

A conventional polymerase chain reaction-based method for the diagnosis of human schistosomiasis in stool samples from individuals in a low-endemicity area

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The aim of this study was to evaluate the efficacy of a polymerase chain reaction (PCR)-based method to detect Schistosoma mansoni DNA in stool samples from individuals living in a low-endemicity area in Brazil. Of the 125 initial stool samples, 80 were ELISA reactive and eggs were identified in 19 of the samples by parasitological examination. For the PCR evaluations, 56 stool samples were selected and divided into five groups. Groups I-IV were scored negative for S. mansoni eggs by parasitological examination. Groups I and II were ELISA reactive, whereas Groups III and IV were ELISA nonreactive. Groups II and III were positive for other intestinal parasites. PCR testing scored eight samples as positive from these four groups. Group V represented the S. mansoni-positive group and it included ELISA-reactive samples that were scored positive for S. mansoni by one or more parasitological examinations (6/19 were positive by Kato-Katz method, 9/17 by saline gradient and 10/13 by Helminex®). PCR scored 13 of these 19 samples as positive for S. mansoni. We conclude that while none of these methods yielded 100% sensitivity, a combination of techniques should be effective for improving the detection of S. mansoni infection in low-endemicity areas.

Key words: *Schistosoma mansoni* - PCR - Kato-Katz - ELISA - saline gradient - Helminex®

Although considered to be among the “neglected tropical diseases” of the world (Molyneux et al. 2005, Hotez et al. 2006), schistosomiasis remains one of the most common parasitic diseases throughout the tropics and subtropics and is a major public health problem in these areas (van der Werf et al. 2003, Steinmann et al. 2006). Laboratory diagnoses are primarily based on the detection of parasite eggs in the stool using the Kato-Katz (KK) method (Katz et al. 1972). This approach is low-cost and easy to perform and provides information on both the prevalence and intensity of the infection. However, a well-known limitation of parasitological (i.e., morphological) examinations is a lack of sensitivity, particularly in low-endemicity areas and in individuals with low parasite loads (Utzinger et al. 2001, Gonçalves et al. 2006, Enk et al. 2008). To overcome this limitation, it is often necessary to analyse larger stool quantities (Bergquist et al. 2009) or to include new diagnostic methods (Cavalcanti et al. 2013).

Other parasitological methods for the detection of schistosomiasis have been described, such as Helminex® or saline gradient methods (Teixeira et al. 2007, Coelho et al. 2009). In the Helminex® method, *Schistosoma mansoni* eggs are concentrated using paramagnetic beads to improve microscopic detection. Somewhat similarly, the saline gradient method is a simple procedure that uses a salt gradient to purify eggs to allow for easier detection with microscopic examination. Serological antibody detection assays, including ELISAs with a variety of antigen preparations (de Noya et al. 2007), have also been used as tools to diagnose schistosomiasis. Although these methods are low-cost, reproducible, objective, rapid and potentially automatable (de Noya et al. 2006), they are poor at distinguishing between past and present infections.

The state of Ceará (CE) in northeastern Brazil has been characterised as a low-prevalence area by the Schistosomiasis Control Program. A study conducted by our group in Planalto do Cajueiro, Maranguape-CE found a disease prevalence of 16% (40 positive individuals out of 250 evaluated) in 2010 using the KK diagnostic method (1 sample, 3 slides). Due to the appreciated lack of sensitivity of this method, we conducted another study in the region using three other parasitological techniques and one serological antibody detection method to better determine the prevalence of this disease. This study demonstrated that the Helminex® technique was more effective than both the KK and saline gradient methods, although certain cases were missed by all three assays (Pinheiro et al. 2012).

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Evaluation of circulating anodic antigen (CAA) and cathodic current antigen (CCA) levels in the serum or urine of infected individuals using ELISA has been proposed as an alternative technique to overcome the problems inherent to antibody detection (Deelder et al. 1989, van Lieshout et al. 1993, van Etten et al. 1994). Studies suggest that the diagnosis of *S. mansoni* through the detection of CCA in urine using reagent strips can provide information on the prevalence of schistosomiasis in high-endemicity areas. Nevertheless, it is of practical importance to assess the efficacy of any new technique for schistosomiasis diagnosis, especially in low-endemicity areas, before applying these techniques for routine diagnostic or epidemiological purposes (Legesse & Erko 2007).

