Review Article

Prevalence of *Schistosoma mansoni* and *S. haematobium* in Snail Intermediate Hosts in Africa: A Systematic Review and Meta-analysis

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Background. Schistosomiasis is caused by Schistosoma mansoni and S. haematobium in Africa. These schistosome parasites use freshwater snail intermediate hosts to complete their lifecycle. Varied prevalence rates of these parasites in the snail intermediate hosts were reported from several African countries, but there were no summarized data for policymakers. Therefore, this study was aimed to systematically summarize the prevalence and geographical distribution of S. mansoni and S. haematobium among freshwater snails in Africa. Methods. Literature search was carried out from PubMed, Science Direct, and Scopus which reported the prevalence of S. mansoni and S. haematobium among freshwater snails in Africa. The pooled prevalence was determined using a random-effect model, while heterogeneities between studies were evaluated by I^2 test. The meta-analyses were conducted using Stata software, metan command. Results. A total of 273,643 snails were examined for the presence of S. mansoni and S. haematobium cercaria in the eligible studies. The pooled prevalence of schistosome cercaria among freshwater snails was 5.5% (95% CI: 4.9-6.1%). The pooled prevalence of S. mansoni and S. haematobium cercaria was 5.6% (95% CI: 4.9-6.3%) and 5.2% (95% CI: 4.6-5.7%), respectively. The highest pooled prevalence was observed from Nigeria (19.0%; 95% CI: 12.7-25.3%), while the lowest prevalence was reported from Chad (0.05%; 95% CI: 0.03-0.13). Higher prevalence of schistosome cercaria was observed from Bulinus globosus (12.3%; 95% CI: 6.2–18.3%) followed by Biomphalaria sudanica (6.7%; 95% CI: 4.5–9.0%) and Biomphalaria pfeifferi (5.1%; 95% CI: 4.1-6.2%). The pooled prevalence of schistosome cercaria obtained using PCR was 26.7% in contrast to 4.5% obtained by shedding cercariae. Conclusion. This study revealed that nearly 6% of freshwater snails in Africa were infected by either S. haematobium or S. mansoni. The high prevalence of schistosomes among freshwater snails highlights the importance of appropriate snail control strategies in Africa.

1. Introduction

Schistosomiasis is one of the neglected tropical diseases (NTD) endemic in 78 countries and infects more than 229 million peoples in tropical and subtropical regions [1]. More than 90% of these cases are concentrated in Africa [2, 3]. The burden of the disease is even more severe in sub-Saharan Africa. Poor environmental sanitation and suitability of the climate conditions for snail intermediate host contribute to the high endemicity of the region. Schistosomiasis ranks second next to malaria from parasitic infection in terms of socioeconomic and health impact in tropics [4].

Human schistosomiasis is caused by *Schistosoma mansoni, S. haematobium, S. japonicum, S. intercalatum, S. mekongi, S. malayensis*, and *S. guineensis* [5–7]. Among these species, *S. mansoni, S. haematobium*, and *S. japonicum* are the major causes of human schistosomiasis globally [8]. *Schistosoma mansoni* and *S. haematobium* are widely distributed and the dominant cause of human schistosomiasis in Africa [5]. The endemicity of the disease in the region is linked with the availability of freshwater snail intermediate hosts.

About 350 species of freshwater snails are known to be medically or veterinary important [9]. Among these diverse

snails, *Biomphalaria, Bulinus*, and *Oncomelania* snails [4] are the dominant snail genera that are involved in the transmission of human schistosomiasis. The *Biomphalaria genus* consists of *B. pfeifferi, B. glabrata, B. sudanica, B. straminea, B. tenagophila, B. alexandarina,* and *B. choanomphala* [10]. *Biomphalaria* snails serve as the intermediate host for *S. mansoni*, which is responsible for intestinal and hepatic schistosomiasis. *Biomphalaria pfeifferi* is the most common and widely distributed snail intermediate host for *S. mansoni* in Africa.

Bulinus consists of 37 recognized species, which is grouped mainly into four species groups, namely, Bulinus africanus, B. forskalii, B. truncates/tropicus, and B. reticulatus [10, 11]. Bulinus snails serve as intermediate hosts for S. haematobium, which is responsible for urinary schistosomiasis. Oncomelania snails consist of only a few species mainly reported from Asia. The most common snail intermediate host for S. japonicum is Oncomelania hupensis, which is found in China, the Philippines, Indonesia, and also Japan [12].

The prevalence of human schistosomiasis is varied greatly in African countries depending on the level of environmental sanitation and the suitability of the area for the snail intermediate hosts, as well as the type of snail in the area. Similarly, the prevalence of schistosomes cercaria in snail intermediate hosts is varied in different locations within the same country and also from country to country in Africa. Several epidemiological studies are available on the types and prevalence of human infecting schistosomes among snail intermediate hosts in Africa. However, up to this time, there has not been any single estimate of the prevalence of S. mansoni and S. haematobium in snail intermediate hosts in Africa that could be used by African policymakers and international organizations working on the prevention and control of schistosomiasis in the continent. Therefore, this study aimed to provide summarized data on the prevalence and geographical variations of S. mansoni and S. haematobium cercaria among freshwater snails in Africa.

2. Methods

2.1. Search Strategies. Relevant literature was systematically searched from online public databases (PubMed Central, Science Direct, and Scopus) using the following key-words: "Schistosomiasis" OR "S. mansoni" OR "S. haematobium" OR "parasitological study" OR "schistosome intermediate host" OR "freshwater snails" OR "malacological survey" OR "Biomphalaria snails" OR "Bulinus snails" AND "Africa". The systematic review and selection of relevant literature were conducted according to PRISMA (Preferred Reporting Items for Systematic Reviews and Meta-analysis) guidelines [13] (Table S1).

2.2. Inclusion and Exclusion Criteria. Literature published in English language from 1979 to June 2020 were extracted from online public databases. Original articles reporting the prevalence of human schistosome cercariae in freshwater

snails in African countries were included in the analysis. The eligibility for the inclusion of a study in our analysis had to fulfill the following criteria: (a) it was published in English, (b) the study was carried out in Africa, (c) the number of examined and infected snails with either *S. mansoni* or *S. haematobium* were clearly stated, and (d) snail species were identified at least to a genus level. Studies that reported nonhuman schistosome and other trematodes species were excluded from the analysis. Besides, review articles and meta-analysis were excluded from the analysis.

2.3. Data Extraction Protocol. The data extraction protocol was prepared and evaluated by all authors. From each published article, we extracted the following information: author information, year of publication, study country, snail species, number of snails (collected, examined and infected), the prevalence of snail infection, the type of schistosomes reported, and methods of schistosome's detection.

