diagnosis, with double reading by the lab technicians of the whole sample with an additional control of a subset of samples by a senior skilled parasitologist. However, this study does has some limitations. First, the STH prevalence in Come hides inter- and intra-specific variations between clusters that will be developed in further analyses once the parent study is unblinded and we have access to those results. It was also necessary to use a staged approach to sampling in order achieve the required number of participants who consented to participate in the longitudinal monitoring cohort, which may have limited its representativeness. Another possible limitation is the reliance on Kato-Katz to detect STH. Kato-Katz is poorly sensitive, particularly for low intensity infections and can be affected by storage and processing time and methods.(50,51) Future analyses using qPCR-based methods will allow for more sensitive detection of of STH in stool.

Conclusion

521 This analysis of the DeWorm3 baseline study data shows that hookworm spp. are the predominant STH in Comé, with a persistent reservoir in adults. This infection reservoir is not addressed by current school-based MDA control measures. These data suggest that community-based MDA may help eliminate STH as a public health problem. Improved unshared sanitation and access to improved water sources are associated with lower prevalence and/or intensity of hookworm infection. Programmatic efforts should pay particular attention to farmers and populations living in poverty in urban, rural and peri-urban environments. The DeWorm3 trial (2017-2022) will determine the feasibility of STH transmission interruption through community-wide MDA given twice-530 a-year for three years in this setting and combined with these results will inform programmatic and policy decisions to improve efforts to eliminate morbidity and infection due to these pervasive infections.

Declaration of competing interest:

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Authors' contribution:

- EFGAA, PH, MA, EA, GC, MI, AJFL, EY contributed to data collection,
- EFGAA performed the statistical analysis and wrote the first draft of the manuscript,
- KHA designed the article statistical methodology and reviewed the analysis and entire
- article draft,
- PH add inputs to the statistical analysis,
- KHA, ARM, JLW, TJL, SG, designed the Deworm3 clinical trial
- EFGAA, MA, KHA, AJFL, MI, AG, JLW reviewed the article draft and the final version.
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Fig 1: Flow diagram of stool sample collection for Benin site DeWorm3 baseline prevalence survey in Comé

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Fig 2: STH unweighted prevalence across age-groups

Fig 3. Age-infection profiles for hookworm. (A) Prevalence (black lines) and intensity (grey lines) of hookworm infection by age for males (solid line and circles) and females (dashed lines and empty circles). Vertical bars represent confidence intervals.

Fig 4. Age-infection profiles for Ascaris. (A) Prevalence (black lines) and intensity (grey lines) of Ascaris infection by age for males (solid line and circles) and females (dashed lines and empty circles). Vertical bars represent confidence intervals.

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Figure 5: Cluster level correlation between prevalence and intensity of hookworm and Ascaris infection in the study population.

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Table 1: Comparison of censused population of the DeWorm3 site and longitudinal monitoring cohort (LMC). \mathbf{r} ÷

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Table 2: Unweighted STH prevalence and intensity of infection by Kato-Katz testing. N=6139

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Table 3 :Factors associated with the the prevalence and the intensity of hookworm infection in Comé, Bénin : findings from a cross-sectional baseline prevalence survey in the DeWorm3 STH–elimination trial.

† Adjusted Generalized logistic mixed model estimating equations with exchangeable correlation structure.

§ 5,366 observations included in fully adjusted model.

‡ Adjusted zero-inflated negative binomial regression model, inflating for sex and age (1–4 years, 5–14 years, 15 years), with an exchangeable correlation matrix.

♣ 5,364 observations included in fully adjusted model.

* Variable dropped from fully adjusted model during model adjustment process using lowest AIC criteria.

** Variable in the final adjusted model but with no significant category

Abbreviation: School Aged Children (SAC), Pre School Aged Children (PSAC), confidence interval (CI), interquartile range (IQR), Moderate Resolution Imaging Spectroradiometer (MODIS)

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Table 4 : Factors associated with *Ascaris* infection prevalence in Comé, Bénin: findings from a baseline prevalence survey using Kato-Katz technique

* Variable dropped from fully adjusted model during model adjustment process using lowest AIC criteria.

** Variable in the final adjusted model but with no significant category

Abbreviation: School Aged Children (SAC), Pre School Aged Children (PSAC), confidence interval (CI), interquartile range (IQR), Moderate Resolution Imaging Spectroradiometer (MODIS)

SUPPLEMENTARY DATA

Table S1: Burden of moderate to high intensity (MHI) STH infection in the study population by age group, during DeWorm3 baseline analysis in Comé, Bénin

[Click here to access/download;Table;Table S2_AJFL](https://www.editorialmanager.com/pntd/download.aspx?id=961593&guid=b408ee81-2527-45d7-941d-1bd875db566e&scheme=1) kha_EA_20201005_Final AJFL.docx

SUPPLEMENTARY DATA

Table S2: Burden of moderate to high STH infection in infected individuals by age group and \overline{z} gender, during DeWorm3 baseline analysis in Comé, Bénin

Table S3: Factors univariately associated with hookworm infection in Comé, Bénin: findings from DeWorm3 cluster randomized trial baseline pre-treatment survey using generalized

SUPPLEMENTARY DATA

Table S3: Factors univariately associated with hookworm infection in Comé, Bénin: findings from DeWorm3 cluster randomized trial baseline pretreatment survey using generalized logistic mixed model.

†Generalized estimating equations with exchangeable correlation structure and logit link applied

All data available displayed for "Infected with Hookworm/ total N (%)"

Acronyms: School Aged Children (SAC), Pre School Aged Children (PSAC) , confidence interval (CI), odds ratio (OR), Moderate Resolution Imaging Spectroradiometer (MODIS)

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SUPPLEMENTARY DATA

Table S4 : Intra-Class Correlation values

*The Intraclass correlation statistics reports two intraclass correlations for this three-level nested model. The first is the level-3 intraclass correlation at the cluster level, the correlation between Hookworm or *Ascaris lumbricoides* infection prevalence in the same cluster. The second is the level-2 intraclass correlation at the household-within-cluster level, the correlation between prevalence of infection with *Ascaris lumbricoides* in the same household and cluster.

** Generalized logistic mixed model with exchangeable correlation matrix