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Methods for quantification of soil-transmitted helminths in environmental media: current techniques and recent advances

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Abstract

Limiting the environmental transmission of soil-transmitted helminths (STH), which infect 1.5 billion people worldwide, will require sensitive, reliable, and cost effective methods to detect and quantify STH in the environment. We review the state of the art of STH quantification in soil, biosolids, water, produce, and vegetation with respect to four major methodological issues: environmental sampling; recovery of STH from environmental matrices; quantification of recovered STH; and viability assessment of STH ova. We conclude that methods for sampling and recovering STH require substantial advances to provide reliable measurements for STH control. Recent innovations in the use of automated image identification and developments in molecular genetic assays offer considerable promise for improving quantification and viability assessment.

Environmental Detection of Soil-Transmitted Helminths for Public Health

Soil-transmitted helminths (STH), including hookworms (*Necator americanus* and *Ancylostoma duodenale*), roundworms (*Ascaris spp.*), and whipworms (*Trichuris trichiura*), impose a substantial burden of disease worldwide. An estimated 819 million people are infected with *Ascaris lumbricoides*, 465 million with *T. trichiura*, and 439 million with hookworms, globally [1]. In addition to a range of acute sequelae, including gastroenteritis, anemia, and intestinal obstruction, chronic helminth infection has been associated with poor physical and cognitive development, and is thought to reinforce poverty [2].

The World Health Organization has launched efforts to reduce the health and economic burden attributable to STH through programs that primarily focus on mass drug administration (MDA) of antihelminthic medicines to school-aged children [3]. These

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campaigns have succeeded in reducing global morbidity due to helminth infections. However MDA is unlikely to break cycles of STH transmission unless coupled with environmental measures to interrupt acquisition of new infections [4-6]. This is in part because STH ova are extremely resistant to environmental stressors, and may survive for years in soils [7]. This hardiness, combined with low infectious doses and high rates of excretion, contributes to the maintenance of STH transmission in the presence of MDA [8, 9]. Quantifying environmental contamination with helminths poses major technical challenges: methods are needed that are both sensitive enough to estimate low—but epidemiologically relevant—concentrations of STH, and cost-effective enough to be deployed in low resource settings where the impact of STH is highest. Here, a systematic review of peer-reviewed and grey literature is presented to assess the state of the art of methods to quantify STH in the environment. Where relevant, information regarding other helminthic parasites with similar biophysical and environmental characteristics (e.g. *Toxocara* spp., *Taenia* spp.) is included.

Literature concerning the four distinct methodological steps involved in quantifying STH in the environment is reviewed below: (i) environmental sampling; (ii) recovering and concentrating STH ova, larvae, or genetic material from the sample matrix; (iii) detecting and quantifying recovered STH or genetic material; and (iv) determining the viability of STH.

Spatial Sampling Regimes For STH

Sampling STH from environmental media requires consideration of their fundamental overdispersion in the environment, with localized clusters of high contamination existing within areas that otherwise exhibit very low STH concentration (Box 1). This follows from the aggregation of high worm burdens in particular individuals, whose feces become a localized source of contamination in the environment [10].

The spatial distribution of STH can be estimated using systematic aligned (Figure 1A - e.g., [11-13]) or unaligned (Figure 1B - e.g., [14, 15]) methods; or walking path transect (Figure 1C - e.g., [16-21]) sampling patterns. A grid-based form of random sampling has occasionally been pursued in which the surveyed area is divided into equal parts that are then randomly selected for sampling (Figure 1D; e.g., [22, 23]). Others have proposed a systematic sampling method combining aspects of grid-based and transect sampling (Figure 1E) [17, 18, 24]. Starting from the corner of a rectangular plot of land, an investigator walks diagonally, turning at the boundaries to create a W-shaped path, and taking samples at regular intervals.

Purposive sampling, which relies on the investigator's judgment to determine appropriate locations from which to draw samples, has been used in some studies to sample from areas where STH are likely to survive, or where human exposures may occur, such as human and animal defectation sites [25, 26], shaded or moist areas [27-29], foot placements around latrine dropholes [30], or areas where children are observed to play [16, 31]. Alternatively, spatial stratified sampling is a method in which a survey site is subdivided into relatively homogeneous areas, from which a share of random or systematic samples is taken based on

their contribution to the total area or spatial variance of the quantity of interest (Figure 1F). Spatial stratified sampling is efficient for sampling highly spatially heterogeneous quantities [32], but has yet to be applied to STH.

