



Development and performance of an automated fecal egg count system for small ruminant strongylids

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ABSTRACT

An automated equine fecal egg count (FEC) system (the Strongly system) was modified for use with small ruminants. Modifications included the incorporation of a shorter flotation step in a flotation system, an adjustment in pre-test sample filtering, and tuning of an image analysis-based egg counting algorithm. In recognition and elimination of false-positive eggs, in preliminary assessments the modified method produced microstrongylid egg counts comparable to manual McMaster analyses of the same samples from both equine and caprine sources. The coefficient of determination (R^2) for the linear correlation between McMaster and automated counts from these samples was 0.94, and there was no significant difference when comparing counts using feces from either sheep or goats. More extensive comparisons of these equine samples split into three groups based on microstrongylid egg counts: Low (0.01–500 EPF), Medium (500–1000 EPF) and High (1000 or greater EPF). Each group contained 5 samples, each of which was used in quadruplicate (divided into two manual 8 times each using both McMaster and the automated method). This, again, showed no difference in accuracy between the techniques, but revealed significantly higher precision, as assessed by coefficients of variation (COV), for the automated method for determining egg counts in the Low and Medium groups. The COV of the McMaster method was 2.2, 2.5 and 2.3 times greater than the automated in the Low, Medium and High groups, respectively. Overall, the automated egg counting system showed good linear agreement with manual egg counts determined with the McMaster method, and demonstrated significantly better precision. This technology allows operators and the results generated from this system to utilize for determination of small ruminant (microstrongylid) fecal egg counts.

1. Introduction

Routine fecal egg counts (FEC) and FEC reduction (FECR) analyses are an integral part of parasite management strategies in small ruminants (Gahan and Woodgate, 2012; Słusarewicz et al., 2019; Burke and Miller, 2020). These have become all the more important since the evolution and spread of anthelmintic resistance in trichostrongyle parasites (Walker, 1994; Jackson and Coop, 2008; Cernuska et al., 2009; Howell et al., 2009; de Graaf et al., 2010; Therios-Missouliou et al., 2017; Ploger and Ewers, 2018; Lambertz et al., 2019). Parasite load has been shown to adversely affect productivity (Miller et al., 2012; Abbas and Hudaib, 2014) and the development of anthelmintic resistance, therefore, represents a growing concern in the small ruminant industry.

Numerous methods are available for conducting FECs and FECRs.

These include the Stoll (Stoll, 1923), McMaster (Gordon and Whitlock, 1939), Wisconsin (Graham and Tyell, 1982), FLOTAC (Oringali et al., 2010) and Mini-FLOTAC (Oringali et al., 2012) methods. These tests can be done on the lower density of wax relative to the non-parasitic elements of the bulk feces, and wax are separated by flotation when a fecal sample is suspended in a dense medium (density between 1.2 and 1.5 g/dl). This separation facilitates the identification and counting of the wax by refractive background produced by the non-parasitic components of the feces.

A common theme in all these tests is the need for manual count of the wax under a microscope in order to obtain quantitative results. As a result, FECs can be affected by the expertise or training status of the analyst (Słusarewicz et al., 2019; Cain et al., 2021). These concerns have been addressed by the development of a variety of new technologies.

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The iDCATM system (Radadi et al., 2018) collects ova in a conical mesh filter (Zinke et al., 2015), where they can be photographed through multiple focal planes by a specialized camera; the composite images are then loaded to a cloud server, where they are manually read remotely by a trained analyst (directly eliminating the need for ova on site), albeit at the cost of a more complex and laborious sample preparation process. Yet other systems have circumvented the possible variability introduced by manual examination by introducing automated counting systems that utilize selected computer vision algorithms produced using deep-learning neural networks (Elghayoury et al., 2020; Nagarsol et al., 2020; Citigözü et al., 2021). These systems, however, require high-resolution images as inputs to their algorithms in order to produce acceptable diagnostic specificity and sensitivity. In practice this requires the collection of a large number of high-resolution images in order to capture the entire sample, and so is a trade-off between optical magnification and field-of-view. As a result, such systems need to take multiple images (using automated stages) to capture the entire sample, which increases test times by between 6 and 42 min (Elghayoury et al., 2020; Nagarsol et al., 2020; Citigözü et al., 2021).

We have developed an alternative automated method that circumvents this dilemma by producing a single low-magnification image with a broad field-of-view that can nevertheless be analyzed computationally (Slosserweitz et al., 2016; Sears et al., 2017). This is achieved by fluorescently labelling eggs by means of the chitin present in their shells using a derivatized secondary chitin-binding protein (Slosserweitz et al., 2016). Imaging in fluorescence mode facilitates the production of high contrast, low background images with sufficient fidelity for automated identification, even at low magnification. As a result, only a single image is needed for each sample, and it is captured in less than a minute, while a four-minute sample preparation and processing procedure is simplified by means of an automated device. Our derivatization of equine stronglyylid larval egg contents, this system has been shown to perform with an accuracy comparable to manual McMaster tests while being significantly more precise (Slosserweitz et al., 2019; Cain et al., 2020).

