

## Review Article

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# Advantages and limitations of diagnostic procedures for zoonotic helminth infections in feces

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## Abstract

Helminthiasis is a common infection in both humans and other animal populations and negatively affects the health of the host, causing a range of morbidity and even mortality, especially in young people and those with weakened immune systems. A variety of diagnostic procedures with a number of modifications are available to identify the sources of infection and to assess the epidemiological situation, the effectiveness of parasite control programs, anthelmintic treatment, and the prevention of drug resistance. These can be simple and cheap methods – for example, stool smear examination – or modern highly sensitive methods – for example, PCR analysis. This review attempts to summarize the advantages and limitations of each of these frequently used methods.

## Introduction

Adequate diagnostic techniques are necessary for the control, prevention, and treatment of common helminth infections in human and other animal populations (Crompton 2000; Utzinger and Keiser 2004). Fecal examination methods are one of the most accessible and simple techniques used for recovery of helminth eggs, larvae, and proglottids. The different fecal examination methods are broadly based on three principles such as floatation of helminth eggs in solutions with high specific gravity, sedimentation of fecal debris, and direct fecal examination. These techniques come with various modifications to increase the sensitivity of the assay such as centrifugation, a combination of different solutions, and straining (Crompton 2000). However, microscopy-based fecal examination tests have low sensitivity and low specificity due to the low concentration of helminth eggs and larvae in feces that can lead to false negative results (Thienpont *et al.* 1979). Moreover, some helminth eggs and larvae such as *Taenia* spp. eggs, Opisthorchiidae eggs, and some lungworm larvae cannot be differentiated by microscopy only (Annoscia *et al.* 2014; Jimenez *et al.* 2010; Rim 2005). Thus, alternative methods of detection of intestinal parasites must be developed. The detection of parasitic antigen in fecal samples and antibody in serum are highly sensitive and specific methods for the detection of individual parasitic infections, though both methods have their own limitations. Recently, a range of DNA-based methods for the detection of intestinal parasites in biological materials, such as feces, blood, and tissues, has been developed as a valuable diagnostic procedure for intestinal parasites. Single, duplex, and multiplex real-time PCR assays have been developed for the detection of many human and animal parasites (Frickmann *et al.* 2015; Ishida *et al.* 2003; Pilotte *et al.* 2016).

This review presents the advantages and limitations of fecal examination methods, ELISA, and PCR diagnostic methods utilized in the setting of helminthiasis detection.

## Fecal examination methods

Fecal examination methods are the most simple and economical procedures for the detection of helminth infection. A variety of modifications are used, each with its own advantages and limitations (Thienpont *et al.* 1979). The most frequently applied procedures include direct fecal smear, floatation, and sedimentation methods. The selection of an appropriate method is highly dependent on the helminth egg's specific gravity. In most cases, this varies between 1.05 and 1.2 g/L. Roundworm and hookworm eggs have a lower specific gravity of around 1.05–1.1 g/L, whereas whipworm and tapeworm eggs are comparatively heavier and have a specific gravity of around 1.13–1.2 g/L. Fluke eggs have a specific gravity of more than 1.2 g/L and precipitate in

most flotation solutions, whose specific gravities usually fall between 1.18 and 2 g/L (David and Lindquist 1982; Harnnoi *et al.* 1998).

### Direct fecal smear

Grassi and Parona first employed a direct fecal smear in 1878 as a diagnostic tool for the detection hookworm infection in humans (Rockefeller 1922). For many years, this method was used for the identification of different parasites with different modifications. For the simple direct fecal smear examination, a small sample of feces is placed on a glass slide, mixed with a drop or two of saline, spread thinly over the slide, and then covered with a glass coverslip. The smear needs to be thin enough to read newsprint through them. Later, Faust (1929) compared the sensitivity of the fecal smear technique without staining and with staining with hematoxylin. He observed that the staining of the fecal smear did not significantly improve diagnostic sensitivity compared to a direct fecal smear without staining. Thick smear technique was introduced by Kato and Miura in 1954 for fecal research (Kato and Miura 1954). They placed 60–70 mg fecal sample on the glass slide, covered it with a cellophane strip soaked with glycerol-malachite green, pressed it with a rubber stopper or finger, and kept it at a room temperature for about 30–60 min. Although the thick smear technique proved more sensitive than a direct fecal smear with a cover glass, it exhibited a low sensitivity for hookworm eggs, which are present in low density in fecal samples and often are destroyed during the long preparation period because hookworm eggs have a delicate consistency, quickly clear with glycerin, and after 30–60 minutes, they cannot be seen (Santos *et al.* 2005; <https://iris.who.int/bitstream/handle/10665/324883/9789240014497-rus.pdf>).