The suitability of this broad range of spatial sampling protocols depends on the investigator's objectives and on the expected distribution of STH within the site. Transect sampling is appropriate for investigating the variance of STH concentration along some other environmental gradient or with distance from a contamination source. Systematic sampling patterns are efficient approaches for arriving at an estimate of two-dimensional spatial distribution of STH, but given the spatial heterogeneity of STH in the environment, spatial stratified sampling may be more efficient, especially when investigators have reason to suspect greater variability in STH concentrations in certain zones within their study sites.

Few studies have investigated the relative performance of environmental sampling strategies for STH. Carabin et al. [43] found that a random sampling method and two purposive sampling methods—selecting areas where animals were thought to defecate and selecting areas where children were seen to play—underestimated *Toxocara* contamination at a daycare center relative to a comprehensive grid-based sampling of the entire area [43]. Thus comprehensive systematic sampling can provide more information to estimate STH contamination across a study site, and may also be significantly more reliable than purposive sampling [43]. Verschave et al. [44] compared the performance of W-route sampling with systematic unaligned sampling of sixteen 0.16 m² plots for estimating local levels of pasture contamination with larval nematode parasites. They found no significant difference in the mean estimate of contamination between the two methods, but noted that the systematic unaligned sampling approach required less time to complete [44].

Recovering STH From Environmental Matrices

In order to quantify STH density in a sample, ova, larvae or their genetic material must be isolated from the environmental matrix and concentrated. Recovery of STH from soils, biosolids, and water samples typically involves five key processes: homogenization, chemical dissociation from the matrix, filtration, sedimentation, and flotation. As an initial step, sample homogenization can yield more reliable estimates of concentration because STHs are often unevenly distributed within environmental samples. Sample homogenization helps to lower variability between samples due to any disproportionate loss of STH associated with material discarded during sample processing.

STH ova tend to adhere to soils and other particles [45], and thus chemical dissociation from the particles in a matrix improves homogenization and prevents loss of ova as matrix particles are removed during subsequent processing steps [39]. Dissociation is usually achieved using ionic detergents such as 7X or Tween, which are thought to displace phosphate anions found on the outermost wall of ova from cationic sites on soil particles (http://www.ewisa.co.za/literature/files/155_107%20Hawksworth.pdf). Recovering STH from samples of plant matter requires a rinsing step (using Tris-buffered saline (TBS), Tween 20, Nacconol, or physiological saline, for instance), after which the rinsate is retained

and processed using membrane filtration, sedimentation and/or flotation as necessary (see, e.g., [40]).

Filtration by sieving helps to remove larger particles that can interfere with STH recovery and detection [39]. When the contents retained on a sieve are to be discarded, there is a risk of loss of ova associated with the discarded material. Homogenization and dissociation of ova from the matrix prior to filtration helps to mitigate this risk. Some protocols use very fine mesh sieves or membrane filters to retain ova and larvae while passing flotation fluid and smaller particles through. Careful matching of mesh size to the target STH is called for in this instance, as meshes small enough to retain *Ascaris* ova, for example, may allow *Trichuris* ova to pass through if they are oriented along their long axis [46, 47].

Sedimentation is used to concentrate STH at the bottom of a sample suspension, resulting in a reduced sample volume that is suitable for examination or further processing. The sedimentation process may be accelerated with the assistance of a centrifuge, or allowed to proceed passively [39]. The passive settling velocities of STH depend on their size and density, as well as the properties of flocs they form with other particles, which may vary strongly between STH species. For instance, the settling of *Ascaris* ova has been found to be more rapid in wastewater compared with tap water (57.6 cm/h vs. 21.6 cm/h) [48], while for *Trichuris* ova slower settling rates were observed in wastewater than in tap water (32.4 cm/h, 54.0 cm/h) [48]. Since organic matter is available for floc formation in wastewater, this result suggests that the flocs formed by *Trichuris* settle more slowly than those formed by *Ascaris* [48].

Flotation refers to the use of high specific gravity (density relative to water, abbreviated as SG) solutions to separate STH from heavier particles. To accomplish this, flotation solutions must be denser than the STH of interest [39] (for densities of various helminth ova, see [49]). Advantages of flotation include increased sensitivity of microscopic and genetic quantification methods through removal of sediment and polymerase inhibitors [39, 46]. Like sedimentation, the partitioning of lighter and heavier particles via flotation may be accelerated through use of a centrifuge [39].