In this paper, we describe the development of a version of this system intended for use with small ruminants, and report on its performance for determining trichostrongylid egg counts in equine and ovine samples compared to the McMaster technique.

2. Materials and methods

2.1. Materials

The properties of the McMaster and automated techniques used in this study are presented in Table 1. The iDCA MED 35.6-9 (98%) is dimethyl sulfoxide medium (specific gravity 1.27) from VWR Ltd. (St. Joseph, Missouri, USA). Trichostrongylid McMaster slides were from FEC Source (Bank, Oregon, USA). An Eclipse E600 microscope (Nikon, Melville, New York, USA) was used for McMaster egg counting.

The Derivatised secondary system was provided by the manufacturer (MFP Equine Solutions, Lexington, KY) and its various components are illustrated in Fig. 1. The system consists of a sample Preparation Tool (SPT), a portable Belgian Dispensing Unit (BDU), disposable Egg Chambers (EGCs), and an Imaging Unit (IU) with an Android tablet running an app that downloads the image from the IU and runs the

egg counting algorithm. The SPT itself consists of a silicone bottle and an interchangeable spring loaded plunger for sample homogenization and a filter cap for sample filtration prior to processing in the BDU. The filter cap contains three filters in a series of decreasing U.S. mesh sizes of 16 (1180 μm), 30 (420 μm), and 70 (210 μm).

Samples were centrifuged using a 15 ml centrifuge tube in a CF-800-1 fixed-angle centrifuge (Hardware Factory Store Inc., Amisa, GA). The fixed-angle centrifuge was chosen to represent a type present in many veterinary practices.

2.2. Samples

Ovine and equine fecal samples were collected during June–July 2020 from various regions of the United States, namely Pennsylvania, Georgia, Arkansas, and California and shipped overnight and on ice from each source. Samples were collected by a mixture of veterinary practices, parasitology research laboratories, and reference laboratories, with some samples being collected rectally and others from freshly deposited feces on the ground. Samples were stored in sealed plastic bags and at 4 °C until analyzed (within 5 days of receipt).

2.3. Algorithm development

Images for algorithm training were produced from the samples described above. Samples were resuspended in iDCA MED at a ratio of 1 g : 9 mL and filtered through two layers of cheesecloth. The filtrate was then diluted into 8 mL portions in 15 ml centrifuge tubes and spun at 2200 g for 3 min. Each supernatant was then processed through the automated system (BDU), and images captured with the IU. Full (18 megapixel) resolution images were then extracted and individual stronglyylid eggs copied into individual images to produce a training set of approximately 5000 images, which were then used to train a TensorFlow-based deep learning neural network (www.tensorflow.org).

The original automated system was designed to count stronglyylid eggs from venines, since the lengths of the minor axes of equine stronglyylid eggs range between 40 and 65 μm (Bosey, 2001; Slosserweitz et al., 2016), they can all be trapped on the 37 μm mesh integrated into the EC that is used to encase eggs in the ovine system. However, ruminant trichostrongylid eggs are significantly smaller, with minor axial lengths ranging between 34 and 50 μm (Forest, 2001). Due to this smaller size, we used equine ECGs fitted with 25 μm meshes to produce images for training. This would prevent possible loss of smaller eggs during sample processing (particularly since the process involves a substantial surface force at several stages to drain reagents between treatment steps), and therefore their possible under-representation in the training set.

2.4. FEC counts

Preliminary comparison to McMaster counts used 13 ovine and 12 equine samples analyzed in triplicate, while further comparison to automated and McMaster counting used 15 samples with a broad range of stronglyylid egg counts to produce fifteen slurries that were then analyzed eight times each by both methods.

To reduce slurry FEC variability when comparing the automated method with the McMaster technique, both tests were conducted on the same slurries rather than on independent slurries produced from different fecal subsamples. Furthermore, slurry subsamples were counted either three or eight times with both methods to, at least partially, reduce subsampling variation. In some cases, samples were analyzed using multiple sequential subsamples of slurry from the SPT, and in other cases slurry from one or multiple SPTs were filtered through the tool and pooled prior to multiple analyses (see below). The McMaster method used in this study has been described previously (Slosserweitz et al., 2019).

The equine automated system makes use of an SPT consisting of a silicone bottle attached to a series of increasingly fine filters designed to

Table 1
Properties of the two fecal egg counting techniques evaluated in the study.