Katz adopted and modified this technique for use in routine epidemiological studies, resulting in its recognition by WHO for the quantitative and qualitative diagnosis of intestinal helminth infections such as *Ascaris lumbricoides*, hookworms, and *Schistosoma mansoni* (Katz *et al.* 1974; Santos *et al.* 2005). For the diagnosis of *S. mansoni*, this method may have low sensitivity. One of the main parameters affecting the sensitivity of the analysis is the minimum infection intensity, which depends on the egg output of one worm. One pair of *S. mansoni* produces about 100 eggs per day, so feces contain many eggs only in the case of intense invasion. Given the small amount of feces analyzed, detection of mild *S. mansoni* infections with only a few female worms is almost impossible (Barenbold *et al.* 2017).

However, fecal smear examination is the simplest and most economical procedure used in laboratories worldwide. The technique has been adopted for the detection of a wide range of parasites, including roundworms, whipworms, hookworms, flukes, and tapeworms. Among the numerous modifications of the direct fecal smear technique, Kato-Katz is recognized as a 'gold standard' and is widely used by both human and veterinary healthcare practitioners (Hong *et al.* 2003).

However, due to the small amount of stool being analyzed (41.7–70 mg), the sensitivity of the method suffers and can give false negative results due to low concentrations of helminth eggs in the stool sample or may appear highly clustered. Sensitivity can be increased by examining multiple Kato-Katz smears prepared from the same stool sample or from multiple samples from the same host or employing a combination of the Kato-Katz smear with a direct fecal smear (Glinz *et al.* 2010; Komiya and Kobayashi 1966). In addition, this method requires rapid sample processing – optimally

within 20–30 minutes after slide preparation. The acceptable processing time for samples is within 24 hours. When stool samples are stored overnight, regardless of temperature, the number of parasite eggs decreases, which makes their detection difficult and does not allow reliably determining the intensity of infection (Bosh *et al.* 2021).

### Concentration method

Microscopic examination of the sedimented fecal matter was one of the first modifications of the fecal smear examination technique (Koutz 1941). Sedimentation relied on the concentration of helminth eggs, larvae, and protozoan cysts at the bottom of a tube to detect parasites or eggs occurring at low densities in feces.

Since 1948, various modifications and simplified improvements have been employed by a number of researchers in clinical laboratories (Blagg *et al.* 1955; Knight *et al.* 1976; Manser *et al.* 2016). The method uses several pieces of apparatus, which must be washed after use with each sample and includes several safety precautions since it involves the use of formalin (an irritant) and ether, which is flammable. These kits are tightly sealed to reduce the hazards of formalin. The major modification of sedimentation method is using the ethyl acetate, a more stable and less flammable alternative, rather than ether (Truant *et al.* 1981; Young *et al.* 1979).

The TF-Test® is another modification of the concentration method for use in routine parasitological surveys. This complex method combines multiple sampling (on three consecutive days), a fixative (SAF-sodium acetate-acetic acid formalin), a concentration method, and a permanent stain (Chlorazol Black dye) (Van Gool *et al.* 2003).

A highly competitive method for field studies is MIF (merthiolate–iodine–formalin). It is a concentration-based method that requires a centrifuge. This technique uses the MIF solution (50 mL formaldehyde at 37%, 10 mL glycerin at 87%, filled to 1 L with distilled water as stock solution I) as a preservative and staining (with 2 g potassium iodide in 10 mL distilled water as stock solution II). Ether is added to dissolve the fecal fats. The MIF method showed higher sensitivity for hookworms, while for *T. trichiura* and *A. lumbricoides*, Kato-Katz performed better. However, MIF detected *Strongyloides stercoralis* for which Kato-Katz method is not specific. Another advantage of this technique is that it allows for the preservation of fecal samples for a long time. Overall, the MIF method is simple and inexpensive, and is suitable for diagnosis and assessment of infection intensity in the field (Incani *et al.* 2016).