Some well-established methods simplify (and circumvent) homogenization, filtration and other steps. For example, STH ova may be recovered from low turbidity water samples using simple membrane filtration, followed by a chemical treatment to render the filter transparent for visual inspection [50]. As another example, hookworm larvae may be recovered using a technique known as the Baermann method: sample material is suspended inside a porous membrane submerged in water, and as larvae migrate out through pores in the membrane, they settle to the bottom of the water column and may be collected for examination [24].

Recovery rates of STH ova from soils and biosolids for various methods reported in the literature vary widely, ranging from less than 10% to over 100% (which indicates measurement error in enumerating seeded and/or recovered ova). The highest reported rates of recovery for each method, as well as the corresponding methods and parameters, are

shown in Tables 1 and 2. Only one study has published recovery rates for the Baermann method for recovery of larvae, which were 80% on average, with a range of 58-99% [26].

Few studies have been published describing the efficiency of recovery of STH from water samples or vegetation. De Souza et al. [51] were able to recover roughly 36% of *Ascaris* ova seeded into wastewater using flotation (ZnSO₄ SG:1.18) and membrane filtration methods [51]. Maya et al. [50] were able to recover 80% and 83% of *Ascaris* ova seeded into well water and wastewater, respectively, and could detect as little as 1 ovum per liter using centrifugal flotation method (ZnSO₄ SG:1.3) with 0.1% Tween 80 as detergent [50]. Using a flotation (ZnSO₄ SG:1.2) and membrane filtration method without detergent, they were able to recover 86% and 63% of ova seeded into well and wastewater, respectively, and could also detect 1 ovum per liter [50]. Rude et al. [52] report an average recovery efficiency of 38.5% for a Nacconol-ether centrifugation method to recover *Ascaris* and *Trichuris* from large aggregate samples of vegetables [52].

Impact of method parameters on recovery from soils and biosolids

The efficiency of flotation is impacted by the properties of the flotation solution, characteristics of the soil matrix, and cross sectional area available for travel of STH in the flotation solution relative to the volume of matrix present [39, 46, 68]. To mitigate against poor recoveries, flotation steps may be repeated to recover any STH trapped between matrix particles during previous flotations. Sequential flotations have been reported to recover up to an additional 10-20% of seeded ova [64].

Soil texture and organic content have a substantial impact on recovery efficiencies. Recovery rates from clay soils are consistently lower and more variable than those from sandy soils [39, 45, 53, 58]. This may be because STH ova are among the lighter elements present in sandy soil, settling on the topmost layer of a suspension, while in suspensions of loam and clay they tend to sediment beneath a layer of lighter particles [69]. Similarly, the makeup of biosolids and sludge can affect recovery; heterogeneous biosolids (e.g., those mixed with soils) have been found to yield more variable recoveries [46]. Chemical treatment applied to biosolids has been found to impact the chemistry of dissociation and flotation processes, affecting recovery efficiency [46].

Properties of the flotation solution have been shown to influence recovery. Generally, higher SG solutions are thought to recover STH ova more efficiently (http://www.susana.org/en/resources/library/details/420). Several comparative studies have supported this assertion [53, 58, 63, 64], but others have reported either no improvement with flotation solutions of higher SG [59], or *superior* recovery with lower SG solutions [56, 62, 63]. Thus, it appears that in some cases, viscosity or chemical interactions with the outer surfaces of specific STH ova, the matrix, or other reagents may be more important determinants of recovery efficiency than SG.

Some authors have reported difficulties using specific flotation solutions. Concentrated NaNO₃ has been observed to crystallize, interfering with examination of microscope slides [59], though the issue has been addressed by adding a drop of glycerol to prevent crystallization [53] and would not matter for methods that separate ova from the flotation

solution with a fine-meshed filter. Concentrated sucrose solution was reported to yield low recoveries, presumably due to the interference of its high viscosity with the movement of ova and sediments during centrifugation and passive flotation [46, 63]. What is more, saturated saccharose has been reported to deform STH ova, potentially interfering with their quantification [70].

The choice of solution used to dissociate ova from soil particles has been found to affect recovery as well. NaOH is commonly used, although some authors have found this detergent to be ineffective [57, 62] or deleterious (e.g., through destruction of ova [59]) for recovering STH. The detergent Tween 40 has been reported to significantly improve recovery of *Toxocara* ova from soil and sand [61, 71], though Tween 20 has been reported as offering no improvement in at least one case [57]. The detergent 7X has been reported to be superior to other dissociation agents, including TritonX 100 and Tween 80 for recovering *Ascaris* ova from biosolids and soil [47], and benzethonium chloride and Tween 80 for recovering *Ascaris* ova in hand rinses [72].