	McMaster	Automated
Volume of Derivatised medium (mL)	54	54
Fecal weight (g)	6	6
Volume of suspension created (mL)	60	60
Chamber size covered	2 × 10 mm X 10 mm	8 mm X 24 mm
Microplate reader	342	25



Fig. 1 Components of the automated system. The system consists of a reusable Sample Preparation Tool (SPT), which itself consists of a silicone bottle, a handle for sample homogenization and a dispensing tip with integrated filter; dispensable egg Chambers (EC) in which samples to be chemically treated and imaged; a Reagent Dispensing Unit (RDU); an Imaging Unit (IU); and an Android tablet with an App for capturing images from the IU and for turning eggs in the images.

removes as much of the fecal debris as possible prior to dropping eggs onto a fine mesh filter (Stamm et al., 2017; Slosarski et al., 2019; Cohn et al., 2020, 2021). Initial attempts to use the ocine system with sheep and goat feces encountered two problems. Firstly, many samples tended to clog the filter system of the SPT, sometimes making complete dispensing of the sample problematic. Secondly, many samples also either clogged the EC, thereby preventing proper coating of chemical reagents during automated processing, or contained sufficient fecal debris to obscure substantial numbers of eggs. We addressed the first problem by removing the (fine) filter from the SPT filter stack, and the second by reconstituting the sample in a sodium nitrate (specific gravity 1.25–1.30) flotation medium (FM) and introducing a post-flotation centrifugation step (see below) to remove the bulk of the fecal particles that could foul the EC mechanism.

The automated equine method was therefore adapted for ruminants as follows. Samples (to 2) were suspended in 54 mL of FICMA M20 using the SPT. In the case of pelletized samples, feces were placed in a plastic bag and shredded in a pasta prior to weighing. Eight mL of suspension was poured into a 15 mL tube through the filter assembly of the SPT and centrifuged at 2200 g for 3 min. The supernatant was then poured into an EC placed on the RDU, coated and then processed with the same cycle used for equine samples. Samples were discarded immediately after centrifugation, because the pellets loosened substantially to the point that significant amounts could dislodge during pouring. When multiple samples were centrifuged simultaneously, all supernatants were discarded immediately afterward into fresh tubes so that they could later be processed as follows.

2.5. Media comparison

Initial experiments sought to determine the extent of any correlation between high-density McMaster and automated counts of the same samples. Upon discovering that small ruminant feces have a propensity to block filters, we were concerned that this might lead to egg loss during filtration. More specifically, using the SPT (even with the finest mesh removed) to produce sub-filters combined slurry for both manual and automated replicate analyses might lower egg counts due to progressive clogging during passage through the meshes. We therefore decided to use coarse cloth as a filter on ocine. However, we were also concerned that results produced using excess cloth may not represent those obtained when using the SPT. We therefore produced a slurry in the SPT, dispensed 8 mL directly into a centrifuge tube, spun the sample, and used this for the filter-automated count. The remaining material in the SPT was also filtered through two layers of cheesecloth, and the filtrate returned to the SPT and harvested through the integrated filter. This

material was then used to conduct all three McMaster and the second and third automated counts.

While the second and third automated counts were, on average, 4.9 and 13.0.8 greater than the first ($n = 26$), respectively, these differences were not significantly different at the 0.05 significance level according to a Wilcoxon signed rank test (see Statistical Analyses, below). Furthermore, the third automated counts were 13.7% greater than the second but also not significantly different. These data indicate that the approach of pre-filtering through cheesecloth prior to the SPT meshes did not produce slurrings with significantly differing counts to the first dispense from the SPT (i.e., the intended in-use condition).

To assess test precision we adopted a similar experimental design to that described previously (Slosarski et al., 2019). Samples were divided into three groups based on trichostrongyle egg count in eggs per gram (EPG): Low (201–500 EPG), Medium (501–1000 EPG) and High (>1000 EPG). Five samples were assigned to each group after screening using a triple-test McMaster counts at the second slurry, and each sample was coded to hide its identity from the analyst. Slurrings were processed using two SPTs and then successively filtered through cheesecloth and into the SPT filter. The pooled filtrate was then counted eight times by both the McMaster and automated methods.

2.6. Statistical analyses

Statistical calculations were done using R version 8.0.0 (The R Foundation) with the “lme4” package installed as with SPSS built 16.0.0.147 (IBM, Armonk, NY). A mixed model was used to determine whether diluting the lower density (LU) pellet had a significant effect on egg count, with sample ID as a random effect. To determine the significance of the difference in relationship between automated versus McMaster test results for ovine and caprine fecal samples, linear regression models were created regressing mean egg count per slide on test method and animal species with an interaction term between test method and animal species. The p value of this interaction term coefficient determined whether animal species had a significant effect on filter slope. Differences between counts at samples prepared by different filtration methods were assessed using a Wilcoxon signed rank test.