2.4. Quality of Individual Study and Assessment of Bias. The quality of studies included in the meta-analysis was assessed by using the Newcastle-Ottawa quality assessment scale (NOS) proposed by Wells et al. [14] (Text S1). The quality assessment tool consists of three parts. First, the selection of study groups graded on a scale containing five stars; mainly deals with methodological qualities of individual study. Second, comparability of groups graded on a scale containing two stars; deals with comparability of studies based on design and analysis. Third, outcomes graded on a scale containing three stars, mainly focused on the assessment of the outcome and statistical analysis. Two authors (TH and EN) independently assessed the quality of individual study, and disagreement was solved by a discussion with the third author (AM). The overall quality of the individual study was categorized as high quality (≥ 8 stars), moderate quality (6-7 stars), and low quality (\leq 5 stars) by the total number stars obtained as described elsewhere [15].

2.5. Publication Bias across Studies. The risks of publication bias across studies were assessed using funnel plot symmetry qualitatively. Egger's and Begg's test were used to determine the presence of publication bias across studies quantitatively.

2.6. Data Analysis. We used a forest plot to estimate the overall pooled effect size with their 95% confidence interval (CI). The heterogeneity among studies used for this metaanalysis was evaluated using the I^2 test [16]. An I^2 - value lower than 25%, between 25% and 50%, and above 50% was regarded as low, moderate, and high heterogeneity, respectively [17]. Because of the high heterogeneity observed among the studies included in the meta-analysis, we used a random-effect model at 95% CI. To sort out the cause of heterogeneity, we used a subgroup analysis, sensitivity test, and meta-regression analysis. The data analysis was conducted using Stata software (version 14, STATA Corp College Station, TX), "metan" command.

3. Results

3.1. Search Results and Eligible Studies. A total of 2,995 relevant studies were screened from online public databases. Out of these studies, 976 articles were removed due to duplications while 1884 articles were excluded based on title and abstract screening. The remaining 135 full-text articles were assessed for eligibility. Of these, a total of 84 articles were excluded from the analysis based on specific exclusion criteria, and the remaining 51 articles were selected for this meta-analysis (Figure 1).

3.2. Characteristics of Subjects in the Eligible Studies. The eligible articles were obtained from 17 African countries: Angola [18], Benin [19], Burkina Faso [20], Burundi [21], Chad [22], Côte d'Ivoire [23, 24], Egypt [25–30], Ethiopia [31–36], Kenya [37–39], Mali [40, 41], Niger [42, 43], Nigeria [44–54], Senegal [55, 56], Sudan [57], Tanzania [58–63], Uganda [64–67], and Zimbabwe [68]. Unfortunately, there were no studies from other African countries that fulfilled the inclusion criteria. Characteristics of the eligible article to this meta-analysis are presented in Table 1.

3.3. *Risk of Bias within Studies.* The Newcastle–Ottawa quality assessment scale indicated that there was no bias within studies. The individual study included in this review was moderate to high-quality score as indicated in Table 1.

3.4. Prevalence of S. mansoni and S. haematobium Cercaria Among Freshwater Snails. A total of 273, 643 snails from Biomphalaria and Bulinus genera were examined for the presence of S. mansoni and S. haematobium cercaria in the 51 eligible studies, respectively (Table 2). Out of these snails, 8,682 of them were infected by either S. mansoni or S. haematobium. The prevalence of schistosome cercaria in the individual study ranged from 0.05% to 58.03% with substantial heterogeneity across studies within and across countries. The pooled prevalence of schistosome cercaria among freshwater snails was 5.5% (95% CI: 4.9–6.1%, $I^2 = 99.4\%$, and p < 0.001) (Figure 2).

3.5. Subgroup Analysis. The highest pooled prevalence of schistosome cercaria was observed among freshwater snails from Nigeria (19.0%; 95% CI: 12.7–25.3%), followed by Ethiopia (15.9%; 95% CI: -5.9–37.5%), Mali (5.2%; 95% CI: -0.3–10.7%), and Tanzania (4.9%; 95% CI: 3.8–6.0%) (Figure 3). We categorized the years of studies into four groups: before 2000, 2001 to 2010, 2011 to 2015, and 2016 to June 2020 to assess the trends on the prevalence of schistosome cercaria in freshwater snails. The pooled prevalence of schistosome cercaria among freshwater snails in years before 2000, 2001–2010, 2011–2015, and 2016–2020 was 1.3% (95% CI: 0.8–1.8%), 2.8% (95% CI: 1.8–3.8%), 6.1% (95% CI: 5.2–7%), and 8.3% (95% CI: 6.6–9.9%), respectively, in Africa (Figure 4).

This meta-analysis targets the two most common and widely distributed *Schistosoma* species (*S. mansoni* and

S. haematobium) in the continent. *Biomphalaria* and *Bulinus* snails were the intermediate hosts for *S. mansoni* and *S. haematobium*, respectively. The pooled prevalence of *S. mansoni* cercaria in *Biomphalaria* snails was 5.6% (95% CI: 4.9–6.3%) while the pooled prevalence of *S. haematobium* cercaria in *Bulinus* snails was 5.2% (95% CI: 4.7–5.7%) (Figure 5).

The pooled prevalence of *S. mansoni* and *S. haematobium* among freshwater snails was varied from country to country. The highest pooled prevalence of *S. mansoni* among *Biomphalaria* snails was observed from Tanzania (16.6%) followed by Ethiopia (15.9%) and Nigeria (14.5%) (Figure S1). On the contrary, the highest pooled prevalence of *S. haematobium* among *Bulinus* snail was observed from Nigeria (19.6%) followed by Angola (14.5%) and Côte d'Ivoire (9.6%) (Figure S2).

Twelve snail species from *Biomphalaria* and *Bulinus* snails were reported in the eligible articles used for this metaanalysis (Table 2). Among these species, *Biomphalaria pfeifferi* was the most common snail species and reported from 19 studies (37.3%) from the total 51 studies included in this study. The pooled prevalence of *S. mansoni* cercaria was 5.1% (95% CI: 4.1–6.2%) among *B. pfeifferi* snails. *Bulinus* snail, particularly *B. globosus* and *B. truncatus*, were the second and third most reported snails species (reported in 14 and 13 studies, respectively) that serve as an intermediate host for *S. haematobium*. The pooled prevalence of *S. haematobium* cercaria was 12.3% (95% CI: 6.2–18.3%) and 5.8% (95% CI: 4.4–7.2%) in *B. globosus* and *B. truncatus* snails, respectively (Table 2).

The studies included in this meta-analysis used shedding of cercariae and PCR-based detection of schistosomes from snail tissue. The pooled prevalence of schistosomes obtained by shedding of cercariae was 4.5% (95% CI: 3.9–5.1%) in contrast to 26.7% (95% CI: 10.5–43.0%) obtained by PCR techniques (Figure 6).

3.6. Publication Bias across Studies. The funnel plot symmetry demonstrates the presence of publication bias among studies included in this meta-analysis (Figure 7). Similarly, Egger's test results (p = 0.02) and Begg's test (p = 407) confirm the presence of publication bias among studies.