### Flotation techniques

The principle of fecal flotation is based on the ability of a solution to allow less dense material (including parasites) to rise to the top (Ballweber *et al.* 2014; Dryden *et al.* 2010).

The specific gravity (SG) of the flotation solution must be higher than the SG of helminth eggs, so commonly used flotation solutions have an SG of around 1.18–1.33 g/L. Common flotation solutions are made by adding a measured amount of sugar (sucrose or dextrose) or salts such as sodium chloride, sodium nitrate, magnesium sulfate, or zinc sulfate to a specific amount of water to achieve the desired SG. Ideally, all helminth eggs would float without loss of morphological structures, while fecal debris would sink. Despite their low cost and simplicity, the flotation methods are highly influenced by several factors, such as

floatation time, tube filling, and precise removal of the coverslip. Flotation methods have many modifications, ranging from simple passive flotation in solution with high SG, the addition of a centrifugation step, and using a chambered slide or special devices that utilize flat-bottomed vials in which the feces/flotation fluid mixture is placed (Ovassay Plus Fecalizer) (Ballweber *et al.* 2014).

The presence of a large amount of fecal debris is a major obstacle for the fecal flotation technique. One modification to alleviate this problem was the addition of a centrifugation step to remove large floating debris. This technique was first introduced by Lane (1924) and later modified by several others (Dryden *et al.* 2010; Zajac *et al.* 2002). The centrifugation time varies from 5 to 20 min depending on the characteristics of the centrifuge used such as the size of the rotor and the relative centrifugation force. For example, Egwang and Slocombe (1982) showed that centrifuging for 4 or 5 min at 264 ×g provided statistically significantly better egg recoveries than shorter (1 min) or longer (20 min) durations. Dryden *et al.* (2010) also observed that a 5-min centrifugation time at 280 ×g showed a significantly higher fecal egg count when compared with the passive flotation method. Another modification of flotation techniques is a fecal egg count method based on microscopy of an aliquot of fecal suspension from a known volume of a fecal sample. This method allows the investigator to express the number of parasitic elements (eggs, larvae) in the fecal sample in terms of eggs per gram of feces and can be used to measure the distribution of infections for epidemiological surveys to detect the presence or build-up of anthelmintic resistance, and to quantify the efficacy of anthelmintic treatment (Dryden *et al.* 2010).

Gordon and Whitlock (1939) proposed the first egg count method using a chambered slide while working with sheep feces, which later came to be known as the McMaster method (Gordon and Whitlock 1939). This procedure used 2 g of feces mixed with 30 mL of flotation solution (Sheater's sugar or saturated sodium chloride), which was then shaken by hand to make a slurry. An aliquot of 1 mL was drawn from the center of the tube and added to three areas of the McMaster counting chamber. Since then, several modifications, including variations in the ratio of feces to fluid, centrifugation time, number of chambers counted, and area of slide counted, have been described. The sensitivity of this technique depends on the weight of the feces examined and the dilution ratio (g of feces/mL of water). In a study by Vadlejch *et al.* (2011), three modifications of the McMaster technique, like Wetzel, Zajíček, and concentration modification according to Roepstorff and Nansen, were compared; the concentration modification according to Roepstorff and Nansen used 4 g of the feces examined and determined that a low dilution ratio (1:14) was more sensitive than when a lower weight of feces (1–2 g) or a higher dilution ratio (1:30) was used (Blagg *et al.* 1955).

A modern modification of the McMaster Technique is FLOTAC®, introduced by Cringoli (2006), which incorporates a centrifugation-enhanced flotation method in a chambered device with a detection limit of 1 or 2 EPG (eggs per gram) (Cringoli 2006; Utzinger *et al.* 2008). A further modification of FLOTAC® was proposed by Barda *et al.* (2013) called Mini-FLOTAC. In this method, a simple device Mini-FLOTAC apparatus comprises two physical components – namely, the base reading disc and two accessories, the key and the microscope adaptor (Barda *et al.* 2014). There are two 1-mL flotation chambers, designed for optimal examination of fecal sample suspensions in a total volume of 2 mL. A major advantage of this method is the lack of a centrifugation step.

## Immunological and molecular analyses

Development of immunoassays and PCR kits provides new highly sensitive and specific methods for the detection of parasitic infections. Coproantigen ELISA assays and PCR tests have been developed for the detection of major parasites of humans and animals in feces.