Differences between coefficients of variation (CVs) were assessed using a log-ratio test. Heteroscedasticity was confirmed both by inspection of Q–Q plots and by using Shapiro-Wilk tests. Levene’s Test for Equality of Variances was used to assess the homogeneity of variance; when variances were determined to be unequal (statistically confirmed when the CVs of all egg level groups were combined), a heteroscedastic test was performed, and in the case of unequal variances, a log-ratio test was used. Differences between egg counts were assessed using

paired *t*-tests after confirming normality of the differences both by an inspection of Q-Q plots and by using Shapiro-Wilk tests.

Differences were considered significant at $P < 0.05$.

3. Results

3.1. Manual equilibration

Centrifugation of samples in flotation medium produced two pellets: a larger one at high density (HD) located at the bottom of the tube and distal to the center of rotation, and a smaller one of LD material towards the top of the slurry meniscus and proximal to the center of rotation (Fig. 2). The latter was also sometimes accompanied by a faint smear of material along the length of the tube.

We were concerned that the LD material might contain some eggs that had adhered to the side of the tube and this result in some egg loss during sample processing. However, we observed that when the supernatant was decanted while the LD pellet faced downward so that the liquid washed over it, the majority of this material was dislodged and deposited into the egg chamber. In contrast, if the tube contents were

decanted after rotating the tube 180 degrees around its longitudinal axis so that the LD pellet faced upward and the supernatant did not flow over it, substantial loss or no material was dislodged (data not presented).

To determine whether dislodging the LD pellet had any effect on the resulting count, three slurries prepared from different samples were rotated using the two panning methods (i.e., with the LD pellet facing upward or downward) to produce 8 counts per slurry per method. There was no significant difference in mean egg count between the two methods. In contrast, the LD pellets captured samples produced CVs that were, on average, 3.4–4.6% larger than the LD-enriched samples (Table 2) and this difference was significant ($P < 0.05$). As a result, all subsequent automated HPLCs were conducted with the LD enrichment procedure.

Once the entire system had been built, we tested the possibility that some ovine eggs might be lost through the pores of the 39 μm mesh of ovine FCs by running quadruplet automated counts of *Trichostrongylus* using FCs that were either 25 or 88 μm screens (Table 2). The counts generated with the larger mesh size screens were, on average, 21.6% lower than those generated with the smaller screens, a difference that was statistically significant ($P < 0.001$). As a result, 25 μm screens were used for the remainder of the study.

3.2. Method performance

The correlation of the McMaster and automated methods was assessed by comparing replicate counts from the same slurry. Both ovine and caprine samples were analyzed in triplicate using manual and automated counts from single slurries of each. The combined manual and automated counts for both species exhibited a strong positive correlation with a coefficient of determination (R^2) of 0.956 and produced a line with a slope approaching unity (1.0; Fig. 3). When treated separately, the ovine and caprine data produced lines with slopes of 0.924 and 1.05, and R^2 values of 0.884 and 0.972, respectively. The difference between the ovine and caprine slopes was not statistically significant.

The precision of the two methods was next compared using only ovine samples (3 groups based on egg count, with 5 samples per group) and each counted 8 times. After averaging replicate counts, these data produced a line with a slope of 0.973 and an even stronger correlation ($R^2 = 0.981$), presumably due to the increased number of replicates (Fig. 4). There was no significant difference between the counts within each group nor in all the groups combined. Precision (as assessed by the CoV) was superior for the automated method in all 3 groups (Table 4), but was significant only in the low and medium count groups ($P < 0.01$ and $P < 0.05$, respectively). The CoV for McMaster was 2.2, 2.5 and 1.3 fold greater than for the automated method in the Low, Medium, and High groups, respectively. In all the groups combined, the McMaster CoV was 2.0-fold higher than that of the automated method, and this difference was also statistically significant ($P < 0.001$).

4. Discussion

The purpose of this study was to modify an automated equine FEC

Table 2

Effect of egg chamber filter pore size on egg counts. A single feed slurry was counted in quadruplicate using egg chambers designed for use with equine samples (LD) or in fluid with a finer mesh (25 μm). Mean results for each group are shown.

	McMaster	25 μm
Mean FEC	91.5	1.60
S.D.	77.8	85.1
CoV	8.5	7.9

S.D. = standard deviation (S.D.).

CoV = coefficient of variation.

FEC = eggs/g.