3.7. Metaregression Analysis and Sensitivity Test. There were clear heterogeneities across studies included in this metaanalysis. We performed a meta-regression analysis to identify the sources of heterogeneity across studies. The metaregression analysis showed that methods of schistosome detection from snails (regression coefficient: 2.55, 95% CI, 1.17–5.54, p = 0.02) might be the source of heterogeneity. The country of study (regression coefficient: 0.99, 95% CI, 0.89–1.11, p = 0.97), years of publication (regression coefficient: 1.41, 95% CI, 0.91–2.18, p = 0.12), snail genus (regression coefficient: 1.19, 95% CI, 0.56–2.55, p = 0.65), and snail species (regression coefficient: 0.92, 95% CI, 0.8 $p = 0.65 \ p = 0.14$) did not contribute for the heterogeneity. Besides, a sensitivity analysis was performed by recalculating the pooled prevalence by sequentially removing one-by-one



FIGURE 1: PRISMA flow diagram of for the inclusion of studies on the prevalence of *S. mansoni* and *S. haematobium* in freshwater snails in Africa.

to assess the effect of individual studies to overall effect. The pooled prevalence remained stable, and the result was not driven by individual studies included in the meta-analysis.

4. Discussion

Schistosomiasis is the second leading cause of infectious diseases next to malaria in Africa. Despite intensive efforts to tackle schistosomiasis, the prevalence is still unacceptably high in many African countries. The control and prevention strategies mainly rely on treatment of infected cases and mass drug administration of school-aged children. In many African countries, snail control strategies are not routinely implemented and sometimes considered as old-fashion approaches [70] despite their vital contributions to the elimination of schistosomes witnessed from Japan, Iran, and Tunisia [71-73]. In addition, enormous progresses have been observed in the elimination program from Morocco, Oman, Lebanon, and Caribbean Islands [73, 74]. The intensity and prevalence of schistosomes' infection among freshwater snails are scarce from many African countries. Summarized information about the prevalence of schistosomes among snail intermediate hosts is important for policymakers to give better attention to snail control strategies in Africa.

The overall pooled prevalence of schistosome cercaria was nearly 6% among freshwater snails in Africa. This finding is slightly lower than 9% reported from freshwater snails in Brazil [75]. In contrast, a lower prevalence of infected snails was observed from Indonesia [76] and Brazil [77]. These differences might be associated with prevalence and intensity of schistosome infection in the community, the level of environmental sanitation, suitability of the climate for the snails, level of existing snail control strategies, level of human exposure to open surface water, methods of schistosome detection, and seasons of snail collection and examination.

The highest pooled prevalence of schistosome cercaria among freshwater snails was observed in Nigeria followed by Ethiopia. In contrast, low prevalences of schistosomes were observed from Benin, Burundi, and Chad. The high prevalence of schistosomes among snail species in Nigeria and Ethiopia might be associated with the high prevalence of schistosomes in the community. The prevalence of schistosomiasis could reach as high as 90% in Ethiopia [78] and 94% in Nigeria [79, 80]. In addition, the difference in the level of environmental sanitation and the suitability of the area for the intermediate host, as well as the types of snail species in the area, may contribute for the difference in infection of snails across countries. Moreover, larger numbers of studies were reported from these two countries. Out of 51 studies, 17 (23.9%) studies included in this review were from the two countries.

Several species of freshwater snails that potentially serve as intermediate hosts for *S. mansoni* and *S. haematobium* have been recently reviewed [81]. *Biomphalaria* and *Bulinus* snails are the common and widely distributed intermediate hosts for schistosomes in Africa. The twelve snail species observed in this review belonged to either *Biomphalaria* or *Bulinus* genus.

Biomphalaria snails are well-known intermediate hosts of *S. mansoni* in Africa. This review showed that 5.6% of *Biomphalaria* snails were infected by *S. mansoni* in Africa. The highest pooled prevalence of *S. mansoni* was observed from Tanzania followed by Ethiopia and Nigeria, while low pooled prevalence was reported from Benin, Burundi, and Chad. These reported differences might be associated with

