Evaluation of real-time PCR for *Strongyloides stercoralis* and hookworm as diagnostic tool in asymptomatic schoolchildren in Cambodia

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Abstract

Diagnosis of soil-transmitted helminths such as Strongyloides stercoralis and

hookworms (Ancylostoma duodenale and Necator americanus) is challenging due to

irregular larval and egg output in infected individuals and insensitive conventional

diagnostic procedures. Sensitive novel real-time PCR assays have been developed. Our

study aimed to evaluate the real-time PCR assays as a diagnostic tool for detection of

Strongyloides spp. and hookworms in a random stool sample of 218 asymptomatic

schoolchildren in Cambodia.

Overall prevalence of 17.4% (38/218) and 34.9% (76/218) were determined by real-time

PCR for S. stercoralis and hookworms, respectively. Sensitivity and specificity of

S. stercoralis specific real-time PCR as compared to the combination of Baermann/Koga

Agar as gold standard were 88.9% and 92.7%, respectively. For hookworm specific real-

time PCR a sensitivity of 78.9% and specificity of 78.9% were calculated. Co-infections

were detectable by PCR in 12.8% (28/218) of individuals.

Strongyloides stercoralis real-time PCR applied in asymptomatic cases showed a lower

sensitivity compared to studies undertaken with symptomatic patients with the same

molecular tool, yet it proved to be a valid supplement in the diagnosis of STH infection in

Cambodia.

Keywords: Real-time PCR, Strongyloides stercoralis, hookworm, sensitivity

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1. Introduction

Soil-transmitted helminths (STH) are the cause of the most neglected diseases worldwide (Hotez et al., 2006). STH are prevalent in tropical and sub-tropical regions with high prevalence rates in areas with poor hygiene and sanitation. In particular *S. stercoralis* and the hookworms *Necator americanus* and *Ancylostoma duodenale* represent clinically relevant members of STH. *Strongyloides stercoralis*, an intestinal nematode, infects an estimated 30-100 million people worldwide (Bethony et al., 2006). In most cases it causes asymptomatic infections. However, severe and life-threatening infections known as hyper-infection syndrome may occur (Marcos et al., 2008). Chronic infection with the hookworm may cause anemia and protein deficiency in both adults and children (Stoltzfus et al., 1997).

Traditionally, diagnosis is accomplished by microscopic examination of stool samples and culture methods. However, in light infected individuals microscopy remains insensitive and culture is time consuming (Agrawal et al., 2009). Sensitivity considerably increases when more than one sample is analyzed (Knopp et al., 2008; Marti and Koella, 1993). Use of consecutive samples on up to four days is therefore recommended (Dreyer et al., 1996). Importantly, parasite density in the majority of *S. stercoralis* cases is low, and therefore it would be desirable to increase the sensitivity of diagnostic methods. Diagnosis of hookworm infection is not as challenging due to higher parasite concentrations. Complicating, multiple infections with different STH can occur rendering simultaneous detection of different helminths a useful expansion of diagnostic tests. Novel molecular methods have been described.() Verweij and colleagues recently presented data of a pilot study conducted in Ghana describing a novel *S. stercoralis* real-time polymerase chain reaction (PCR) and a novel hookworm specific real-time PCR (Verweij et al., 2007; Verweij et al., 2009). These assays displayed a high analytical

sensitivity and proved their principle. We validated a described *Strongyloides* spp. and hookworm real-time PCR assay in asymptomatic Cambodian children using a gold standard of microscopy and culture techniques.

2. Materials and Methods

2.1 Ethical consideration

The study was integrated into a larger study on *S. stercoralis* infection and risk factors approved by the ethics committee of the cantons of Basel-Stadt and Basel-Land (EKBB, number 21/09, dated 29 January 2009), Switzerland, and the National Ethics Committee for Health Research (NECHR, number 033, dated 20 March 2009), Ministry of Health, Cambodia. Children with a positive diagnostic result for *S. stercoralis* and hookworm by microscopy were treated with ivermectin (200 µg/kg BW divided in two doses, over two days) and mebendazole (500 mg, single dose), respectively.