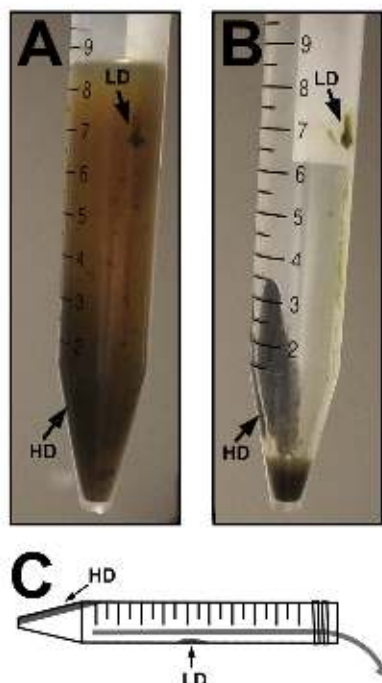


Fig. 2. Sample separation procedure. (A) Sample centrifugation produces a large, high density (HD) pellet at the bottom of the slurry (to the distal to the axis of rotation) and a smaller low density (LD) pellet on the opposite side of the tube, that was often accompanied by a faint smear of material below it. (B) Appearance of the same tube after carefully decanting supernatant with the LD pellet facing upward to minimize its disturbance. (C) Schematic diagram of this system for the decanting operation while maximizing retrieval of the LD pellet material.

Table 3

Effect of post-centrifugation enrichment of the low density (LD) pellet on egg count. Samples collected from three fecal samples (A, B and C) were counted after detaching the eggs onto a slide, which is easier for LD pellet egg counts per group. Results are reported as mean counts per fecal group

Sample	enriched LD pellet			wash to pellet		
	A	B	C	A	B	C
Mean (SD)	520.3 (227.5)	527.5 (215.5)	441.6 (194.0)	1049.4 (312.4)	1322.4 (316.1)	1163.1 (252.2)
CV	43.7	40.9	44.1	29.8	23.9	21.7

S.D. standard deviation (SD)

CV = coefficient of variation

EPG = eggs/g

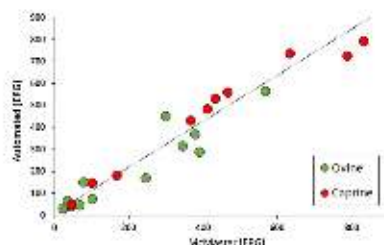


Fig. 3 Correlation of triplicate ovine and caprine automated counts with a single McMaster count. Thirteen ovine and 12 caprine samples were used to generate counts, but were then counted in duplicate by the McMaster and automated methods. The averaged results were then plotted together. The line represents the best least squares fit to the combined data. The slope is 1.05 with a coefficient of determination (R^2) of 0.958

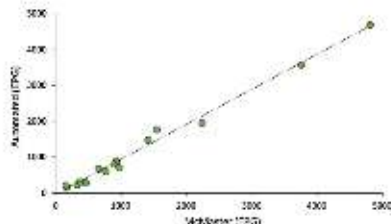


Fig. 4 Correlation of triplicate ovine automated counts with a single McMaster count. Five caprine fecal samples with a broad range of counts were used to produce sludges that were then counted either times by the automated and McMaster methods. The averaged results were then plotted together. The line represents the best least squares fit to the data. The slope is 0.971 with a coefficient of determination (R^2) of 0.991

system for use with small quantities and to compare its performance to manual McMaster counting of trichostrongyle eggs. We found that when analyzing ovine samples, the modified system performs comparably to the McMaster with respect to accuracy while operating with a higher precision.

It should be acknowledged that this study did not differentiate the strongyle eggs by species or species, and more work is needed to ensure that the algorithm will reliably recognize and count all eggs of interest. While we assume that a majority of eggs present in the sample material were of trichostrongyle origin, some could be from strongyle species,

such as *Oesophagostomum* spp., and it is possible that this could add to variability in size and shape.

Sample processing in small amounts was very different than that for equines in that the filters used for the latter were rapidly clogged when using fecal material from the former. We attribute this behavior to a difference in particle sizes in the feces of small ruminants versus equines (Clausen et al., 2013). The former engage in low-gut fermentation (rumination), which is both more efficient and time consuming than the high-gut fermentation of the latter, leading to more complex degradation of plant matter and therefore a sizeier particle size distribution in their feces. In fact, the reported mean particle sizes for ovine and caprine feces are smaller than pores of the first two filters of the SPT (Clausen et al., 2013), meaning that many of the fecal particles could potentially gain access to the final filter where they could over-whelm and eventually block it. Furthermore, the greater relative abundance of smaller particles means that a greater number could eventually be deposited on the 10- μ m mesh, thereby clogging this too. While the former problem could be addressed by modifying the pore sizes in the filter stack of the SPT, the latter could not. We ameliorated SPT clogging by removing the final, finest mesh filter from the SPT and we also addressed both problems with a single solution – that is the use of a tried, high speed centrifuge step as a filtration medium (to separate the bulk of the fecal debris from the eggs). It should be noted that, unlike FFC methods such as the Sieff and Wainman, the purpose of this step is not to float the eggs to the surface of the liquid, but rather to sediment the fecal particles to the bottom of the tube. We used an FM consisting of sodium chloride, because both zinc and magnesium sulfate, which are commonly used for this, would precipitate as metal oxides during the bleaching step of sample processing and block the FFC mesh, thereby preventing further colour exchange. Other FFCs such as aluminum solution, salt and sugar/salt mixtures that do not contain divalent cations whose oxides are insoluble could also be used for this purpose.