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				Snails		In	fected snails			I	otal		
Author	Year	Country	Snail species	examined (n)	Cercaria shedding (n)	Prevalence (%)	Screened by PCR	Positive (n)	Prevalence (%)	Infected snails (<i>n</i>)	Prevalence (%)	<i>Schistosoma</i> species	Quality score
Sturrock et al. [39]	1979	Kenya	B. pfeifferi	938	65	6.93		I	I	65	6.93	S. mansoni	4
King et al. [30]	1982	Egypt	Bullines spp.	4312	6	0.21	Ι	Ι		6	0.21	S. haematobium	8
Lwambo [60]	1988	Tanzania	B. nasutus	17646	156	0.88		I		156	0.88	S. haematobium	6
Gryseels [21]	1991	Burundi	B. pfeifferi	29199	249	0.85				249	0.85	S. mansoni	6
Bado et al. [40]	1997	Mali	B. truncatus B. globosus	266 90	- 14	- 7.80	266 —	22 —	8.30 —	22 7	8.3 7.80	S. haematobium S. haematobium	8
Kariuki et al. [37]	2004	Kenya	B. nasutus	11000	122	1.20	I	I	I	122	1.20	S. haematobium	8
Kazibwe et al. [66]	2006	Uganda	B. stanleyi B. sudanica	21715 8452	949 296	4.40 3.50				949 296	4.40 3.50	S. mansoni S. mansoni	6
Hussien et al. [29]	2007	Egypt	B. alexandrina	2070	10	0.48	I		I	10	0.48	S. mansoni	8
Labbo et al. [43]	2007	Niger	B. forskalii B. truncatus	9450 27125	5 509	0.05 1.90				5 509	0.05 1.90	S. haematobium S. haematobium	×
Hussien et al. [25]	2008	Egypt	B. truncatus	837	14	1.67			I	14	1.67	S. haematobium	7
Odongo-		,	B. choanomphala	9194	164	1.78	I		I	164	1.78	S. mansoni	
Aginya et al. [64]	2008	Uganda	B. pfeifferi	4173	75	1.79	I		I	75	1.79	S. mansoni	6
Ayanda [47]	2009	Nigeria	B. globosus B. pfeifferi	392 256	72 52	18.37 20.31				72 52	18.37 20.31	S. haematobium S. mansoni	~
lbikounlé et al. [19]	2009	Benin	B. pfeifferi	357	2	0.56	I		I	2	0.56	S. mansoni	8
Zongo et al. [20]	2009	Burkina Faso	B. truncatus	27	2	7.40	I	I	I	2	7.40	S. haematobium	4
Akinwale et al. [53]	2011	Nigeria	B. truncatus	138	I	I	138	41	29.70	41	29.70	S. haematobium	2
Lotf et al. [26]	2011	Egypt	B. alexandrina	277	17	6.10	I		I	17	6.10	S. mansoni	7
Mengistu et al. [34]	2011	Ethiopia	Biomphalaria spp.	560	325	58.00	I		I	325	58.00	S. mansoni	4
Onisa et al.			B. pfeifferi	425	7	1.60				7	1.60	S. mansoni	
[38]	2011	Kenya	B. sudanica	407		1.70			I	Γ ι	1.70	S. mansoni	6
Aba et al [50]	2012	Nigeria	B. globosus R truncatus	777	υ c	2.20 44.64				υ Ľ	2.20 44.64	S. haematobium S. haematohium	L
Iboh et al. [48]	2012	Nigeria	B. globosus	120	13	10.80	I		I	13	10.80	S. haematobium	~ ∞
Mekonnen et al. [35]	2012	Ethiopia	B. pfeifferi	80	2	2.50	I		I	2	2.50	S. mansoni	7
Kinanpara [23]	2013	Côte d'I <i>v</i> oire	B. globosus B nfeifferi	189 1409	23 25	12.17 1 77				23 25	12.17 1 77	S. haematobium S. mansoni	~
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						TABLE 1: CO	ntinued.						
				Snails		Inf	fected snails			To	ıtal		
Author	Year	Country	Snail species	examined (n)	Cercaria shedding (n)	Prevalence (%)	Screened by PCR	Positive (n)	Prevalence (%)	Infected snails (<i>n</i>)	Prevalence (%)	<i>Schistosoma</i> species	Quality score
Alebie et al. [31]	2014	Ethiopia	B. pfeifferi	301	32	10.60	I	I	I	32	10.60	S. mansoni	7
Angelo et al. [62]	2014	Tanzania	B. sudanica	1470	11	0.75		I		11	0.75	S. mansoni	8
Ivoke et al. [49]	2014	Nigeria	B. globosus	308	62	20.10			I	62	20.10	S. haematobium	8
Moser et al. [22]	2014	Chad	B. truncatus B. pfeifferi	4119 108	1 1	0.80 0.90				1 1	0.02 0.90	S. haematobium S, mansoni	×
Akinwale et al. [54]	2015	Nigeria	B. globosus B. forskalii B. camerunensis	109 22 7			109 22 7	38 8 8 4	34.80 36.40 57.00	86 8 4	34.80 36.40 57.00	S. haematobium S. haematobium S. haematohium	6
- - -			B. senegalensis	11	Ι	I	11	5	18.20	5	18.20	S. haematobium	
Amsalu et al. [33]	2015	Ethiopia	B. pfeifferi	31	1	3.20			I	1	3.20	S. mansoni	7
Bado et al. [41]	2015	Mali	B. truncatus B. pfeifferi	324 189	11 2	3.40 1.10				11 2	$3.40 \\ 1.10$	S. haematobium S. mansoni	8
Gashaw et al. [32]	2015	Ethiopia	B. pfeifferi	14	2	14.30		I	I	2	14.30	S. mansoni	7
Rowel et al., [67]	2015	Uganda	Biomphalaria spp	19355	127	0.70		I	I	127	0.70	S. mansoni	6
Senghor [55]	2015	Senegal	B. senegalensis B. umbilicatus	7333 339	8 22	0.11 6.51				8 22	0.11 6.51	S. haematobium S. haematobium	6
Aboelhadid et al. [27]	2016	Egypt	B. alexandrina B. truncatus	822 423	22 9	22.80 2.10				22 9	2.68 2.10	S. mansoni S. haematobium	2
Farghaly et al. [28]	2016	Egypt	B. alexandrina	400	б	0.8	400	13	3.3	13	3.3	S. mansoni	6
Mohammed et al. [57] Pennance [63]	2016 2016	Sudam Tanzania	B. truncatus B. pfeifferi B. globosus	1403 5100 1111	2 82 26	0.10 1.60 2.30				2 82 26	0.10 1.60 2.30	S. haematobium S. mansoni S. haematobium	6 8
Alemayehu et al. [36]	2017	Ethiopia	B. pfeifferi	111	7	6.30			I	7	6.30	S. mansoni	8
Allan et al. [18]	2017	Angola	B. globosus	173	25	14.50			I	25	14.50	S. haematobium	8
Bakuza et al. [69]	2017	Tanzania	B. pfeifferi	235	29	12.30	219	103	47.00	103	43.8	S. mansoni	8
Gouvras et al. [61] Otaba and	2017	Tanzania	B. sudanica B. choanomphala	35,910 6906	439 61	$1.21 \\ 0.88$				439 61	$1.21 \\ 0.88$	S. mansoni S. mansoni	~
Ubachukwu	2017	Nigeria	B. pfeifferi	460	20	4.30		I	I	20	4.30	S. mansoni	×
Stanton et al. [65]	2017	Uganda	Biomphalaria spp.	499	5	1.00	118	22	18.64	27	5.4	S. mansoni	8

	,	Quality score		8		0	0	8	6	c	۷	c	٧	7	6	σ	`
	,	<i>Schistosoma</i> C species	S. mansoni	S. haematobium	S. haematobium	S. haematobium	S. haematobium	S. mansoni	S. haematobium	S. haematobium	S. mansoni	S. haematobium	S. haematobium	S. mansoni	S. mansoni	S haematohium	0. 11401111110011111
	otal	Prevalence (%)	26.00	23.90	24.30	19.20	17.00	7.45	4.50	0.20	3.40	0.10	2.00	2.20	35.4	1 90	0/1
	Tc	Infected snails (n)	156	1421	2166	34	18	16	5	24	79	2	5	6	279	30	2
		Prevalence (%)	I	I	I			7.45	I	I		I			35.40		
		Positive (n)	I	I	I	Ι	I	16	I	I	Ι	I	I	I	279	l	
ntinued.	ected snails	Screened by PCR		I	I			212	I	I		I		I	788		
Table 1: Coi	Inf	Prevalence (%)	26.00	23.90	24.30	19.20	17.00		4.50	0.20	3.40	0.10	2.00	2.20	I	1 90	7.77
		Cercaria shedding (n)	156	1421	2166	34	18		5	24	79	2	5	6	I	30	2
	Snails examined (n)		592	5942	8894	177	106	212	112	11989	2290	1772	247	407	788	1547	71.0T
		Snail species	B. pfeifferi	B. truncatus	B. globosus	B. globosus	B. truncatus	B. pfeifferi	B. globosus	B. forskalii	B. pfeifferi	B. truncatus	B. globosus	B. pfeifferi	B. sudanica	R alabour	D. Zwwwa
		Country		Nigeria		Miccourto	INIGETIA	Nigeria	Nigeria	NE	INIGE	Côte	d'Ivoire	Senegal	Tanzania	Zimhahwa	
		Year		2018		0100	6107	2019	2019	0100	6107	0100	6107	2020	2020	0000	0404
		Author		Aliyu et al. [51]		Afiukwa et al.	[46]	Okeke et al. [45]	Peletu et al. [44]	Rabone et al.	[42]	Tian-Bi et al.	[24]	Catalano et al. [56]	Fuss et al. [59]	Mutsaka- Malanyaza	et al. [68]