Ova may be lost if they adhere to the walls of pipettes, tubes and beakers. To reduce this possibility, some protocols recommend treating labware with organosilane (Rain-X) to reduce adhesiveness. However, Jeandron et al. [71] found that organosilane treatment reduced yields from both glass and plastic materials, and recommended that non-coated pipettes and falcon tubes be used in protocols to recover helminth ova [72]. Adhesion can also be exploited for ova recovery, such as in the coverslip method commonly used for recovering *Toxocara*. In this method a test tube containing the sample is filled to the brim with flotation fluid in order to create a meniscus. A glass slide or coverslip is then placed so that ova traveling upwards via passive or centrifugal flotation may adhere to it and be counted (e.g., [61, 62, 71]). The relative sensitivity of this technique has not been compared with methods in which the entire final sample volume is examined for STH.

Some researchers, especially those conducting surveillance for *Toxocara*, pre-process samples by filtering them through coarse 4 mm² sieves (e.g. [12, 64]), or fine 150 μ m sieves (e.g. [21, 25, 27, 28]), prior to homogenization and other processing in order to remove twigs, rocks, and larger soil particles. The use of the finer sieve, in particular, prior to sample homogenization, seems likely to result in some loss of ova associated with retained materials. Nonetheless, one group has claimed that by drying a 200 gram soil sample, then sieving it down to 2 grams of powdery sand using the 150 μ m sieve, up to a maximum of 40% efficient recovery may be achieved, and so this method may provide greater throughput for soil samples at the expense of some sensitivity [29]. Others have reported that mechanical blending to disrupt large particles provides better results than sieving [62].

Recovery methods for biosolids and sludge sometimes include a phase extraction step in which the sample is partitioned between an acidic aqueous phase, where ova sediment, and a lipophilic phase [70]. However, one group of researchers reported decreased recovery efficiency and lower *Ascaris* ova viability when using an acid-alcohol phase extraction compared with a method using a fine mesh sieve to retain ova after flotation [47]. Exposure of ova to an acid-alcohol extraction mixture for more than 30 minutes yielded reduced

viability, and thus any acid-alcohol extraction procedure should be limited to short durations [47].

Storage conditions are important determinants of the recovery of viable STH ova and larvae. Ova of *Ascaris* and *Trichuris* are durable and can be stored at 4° C or room temperature for weeks or days, respectively, without loss of viability. However, hookworm ova are less resilient and have been shown to degrade within hours at room temperature if kept in a low moisture environment [73]. Hookworm larvae are even less hardy, and samples should be stored for no more than 7 days at 4° C to ensure a reliable assessment of viability [24]. It is important to note that differences in the physiochemical properties of ova across helminth species may necessitate adjustments to optimize other aspects of species-specific recovery methods, including flotation, dissociation, and sedimentation protocols.

Quantification Of STH Larvae And Ova

Light microscopy

After recovery from the sample matrix, STH have traditionally been quantified via light microscopy, a time-consuming and fatiguing process that requires expert knowledge of parasite morphology and is prone to human error, particularly when environmental processes or sampling and isolation methods distort parasite morphology. Furthermore, some species of STH, such as *Ascaris lumbricoides* and *Ascaris suum*, are nearly or completely indistinguishable by visual inspection. As an alternative to visual inspection, automated analysis of light microscopy images by computer could yield more rapid and reliable STH quantification. Such an approach has been described for detection of helminth ova isolated from cow feces [74]; identification of helminth ova isolated from human feces [75]; and quantification of different developmental stages of cultured *Trichuris* ova [76]. In concert with file sharing services, automated image identification has the potential to rapidly accelerate and standardize quantification of STH in environmental samples.

Nucleic acid-based methods

Genetic techniques for identifying and quantifying STH also have the potential to yield faster and more reliable sample analysis (see summary of genetic assay sensitivity for STH in Table 3). Genetic methods such as loop mediated isothermal amplification (LAMP) and polymerase chain reaction (PCR) have been used to detect STH in a wide range of contexts. Genetic methods can be species- or subspecies-specific, are highly sensitive (many assays target DNA sequences with multiple copies per cell, and can detect the equivalent of well under one mature ovum per gram of stool), and unlike microscopic or immunological assays, do not depend on parasite morphology remaining unaltered by environmental or laboratory processes [77]. Nonetheless, assays that rely on microscopic analysis are likely to be more practical for low-resource settings, where equipment for thermocycling and quantification of STH in low-resource settings, since it can be performed with simple equipment such as a water bath (even a thermos of hot water [78]), and produces enough magnesium pyrophosphate precipitate that a positive reaction can be discerned with the

naked eye [79]. Multiplex procedures to simultaneously detect multiple STH are available for PCR, but not yet for LAMP[80].

