



Evaluation of banked urine samples for the detection of circulating anodic and cathodic antigens in *Schistosoma mekongi* and *S. japonicum* infections: A proof-of-concept study

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ABSTRACT

In Asia, *Schistosoma japonicum* is the predominant schistosome species, while *Schistosoma mekongi* is confined to limited foci in Cambodia and Lao People's Democratic Republic. While the People's Republic of China has been successful in controlling schistosomiasis, the disease remains a major public health issue in other areas. In order to prioritise intervention areas, not only accurate diagnosis is important but also other factors, such as practicality, time-efficiency and cost-effectiveness, since they strongly influence the success of control programmes. To evaluate the highly specific urine-based assays for the schistosome circulating cathodic antigen (CCA) and the circulating anodic antigen (CAA), banked urine samples from Cambodia ($n = 106$) and the Philippines ($n = 43$) were examined by the upconverted phosphor lateral flow (UCP-LF) CAA assay and the point-of-care (POC)-CCA urine assay. Based on 250 μ l urine samples, UCP-LF CAA sensitivity outcomes surpassed a single stool examination by the Kato-Katz technique. The banked urine samples in the current study did not allow the evaluation of larger volumes, which conceivably should deliver considerably higher readings. The sensitivity of a single urine POC-CCA was in the same order as that of a single Kato-Katz thick smear examination, while the sensitivity approached that of triplicate Kato-Katz when a combination of both CAA and CCA assays was used. The promising results from the current proof-of-concept study call for larger investigations that will determine the accuracy of the urine-based CCA and CAA assays for *S. mekongi* and *S. japonicum* diagnosis.

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1. Introduction

An estimated 230 million people are infected with one of the six *Schistosoma* species that cause schistosomiasis, a chronic and debilitating disease of the tropics and subtropics (Colley et al., 2014). More than 90% of the global schistosomiasis burden is concentrated in Africa, where the infections are mainly due to *Schistosoma haematobium* and *Schistosoma mansoni* (Murray et al.,

2012; Utzinger et al., 2009). In Asia, *Schistosoma japonicum* is found in the People's Republic of China (P.R. China), the Philippines (Olveda et al., 2014; Zhou et al., 2010) and Indonesia (Satrija et al., 2015), while the infection is caused by *Schistosoma mekongi* in Cambodia and Lao People's Democratic Republic (Lao PDR) (Muth et al., 2010).

Accurate determination of prevalence and intensity of infection are important for prioritising areas for interventions (Brooker et al., 2009). As long as the prevalence is medium or high (>10%), microscopy works reasonably well (Bergquist et al., 2009; Knopp et al., 2013; Utzinger et al., 2011), although multiple specimens may need to be tested. However, areas characterised by low prevalence (<5%) and low intensity of infection present a challenge, particularly when the overall prevalence falls below 1%, as many positive

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cases are missed if not diagnostics of considerably higher sensitivity are applied (van Dam et al., 2015). Reflecting this need, diagnostics development has concentrated on techniques based on the detection of the pathogen DNA. Such molecular methods have successfully been used for the detection of DNA from various schistosome species, including *S. japonicum* and *S. mekongi* (Clerinx et al., 2013; Wang et al., 2011). Additionally, similar levels of sensitivity have also been achieved with reagents active against the schistosome circulating cathodic antigen (CCA) and the circulating anodic antigen (CAA) in the blood and urine of infected hosts (Colley et al., 2013; Corstjens et al., 2014; van Dam et al., 2014). While the latter tests have been widely applied for the diagnosis of African schistosomiasis (*S. mansoni* and *S. haematobium*) as well as for *S. japonicum* infections (van Lieshout et al., 2000), detection of CAA and CCA in *S. mekongi* infections has not yet been studied. Besides the diagnostic shortcomings of stool microscopy, compliance to provide (multiple) faecal samples is a challenge, specifically in areas of lower prevalence and morbidity (Bergquist, 2013). Using urine as a diagnostic specimen would greatly enhance the possibility of reliable case finding.

While P.R. China has been uniquely successful in reducing prevalence and disease burden of *S. japonicum* infections over the past 60 years (Uttinger et al., 2005; Xu et al., 2015; Zhou et al., 2010), schistosomiasis is still a major problem in the Philippines (Bergquist and Tanner, 2010; Olveda et al., 2014). Presence of *S. mekongi* infections is limited to a few foci in northern Cambodia and southern Lao PDR (Muth et al., 2010).