Initial experiments uncovered that two types of pellet formed during centrifugation (Fig. 2). One, containing the bulk of the fecal debris sedimented in the bottom of the tube and therefore retained a normal rate was denser than the FM (approximately 1.25 g/ml). However, a smaller, but significant amount of material that was less dense than the FM floated instead during centrifugation and was deposited along the inside wall of the tube. We found that the stability of the lower density pellet was dependent on the orientation of the tube when the supernatant was decanted, and the best position was to decant when the pellet was washed into the FFC prior to the rest, suggesting that it may have contained some trapped eggs whose presence or absence have an effect on the final result. However, these observations were made on a limited number of samples and so any conclusions on this point should remain tentative pending more rigorous investigation. Such investigations could be warranted because the possibility of egg loss in the lower density pellet could be relevant to both practitioners and researchers utilizing FFC-angle correctors.

As with the equine test, the automated small ruminant system produced counts that exhibited a strong positive correlation with manual counting and produced counts comparable to the McMaster (Sitarzewicz et al., 2019; Cain et al., 2020). Since the algorithm had been trained with a mixture of both ovine and caprine trichostrongyle eggs, it could not tell apart with apparent equal accuracy, since the slopes of the lines of automated vs. McMaster counts were not statistically different.

The precision of the method was, in broad terms, similar to that of the equine method, as reported previously (Sitarzewicz et al., 2019; Cain et al., 2020), in that the automated method performed with approximately twice the precision of the McMaster (half the CV) in the low and medium egg level groups. In contrast, the automated method exhibited only a 30 % improvement in precision in the high group, a difference that was not statistically significant, while in previous studies this difference was also resolved and significant. This may have been partially due to an over performance of McMaster relative to previous

Table 4

Precision of manual and automated methods using ovine samples. Fifteen ovine fecal samples were placed into three groups (high to medium to low) based on their egg content (5 per group), and were used to produce counts that were then counted, stage times used by the automated and McMaster methods. The mean results in eggs/g group for each group of eight counts are shown to the left, as are the observed coefficients of variation. The average count and CV for each group are shown in the right (upper numbers) along with the 95% confidence intervals for each mean (lower ranges).

	Manual CV		CV		Mean (CV)/95% CI		Mean CV/95% CI	
	McMaster	Automated	McMaster	Automated	McMaster	Automated	McMaster	Automated
low	391.5	391.6	35.0	15.0	253.0	245.0	22.0	14.0
	482.7	252.2	14.2	12.4				
	767.1	767.1	37.5	9.9				
	1541.0	134.2	33.4	9.0				
	274.6	224.8	22.4	22.2	1.774–4087	192.1–291.8	20.9–41.1	51–18.7
Medium	783.9	316.0	43.4	9.8	529.2	744.7	25.5	13.2
	349.4	379.0	22.3	6.2				
	288.2	288.7	25.4	8.2				
	582.5	120.0	15.5	13.4				
	189.9	711.9	14.2	13.8	244.6–181.7	845.2–444.4	14.3–15.0	7.2–11.2
High	1475.0	1452.6	15.2	11.4	2751.4	2299.0	15.1	11.2
	1244.2	1749.0	11.8	17.2				
	3728.7	3728.2	14.0	13.7				
	2238.2	1936.6	17.0	8.4				
	4812.5	4816.6	11.2	8.5	1486.1–4048.7	1491.1–2928.5	11.9–13.4	8.1–14.4

CV = coefficient of variation.

CI = 95% confidence interval.

DFG = eggs/g

samples because the high group contained two samples with substantially higher egg counts than in previous equine studies (approximately 3500 and 4000 DFG). Such high counts are not uncommon in sheep (Nisenzon et al., 1998; Le Gallouec et al., 2007), and these samples also produced the lowest McMaster CVs, presumably because stochastic subsampling variation decreases with increasing egg concentration (Jorgensen et al., 2012). Further studies using sample groups stratified at higher egg count levels could clarify whether manual and automated count precision continue to converge at extremely high egg counts. However, it should be acknowledged that an accurate and precise determination of these high counts probably has limited clinical implications, whereas performance in the lower count ranges would be more relevant, especially for older animals/sores.

In this study, precision was investigated only using ovine fecal samples. However, the facts that the algorithm was trained on a combination of ovine and equine samples and that equine and ovine samples formed statistically indistinguishable lines when compared to McMaster suggests that the ovine results presented here would be reproducible when using equine samples, although this would need to be confirmed by a specific study. Furthermore, in this study did not assess test performance on eggs from large mammals, which will be the subject of future studies. Furthermore, the algorithm can be trained to also recognize other parasite egg types such as *Neomonaxia* and *Trichostrongylus*, which should be pursued as well.