### Coproantigen detection

Highly sensitive and specific immunological assays employing antibodies have been developed to identify helminth parasite antigens released in the host feces. The helminth coproantigens are detected by enzyme-linked immunosorbent assays (ELISAs). One major advantage of such coproantigen ELISAs over serum antibody assays is that this method indicates only the current infection and avoids handling of serum that may contain parasitic products from previous infestations. The first such assay using agar gel diffusion was developed for the detection of *Echinococcus granulosus* (Babos and Nemeth 1962). Other coproantigen assays have been developed for many human and animal cestode parasites to identify of major flatworm infections such as *Opisthorchis viverrini*, *Fasciola hepatica*, *F. gigantica*, and *Echinostoma capronii* (Abdel-Rahman *et al.* 1998; Estuningsih *et al.* 2004; Fraser and Craig 1997; Mezo *et al.* 2004; Watwiengkam *et al.* 2013).

Recently, an ELISA developed with monoclonal mouse IgG antibodies against *Clonorchis sinensis* was used for its detection in experimentally infected rats (Rahman *et al.* 2012). However, only a handful of coproantigen assays have been reported for nematodes, including assays for the detection of a number of species like *Haemonchus contortus* (Ellis *et al.* 1993), *Ascaris suum* (Schnierer 1995), *Strongyloides ratti* (Nageswaran *et al.* 1994), and *Heligmosomoides polygyrus* (Johnson *et al.* 1996). These tests have been shown to be relatively sensitive and species-specific.

### PCR based detection methods

The first polymerase chain reaction (PCR) based assay for the identification of parasite DNA from eggs in feces was demonstrated by Flisser *et al.* (1988, 1990) and then by Bretagne *et al.* (1993). Common techniques such as conventional PCR, multiplex PCR, and real-time PCR are used for detecting parasite DNA in host feces (Bergquist *et al.* 2009). Conventional PCR involves the amplification of a sequence of target DNA using a primer pair to detect foreign DNA belonging to any parasite stage in either tissue, blood, or urine. The conventional PCR technique is slightly modified to perform a multiplex PCR that allows amplification of multiple target DNA sequences simultaneously (Gordon *et al.* 2011; Toze 1999). Multiplex PCR can also be used for the detection of multiple parasites species from the same individual's fecal sample or blood.

The development of the real-time PCR introduced researchers to a powerful tool to study gene expression profiles as a function of the relative abundance of the target gene in the sample. Real-time or quantitative PCR (qPCR) does this by measuring the fluorescence released as a by-product of the PCR reaction and presenting it as a graph of fluorescence intensity relative to time. Thus, samples with a higher concentration of the target DNA show a peak in fluorescence at an earlier time point during the PCR (Frickmann *et al.* 2015; Heid *et al.* 1996). PCR, however, has a number of limitations, such as the risk of contamination, false positive results due to the presence of naked nucleic acids and non-viable microorganism, and difficulty in quantification in water and wastewater (Toze 1999).

Modern diagnostic molecular-based techniques includes DNA sequencing, DNA barcoding and Loop-Mediated Isothermal Amplification (LAMP).

DNA barcoding was established as a rapid, powerful method for taxonomic research. DNA barcoding uses molecular data for identification and differentiation of species. PCR uses for amplification of short DNA fragments which compared to a public DNA databases for the possible sequences matches. *Nuclear and mitochondrial genes, such as 18S rRNA, 16S rRNA, ITS regions, and cox1, are used for DNA barcoding of eukaryotic organisms.*