2.2 Sample collection

During a field study in the semi-rural province of Kandal, south of Phnom Penh in Cambodia in 2009, 501 stool samples were collected from children from four different schools. The stool samples were transported to the laboratory within 2 hours after collection and subsequently analyzed by Baermann (Baermann, 1917), Koga Agar (Koga et al., 1991) and Kato-Katz (Katz et al., 1972) standard techniques. Kato Katz slides were read starting 30 minutes after preparation, to allow clearing time for the hookworm eggs. Koga Agar slides were read after incubation at 28°C for 48 hours. In addition, an aliquot of stool was taken and immediately frozen at -20°C and shipped to Basel, Switzerland, for further analysis. A total of 218 samples were randomly selected (114; 52.3% male versus 104; 47.3% female, average age 10.6 years) and analyzed by real-time PCR assays.

2.3 DNA extraction from stool samples

For isolation of nucleic acids from stool samples the QIAmp DNA stool kit (Qiagen, Hilden, Germany) was used according to the manufacturer's instructions. 200 mg of stool was used as sample input volume, elution volume was 200 µl.

2.4 Real-time PCR assays

Real-time-PCR assays for *S. stercoralis*, *A. duodenale* and *N. americanus* were done according to Verweij et al. with minor modifications (Verweij et al., 2007; Verweij et al., 2009). In brief, a 25 µl monoplex-reaction contained 5 µl DNA template, 12,5 µl HotStarMastermix (Qiagen), 3mM MgCl₂, 400 nM of each sense and antisense primer (TIB-Molbiol, Berlin, Germany) and 200 nM of the respective probe (TIB-Molbiol). All probes were labelled with the dye FAM and BBQ as quencher. Thermal cycling on an ABI 7500 Realtime PCR System (Applied Biosystems, Weiterstadt, Germany) comprised denaturation at 95°C for 10 min, followed by 40 cycles of 95°C for 15 s and 60°C for 30 s. Fluorescence was measured during the 60°C step.

2.5 Construction of DNA standards

A 101 base pair (bp) real-time amplicon of the 18S rRNA gene of *S. stercoralis*, a 70 bp fragment of the ITS-2 gene of *A. duodenale* and a 100 bp fragment of the ITS-2 gene of *N. americanus* was ligated into pCR2.1 plasmid vector and cloned into *Escherichia coli* using the pCR 2.1 TOPO TA cloning kit (Invitrogen, Karlsruhe, Germany) as described (Panning et al., 2008). Resulting plasmids of the correct length were purified by means of a QIAprep Spin Miniprep kit (Qiagen) and DNA concentration was spectrophotometrically measured. Each of the plasmids was used as a run control at a concentration of 100 DNA copies/PCR reaction. No-template controls were included in each run.

2.6 PCR inhibition control

Inhibition was monitored by spiking the *S. stercoralis* plasmid at a concentration of 100 copies/real-time PCR reaction in each duplicate sample.

2.7 Cross reactivity panel

DNA of the following parasites was tested with each real-time PCR to assess a possible cross reaction: A. duodenale, N. americanus, Trichuris trichiura, Ascaris suum, Trichinella spiralis, Toxocara canis, Wuchereria bancrofti, Loa loa, Dirofilaria immitis, Echinococcus granulosus, Schistosoma mansoni, S. stercoralis.

2.8 Data analysis

Results of parasitological and real-time PCR examinations were double-entered into EpiData version 3.1 (EpiData Association, Odense Denmark) and Microsoft Excel, respectively. After validation, analysis was performed with Stata version 10 (Stata Corporation, Texas USA).

S. stercoralis real-time PCR were compared with results of Baermann and Koga Agar methods (gold standard) of the same stool sample. Results of Kato-Katz technique were regarded as gold standard for the hookworm real-time PCR assays. Standard statistical techniques (including McNemar test) were applied where appropriate. A *p*-value of less than 5% was regarded as statistically significant.

3. Results

All real-time PCR assays were optimized with respect to sensitivity and specificity. 10 DNA plasmid copies per reaction of *A. duodenale, N. americanus, Trichuris trichiura, Ascaris suum, Trichinella spiralis, Toxocara canis, Wuchereria bancrofti, Loa loa, Dirofilaria immitis, Echinococcus granulosus, Schistosoma mansoni, S. stercoralis were detectable on a regular basis by each individual real-time PCR. No unspecific amplification with other related parasites was detectable (data not shown).*

3.1 Validity of S. stercoralis PCR

Overall, real-time PCR detected *Strongyloides* DNA in 38 of 218 (17.4%) samples whereas Baermann and Koga Agar combined detected *S. stercoralis* larvae in 41 of 218 (18.8%) samples (p=0.851). 25 of 38 (65.8%) real-time PCR positive samples were also positive by Baermann and/or Koga Agar methods (Table 1). Real-time PCR failed to demonstrate DNA in two Baermann/Koga Agar positive and in one Baermann positive patient. In 17 Koga Agar positive samples only 4 (23.5%) were also positive by real-time PCR. Stool samples of 164 study participants were negative in all three methods. None of the sample contained substances inhibitory to real-time PCR as demonstrated by amplification of the spiked external inhibition control. Sensitivity and specificity of real-time PCR as compared to the combination of Baermann/Koga Agar methods as gold standard were 88.9% and 92.7%, respectively (Table 1).