7

Servil assess	Smail amonion	Studios (4)	Examined ensite (u)		Infected snails
shan genus	Shall species	studies (n)	Examined shalls (n)	п	pp (95%CI)
	B. pfeifferi	19	46480	987	5.10 (4.05-6.15)
	B. sudanica	5	47027	1032	6.73 (4.46-9.01)
D:	B. alexandrina	4	3569	62	2.81 (0.75-4.87)
вютрпагата	B. choanomphala	2	16,100	225	1.58 (0.44-2.21)
	B. stanleyi	1	21,715	949	4.37 (4.09-4.64)
	Unclassified Biomphalaria snails	3	20414	479	21.28 (-1.78-44.34)
	B. globosus	14	13691	2511	12.25 (6.23-18.27)
	B. truncatus	13	42538	2077	5.78 (4.36-7.20)
	B. nasutus	2	28648	278	0.98 (0.76-1.20)
D1:	B. senegalensis	2	7344	10	4.73 (-10.72-20.18)
Dulinus	B. umbilicatus	1	339	22	6.49 (3.71-9.27)
	B. forskalii	3	21461	37	0.13 (0.08-0.34)
	B. camerunensis	1	7	4	57.14 (21.30-92.99)
	Unclassified Bulinus snails	1	4312	9	2.1 (0.10-3.90)
	Total	51	273 643	8682	5.51 (4.95-6.07)

TABLE 2: The pooled prevalences of S. mansoni and S. haematobium infection among Biomphalaria and Bulinus snails of Africa.

the level of environmental sanitation, the abundance of *Biomphalaria* snails, and seasons of snail collection and examination. The highest prevalence of *S. mansoni* among *Biomphalaria* snails was reported during the dry season or just before the beginning of the rain seasons [47, 82].

Five species of *Biomphalaria* snails (*B. pfeifferi*, *B. sudanica*, *B. choanomphala*, *B. alexandrina*, and *B. stanleyi*) were included in the eligible articles for this review. As reviewed by Abe et al. [81], *B. pfeifferi*, *B. sudanica*, *B. choanomphala*, and *B. alexandrina* were the common intermediate hosts of *S. mansoni* in Africa. The pooled prevalence of *S. mansoni* varied from 1.3% to 6.7% among these snail species.

Biomphalaria pfeifferi were the most common snails infected by schistosome parasite. About 40% of the eligible studies included in this meta-analysis reported *B. pfeifferi*. The role of *B. pfeifferi* as an intermediate host of *S. mansoni* varied from country to country. *Biomphalaria pfeifferi* is the sole intermediate host for *S. mansoni* in Côte d'Ivoire [83] and Senegal [84] and the dominant intermediate host in many other African countries [85–88].

Biomphalaria sudanica was the second common *Biomphalaria* snails that serve as an intermediate host for *S. mansoni* as observed in this study. *Biomphalaria sudanica* is an intermediate host for *S. mansoni* in Kenya [47] and Tanzania [19, 89]. *Biomphalaria sudanica* is also reported from Ethiopia but limited to around Lake Ziway [90] and Tikur Wuha [87].

Biomphalaria alexandrina was the third common intermediate host for *S. mansoni* observed in this study. However, its contribution as an intermediate host for *S. mansoni* is restricted in geographical distribution, mainly reported from Egypt [91–93]. *Biomphalaria choanomphala* was another intermediate host for *Schistosoma* mansoni reported from Uganda [64] and Tanzania [61]. This snail species is widely distributed around Lake Victoria, which is divided among three countries (Kenya, Tanzania, and Uganda). *Biomphalaria choanomphala* is also reported from Kenya [94], but the infection intensity was not determined. *Bulinus* snail is the common intermediate host for *S. haematobium* in Africa. This study revealed that about 5.2% of *Bulinus* snails were infected with *S. haematobium* in Africa. The highest pooled prevalence of *S. haematobium* infection among *Bulinus* snail was observed from Nigeria followed by Angola and Côte d'Ivoire. In contrast, there was a low prevalence of *S. haematobium* infection among *Bulinus* snails from Chad, Niger, Senegal, and Sudan. These variations might be associated with the difference in the level of endemicity of *S. haematobium* in the countries. A recent review indicated that about one-third of the populations of Nigeria were infected by *S. haematobium* [95, 96]. The higher infection intensity in the population might lead to a high level of environmental contamination that resulted in higher snail infection in Nigeria.

The eligible studies included in this review report seven species of *Bulinus* snails (*B. truncates B. globosus, B. forskalii, B. senegalensis, B. nasutus, B. camerunensis, and B. umbilicatus*) from African countries. According to Abe et al. [81], *B. truncates, B. africanus, B. forskalii, B. senegalensis,* and *B. camerunensis* were the predominant *Bulinus* snails that serve as an intermediate host for *S. haematobium*.

Among the *Bulinus* snails, *B. globosus* was reported from 14 studies in 7 African countries included in this review. The present study revealed that 12.3% of *B. globosus* was infected by *S. haematobium*. Similarly, high pooled prevalence (18%) of *S. haematobium* among *B. globosus* was recently reported in a meta-analysis from Nigeria [96]. *Bulinus truncatus* are the other important *Bulinus* snails that serve as an intermediate host for *S. haematobium* was 5.9% among *Bulinus truncatus* snails in Africa. In contrast to our result, 19% prevalence of *S. haematobium* was reported from *B. truncates* snails in Nigeria [31]. Similarly, *B. truncatus* is the predominant intermediate host for *S. haematobium* in Niger [42] and Côte d'Ivoire [23, 24].

Detection of schistosome infection is determined by shedding of cercariae and/or PCR based approaches from snail tissue. There was a significant difference in the pooled