Ultrasensitive diagnosis is a prerequisite for delineating the spatial distribution of *S. mekongi* and *S. japonicum* infections. Furthermore, sensitivity is imperative in the elimination stage, where the role of low-level infections play a critical role (Knopp et al., 2013; Rollinson et al., 2013; van Dam et al., 2014). This has convincingly been shown in Africa (Colley et al., 2013) and recently in P.R. China (van Dam et al., 2015). In order to extend the proof-of-principle not only by investigating various intensities of *S. japonicum* infection but also exploring the distribution of *S. mekongi* more exactly, sets of banked urine samples were used to evaluate the sensitivity of assays based on circulating antigens in areas where presently available diagnostic tools have proved inadequate.

2. Materials and methods

2.1. Study area

The current study was facilitated by banked urine samples collected some 20 years ago in schistosome-endemic areas of longstanding foci in Cambodia and the Philippines. In brief, urine samples from Cambodia were collected in 1995 from school-aged children in Krakor, a village situated approximately 5 km north of Kratié (northern Cambodia). Krakor is one of 20 villages along the Mekong River in the districts of Kratié and Sambo of Kratié province that had been subjected to *S. mekongi* risk profiling and subsequent control by the Cambodian Ministry of Health (MoH), in collaboration with Médecins Sans Frontières (Stich et al., 1999).

Urine samples from the Philippines were collected as part of a cohort study pertaining to human water contact patterns performed in 1993–1995 in the island of Leyte. The study was originally designed to investigate exposure and infection patterns in combination with cellular immunological mechanisms and evaluation of diagnostics (Acosta et al., 2002a,b; Webster et al., 1997).

Both studies were approved by the responsible authorities and consent was obtained according to the prevailing and acceptable procedures at that time, as detailed in the cited papers (Stich et al., 1999; Acosta et al., 2002b).

2.2. Sample collection and storage

In Cambodia, stool and urine samples were collected from 126 schoolchildren in the primary school of Krakor. Children were invited to provide three faecal samples over a period of five consecutive days. Additionally, on the first examination day, children provided a urine sample. Stool and urine samples were transferred to a central laboratory in Kratié. Faecal samples were subjected to the Kato-Katz technique using a single 41.7 mg thick smear (Katz et al., 1972). *S. mekongi* and other helminth eggs were counted and recorded separately for each sample with the geometric means calculated for egg counts from multiple samples. A small aliquot of urine (5–10 ml) was frozen and kept at -20°C in the hospital laboratory in Kratié prior to transfer to the Department of Parasitology, Leiden University Medical Center (LUMC) in the Netherlands pending further analysis. Complete data records (i.e. at least 1 faecal sample and 1 urine sample examination) were finally available from 106 children.

In the Philippines, the stool samples were transferred to central laboratories and processed the same day according to Katz et al. (1972) using 2 slides of 50 mg faeces per sample. Urine samples were obtained from a sub-cohort from the larger study, frozen and transported to the LUMC for circulating antigen testing. Overall, a panel of 43 *S. japonicum* egg-positive individuals (according to Kato-Katz results) was selected for diagnostic evaluation by the CAA and CCA assays. Additionally, 10 urine samples for negative endemic controls were evaluated for specificity.

2.3. Urine examination

All urine samples were frozen and transferred to LUMC in Leiden, where they were stored at -20°C until detailed follow-up. Overall, 106 urine samples from Cambodia and 43 urine samples from the Philippines were examined by an upconverted phosphor lateral flow (UCP-LF) assay for schistosome CAA. Samples were investigated both by the conventional as well as a concentration assay (UCAA10 and UCAA250 with wet reagents, as described elsewhere (Corstjens et al., 2014). Briefly, samples were diluted with an equal volume of 4% (w/v) tri-chloro-acetic acid (TCA) and centrifuged and assayed as such (UCAA10). Alternatively, the clear supernatants were reduced to amounts of 20–30 μl using a Millipore Ultra-0.5 device (UCAA250). The latter was only applied if, in the UCAA10, CAA levels were below 30 pg/ml (i.e. low or negative). After incubation with the UCP-antibody conjugate, strips were added and the samples allowed to run as described previously (van Dam et al., 2013). Following drying overnight, the strips were scanned for bound UCP using a Packard FluoroCount microtiter plate reader adapted with an IR laser (980 nm) modified to scan LF strips (Niedbala et al., 2001).