Certain compounds present in environmental matrices may inhibit LAMP and PCR reactions, including clays, humic and fulvic acids, polysaccharides, salts, heavy metals, and various organic molecules [39], though the BstDNA polymerase used in LAMP tends to be more resistant to inhibitors than Taq polymerases used in PCR [80]. Removal of inhibitors can be accomplished in part by using centrifugal flotation to recover STH ova [81], though even after flotation, adding anti-inhibition agents has been found to increase the sensitivity of genetic assays for STH [82]. To determine whether inhibitors are affecting reactions in a genetic assay, internal control targets can be used (e.g., [83]).

While most researchers who use genetic methods to detect sth in environmental matrices perform nucleic acid extraction after recovering ova by flotation [66, 84-86], some have successfully performed extraction directly on the environmental matrix [87, 88]. Some protocols call for disrupting the outer shell of certain helminth ova (primarily *trichuris* and *baylisascaris procyonis*) in order to efficiently extract genetic material; this can be accomplished through freeze-thaw cycles [85, 86], heating/boiling [85], shaking with glass beads [84, 85], or digestion with proteinase k [84, 86].

Determination Of STH Viability

Traditionally, STH ova viability has been assessed microscopically using an embryonation assay: after incubation under favorable conditions for 10-21 days, ova containing visibly developed larvae are recorded as viable, and undeveloped ova are assumed to be non-viable. The minimum time required for embryonation has been reported as 13 days for *Ascaris* and 16 days for *Trichuris* when incubated at 30 °C under constant aeration [94]. Hookworm ova hatch in soil within 1-2 days of excretion, and larvae survive in the environment for a few weeks at most (http://www.cdc.gov/dpdx/hookworm/index.html). Therefore, there is a risk that quantification of viable *Ascaris* and *Trichuris* ova via an embryonation assay may not generate accurate estimates of hookworm contamination in a sample – by the time *Ascaris* or *Trichuris* ova have matured, hookworm ova will have hatched and the larvae may have died and decomposed.

Determination of STH viability by microscopy is laborious and prone to human error, and as for STH detection and quantification generally, developments in automated image analysis —particularly when used with dyes that stain only live or dead cells—have the potential to improve the efficiency and reliability of microscopy-based viability assessment. Dabrowska et al. [95] were able to differentiate *Ascaris, Toxocara*, and *Trichuris* ova with high efficiency using SYTO 9 to stain live cells and propidium iodide to stain dead cells. The sensitivity of the assay decreased slightly when applied to sewage sludge samples, most likely due to interference of bile salts with the uptake of fluorescent dyes [95].

Genetic assays also show promise for establishing the viability of STH ova. In one such approach, ITS-1 DNA (internal transcribed spacer region of ribosomal DNA, present in multiple copies per genome) is amplified by quantitative PCR in order to track the development and viability of *Ascaris* ova [92, 96]. Raynal et al. [96] determined that ITS-1

DNA signals in embryonated ova decreased to negligible levels 10 days after inactivation, but persisted in viable ova [96]. Pecson et al. [92] proposed that persistent DNA in inactivated ova could be removed by treatment with proteinases and nucleases prior to PCR, and demonstrated that RTPCR targeting ITS-1 mRNA detected only viable ova, though the RNA assay was not sensitive nor consistent enough to be quantitative [92]. Other possible means of quantifying viability using genetic assays include using vital stains such as ethidium mono azide as inhibitors that only affect dead cells [97], and using quantitative RT-PCR to measure heat shock protein mRNA after exposure to elevated temperatures [39], but neither approach has been published for STH.

Concluding Remarks

Methods for reliably, rapidly, and cost-effectively assessing environmental contamination with STH will be crucial for understanding and disrupting STH transmission. Dominant methodologies are time and labor intensive, unreliable, and unstandardized in their approach. Key unresolved issues remain regarding optimal environmental sampling methods for STH, and optimal parameters for recovery, quantification, and viability assessment protocols (see Outstanding Questions Box). Future research examining sampling error under various spatial sampling strategies for STH and the effect of systematic variations in method parameters on recovery efficiency, will be critical to establishing standardized, efficient and reliable STH detection methods. To begin to address the many gaps in our understanding of the optimal strategies for recovering STH ova from environmental samples, some have called for universal use of internal process controls during analysis of environmental samples [98]. Stained control ova can be introduced to environmental samples, providing a standardized means of estimating efficiency of STH recovery for each published assay. Such a strategy could help to accelerate a consensus on optimal techniques.