In summary, we have described the development of a rapid, automated egg counting procedure for use with small ruminants and evaluated its performance relative to manual McMaster counting using sheep feces. There was no significant difference in the accuracy of the manual and automated methods (i.e., they produced, on average, equivalent counts). However, the automated method was significantly more precise than McMaster counting at egg levels below 1000 DFG, and equally precise at higher counts.

CRDIT authorship contribution statement

M.K. Nielsen and P. Słusarczyk conceptualized the study, administered and supervised the project, P. Słusarczyk collected and counted the data, and J. Słusarczyk conducted the statistical analyses. P. Słusarczyk wrote the manuscript with input from all co-authors.

Declaration of Competing Interest

PS and MKS both hold stock in MEF Equine Solutions, LLC and PS is an employee of this company, which is commercially an automated image-based parasite egg counting technology.

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References

- Abdel, S., Hothoth, M., 2014. *Use of artificial intelligence and optical fiber detecting parasite eggs from feces in the fecal flotation of *Blasiusia* in ovine eggs to find*. *Ameghino Vet. Parasitol.* 269, 20–24.
- Berke, J.M., Miller, J.S., 2020. Distributional approach to parasite control in ruminant livestock. *Westvax Veterinary Clinics Field Manual Parasit* 36, 89–107.
- Chen, J.L., Simonsen, L., Hjaltny, H.L., Moseby, M.E., Walden, R.M., Kyndy, L.M., McElroy, J.M., Ross, J.H., Stone, A.F., Nielsen, M.P., 2020. The genetic performance of *Blasiusia*, *Trichostrongylus* and automated egg counting techniques for measurement of equine strongyle eggs in fecal samples. *Vet. Parasitol.* 254, 101793.
- Chen, J.L., Ross, A.L., Sun, P., Slobin, A., Furlong, R.W., Kyndy, R.M., 2021. The effect of ambient humidity on fecal egg counting variability. *Parasitol. Res.* 120, 1282–1290.
- Correia, D., Vercy, M., Corbe, E., 2016. A survey on anti-parasitic resistance in automated pastures of sheep in an Slovak Republic. *Vet. Parasitol.* 129, 39–49.
- Correia, M., Vercy, P., Hájek, P., Hájek, E., Kment, A., Vercy, J., Antonic, S.L., Hájek, J., 2015. *Effect of climate and climate change on the prevalence of parasitic diseases*. *Comp. Contin. Parasitol. A Res. Image J Parasitol.* 129, 182–192.
- Corbe, E., Long, E.L., White, C.S., Gross, S.L., Somers, S.L., 2016. Analysis of method related on cases for single field of view parasite egg microscopy. *J. Parasitol.* 297, 512–521.
- Cox, D.D., Todd, A.C., 1982. Survey of gastrointestinal parasites in Wisconsin dairy cattle. *J. Am. Vet. Med. Assoc.* 121, 706–709.
- D'Amico, G., Khalil, L., Mura, M.P., Lutzini, L., 2013. *ELISA and molecular biology techniques for qualitative and quantitative diagnosis of giardiasis of parasites in animals and humans*. *Hum. Parasit.* 4, 202–215.
- D'Amico, G., Mura, M.P., Lutzini, L., Tassi, A., Taramino, L., Lutzini, L., Khalil, L., 2013. *The Mini-ELISA test for the diagnosis of lamblia and giardia infections in humans and animals*. *Am. J. Parasitol.* 11, 1253–1211.
- D'Amico, G., Mura, M.P., Mura, M.P., Lutzini, L., Tassi, A., Tassi, A., Khalil, L., Khalil, M., Khalil, M., 2015. *Automated fecal flotation microscope (AFM): a new device for reference for helminth egg counts*. *Parasitology* 145, 437–451.