Ct-values obtained for the real-time PCR positive samples were compared between Baermann/Koga Agar positive samples, Baermann positive/Koga Agar negative samples, Baermann negative/Koga Agar positive and Baermann/Koga Agar negative samples, respectively. Ct-values were low in samples with parasitological positive results and high

in samples with negative results: Mean and median Ct-values were 31.9, 38.2, 35.0 and 39.1 and 33.0, 39.0, 34.5 and 39.0, respectively. A statistically significant difference (p<0.05) was observed between Baermann/Koga Agar positive and Baermann positive only samples and for Baermann/Koga Agar positive and Baermann/Koga Agar negative samples (p=0.012; Figure 1).

3.2 Validity of hookworm PCR

A total of 76 of 218 (34.9%) samples yielded a positive result by real-time PCR of which 52 (23.9%) samples were positive by the Kato-Katz method (p=0.016). 41 of 52 (78.8%) of the Kato-Katz positive samples were positive by real-time PCR. Interestingly, 35 of 218 (16.1%) samples yielded a positive real-time PCR result without detection of eggs. Of note, 16 of these 35 (45.7%) samples yielded a Kato-Katz positive result when two further stool samples of these children were examined. Finally, 131 of 218 (60.1%) samples demonstrated to be negative by all methods used. The overall analysis yielded a sensitivity of 78.9% and specificity of 78.9% of real-time PCR.

Ct-values of *N. americanus* real-time PCR positive samples were lower in Kato-Katz positive (median Ct-value 36.5) than Kato-Katz negative samples (median Ct-value 37.0), however no statistical difference was recorded (p=0.17). Ct-values of *A. duodenale* real-time PCR positive samples were not done due to its low prevalence.

Real-time PCR was used for the identification of hookworm species. A total of 56 of 76 (73.7%) hookworm real-time PCR positive samples were positive for *N. americanus* specific real-time PCR only. In particular five of 76 (6.6%) were positive for *A. duodenale* specific real-time PCR only and 15 of 76 (19.7%) samples yielded positive results for both *N. americanus* and *A. duodenale* specific real-time PCR.

Among the 56 *N. americanus*-DNA positive samples half (28, 50.0%) were concomitantly positive by microscopy. Only 1 of 5 *A. duodenale*-DNA positive sample (20.0%) was also positive by microscopy.

Twelve of 15 (80.0%) samples with positive results for *N. americanus* and *A. duodenale*-specific real-time PCR were also positive by Kato-Katz (Table 2)

3.3 Samples with multiple detections

Strongyloides spp. and hookworm co-infections were detected in 26 of 218 (11.9%) samples by classical parasitological methods. Twenty-three of these positive samples (88.5%) yielded a real-time PCR positive result. In detail, 13 (50.0%) yielded positive results by both real-time PCR for *S. stercoralis* and hookworms; 2 (7.7%) were positive by *S. stercoralis* real-time PCR only; 8 (30.8%) samples were positive by hookworm real-time PCR only; 3 (11.5%) were negative by both real-time PCR.

Interestingly, 11 of 15 (73.3%) Koga Agar positive samples for *S. stercoralis*, which tested negative by *Strongyloides* real-time PCR yielded a hookworm positive result in the Kato-Katz method. In addition, 4 of 11 (36.4%) hookworm positive samples *Strongyloides* larvae were found in Koga Agar method.