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Box 1. Characteristics of the spatial and temporal distribution of STH in the environment

STH ova exhibit the longest survival times in moist environmental conditions with little sunlight [33], and STH embryos develop most efficiently in aerobic environments [34]. Thus ova within sandy or loose soils, which retain water poorly, have been shown to be less resistant to desiccation and ultraviolet radiation [12, 35], while soils that are impermeable and anoxic have been observed to slow ova maturation [35]. Optimal conditions for hookworm larvae are similar, except that spacing between soil particles facilitates their migration through the soil column. Infective hookworm larvae have been observed more frequently in moist, shaded, sandy soils [36].

The vertical distribution of STH in the soil column is poorly understood. While exposure to viable STH ova that have passed below the surface layer is presumed unlikely, vertical transport by earthworms and disturbance of soil by rooting animals and human activity may return ova from deeper layers to the soil surface [37, 38]. Hookworm larvae actively move within the soil column, and are capable of migrating between the surface and depths of up to 20 cm to avoid dry conditions [22].

Among environmental media, STH tend to be sparsely concentrated in surface waters where ova settle rapidly out of the water column. In wastewater, however, STH may be present in high concentrations and relatively evenly dispersed due to mixing of numerous fecal sources [39]. Food crops grown close to, but above the ground are most frequently and heavily contaminated with STH [40]. Root crops are also prone to STH contamination [40], and the large surface area of leafy greens facilitates STH attachment and provides protection from drying and UV radiation, making these some of the most frequently contaminated crops [40, 41].

Seasonal fluctuations in temperature, moisture, and infection prevalence (due to MDA campaigns) are known to affect the seasonal distribution of STH in soils. Generally, more frequent soil contamination is found during wet seasons [19, 26-28, 42]. Some researchers have reported higher occurrence of STH, but lower STH viability, in soils during dry seasons, perhaps due to lack of rain to wash ova away [13, 33]. Temporal sampling regimes for STH often reflect these seasonal trends, with samples drawn during both wet and dry seasons to capture high and low contamination conditions [13, 19, 20, 27, 28].

Outstanding questions

Despite the long history of methods for the environmental detection of STHs in soils and other media, key methodological issues remain:

- What spatial sampling strategies are most efficient for estimating the environmental distribution of STHs in soil, surface water, and vegetation?
- Few studies have taken a systematic approach to environmental sampling for STHs, and thus optimal sampling strategies that reduce measurement error and uncertainty for specific STHs in particular environmental media remain poorly understood.
- What aspects of recovery protocols are most important for maximizing STH recovery for specific species and environmental media?
- Limited comparative studies have been conducted examining the efficiency of recovery methods conducted with controlled differences in procedural parameters. Parameters whose effects need clarification include the choice of flotation solution, its specific gravity or exact chemical nature, the choice of dissociation agent, lengths of sedimentation, and preprocessing and lipid extraction steps. There is a particular absence of data on the efficiency of methods to recover STHs from plant matter.

Trends

- The state of the art and key developments in environmental methods for sampling, recovery and concentration, quantification and viability assessment of soil transmitted helminths (STHs) are reviewed.

- Optimal protocols for sampling and recovery of STHs from environmental samples have not been developed, and systematic investigation is needed.

- Recent advances in genetic assays and automated image analysis for quantification and viability assessment offer improved sensitivity, reliability, and sample throughput.



Figure 1.

Spatial sampling regimes for soil transmitted helminths (STH) in soil and vegetation. (A) Systematic aligned sampling. (B) Systematic unaligned sampling. (C) Walking path transect sampling. (D) Random sampling of a grid. (E) W-route sampling. (F) Spatial stratified sampling with random samples designated according to expected variance within each stratum.