All samples were examined in one series, including two standard curves of serial dilutions of *S. mansoni* in a TCA-soluble fraction of adult worm antigen (AWA) containing 3% CAA (AWA-TCA) added to negative urine to allow quantitative CAA determination (pg/ml urine) as described elsewhere (Polman et al., 2000). The assay cut-offs were decided in accordance with Corstjens et al. (2014) using 10 pg/ml urine (UCAA10) and 1 pg/ml urine (UCAA250) as the lower limits of detection (LOD), and 5 pg/ml and 0.5 pg/ml urine as the LOD if the assay would have been performed with multiple samples under ideal laboratory conditions. The region between the two LODs was designated as 'indecisive', indicating that samples were suspected to be positive but that retesting would be required to truly ascertain the score, preferable using a higher sample volume.

Additionally, the commercially available point-of-care (POC)-CCA urine assay (Rapid Medical Diagnostics; Pretoria, South Africa) was performed, according to the manufacturer's instructions. The scoring of the test lines was done similarly to other groups who

have worked with the POC-CCA for the diagnosis of *S. mansoni* in different African settings (Colley et al., 2013; Coulibaly et al., 2011) with negative, 'trace', and three increasing intensity grades of positivity.

2.4. Statistical analysis

Data were entered in an Excel 2010™ spreadsheet. Statistical analysis was done with Excel and SPSS version 20 (IBM Corp.; Armonk, USA). Non-parametric statistics were used for correlation between test results. The sensitivity of the diagnostic tests was calculated and used for indication of test performance. For this reason, we applied a combined infection positive ('gold') standard, defined as being positive if a sample from an individual presented with *Schistosoma* spp. eggs at least once out of three faecal samples and/or found positive in the UCP-LF CAA assay and/or the POC-CCA. This approach implied that, by definition, the specificities of the individual tests were 100%. This was considered valid because of the very high specificities of the circulating antigen assays used (Midzi et al., 2009). Following the approach described previously (Coulibaly et al., 2013), an 'indecisive' or 'trace' category was defined and analysed separately, both for the UCAA and the POC-CCA results.

3. Results

3.1. Banked urine samples for *S. mekongi* diagnosis

Among the 106 children (age range 9–16 years, median 13 year) with complete data records (i.e. at least one stool and one urine sample examined), 62 were found positive for *S. mekongi* eggs in stool (58%); the infection intensity ranged between 0 and 480 eggs per 1 g of stool (EPG) with a median of 4 EPG. On the three consecutive days, the positive rates were 40%, 43% and 44%, respectively (although not all individuals supplied a stool sample each day). In addition to the 62 egg-positive cases, six positives were found by CAA and one by POC-CCA, resulting in a final positive rate of 65%. The sensitivities of the various assays, including the separate analysis with trace/indecisive results being considered as negative or positive versus our pre-defined 'gold standard', are shown in Table 1. There were 58 cases with complete egg count data and out of these the sensitivities of a single stool examination with one Kato-Katz thick smear versus triplicate Kato-Katz ranged between 38% and 45%. The 'true' prevalence, based on a readily available pocket-chart model would have been about 73% (de Vlas et al., 1993).

Three stool examinations with single Kato-Katz thick smears each could detect 90% of the positive individuals. The POC-CCA sensitivity was 39% (including trace positives, the sensitivity was 49%), the UCAA250 sensitivity was 57% (including indecisive positives, the sensitivity was 70%) and POC-CCA and/or CAA 68% (including trace positives, the sensitivity was 80%). In those corresponding to egg counts >100 EPG ($n=18$), one case was missed by POC-CCA (trace positivity only) and two cases by the UCP-LF CAA test (sensitivity 94% and 89%, respectively). However, the combined POC-CCA/CAA results detected all cases with more than 100 EPG. Based on the 10 endemic negatives, the UCP-LF CAA high specificity cut-off was 0.23 pg/ml.

Fig. 1 depicts the correlations between the CAA concentrations and egg counts (panel A, Spearman's $\rho=0.68$, $p<0.001$) and POC-CCA intensity scores (panel B, Spearman's $\rho=0.60$, $p<0.001$) as well as between egg counts and POC-CCA (panel C, Spearman's $\rho=0.64$, $p<0.001$). Panel A shows that the eight samples with CAA concentrations in the indecisive range from 0.5 (dotted line in Fig. 1) to 1 pg/ml were confirmed by positive egg counts. The data