The cytochrome *c* oxidase (*cox1*) is the most slowly evolving gene, acknowledged as the 'gold standard' for DNA barcoding of eukaryotic organisms. For the determination of partial *cox1* sequences of Platyhelminthes, the primer set of JB3 (5'-TTT TTT GGG CAT CCT GAG GTT TAT-3') and JB4.5 (5'-TAA AGA AAG AAC ATA ATG AAA ATG-3') (Bowles *et al.* 1993) was widely used for investigating the inter- and intra-species variations of trematodes and cestodes. A PCR, followed by sequencing, on the mitochondrial genes *cox1* and *nad1* with primers published by Bowles and co-authors was the standard test at our laboratory for addressing both the *Taenia* sp. determination as the detection of *Echinococcus* sp. with subsequent species determination (Bowles *et al.* 1992; Bowles and McManus 1993; Gasser *et al.* 1999). The polymorphisms in these genes, which are so useful for typing, also interfere with the PCR. The annealing site for these primers is polymorphic as well. With several specimens of cestodes, the PCRs – most often the *nad1* PCR – failed to amplify one of the two targets. Many different PCR based approaches to detect and type various Cestodes have been described (Abbasi *et al.* 2003; Al-Sabi & Kapel 2011; Bart *et al.* 2004; Boubaker *et al.* 2013; Dinkel *et al.* 2004; Gonzalez *et al.* 2002; Jeon *et al.* 2009; Knapp *et al.* 2014; Maurelli *et al.* 2009; Mayta *et al.* 2008; Schneider *et al.* 2008; Siles-Lucas and Gottstein 2001; Stefanic *et al.* 2004; Trachsel *et al.* 2007; van der Giessen *et al.* 1999; von Nickisch-Rosenegk *et al.* 1999; Yamasaki *et al.* 2004). Most of these studies, however, use different PCRs for the detection of different cestodes. The PCRs are not specifically designed for human diagnostics. For routine diagnostics, a single test that is able to detect and type a range of cestodes is preferable. A widely published PCR target is the 12SrRNA gene, hereafter referred to as 12S. A number of different primers have been designed for the amplification of this gene because different research groups had different purposes (Boufana *et al.* 2008; Dinkel *et al.* 2004; Stefanic *et al.* 2004; Trachsel *et al.* 2007; van der Giessen *et al.* 1999). The primers on 12S were designed to amplify a broad range of cestodes. The reverse primer is essentially the same as one of the primers published by Trachsel and co-authors for *Echinococcus* sp. and *Taenia* sp. (Trachsel *et al.* 2007). In addition to the cestode 12S primers, the *Taenia* sp. specific primers designed by Trachsel and co-authors were also used.

Additionally, the primers for the *nad5* gene specifically for amplification of *Echinococcus* sp. were designed. Both the 12S and *nad5* genes are mitochondrial, which ensures high sensitivity. The genes are also genetically highly variable, which is especially useful for species determination within the genus *Echinococcus* sp. We investigated the sensitivity and specificity of the PCRs on these genes. Furthermore, material from suspected echinococcosis patients was tested with *cox1*, *nad1*, 12S, and *nad5* PCRs to compare sensitivity in clinical samples. To compare the targets for their use in species determination, we tested a variety of different *Echinococcus* sp. samples.

DNA barcoding has emerged as a powerful, rapid, molecular-based method with significant contributions to both taxonomic and

biodiversity research (Hajibabaei *et al.* 2007; Hebert *et al.* 2003; Pereira *et al.* 2013; Weigt *et al.* 2012). Organisms can be accurately identified at the species-level by this process, which uses polymerase chain reaction (PCR) to amplify short DNA fragments corresponding to standardized regions of the genome with associated discriminatory sequence variations that are then subject to DNA sequencing. These sequences are then compared to a public DNA database of all the possible sequence matches or to a defined reference library populated with DNA barcoding sequences of voucher specimens (e.g., BOLD Systems v3; <http://www.boldsystems.org/>).

Nuclear (e.g., 16S rRNA, 18S rRNA, ITS regions), mitochondrial (e.g., cytochrome *b*, mitochondrial control regions), and some chloroplastic genes have all been used in barcoding eukaryotic organisms (Patwardhan *et al.* 2014). The more slowly evolving gene of the mitochondrial genome, cytochrome *c* oxidase I (*cox1*), is acknowledged as the 'gold standard' for species identification and DNA barcoding of animals (Hebert *et al.* 2003). However, for parasite identification, discovery, and diversity research, the nuclear 18S small subunit (SSU) rRNA gene has been more commonly used (e.g., Harris & Rogers 2011; Maia *et al.* 2012; Netherlands *et al.* 2014; Perkins & Keller 2001; Wozniak *et al.* 1994).

Notomi and co-authors, in 2000, described a new method of DNA amplification called Loop-Mediated Isothermal Amplification (LAMP) that has since been used for identification of major human and animal parasites and demonstrated comparable sensitivity and specificity to that of qPCR (Notomi *et al.* 2000).