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Table 1

Reported p	arameters	yielding maximu	m rates of reco	very of STF	H from soils.			
Organism	Soil type	Seeding level (EPG a)	Method type	Detergent	Flotation solution (SG)	Percent recovery	Estimated processing time b	Refs.
	Clay	4				15.3%		
Ascaris	Sandy	20	Passive flotation	Bleach (20%)	NaNO ₃ (1.3)	24.32%	2 hours	[53]
	Loamy	20				25.04%		
Ascaris	NR ^C	1452- 32393	Centrifugal flotation	NaOH (0.5 N)	Glucose monohydrate (1.27)	82%	2 days	[54]
				Tween40 (0.5%)	NaNO ₃ (1.35)	47%	<1 hour	
Ascaris	NR	10	Centrifugal flotation	Tween80 (1%)	NaNO ₃ (1.35)	66.5%	<1 hour	[55]
				None	ZnSO4 (1.2)	9.5%	3 hours	
Trichuris	NR	1452- 32393	Centrifugal flotation	NaOH (0.5 N)	Glucose monohydrate (1.27)	80%	2 days	[54]
Trichuris	NR	-	Centrifugal flotation, recovery by adherence to 4 coverslips	None	MgSO4 (1.28)	96.4%	1-2 hours	[56]
				Tween40 (0.5%)	NaNO ₃ (1.35)	39.33%	<1 hour	
Ancylostoma	NR	10	Centrifugal flotation	Tween80 (1%)	NaNO ₃ (1.35)	50%	<1 hour	[55]
				None	ZnSO4 (1.2)	%6	3 hours	
Toxocara	Sandy	7	Centrifugal flotation, recovery by adherence to 6 coverslips	Tween80 (NR)	NaNO ₃ (1.22)	15%	2.3 hours	[11]
		2	Passive flotation, recovery by adherence to 4 coverslips	Tween40 (NR)	ZnSO4 (1.2)	15%		
Toxocara	Sandy	543	Passive flotation	None	Saccharose (1.27)	99.98%	<1 hour	[57]
	Silty clay					7.5%		
Tavaoma	Clay silt	000	Centrifugal	NaOH	NanCraO ₂ (1.35)	13.9%	/1 hour	[58]
1 OXOCUTU	Sandy	7007	flotation	(0.1 N)	10011/1021020VI	38%	mou 1>	[oc]

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62.5%

Sand

Organism	Soil type	Seeding level (EPG ^{a})	Method type	Detergent	Flotation solution (SG)	Percent recovery	Estimated processing time b	Refs.
Toxocara	Sandy	200	Centrifugal flotation, recovery by adherence to coverslip	Tween80 (0.2%)	ZnSO4 (1.2)	10.75%	2 days	[59, 60]
				NaOH (0.1 N)		8.8%		
Toxocara	NR	-	Centrifugal flotation, recovery by adherence to 4 coverslips	None	MgSO ₄ (1.28)	47.2%	1-2 hours	[56]
			Centrifugal	Tween40 (0.5%)	NaNO ₃ (1.35)	46.83%	<1 hour	
Toxocara	NR	10	different methods: ([12,	Tween80 (1%)	NaNO ₃ (1.35)	71%	<1 hour	[55]
			00, 02])	None	ZnSO4 (1.2)	7.67%	3 hours	
Toxocara	Clay	210	Centrifugal flotation X $3d$, recovery by adherence to $4-5$ coverslips	NaOH (0.1 N)	ZnSO4 (1.18)	66.19%	1-2 hours	[62]
	Silt				NaCl (1.25)	45.3%		
Toxocara	Sand	20	Centrifugal flotation	Tween80 (0.1%)	NaCl (1.25)	100.8%	1-2 hours	[63]
	Laterite				ZnSO4 (1.52)	56.8%		
Toxocara	${}^{\rm NR}{}^d$	0.48	Centrifugal flotation X 4	Tween80 (0.0025%)	MgSO4 (1.275)	82.5%	1-2 hours	[64]
Toxocara	NR	NR	Centrifugal flotation, recovery by adherence to 2 coverslips	Tween80 (1%)	NaNO3 (1.35)	69.8%	1-2 hours	[12]
Echinococcus	NR	100	Centrifugal flotation, recovery by adherence to coverslip	Tween80 (0.05%)	Sucrose (1.27)	10%	2-3 hours	[65]
Echinococcus	Sand	NR	Centrifugal flotation, recovery by adherence to coverslip	Tween80 (0.05%)	ZnCl ₂ (1.4)	NR	1-2 hours	[66]
¹ EPG - Eggs pe	y gram.							

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b Processing times estimated based on given times for incubation and centrifugation steps, and do not include time needed to examine the sample after recovery.

 c NR - not reported.

Anthon Manuel (e.g. X 4 denotes four sequential flotations).

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Reported parameters yielding maximum rates of recovery of Ascaris from biosolids and sludges.