Recent studies have shown that indirect diagnostic tests (e.g., point-of-care circulating cathodic antigen) are valuable alternatives to direct parasitological methods for the diagnosis of *S. mansoni* infections (Coulibaly et al. 2011, Stothard et al. 2011). Although schistosome antigens, such as CCA and CAA, can be detected in the serum and urine of infected individuals (Coulibaly et al. 2013), the sensitivity and specificity of these assays have not been determined in low-endemicity areas (Colley et al. 2013).

Techniques for identifying circulating *S. mansoni* DNA have been widely used and such methods have demonstrated that it is possible to detect infections as early as one day after parasite exposure (Kato-Hayashi et al. 2010, Sulbaran et al. 2010, Gentile et al. 2011). PCR-based diagnoses have been shown to be highly sensitive and specific and should therefore be considered as alternative methods for the diagnosis of *S. mansoni* infections. Such methods have already been used to detect *S. mansoni* DNA in samples of stool (Pontes et al. 2002, Espírito-Santo et al. 2012), blood serum (ten Hove et al. 2008, Haggag & Abdullah 2011), blood plasma (Wichmann et al. 2009) and urine (Sandoval et al. 2006, Enk et al. 2012). Real-time polymerase chain reaction (PCR) has also been used to diagnose schistosomiasis (ten Hove et al. 2008) and this technique can be used as a potential therapeutic cure control method, as has been demonstrated in animal models of experimental infection (Gentile et al. 2011).

In this study, we evaluated the efficacy of a conventional PCR-based method for the diagnosis of human schistosomiasis in individuals from a low-endemicity area in CE.

SUBJECTS, MATERIALS AND METHODS

Study population - The Planalto do Cajueiro is a small community composed of approximately 903 inhabitants distributed across five urban blocks, according to a survey conducted by the city hall. The area is surrounded by two streams containing *Biomphalaria straminea*, the intermediate host of *S. mansoni*. The main source of income for the residents of this town is subsistence agriculture, with the addition of some textile and footwear manufacturing. Due to the poor infrastructure and sanitation, this population is at risk for the transmission of a number of diseases, including schistosomiasis, leishmaniasis, dengue and others.

In total, 125 stool samples were collected from individuals living in Planalto do Cajueiro, Maranguape, a low-endemicity area for schistosomiasis, and were analysed using the parasitological KK - Helm-Test® (hereafter KK) and ELISA.

Based on the results of serological and parasitological techniques (Carneiro et al. 2012, Pinheiro et al. 2012), 56 stool samples were selected from individuals aged between four-72 years, with a mean age of 30.49 years [standard deviation (SD) 17.78] and a median age of 26 years. This sample was divided into the following five groups for PCR analysis: Group I - Ten stool samples from individuals with reactive ELISA results, but without parasitological confirmation of *S. mansoni* eggs by KK. These stool samples were also negative for other parasitic diseases; Group II - Ten stool samples from individuals with reactive ELISA results, but without parasitological confirmation of *S. mansoni* eggs by KK. These stool samples were also positive for other parasitic diseases [Hookworm (5), *Trichuris trichiura* (5), *Strongyloides stercoralis* (2) and *Taenia* spp (1)]; Group III - Seven stool samples from individuals with non-reactive ELISA results and without parasitological confirmation of *S. mansoni* eggs by KK. These stool samples were also positive for other parasitic diseases [Hookworm (1), *T. trichiura* (1), *S. stercoralis* (1), *Entamoeba coli* (2), *Entamoeba histolytica/Entamoeba dispar* complex (3), *Ascaris lumbricoides* (1) and *Enterobius vermicularis* (1)]; Group IV - Ten stool samples from individuals with non-reactive ELISA results, negative results for *S. mansoni* eggs by KK and negative stool tests for other parasitic diseases; Group V - Nineteen stool samples from individuals with reactive ELISA results and who tested positive for the presence of *S. mansoni* eggs by parasitological examination. Stool samples from Group V in sufficient amounts were also tested using two additional parasitological methods: the saline gradient method (n = 17) and Helmintex® (n = 13).

Parasitological methods - The KK Kit [Bio-Manguinhos, Oswaldo Cruz Foundation, Rio de Janeiro (RJ), Brazil] was used according to the manufacturer's specifications and the saline gradient and Helmintex® methods were performed as described in previous studies (Pinheiro et al. 2012). One stool sample was collected from each individual and the KK exams were performed by analysing three slides per sample. For the identification of parasites other than *S. mansoni*, the Hoffman method (Hoffman et al. 1934) was used.