Author (year of publication)	ES (95% CI)	% Weight
Sturrock et al. (1979)	0.07 (0.05, 0.09)	2.28
King et al. (1982)	0.00 (0.00, 0.00)	2.87
Lwambo (1988)	0.01 (0.01, 0.01)	2.88
Gryseels (1991)	0.01 (0.01, 0.01)	2.88
Bado et al. (1997)	0.08 (0.06, 0.11)	1.58
Kariuki et al. (2004)	0.01 (0.01, 0.01)	2.87
Kazibwe et al (2006)	0.04(0.04, 0.04)	2.87
Hussein et al. (2007)	0.00(0.00, 0.01)	2.83
Labbo et al. (2007)	0.01 (0.01, 0.02)	2.87
Hussien et al. (2008)	0.02 (0.01, 0.03)	2.66
Odongo-Aginya et al (2008)	0.02(0.01, 0.02)	2.86
Avanda (2009)	0.19(0.17, 0.22)	1.67
bikoundé et al (2009)	0.01 (0.00, 0.02)	2.65
	0.07 (0.01, 0.24)	0.21
Akinwale et al (2011)	0.30(0.22, 0.38)	0.43
Lotfetal (2011)	0.06(0.04, 0.10)	1.57
Menority et al (2011)	0.58 (0.54 0.61)	1.29
		2 70
Aba et al (2017)	0.02 (0.01, 0.03)	0.16
The tail (2012)	0.45 (0.51, 0.59)	0.10
	0.03 (0.00, 0.09)	1.09
	0.03(0.00, 0.09)	2.60
Alabia a (2014)	0.03(0.02, 0.04)	1.30
Anorde et al. (2014)	0.11(0.07, 0.13)	2.92
Angelo et al. (2014)	0.01(0.00, 0.01)	2.65
	0.20 (0.10, 0.23)	0.97
Moser et al. (2014) Altimute at al (2015)	0.00(0.00, 0.00)	2.88
Akniwate et al. (2015)	0.33(0.27, 0.43)	0.42
Amsau et al. (2015)	0.03 (0.00, 0.17)	0.39
	0.03 (0.01, 0.04)	2.40
Gasnaw et al. (2015)	0.14 (0.02, 0.43)	0.07
	0.01 (0.01, 0.01)	2.87
Sengnor (2015)	0.00 (0.00, 0.01)	2.87
Aboeinadid et al. (2016)	0.02 (0.02, 0.04)	2.67
rargnaily et al. (2016)	0.03 (0.02, 0.05)	2.16
Monammed et al. (2016)	0.01 (0.01, 0.02)	2.86
Pennance (2016)	0.02 (0.02, 0.03)	2.66
Alemayehu et al. (2017)	0.06 (0.03, 0.13)	0.87
Ailan et al. (2017)	0.14 (0.10, 0.21)	0.76
Bakuza et al. (2017)	0.44 (0.37, 0.50)	0.59
Goldvas et al. (2017)		2.88
Okeke & Ubachukwu (2017)	0.04 (0.03, 0.07)	2.12
Stanton et al. (2017)	0.05 (0.04, 0.08)	2.05
Airyu et al. (2018)	0.24 (0.24, 0.25)	2.76
Ahukwa et al. (2019)	0.19(0.16, 0.22)	1.53
Okeke et al. (2019)	0.07 (0.04, 0.12)	1.24
Peletu BJ et al. (2019)	0.05 (0.01, 0.10)	1.07
Rabone et al. (2019)	0.01 (0.01, 0.01)	2.87
1 ian-Bi et al. (2019)	0.00 (0.00, 0.01)	2.86
Catalano et al. (2020)	0.02 (0.01, 0.04)	2.35
russ et al. (2020)	0.35 (0.32, 0.39)	1.39
(1)	0.02(0.01, 0.03)	2.75
Overall (1-squared = 99.4%, $p \le 0.001$)	0.06 (0.05, 0.06)	100.00
NOTE: weights are from random effects analysis	Т	
-0.612 I	 0.612	

FIGURE 2: Forest plot diagram showing the prevalence of human infecting schistosomes (*S. mansoni* and *S. haematobium*) examined in snails in Africa. Each square represent effect size (ES) of individual studies, and the horizontal line represents the 95% CI. The diamond indicates the pooled effect and the vertical dash lines indicate the overall estimate.

prevalence of schistosome results between shedding of cercaria (4.5%) and PCR (26.7%) among snail species in Africa. Similar differences (1.56% vs. 39.8%) are seen in the prevalence of schistosome infection between cercarial shedding and PCR methods among snails as reported from Kenya [97]. This difference is associated with the sensitivity of PCR to detect schistosome infection from snail tissue. Cercarial shedding is suffered by several limitations such as low parasite burden; snails may not shed cercariae during

the prepatent period; time-consuming, and labour-intensive [98]. PCR-based detection of schistosome infection from snails is generally rapid, efficient, sensitive, and cost-effective for large-scale detection [99, 100].

Despite the ongoing schistosomiasis control strategies in many African countries, the pooled prevalence of schistosomes among freshwater snails had increased over time from 1.3% (before 2000) to 8.3% between 2016 and 2020. This might be attributed to the large number of epidemiological



FIGURE 3: Geographical distribution and pooled prevalence of *S. mansoni* and *S. haematobium* among *Biomphalaria* and *Bulinus* snails, respectively, in African countries.

studies conducted and reported recently. Besides, molecular based detection of schistosome infection from snail tissue received attention in the recent years. These situations may contribute to the increased prevalence of schistosomes among freshwater snails recently.

4.1. *Limitation of the Study*. Although this systematic review generated valuable data on the prevalence of *S. mansoni* and

S. haematobium among freshwater snails in Africa, it also has limitations. First, information on the prevalence of schistosome cercaria among snail species was not obtained from all African countries. Prevalence data were available only from 17 African countries. The pooled prevalences of schistosomes in this review may not fully represent the prevalence of *S. mansoni* and *S. haematobium* among freshwater snails of Africa. Second, the numbers of published studies were not evenly distributed even in the 17

Author (year of publication)	ES (95% CI)	% Weight
1979 to 2000		
Sturrock et al. (1979)	0.07 (0.05, 0.09)	2.28
King et al. (1982)	0.00 (0.00, 0.00)	2.87
Lwambo (1988)	0.01 (0.01, 0.01)	2.88
Gryseels (1991)	0.01(0.01, 0.01)	2.88
		1.58
Subtotal (1-squared = 97.2% , $p \le 0.001$)	0.01 (0.01, 0.02)	12.49
2001 to 2010		
Kariuki et al. (2004)	0.01 (0.01, 0.01)	2.87
Kazıbwe et al. (2006)	0.04 (0.04, 0.04)	2.87
Hussein et al. (2007)	0.00(0.00, 0.01)	2.83
Labbo et al. (2007)	0.01(0.01, 0.02)	2.87
Odongo Aginya et al. (2008)	0.02(0.01, 0.03)	2.00
Avanda (2000)	0.02(0.02, 0.02)	1.67
Ibikounlé et al. (2009)	0.19(0.17, 0.22)	2.65
Zongo et al. (2009)		0.21
Subtotal (Leavared = 98.9% th ≤ 0.001)	0.03(0.02,0.04)	21.40
Subjoint (1-squarea $-30.376, p \le 0.001$)	0.03 (0.02, 0.04)	21.49
2011 to 2015	0.20 (0.22, 0.28)	0.42
Akinwale et al. (2011)		0.43
Lott et al. (2011)	0.58(0.54, 0.10)	1.57
Opios et al. (2011)	- 0.58 (0.54, 0.61)	1.29
Abe et al. (2012)	0.02(0.01, 0.03)	2.70
Those et al. (2012)	◆ 0.15 (0.57, 0.57) ◆ 0.11 (0.06, 0.18)	0.67
Mekonnen et al. (2012)	0.03 (0.00, 0.09)	1.09
Kinanpara (2013)	0.03(0.02, 0.04)	2.69
Alebie et al. (2014)	• 0.11 (0.07, 0.15)	1.30
Angelo et al. (2014)	0.01 (0.00, 0.01)	2.83
Ivoke et al. (2014)	• 0.20 (0.16, 0.25)	0.97
Moser et al. (2014)	0.00 (0.00, 0.00)	2.88
Akinwale et al. (2015)		0.42
Amsalu et al. (2015)	0.03 (0.00, 0.17)	0.39
Dabo et al. (2015)	0.03 (0.01, 0.04)	2.40
Gashaw et al. (2015)	• 0.14 (0.02, 0.43)	0.07
Rowel et al. (2015)	0.01 (0.01, 0.01)	2.87
Senghor (2015)	0.00 (0.00, 0.01)	2.87
Subtotal (I-squared = 98.8%, $p \le 0.001$)	0.06 (0.05, 0.07)	27.60
2016 to 2020		
Aboelhadid et al. (2016)	0.02 (0.02, 0.04)	2.67
Farghaly et al. (2016)	0.03 (0.02, 0.05)	2.16
Monammed et al. (2016)	0.01(0.01, 0.02)	2.86
Alemance (2016)	0.02(0.02, 0.03)	2.00
Allan et al. (2017)	- 0.00 (0.03, 0.13)	0.87
Bakuza et al. (2017)		0.59
Gouvras et al. (2017)	0.01 (0.01, 0.01)	2.88
Okeke & Ubachukwu (2017)	0.04 (0.03, 0.07)	2.12
Stanton et al. (2017)	0.05 (0.04, 0.08)	2.05
Aliyu et al. (2018)	• 0.24 (0.24, 0.25)	2.76
Afiukwa et al. (2019)	•• 0.19 (0.16, 0.22)	1.53
Okeke et al. (2019)	- 0.07 (0.04, 0.12)	1.24
Peletu BJ et al. (2019)	- 0.05 (0.01, 0.10)	1.07
Rabone et al. (2019)	0.01 (0.01, 0.01)	2.87
Tian-Bi et al. (2019)	0.00 (0.00, 0.01)	2.86
Catalano et al. (2020)	0.02 (0.01, 0.04)	2.35
Fuss et al. (2020)	• 0.35 (0.32, 0.39)	1.39
Mutsaka-Makuvaza et al. (2020)	0.02 (0.01, 0.03)	2.75
Subtotal (I-squared = 99.7%, $p \le 0.001$)	0.08 (0.07, 0.10)	38.42
Overall (<i>I</i> -squared = 99.4%, <i>p</i> ≤ 0.001)	0.06 (0.05, 0.06)	100.00
NOTE: weights are from random effects analysis		
 _0.612 0	0.612	
0.012 0	0.012	