4. Discussion

STH have a significant impact on public health in affected countries (Bethony et al., 2006) (Olsen et al., 2009). Many parts of Southeast Asia are ideal ecological and socioeconomical settings for *S. stercoralis* and hookworms. Reports from Cambodia and Laos PDR showed prevalence rates as high as 20.2% for *S. stercoralis* (Chhakda et al., 2006) and 76.8% for hookworms (Sayasone et al., 2011). Therefore, precise and reliable diagnostic methods remain crucial for public health interventions. Recently, molecular methods have been established for *Strongyloides* spp. and hookworms.(Verweij et al., 2009). Using the *S. stercoralis* positive results of Baermann and/or Koga Agar, our validation of real-time detection of DNA for *Strongyloides* spp. and hookworms revealed a sensitivity and specificity of 61.0% and 92.7% and 78.9% and 78.9%, respectively. These values for sensitivity of *S. stercoralis* detection are similar to the values obtained by Verweij and colleagues (61.1% and 92.4%) (2007, 2009).

When comparing the coprological diagnostic methods with the real-time PCR, different factors have to be considered. First, the sample input volume of culture methods and of real-time PCR assays differs by a factor of 10, which considerably increases sensitivity for the culture method. Increasing input volume for real-time PCR is not an option since stool contains substances which may be inhibitory to real-time PCR. However, real-time PCR inhibition was not an issue in our samples as no significant inhibition of the external amplification control was observed. As reported earlier and confirmed in our study use of consecutive samples increased sensitivity by a factor of 1.5 for *Strongyloides* as well as for the hookworms by conventional methods (Marti and Koella, 1993). It is likely that repeated testing of consecutive sample by real-time PCR will also increase the detection rate. However, this is not an option in resource limited settings but might be considered in in-depth epidemiological studies.

Second, in our study only asymptomatic patients were included. It is reasonable to assume that these patients do not excrete *S. stercoralis* larvae and hookworm eggs in high concentrations. Symptomatic infections are more likely to be of higher parasite burden, thus increasing the number of detectable cases. This is also reflected by the overall high Ct-values in our study indicating rather low parasite burden.

Technically, for maximum sensitivity we used a monoplex real-time PCR approach in contrast to Verweji et al. This further minimizes the risk of false-positive results as multiplex real-time PCR assays are prone to complex oligonucleotide/probe interactions. Analytical sensitivity, as demonstrated on dilution series of plasmid DNA, was comparable to other assays used for parasite detection. However, we can not rule out that infections with low parasite numbers were missed by this approach due to clumped occurrence (Poisson distribution) of the parasites. Interestingly, Wichmann and colleages (2009) have recently demonstrated that detection of free parasite DNA in serum is possible for *Schistosoma* spp. and might constitute a feasible alternative for pathogen detection (Wichmann et al., 2009). Given the life cycles of *Strongyloides* and hookworm infections this might also be a feasible approach and deserves further studies.

Although real-time PCR in particular seems to be a valid technique for diagnosis its application in resource poor settings remains limited due to rather expensive laboratory equipment. Koga Agar culture and/or Baermann technique can be performed at the village level in the field. They have yielded superior performance in this study. However, given the importance of multiple parasitic infections in endemic countries molecular techniques become more advantageous. Especially multiplex real-time PCR methods as proposed by Verweij and colleagues seem to be a valid and cost effective way in diagnosing multiple infections (Verweij et al., 2007). Of note, experience with microscopy and conventional methods is waning in many laboratories in industrialized countries. In

this respect, for *S. stercoralis* sensitivity of real-time PCR in comparison to the Koga Agar method was disappointing. One reason might by a possible misidentification of larvae. Further studies are needed if molecular methods can compensate for this.

To conclude, we could demonstrate that real-time PCR assays can be applied in asymptomatic cases but sensitivity was slightly lower compared to culture and/or microscopy. Further studies are needed to determine the applicability of real-time PCR in the clinical as well as in a research setting.

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Table 1: Comparison of results obtained by Baermann and Koga Agar and real-time PCR methods for the detection of *Strongyloides stercoralis*

Microscopy			PCR		Sensitivity	Specificity
n	Baermann	Koga Agar	Positive	Negative	%	%
18	+	+	16	2	88.9	
6	+	-	5	1	83.3	
17	-	+	4	13	23.5	
41	+/-	+/-	25	16	61.0	
177	-	-	13	164		92.7

Table 2: Comparison of results obtained by Kato-Katz and real-time PCR methods for the detection of hookworms.

n	Microscopy	PCR		Sensitivity	Specificity
	Kato-Katz	Positive	Negative	%	%
52	+	41	11	78.9	
166	-	35	131		78.9

Figure 1: Comparison of Ct-values of *Strongyloides stercoralis* real-time PCR results with Baermann and/or Koga Agar positive and negative samples, respectively. Y-axis denotes Ct-values of real-time PCR.