Serological assay for *S. mansoni* antibody detection (IgG ELISA) - Serum samples were tested using an immunoenzymatic reaction to measure the levels of IgG antibodies specific to *S. mansoni*, according to the protocol described by Colley et al. (1977), with a few modifications. A total extract from *S. mansoni* adult worms (5 µg/mL) was used as the antigen (provided by Prof Dr Alfredo Góes, Institute of Biological Sciences, Federal University of Minas Gerais). Peroxidase-labelled sheep anti-human immunoglobulin G antibody (1:1500) (Sigma-Aldrich, St. Louis, MO, USA) was used as the conjugate and ortho-phenylenediamine (1 mg/mL) with H₂O₂

(0.1 mL/mL) in 0.2 M citrate buffer (pH 5.0) was used as the substrate developer. All reactions were performed in a final volume of 100 µL. The cut-off value was determined to be 2 SD greater than the mean optical density value derived from 35 negative control samples tested on the same day as the experimental samples; these samples were obtained from apparently healthy individuals with no detectable parasitic diseases living outside of *S. mansoni* endemic areas.

DNA detection using PCR analyses - Control stool samples - DNA was extracted from *S. mansoni* eggs and from two stool samples from individuals with proven parasitic infections from an endemic area in RJ, which were used as positive controls. One stool sample had greater than 400 eggs/g and the other had fewer than 100 eggs/g. In addition, 10 samples obtained from individuals living in non-endemic areas were used as negative controls.

DNA extraction - Total DNA was extracted from 300 µL of each stool sample using the FastDNA Spin Kit (MP Biomedicals, Vista, CA, USA), according to the manufacturer's instructions. Eggs were disrupted using the FPI20 Fast Prep Cell Disrupter (MP Biomedicals) for 10 sec at a speed of 5.5 ms⁻¹. The DNA was then purified using the QIAquick PCR Purification Kit (QIAGEN, Hilden, Germany) and was stored at -20°C.

PCR - The following PCR parameters were used, according to the protocol described by Pontes et al. (2002), with the exception that the reverse primer sequence was modified by one nucleotide to yield 100% homology to the *S. mansoni* genome (SMFPONTES: 5'-GATCTGAATCCGACC CG-3' and SMRPER: 5'-TAT-TAACGCCACGCTCTCG-3'). The oligonucleotide primer pairs were designed to be specific to a highly repetitive region of the *S. mansoni* genome described by Hamburger et al. (1991). The reactions were performed in a final volume of 50 µL and included 48 µL Master-Mix [45 µL SuperMix (Invitrogen, Carlsbad, CA, USA), 1 µL forward primer F (10 µM), 1 µL reverse primer (10 µM), 1 µL magnesium chloride (1.5 mM)] and 2 µL purified DNA.

Template DNA was amplified using 35 cycles the following: a denaturation phase at 94°C for 30 sec, a hybridisation phase at 55°C for 30 sec and an extension phase at 72°C for 30 sec, followed by a final extension phase at 72°C for 7 min. A GeneAmp PCR System 2400 thermocycler (Applied BioSystems, CA, USA) was used. Amplified products (110 bp) were analysed using a photodocumentation system (Bio-Rad, Hercules, CA, USA) following electrophoresis through 2% agarose gels (Invitrogen) containing ethidium bromide.

Ethical considerations - The study was approved by the Ethical Review Board of the Faculty of Medicine of the Federal University of Ceará (165/09). Informed written consent was obtained from each study participant or, in the case of minors, from their legal guardians. In cases of *S. mansoni* infection, a single dose of praziquantel (60 mg/kg body weight) was administered. Participants with positive faecal examinations for intestinal helminths were treated with albendazole (400 mg/dose).

RESULTS

Stool and serum samples from 125 individuals were initially analysed using KK and ELISA tests. Only 5.6% (7) of the individuals tested positive for the presence of *S. mansoni* eggs in their faeces based on the KK tests. One sample was deemed positive for eggs based on one slide, one sample was deemed positive based on two slides and an additional five samples were deemed positive based on all three slides. Of the 125 serum samples, 64% (80) were reactive in the *S. mansoni* ELISAs, including the seven KK-positive individuals. However, one of the KK-positive individuals did not provide a sufficient amount of stool for further parasitological examinations or PCR analysis.