FIGURE 4: The pooled prevalence of schistosome cercaria among freshwater snails in Africa based on year of publication.

countries (varied from 1 study to 11 studies in a country). Third, the studies included in this review were published in English, and we did not include studies published in other

languages such as French due to language barriers and translation-related challenges. Fourth, most of the studies included in this review used cercarial shedding rather than

Author (year of publication)	ES (95% CI)	% Weight
Biomphalaria Snails		
Sturrock et al. (1979)	• 0.07 (0.05, 0.09)	1.97
Gryseels (1991)	0.01 (0.01, 0.01)	2.51
Kazibwe et al. (2006)	0.04 (0.04, 0.04)	2.50
Hussein et al. (2007)	0.00 (0.00, 0.01)	2.47
Odongo-Aginya et al. (2008)	0.02 (0.02, 0.02)	2.50
Ayanda (2009)	$- \bullet - 0.20 (0.16, 0.26)$	0.71
Ibikounie et al. (2009)		2.30
Lott et al. (2011) Managistra et al. (2011)	- 0.06 (0.04, 0.10)	1.55
Opica et al. (2011)		2.31
Mekonnen et al. (2012)	= 0.03 (0.00, 0.09)	0.92
Kinanpara (2013)	0.02 (0.01, 0.03)	2.39
Alebie et al. (2014)		1.09
Angelo et al. (2014)	0.01 (0.00, 0.01)	2.46
Moser et al. (2014)	0.01 (0.00, 0.05)	1.55
Amsalu et al. (2015)	0.03 (0.00, 0.17)	0.32
Dabo et al. (2015)	0.01 (0.00, 0.04)	1.90
Gashaw et al. (2015)	• 0.14 (0.02, 0.43)	0.06
Rowel et al. (2015)	0.01 (0.01, 0.01)	2.51
Aboelhadid et al. (2016)	0.03 (0.02, 0.04)	2.22
Farghaly et al. (2016)	0.03 (0.02, 0.05)	1.86
Mohammed et al. (2016)	0.02 (0.01, 0.02)	2.49
Alemayehu et al. (2017)		0.73
Bakuza et al. (2017)	0.44(0.37, 0.50)	0.49
Okeke & Libechukuu (2017)		2.51
Stanton et al. (2017)	0.05(0.04, 0.05)	1.82
Alivu et al. (2018)		1.10
Okeke et al. (2019)	• 0.08 (0.04, 0.12)	1.04
Rabone et al. (2019)	0.03 (0.03, 0.04)	2.38
Catalano et al. (2020)	0.02 (0.01, 0.04)	2.03
Fuss et al. (2020)	→ 0.35 (0.32, 0.39)	1.18
Subtotal (I-squared = 98.9%, $p \le 0.001$)	0.06 (0.05, 0.06)	54.50
Bulinus Snails		
King et al. (1982)	0.00 (0.00, 0.00)	2.51
Lwambo (1988)	0.01 (0.01, 0.01)	2.51
Dado et al. (1997) Koriuli et al. (2004)		1.54
Labbo et al. (2007)	0.01(0.01, 0.01)	2.51
Hussien et al. (2008)	0.02(0.01, 0.02)	2.32
Avanda (2009)		0.99
Zongo et al. (2009)	0.07 (0.01, 0.24)	0.17
Akinwale et al. (2011)	0.30 (0.22, 0.38)	0.35
Opisa et al. (2011)	0.02 (0.01, 0.04)	1.91
Abe et al. (2012)	• 0.45 (0.31, 0.59)	0.13
Iboh et al. (2012)	— 0.11 (0.06, 0.18)	0.56
Kinanpara (2013)	• 0.12 (0.08, 0.18)	0.75
lvoke et al. (2014)		0.81
Moser et al. (2014)	0.00 (0.00, 0.00)	2.52
Akinwale et al. (2015) Daba at al. (2015)	0.35(0.27, 0.43)	0.35
Senghor (2015)	0.00(0.02, 0.00)	2.51
Aboelhadid et al. (2016)	0.02 (0.01, 0.04)	2.06
Mohammed et al. (2016)	0.00 (0.00, 0.01)	2.50
Pennance (2016)	0.02 (0.02, 0.03)	2.32
Allan et al. (2017)	→ 0.14 (0.10, 0.21)	0.63
Aliyu et al. (2018)	• 0.24 (0.23, 0.25)	2.40
Afiukwa et al. (2019)	→ 0.18 (0.14, 0.23)	0.80
Peletu BJ et al. (2019)	0.05 (0.01, 0.10)	0.89
Kabone et al. (2019)	0.00 (0.00, 0.00)	2.51
Itan-Bi et al. (2019)	0.00(0.00, 0.01)	2.50
IVILISARA-IVIAKUVAZA ET AI. (2020)	0.02 (0.01, 0.03)	2.39 45 50
· · · · · · · · · · · · · · · · · · ·		45.50
Overall (<i>I</i> -squared = 99.3%, $p \le 0.001$)	0.06 (0.05, 0.06)	100.00
NOTE: weights are from random effects analysis		
-0.612 0	0.612	

FIGURE 5: Forest plot diagram showing the estimated effect size of S. mansoni in Biomphalaria snails and S. haematobium Bulinus snail in Africa.