Organism	Type	Seeding level (EPG a)	Method type	Detergent	Flotation solution (SG)	Percent recovery	Estimated processing time $m{b}$	Reference
	Acid-treated					80.5%		
-	Anaerobically composted	¢	Centrifugal	XL	MgSO4	75.5%		72
Ascarts	Soil-blended	NR	flotation	(NR)	(1.2)	75.5%	stad c	[0+]
	Alkaline- treated	I			-	58.5%		
	Urine-diverting		Centrifugal flotation,	NH4HCO3 (NR)	Posuz	77.28%	2-3 hours, significant time to read slides due to debris	
Ascaris	toilet biosolids	0071	Centrifugal flotation with sequential sieving	None	(1.4)	57.14%	<1 hour	(nd, mowsaweau2.8%) u L. cc1/sein/sumeranne.ac.ac.ac.ac.ac.ac.ac.ac.ac.ac.ac.ac.ac.
Ascaris	Sewage Sludge	125 (per gram total solids)	Centrifugal flotation	7X (1%)	MgSO4 (1.2)	82.00%	5 days	[67]
a _{FPG - Fo}	os ner oram							

г ч – Eggs per gram

b Processing times estimated based on given times for incubation and centrifugation steps, and do not include time needed to examine the sample after recovery.

 c NR – not reported.

Table 3

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Reported sensitivities of genetic assays for STH detection and quantification.

Organism	Type	Life stage	Target gene(s)	Sample matrix	Extraction kit	Detection method	Limit of detection ^a
Echinococcus multilocularis	PCR	Ova	12S rDNA	Soil	Sherlock AX Kit (A&A Biotechnology)	Electrophoresis	80% detection at 0.25 EPG^{b}
Strongyloides stercoralis	LAMP	Larva	28S rRNA	Stool	Powersoil DNA Isolation Kit (Mobio)	Syto-82 fluorescent dye	0.4 larvae / mL
Taenia solium Taenia asiatica Taenia saginata	Multiplex PCR	Ova	Valine tRNA NADH dehydrogenase	Stool		NR ^c	10 EPG
Taenia solium Taenia asiatica Taenia saginata	LAMP	Ova	COX1 CLP	Stool	QI Aamp DNA stool Minikit (QIAGEN)	Electro-phoresis	88.4% detection of single genome
Taenia spp	Multiplex PCR	Ova	COX1	Stool	QIAamp DNA Stool Minikit (QIAGEN)	Electro-phoresis	50 EPG
Toxocara canis		Ova	ITS-2	170 S	P		33% detection at 2 EPG
Ancylostoma caninum	LUN	(1 cell)	rDNA	1100	NA		0.4 EPG
Toxocara canis Toxocara cati	LAMP	Ova	ITS-2 rDNA	Sand	NA	Electro-phoresis	0.6 EPG
Strongyloides stercoralis	Multiplex quantitative PCR	Larvae	18S rRNA	Stool	QIAamp DNA Stool Minikit (QIAGEN)	Fluorescent Probes	0.4 larvae per gram
Trichuris spp			ITS-1 rDNA				
Necator/ Ancylostoma spp	Quantitative PCR	Ova	ITS-2 rDNA	Stool	NA ²	EvaGreen fluorescent dye	1 EPG ³
Toxocara spp		-	23S rDNA				
Ascaris lumbricoides	Quantitative PCR	Ova	ITS-1 rDNA	Sterile water	UltraClean microbial DNA and RNA kits (Mobio)	Fluorescent probes	3,750 copies or ~ 4 cells
Toxocara canis Toxocara Cati	Multiplex quantitative PCR	Ova	ITS-2 fRNA	Soil	PowerMax [®] soil DNA isolation kit (Mobio)	Fluorescent probes	2 EPG

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Refs [99] [87]

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Organism	Type	Life stage	Target gene(s)	Sample matrix	Extraction kit	Detection method	Limit of detection ^a	Refs
Ancylostoma duodenale		Ova	ITS-2 rDNA				1 copy per gram	
Necator americanus	Multiplex	Ova	ITS-2 rDNA	1000	QIAamp DNA Stool Mini kit		100 copies per gram	
Ascaris lumbricoides	PCR	Ova	ITS-1 rDNA	10016	(QIAGEN)	Filorescent proces	1 copy per gram	[06,00]
Strongyloides stercoralis		Larvae	18S rDNA	_			1 copy per gram	

^a Limits of detection are the lowest tested concentration at which 100% recovery was reported, unless otherwise stated.

 $b_{
m EPG-eggs}$ per gram.

 c NR – not reported.

 d_{MA} – authors used their own in-house nucleic acid extraction techniques.