When comparing the KK and PCR results from 56 *S. mansoni*-positive stool samples, we found that the KK method correctly identified positive samples in six (10.7%) cases and that PCR correctly identified positive samples in 21 (37.5%) cases. Thirteen of the PCR-positive samples were from Group V and the other PCR-positive samples were as follows: four/10 from Group I, two/10 from Group II and one positive sample each from Groups III and IV. The samples from Groups II and III revealed the presence of other parasites and no correlations were observed between these groups and the groups without parasite infection that also had positive PCR results (Table I).

Considering the 19 Group V samples, which were proven positive for *S. mansoni* using parasitological methods other than KK, six/19 were also positive by KK, nine/17 were positive by saline gradient and 10/13 were positive by Helmintex®. PCR analysis detected the presence of parasite DNA in 13 of these 19 samples (68.4%). When comparing the KK and PCR results for the 19 Group V samples, five were scored positive for the presence of *S. mansoni* by both methods, eight were positive only by PCR, one was positive only by KK and five were scored negative by both tests. The proportion correctly detected by PCR was significantly higher ($p = 0.002$) than the proportion detected by the KK method (Table II).

Both the saline gradient method and PCR were performed on 17 samples in Group V. Seven samples were scored positive by both assays, four were positive only by PCR and two were positive only by the saline gradient method. There was no statistically significant difference between these methods ($p = 0.688$) (Table III).

The Helmintex® method and PCR were performed on 13 samples in Group V. Seven samples were scored positive by both assays, three were positive only by Helmintex® and one was positive only by PCR. The proportion of samples correctly detected by PCR was statistically similar ($p = 0.125$) to the proportion detected by Helmintex® (Table IV).

The relationships between the results of the three parasitological tests and the PCR-based analysis are presented in Table V.

To verify the presence of inhibitory substances in the amplification reactions, all six stool samples from individuals without *S. mansoni* eggs and who tested negative by PCR were contaminated with 5 fg *S. mansoni*

DNA. Amplification was not observed in two of these six samples, suggesting the presence of reaction inhibitors in these samples.

DISCUSSION

In this study, we examined the efficacy of a PCR-based method to diagnose the presence of *S. mansoni* eggs in the stool samples of individuals with low-intensity infections. To adequately monitor and control schistosomiasis, more accurate diagnostic methods are needed, which is especially pressing due to the high prevalence and incidence of this disease in some regions (Melo 2006). The control of schistosomiasis is now possible due to the development of a single-dose oral medication that can be easily administered to infected patients. Indeed, the treatment of entire communities can decrease morbidity and reduce egg excretion, although certain individuals will usually remain infected. Despite experimental data suggesting that praziquantel resistance can arise, there is currently no clinical evidence for this phenomenon (Doenhoff et al. 2008, Greenberg 2013). The most reasonable explanation for positive exams following treatment is re-infection due to living in an endemic area. Furthermore, the widespread impacts of chemical-based and other treatment measures have been difficult to determine because the sensitivity of parasitological exams decreases dramatically when egg excretion is

low or absent, particularly in areas of low transmission and/or endemicity. Therefore, more sensitive diagnostic methods are needed to combat schistosomiasis effectively (Gomes et al. 2010, Cavalcanti et al. 2013).

Relatively few studies have addressed the detection of *S. mansoni* DNA in clinical samples (Pontes et al. 2002, 2003, Sandoval et al. 2006, Oliveira et al. 2010). The usefulness of PCR for detecting *S. mansoni* DNA in human faecal samples was first described by Pontes et al. (2002). They demonstrated the detection of *S. mansoni* DNA in stool samples at 2.4 eggs per gram (EPG) of faeces, which indicated a 10-fold greater sensitivity than KK examination; the detection limit was shown to be 1 fg of pure *Schistosoma* spp DNA.

A recent study by Gomes et al. (2010) assessed 206 individuals living in the town of Pedra Preta, state of Minas Gerais (MG), an endemic area for schistosomiasis. Using the KK technique, they determined an *S. mansoni* infection rate of 18% when examining 12 slides for each stool sample. In contrast, using a PCR-ELISA technique that was standardised for the detection of *S. mansoni* DNA, they determined a 30% infection rate. These results demonstrate that diagnostic sensitivity can be increased using this molecular technique.