Author (year of publication)		ES (95% CI)	% Weight
Cercaria shedding			
Sturrock et al. (1979)	•	0.07 (0.05, 0.09)	2.28
King et al. (1982)	•	0.00 (0.00, 0.00)	2.87
Lwambo (1988)		0.01 (0.01, 0.01)	2.88
Gryseels (1991)		0.01 (0.01, 0.01)	2.88
Bado et al. (1997)	<u> </u> ● −	0.08 (0.06, 0.11)	1.58
Kariuki et al. (2004)		0.01(0.01, 0.01)	2.87
Kazibwe et al. (2006)		0.04 (0.04, 0.04)	2.87
Labbo et al. (2007)		0.00(0.00, 0.01)	2.85
Hussien et al. (2007)		0.01(0.01, 0.02) 0.02(0.01, 0.03)	2.67
Odongo-Aginya et al. (2008)		0.02 (0.02, 0.02)	2.86
Avanda (2009)		0.19 (0.17, 0.22)	1.67
Ibikounlé et al. (2009)	•	0.01 (0.00, 0.02)	2.65
Zongo et al. (2009)		0.07 (0.01, 0.24)	0.21
Lotf et al. (2011)	+	0.06 (0.04, 0.10)	1.57
Mengistu et al. (2011)	-		1.29
Opisa et al. (2011)	•	0.02 (0.01, 0.03)	2.70
Abe et al. (2012)	↓ ↓	- 0.45 (0.31, 0.59)	0.16
Iboh et al. (2012)		0.11 (0.06, 0.18)	0.67
Mekonnen et al. (2012)	◆	0.03 (0.00, 0.09)	1.09
Kinanpara (2013)		0.03 (0.02, 0.04)	2.69
Angele et al. (2014)		0.11(0.07, 0.15)	1.30
Jucke et al. (2014)		0.01(0.00, 0.01) 0.20(0.16, 0.25)	2.83
Moser et al. (2014)		0.20(0.10, 0.23) 0.00(0.00, 0.00)	2.88
Amsalu et al. (2015)	Ĩ <u>.</u>	0.03(0.00, 0.17)	0.39
Dabo et al. (2015)		0.03 (0.01, 0.04)	2.40
Gashaw et al. (2015)		0.14 (0.02, 0.43)	0.07
Rowel et al. (2015)		0.01 (0.01, 0.01)	2.87
Senghor (2015)	•	0.00 (0.00, 0.01)	2.87
Aboelhadid et al. (2016)		0.02 (0.02, 0.04)	2.67
Mohammed et al. (2016)	•	0.01 (0.01, 0.02)	2.86
Pennance (2016)		0.02 (0.02, 0.03)	2.66
Alemayehu et al. (2017)		0.06 (0.03, 0.13)	0.87
Alian et al. (2017)		0.14 (0.10, 0.21)	0.76
Olvelse & Libechylguy (2017)		0.01(0.01, 0.01)	2.88
Alivn et al. (2018)		0.04(0.03, 0.07) 0.24(0.24, 0.25)	2.12
Afjukwa et al. (2019)		0.24(0.24, 0.23) 0.19(0.16, 0.22)	1.53
Peletu BI et al. (2019)		0.05 (0.01, 0.10)	1.07
Rabone et al. (2019)	•	0.01 (0.01, 0.01)	2.87
Tian-Bi et al. (2019)	•	0.00 (0.00, 0.01)	2.86
Catalano et al. (2020)		0.02 (0.01, 0.04)	2.35
Mutsaka-Makuvaza et al. (2020)		0.02 (0.01, 0.03)	2.75
Subtotal (I-squared = 99.4%, $p \le 0.001$)	◊	0.05 (0.04, 0.05)	91.72
PCR			
Akinwale et al. (2011)		0.30(0.22, 0.38)	0.43
Akinwale et al. (2015) Okolega et al. (2010)	→ →	0.35(0.27, 0.43) 0.07(0.04, 0.12)	0.42
Fuse et al. (2019)		0.07 (0.04, 0.12) 0.35 (0.32, 0.39)	1.24
Subtotal (I-sauared = 97.6% , p < 0.001)		0.33(0.32, 0.3)) 0.27(0.10, 0.43)	3.48
		0.27 (0.10, 0.10)	5.10
Cercaria shedding & PCR			
Farghaly et al. (2016)	*	0.03 (0.02, 0.05)	2.16
Bakuza et al. (2017)	_ →	0.44 (0.37, 0.50)	0.59
Stanton et al. (2017)	*	0.05 (0.04, 0.08)	2.05
Subtotal (I-squared = 98.6%, $p \le 0.001$)		0.17 (0.04, 0.30)	4.80
Overall (<i>I</i> -squared = 99.4%, <i>p</i> = 0.000)		0.06 (0.05, 0.06)	100.00
NOTE: weights are from random effects analysis			
	1		
-0.612	0	0.612	

FIGURE 6: Forest plot diagram showing the pooled prevalences of *S. mansoni* and *S. haematobium* among freshwater snails examined by cercarial shedding and PCR.

PCR for the detection of schistosome infection. Cercarial shedding is less sensitive for the detection of schistosomes due to its inherent limitation that may result in low prevalence of schistosomes among infected snails in Africa. The pooled prevalence of schistosome cercaria among freshwater snails of Africa observed in this review might be below the actual infection intensity. Fifth, this review showed that there was high heterogeneity across

0 0.5 1 1.5 -2 -8 -6 $^{-4}$ 0 loges

FIGURE 7: Funnel plot with 95% confidence limit showing publication bias across studies on the prevalence of S. mansoni and S. haematobium among freshwater snails of Africa.

studies included in this review. These might be associated with study design, seasons of snail collection, method of detection, and variation of endemicity of schistosomes across countries.

5. Conclusions

This review showed that nearly 6% of freshwater snails in Africa were infected by either S. haematobium or S. mansoni. The pooled prevalences of schistosome cercaria among freshwater snails have increased in the recent years in many African countries. The higher and increased trends in the prevalence of schistosomes among freshwater snails highlight the need for appropriate snail control strategies in the region. Policymakers should give better attention about integration of snail control strategies to the ongoing treatment-based prevention of schistosomiasis in Africa.

Abbreviations

CI:	Confidence interval
I^2 :	Inverse variance index
NOS:	Newcastle-Ottawa quality assessment scale
PRISMA:	Preferred Reporting Items for Systematic
	Reviews and Meta-analysis
PCR:	Polymerase chain reaction
NTD:	Neglected tropical disease.

Data Availability

All the datasets are included in the manuscript.

Conflicts of Interest

The authors declare that there are no conflicts of interest.

Authors' Contributions

TH was involved in the design, conducted the review, data analysis, and interpretation of the findings, and drafted the manuscript. EN and AM were involved in drafting the manuscript, reviewing, and editing the manuscript.

Supplementary Materials

Table: PRISMA checklist. S1 **S**1 Text: NEWCASTLE-OTTAWA QUALITY ASSESSMENT SCALE. S1 Fig: forest plot showing the pooled effect estimate of S. mansoni among Biomphalaria snails in Africa. S2 Fig: forest plot showing the pooled prevalence estimate of S. haematobium among Bulinus snails in Africa. (Supplementary Materials)

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Funnel plot with pseudo 95% confidence limits

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