LAMP employs a DNA polymerase with high strand displacement activity and a set of four primers that recognize six distinct sequences of the target DNA, giving it very high specificity. Although relatively new in the category of diagnostic techniques, the LAMP has distinct advantages over both conventional PCR and qPCR, such as high specificity of DNA amplification in the presence of non-target DNA, cost-effective amplification at isothermal conditions since no thermal cyclers is necessary, and simple detection of amplification products using fluorescent dyes. However, the lack of data with this technique hinders the development of LAMP kits and their use in mass screening of helminth infections (Biswal 2016).

### Which method is the best?

The sensitivity and specificity of diagnostic tests significantly influence the results of epidemiological surveys for helminth infections and the particular efficacy of anthelmintic treatment or drug resistance. The sensitivity of the diagnostic tool used is also important in helminth control programs; a low sensitivity leads to false negative results that in turn may lead to premature treatment cessation. Among the different diagnostic techniques described above, the fecal smear examination methods are the most frequently used in epidemiological surveys. The World Health Organization recommends the Kato-Katz method of duplicate slides for the detection of soil-transmitted helminth infections (*Ascaris lumbricoides*, *Trichuris trichiura*, *Ancylostoma duodenale*). The Faust method with zinc sulfate solution is one of the most common methods used in veterinary studies (Faust *et al.* 1929). Other methods such as sedimentation, McMaster, and FLOTAC are also used in parasitological studies (Biswal 2016). Comparative data on fecal examination methods are controversial since each procedure varies due to helminth species and study methodology. Moreover, the similar examination methods can have different sensitivities for the same parasite species in different studies. However, a summary of recently published data on the comparison of different fecal

examination methods FLOTAC and mini-FLOTAC shows higher sensitivity for most parasites when compared with the gold-standard Kato-Katz method, McMaster egg count method, and sedimentation techniques. PCR-based methods have a higher sensitivity than traditional coproscopical examination techniques, ranging from 78.9% to 100% (Knopp *et al.* 2009; Taniuchi *et al.* 2011) for different parasitic species. Coproantigen ELISA tests are also highly sensitive methods, especially for identification of tapeworm infections – for example, sensitivity of the coproantigen test for the detection of *Taenia pisiformis* is 87.5% (Schär *et al.* 2013), and recently developed coproantigen assays for detection of fluke worm infections such as *Opisthorchis* and *Clonorchis* range in sensitivity from 93.3% to 100% (Craig *et al.* 1995; Teimoori *et al.* 2017).

In terms of specificity, coproantigen tests rank lower than PCR tests, ranging from 54.2% to 100% (Rahman 2012; Watwiengkam 2013). In a study by Nageswaran *et al.* (1994), the coproantigen test for detection of *S.ratti* had cross-reactions with *Necator americanus* and *S. muris* (Nageswaran *et al.* 1994).

One major advantage of PCR analyses over ELISAs is a significant decrease in the detection limits. For example, positive results have been obtained with a concentration 0.11–0.35 ng of parasite DNA (Ai *et al.* 2010), whereas ELISA coproantigen tests typically require around 0.3–0.6 ng of parasite antigens (Mezo *et al.* 2004).

Coproscopical examination methods are still favored by diagnostic clinics owing to their simple and economical set-ups. Without the use of costly equipment like thermal cyclers or spectrophotometers, coproscopical examinations can be performed in small laboratories or field studies. In order to circumvent the possibility of false negative results due to low concentrations of helminth eggs, low detection limit techniques like the Wisconsin and FLOTAC methods (detection limit 1 EPG), the McMaster method (10 EPG), and Kato-Katz (24 EPG) are recommended (Glinz *et al.* 2010; Knopp *et al.* 2009; Levecke *et al.* 2012). The sensitivity can be augmented by multiple stool examinations (Bogoch *et al.* 2013).

The choice of an appropriate examination method is also dependent on the study objectives, including host species, estimation of overall parasite prevalence or individual parasite infection, the intensity of parasite infection, and efficacy of anthelmintic treatment or drug resistance. Moreover, recently published reports claim an absence of the so-called ‘gold standard’ of helminth infections diagnostics (Nikolay *et al.* 2014; Tarafder *et al.* 2010). Thus, different methods can be employed in parasitological studies.

## Conclusions

For the detection of helminth infestations, different methods, based on the relative ease of use, the purpose of the study, the availability of laboratory equipment, and the practitioner’s qualifications, must be tested. Attaining an adequate accuracy using each methodology should be the goal of selecting a particular examination method.

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