In the present study, we analysed 56 samples and compared the results from a conventional PCR-based method with results from serological and parasitologi-

TABLE I
Results of Hoffman and polymerase chain reaction (PCR) methods from stool samples of Groups II and III

Groups	Individuals	Hoffman	PCR
II	A1	<i>Strongyloides stercoralis</i> <i>Trichuris trichiura</i>	N
	A2	<i>Ancylostoma</i> spp	N
	A3	<i>T. trichiura</i>	P
	A4	<i>Ancylostoma</i> spp	N
	A5	<i>T. trichiura</i>	P
	A6	<i>Ancylostoma</i> spp	N
	A7	<i>S. stercoralis</i>	N
	A8	<i>Taenia</i> sp.	N
	A9	<i>Ancylostoma</i> spp <i>T. trichiura</i>	N
	A10	<i>Ancylostoma</i> spp <i>T. trichiura</i>	N
III	C1	<i>S. stercoralis</i>	N
	C2	<i>Ancylostoma</i> spp	N
	C3	<i>T. trichiura</i> , <i>Entamoeba histolytica</i> / <i>Entamoeba dispar</i> , <i>Escherichia coli</i>	N
	C4	<i>Enterobius vermicularis</i>	N
	C5	<i>E. coli</i> <i>E. histolytica</i> / <i>dispar</i>	N
	C6	<i>Ascaris lumbricoides</i>	N
	C7	<i>E. histolytica</i> / <i>dispar</i>	P

N: negative; P: positive.

cal tests. Among the 19 samples comprising Group V, *S. mansoni* was detected by the KK technique (reading 3 slides) in 33.9% of cases. In contrast, the PCR-based method was able to correctly score the samples in 68.4% of cases - a more than two-fold increase in sensitivity. However, these results differ from those obtained by Pontes et al. (2003) in a study conducted in Comercinho,

MG. In that study, the authors reported a smaller difference between the results of parasitological examination and their PCR-based technique (30.9% and 38.1%, respectively), detecting only a 0.8-fold increase in sensitivity using PCR. This discrepancy may be explained by the use of three stool samples in the parasitological analysis or because the study was performed in a highly endemic area for *S. mansoni*.

Of the 17 serum samples in negative control groups (Groups III and IV), PCR yielded two false positives. The presence of other parasites did not influence this result, as the number of PCR-positive samples in Group III (with other parasites) was similar to the number detected in Group IV (no other parasites). Although the potential for sample or reaction contamination did exist, the experiments were all conducted in duplicate and all conditions were properly controlled to minimise the possibility of contamination. When analysing the results from Groups II and III (with other parasites), we did not observe any relationships between the parasite species that were present and the results of the PCR analyses (Table I). In particular, only three samples were scored positive by PCR, two of which had *T. trichiura* and one of which had an *E. histolytica/dispar* complex. Indeed, these results may suggest past *S. mansoni* infections, as molecular techniques have been shown to display

TABLE II

Comparison between the reference method Kato-Katz (KK) and polymerase chain reaction (PCR) method for schistosomiasis detection

PCR	KK n (%)		Total n (%)
	P	N	
P	5	8	13 (68.4)
N	1	5	6 (31.6)
Total	6 (31.6)	13 (68.4)	19 (100)

N: negative; P: positive.

TABLE III

Comparison between the polymerase chain reaction (PCR) and saline gradient methods for schistosomiasis detection

PCR	Saline gradient n (%)		Total n (%)
	P	N	
P	7	4	11 (64.7)
N	2	4	6 (35.3)
Total	9 (52.9)	8 (47.1)	17 (100)

N: negative; P: positive.

TABLE IV

Comparison between the polymerase chain reaction (PCR) and Helmintex® methods for schistosomiasis detection

PCR	Helmintex® n (%)		Total n (%)
	P	N	
P	7	1	8 (61.5)
N	3	2	5 (38.5)
Total	10 (76.9)	3 (23.1)	13 (100)

N: negative; P: positive.

TABLE V

Results of Kato-Katz (KK), saline gradient, Helmintex® and polymerase chain reaction (PCR) methods from stool samples of Group V for the presence of *Schistosoma mansoni*

Individuals	Parasitological methods (EPG)			
	KK	Saline gradient	Helmintex®	PCR
E1	N	P (10.0)	N	P
E2	N	N	P (0.10)	N
E3	N	P (2.0)	P (0.13)	P
E4	N	N	P (0.17)	P
E5	P (8.0)	P (10.0)	P (0.47)	P
E6	N	P (6.0)	P (0.13)	P
E7	P (8.0)	NR	P (4.77)	P
E8	N	P (2.0)	N	N
E9	N	P (6.0)	NB	P
E10	N	P (2.0)	NB	P
E11	P (8.0)	NR	NB	P
E12	P (8.0)	N	NB	P
E13	N	N	P (0.20)	P
E14	P (8.0)	N	N	N
E15	N	P (2.0)	NB	N
E16	P (8.0)	N	P (0.20)	P
E17	N	N	P (0.20)	N
E18	N	P (2.0)	NB	P
E19	N	N	P (0.30)	N

EPG: eggs per gram; N: negative; NR: not realised; P: positive.

higher sensitivity when samples are collected from individuals living in endemic areas (Gomes et al. 2010, Oliveira et al. 2010).