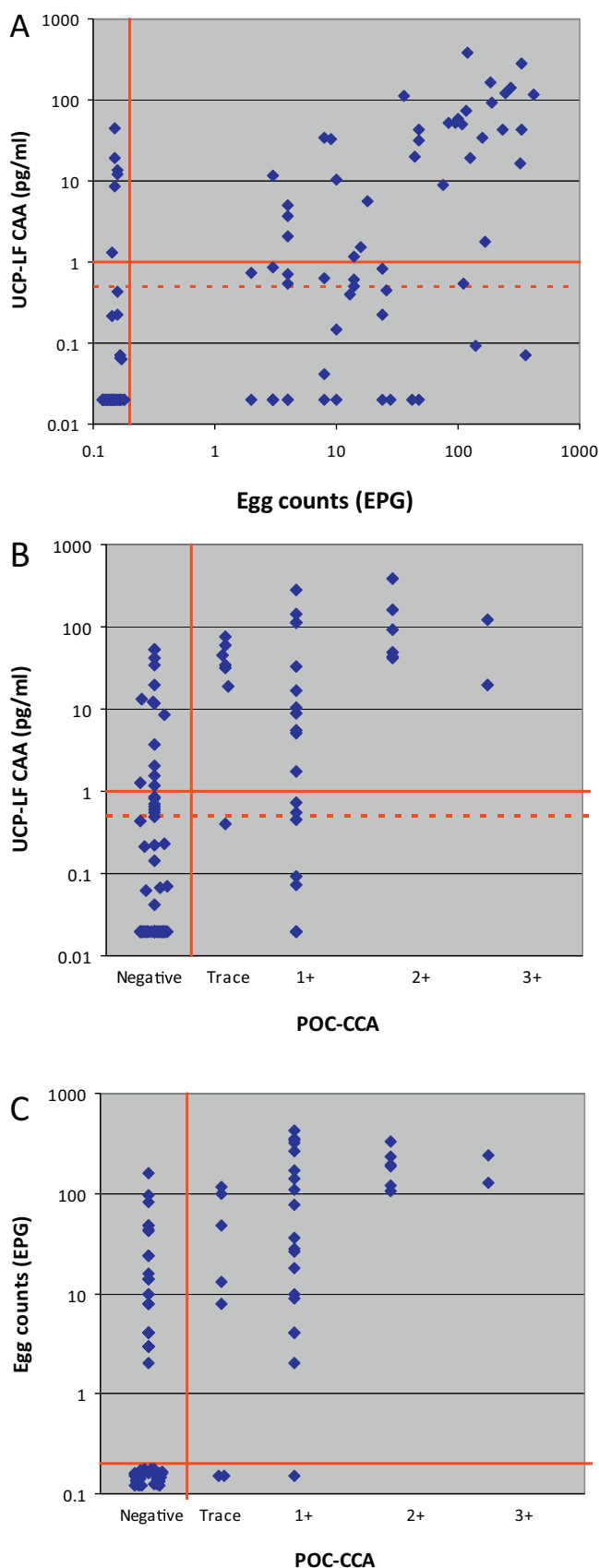


Fig. 1. Correlations of *S. mekongi* urine CAA levels (pg/ml) determined by the UCP-LF CAA test with (panel A) stool egg counts (mean EPG of 3 slides, 1 slide per stool), and with (panel B) POC-CCA intensity scores; panel C shows correlations between EPG and POC-CCA. The solid line represents the high specificity cut-off for the UCP-LF CAA assay, while the region between the dotted line and the solid line is classified as 'indecisive'.

(measured by Kato-Katz stool examination) in most endemic villages are below 5%, which differs from the endemic villages in Lao PDR where infection rates of 20% and more are frequently observed (Lovis et al., 2012; Sayasone et al., 2012). Furthermore, co-infection with other trematode species might influence test performance. In particular, the fish-borne trematode *Opisthorchis viverrini* is highly endemic in *S. mekongi*-endemic areas in Lao PDR and might therefore influence test outcomes (Lovis et al., 2012; Sayasone et al., 2011, 2012).

In conclusion, an adequate diagnostic approach is required with 'surveillance and response' approaches taking over from extensive monitoring and evaluation (Tambo et al., 2014). Highly sensitive diagnostic tools must be in place to confirm elimination or the potential beginning of resurgence of infection (Bergquist et al., 2015). These tools should follow the ASSURED characteristics as defined by the World Health Organization (i.e. affordable, sensitive, specific, user-friendly, rapid and robust, equipment-free, and deliverable to users). This proof-of-concept study, although performed on a limited number of samples collected some 20 years ago, underscores that CCA- and CAA-based assays would hold high expectations as promising candidates fulfilling these needs and requirements.

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