- de Lima, D.G., de Bortoli, L.G., Araújo, N.A., Palomares, J.L., Oliveira, Jr., A.C., de Melo, M.F., Simões, C.B.: 2014, 'Ameliorating efficiency and environmental practices in deep-sea fishing in state of Rio de Janeiro, Brazil', *Environ Biol Fish* 100, 100–104.
- Hilborn, R., Waldman, J., Borchert, B., Torres, J.S., de Melo, V., de Paula, T., de Sousa, P., Costa, Z., Sousa, Z., Sobral, B., de Ward, J.: 2009, 'Fishery management for small, wide-spread, low-value fishes for small-scale fisheries in small-scale fisheries', *Environ Biol Fish* 102, 1599–1605.
- Jenny, M.J.: 2004, 'Fishery Management Reference Manual', 5th edition, Iowa State University Press.
- Jordan, R.H., Winfield, S.C.: 1999, 'A new technique for counting otoliths: egg and embryo counts', *J. Great Lakes Res.* 25, 50–52.
- Marino, M., Zaccaro, F.A., Di Stefano, G.F., Scudillo, G., Giamberini, S., Giamberini, F., Simeone, A.P.: 2012, 'Spatio-temporal dynamics of population with multiple recruitment pulses: the case of the bay of Naples (southern Italy)', *Environ. Biol. Fish.* 9, 29–39.
- Moreira, S., Jorde, J.M., Miller, A.L., Patti, T.L., Gascón, R., Williams, K.J., Wilson, L.R., Zito, A.M., Eklöv, P.M.: 2005, 'Resilience of recruitment response to deep-sea fishery in the southern United States', *J. Am. Wetl. Manag.* 25, 1234–1242.
- Ockler, F., Corp, S.A.: 2010, 'The development of multivariate resource-use deep-sea fisheries', *Oceanology* 150 (suppl), 879–884.
- Ota, J.T., Wengrow, B.G.: 2012, 'Invertebrate resource management in the deep-sea by the Hawaiian deep-sea industry', *Sci. Nat.* 100, 35–41.
- Samuel, C., Jaramilla, S., Hernandez, K., Garcia, M.: 2014, 'Spatio-temporal dynamics and environmental responses in young cohort of the resource fishery, *Merluccius* sp. in southern Chile', *BMC Ecol. Evol.* 15, 203–222.
- Ta, Jernan, J.H., Borchert, S., Davis, C.J., Hertzler, L.M., Custer, G.S.: 2007, 'Adopting an egg count for total indices in deep-sea fisheries', *ICES J. Mar. Sci.* 64, 148–155.
- Witte, D.W., Vaughan, L.S., Lewtasick, D.M., Conry, P.M., Doherty, G.M., Warner, G.S.: 2013, 'The genetic diversity of *Merluccius* in the Irish Sea', *Environ. Biol. Fish.* 106, 379–384.
- Aguiar, V., de la Sotilla, E., Ostano, A., Palma, A., Cruz, J., Ferrero, R., Lopez, R.N., Soto, E.L., Gonzalez, B.: 2015, 'Evaluation of the VTKGAN (WAGAN) as a multi-scale and time-fair genetic diversity system integrated with the existing algorithm', *Environ. Genom.* 11, 200–210.
- Almeida, J.P., Sidorowicz, M.A., Wiegman, G.C.: 2015, 'Population genetic structure and genetic diversity of the Atlantic haddock (*Merluccius merluccius*) and *Merluccius aeglefinus* in the North Atlantic', *Sci. Nat.* 103, 41–48.
- Borner, M.H., Steiner, M.A., Wawry, S., Menni, G., Campbell, S.D., Vaughan, J.L., Jellie, A.: 2010, 'Comparison of mtDNA and BEZSN/2012 methods for counting otoliths in the larvae of haddock', *Environ. Biol. Fish.* 11, 229–237.
- Brody, L., Coull, A., Brannon, E., Faxon, A., Chiriboga, M., Leibel, A., Mee, J., M.F., Simons, G., Shuter, B., Grigg, S., Jones, P., Guitton, G.: 2013, 'Deep-sea fisheries: a global overview and future prospects', *Environ. Biol. Fish.* 98, 299–300.
- Costa, J.A., Simões, C.B., P. 2012, 'Merluccius', *Environ. Biol. Fish.* 2012, 'Evolutionary history and genetic structure of the eastern Atlantic haddock, *Merluccius merluccius*, based on mitochondrial DNA D-loop analysis', *Sci. Nat.* 100, 41–49.
- Chakravarti, P., Pagani, S., Miller, G., Papp, G., Chow, E.K., Nandakumar, M., Sedgwick, W., Shaban, R.K.: 2013, 'Autosomal parent-child egg-counting using the common Y-chromosomal marker system and complementary X-chromosomal loci', *Environ. Biol. Fish.* 98, 409–417.
- Chakravarti, M., Chakravarti, P., Nanda, M.K.: 2010, 'The error in counting markers on chromosomes with a marker set polymorphic', *Stat. Probab. Lett.* 82, 1565–1570.
- Mill, R.H.: 1922, 'Investigations on the natural history of the haddock, *Merluccius merluccius* (L.)', *Environ. Biol. Fish.* 102, 1599–1605.
- Torgersen, P.R., Paul, M., Jones, P.J.: 2012, 'The contribution of genetic diversity sampling to observed variability in larval egg counts', *Environ. Biol. Fish.* 101, 399–400.
- Witte, P.J.: 1981, 'The development of variability evidence in *Merluccius* (haddock) stocks', *ICES J. Mar. Sci.* 40, 223–232.