In a study conducted among schoolchildren in Egypt by Allam et al. (2009), the researchers sought to demonstrate decreased detection of schistosomiasis in low-endemicity areas. The authors showed that *S. mansoni* eggs were detected by the Percoll technique in 11% of individuals who were scored negative by the KK technique (Dalton et al. 1997) and they also showed that the PCR technique could detect *S. mansoni* DNA in 23% of the KK-negative individuals. Interestingly, although all of the KK-positive cases were also scored positive by the Percoll method, the PCR technique yielded several negative results among the cases deemed positive by the parasitological methods.

Oliveira et al. (2010) performed another study that included stool samples from individuals in the low-endemicity area of Paracambi, RJ. The authors demonstrated that of 16 samples from a group of individuals with parasite loads less than 10 EPG of faeces, PCR yielded positive results in nine of these cases (56%); this study used the same primer pair as in our study. In our study, 13 samples were scored positive by PCR among the 19 Group V samples, yielding a detection rate of 68.4%.

Among the 39 ELISA-reactive samples, 19 were scored positive by PCR. In other words, using a molecular technique, we measured the presence of parasite DNA - indicating active infection - in 48.7% of the seropositive samples. Indeed, ELISA techniques using egg extracts as antigen have been shown to display cross-reactivity when performed in endemic areas (de Noya et al. 2007, Grenfell et al. 2013), which could explain the results for the 14/20 (Groups I and II) ELISA-positive samples from individuals who did not test positive for *S. mansoni* infection by parasitological detection or PCR. Another possibility is that the ELISA technique can detect circulating antibodies from past infections in individuals who do not currently have eggs in their faeces.

PCR failed to detect DNA in six samples that were confirmed to be positive by parasitological examination and ELISA. Of these, only one sample was scored positive by the KK method (8 EPG) and two were scored positive by the saline gradient method (2 EPG); the other three samples were only scored positive by the Helminx[®] assay (0.10, 0.20 and 0.30 EPG). Material from these six samples was tested for the presence of PCR inhibitors, which was indeed present in two of the six samples that were analysed. It is possible that in the other four samples, the quantity of stool used may not have contained the minimum number of parasite eggs required to obtain DNA during the extraction step, perhaps because the samples were taken from individuals from low-endemicity areas that had low parasite loads. It is also possible that variations in egg production and consequent uneven egg distribution within the faeces may have affected the results. Uneven egg distributions are typically minimised through faecal homogenisation; however, the quantity removed from the sample for parasitological examination may have removed the few eggs present in the sample.

The inhibition of Taq polymerase by certain substances found in clinical samples is already widely known and it presents a serious problem for the sensitivity of PCR-based methods. DNA extraction methods that minimise this effect have been described and have improved the efficiency of these techniques. Changes to PCR protocols and the inclusion of certain substances in DNA solutions, such as bovine serum albumin, have also minimised these effects. However, in some samples, the amounts or types of inhibitors present continue to interfere with PCR assays, compromising the sensitivity of the methods. Indeed, some laboratories have introduced structural modifications into Taq polymerase to make it more resistant to inhibitors, which may be helpful in these types of studies.

The use of PCR-based methods for detecting *Schistosoma* spp DNA in epidemiological surveys has also been limited because DNA extraction and PCR amplification can be relatively cumbersome and expensive compared with other methods. However, advances in DNA isolation and PCR techniques now make such assays a viable alternative (Verweij et al. 2004). Based on the literature, PCR appears to be an extraordinarily sensitive and specific tool for diagnosing schistosomiasis and it can be used to detect DNA in samples from individuals with as little as 2.4 EPG in their faecal matter (Hamburger et al. 1991). Furthermore, when the objective is disease control, the cost of a molecular technique should be weighed against the benefits obtained, which may be quite significant. Although none of the methods exhibited 100% sensitivity, PCR in combination with Helminx[®] detected the presence of *S. mansoni* in 11 of the 13 samples we analysed. Based on these results, we conclude that of the use of two or more diagnostic tests in combination may be necessary for accurate diagnosis of this disease.

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