

## Comparison of Direct Microscopy and Molecular Method to Detect Amoebiasis Cases from Stool Specimen and Also Identify the *Entamoeba* Species Involved In Infection: A Study of Nepal

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### ABSTRACT

Direct microscopic examination of stained or unstained wet mount preparations or fixed stained smears of clinical material can often provide the etiological diagnosis of an infectious process. A total of 266 human stool specimens from children fewer than 15 years were examined for the presence of amoebiasis by a combination of microscopic examination and molecular method. In molecular method, nested polymerase chain reaction (Nested-PCR) targeting genomic *Entamoeba* species was used. Stool specimens were collected from 266 participants in Southern plains of Nepal and analyzed at Kathmandu Centre for Genomics and Research Laboratory. The stool specimens were processed by wet mount method using saline as well as iodine staining and examined via microscopy for the presence of *Entamoeba* cysts or trophozoites. Furthermore, the stool specimens were characterized using Nested-PCR targeting genomic *Entamoeba* species. Based on microscopic examination, overall prevalence of *Entamoeba* infection was 17.6% (47/266). The PCR results showed that 52(19.5%) out of 266 stool specimens successfully generated species-specific amplicons. Males (21.7% in PCR) were more commonly infected compared to females (16.6% in PCR). Comparison by age groups showed that 10-15 years age-group (26.6% in PCR) had higher infection than age-group 5-10 years (16.6%) years and 1-5 years (15.2%). The infection with *E. histolytica* (100%; 52/266) was the predominant cause of amoebiasis, while infection with *E. dispar* and *E. moshkovskii* was not found. PCR is a more effective method for the identification of *Entamoeba* infection than microscopy. This study also suggests that older children especially male are more prone to *Entamoeba* infection.

**Key words:** Amoebiasis, wet mount method, microscopy, nested-PCR, Nepal,

### INTRODUCTION

*Entamoeba histolytica* is the causative agent of amoebiasis and this is one of the leading parasites that cause human mortality (Haque *et al.*, 2003, Haque *et al.*, 2006 and WHO 1997). Each year, the parasite invades approximately 50 million people resulting in 100,000 deaths (WHO 1997). Likewise, the productivity of animals is constrained due to *Entamoeba* infection that can have extensive consequences, ranging from reduced animal performances to mortality (Sykes, 1994; Waller, 1999).

Although the parasite is distributed worldwide; the prevalence rate is higher in the developing countries (Stanley, 2003). *Entamoeba* infection spreads through fecal-oral route (Blessmann *et al.*, 2003 and Fotedar *et al.*, 2007). Open defecation and improper disposal waste water often lead to contamination of food and water sources in developing countries and this is a serious issue contributing to human morbidity (Estevenz *et al.*, 1983 and Gyawali *et al.*, 2009).

Similarly, grazing management has been a serious issue behind GI infection in animals since pastures are usually not provided or not properly managed.

*Entamoeba histolytica* causes gastrointestinal (GI) diseases in farm resulting in reduced productivity, weight gain as well as milk production (Jittapalapong *et al.*, 2011).

Moreover, in some parts of the world, wastewater and human and animal excreta are used as fertilizer to increase the crop yield. This is normally practiced in China, South East Asia as well as various areas in Africa (Drechsel *et al.*, 2011; Cross, 1985 and Timmer and Visser, 1998) with severe water scarcity. Such practices can easily contaminate food sources and spread GI infection to humans and animals. Furthermore, lack of hygiene and poor hand washing practices among the farm workers play a major role in increasing the infection rate. Generally, *E. histolytica* is the common pathogen that can cause such GI diseases.

The diagnosis of an intestinal infection with *E. histolytica* is commonly made by microscopic examination of stool, in which one must recognize and differentiate the cysts or trophozoites of *E. histolytica* from those of morphologically different species. The identification of these cysts and trophozoites requires a lot of skill and patience by the microscopist. To overcome the difficulty in recognizing the pathogens and also know the species of the pathogen, molecular methods are increasingly being used for research purposes. Some molecular methods, including conventional and real-time PCR, have been developed for the detection and differentiation of *Entamoeba* at species level from clinical samples (Hamzah *et al.*, 2006).

Polymerase Chain Reaction (PCR) assay allows the in vitro amplification of a specific DNA fragment in a cyclic process of denaturation, hybridization of primers, and elongation of the DNA strand using a thermostable DNA polymerase. In nested PCR, amplicons from a PCR are used as the template in a second PCR using one primer or two primers different from those used in the initial PCR and located within the sequence amplified by the first primer set. Nested PCRs are used to increase sensitivity and specificity (Messmer *et al.*, 1997).

The study is supportive to describe the current status of amoeba infection in the endemic areas. The study has few general and specific objectives, in general it detect amoeba infection using microscopy and Polymerase Chain Reaction and also identify the *Entamoeba* species in the stool specimen and in particular it compares the results of PCR and microscopy for the detection of *Entamoeba* infection also find the *Entamoeba* species in stool specimen.

## MATERIALS AND METHODS

This cross sectional study was carried from 15 July to 18 January 2017. For this study, stool specimens were collected from hospitals and health centers in southern plains of Nepal and analyzed in Kathmandu Centre of Genomics and Research Laboratory (KCGRL), Lalitpur, Nepal. Nepal lies in sub-tropical zone of Asia sharing its open border with India. Due to compromise in drinking water

supply and unmanaged human settlements in southern plains, prevalence of amoebiasis is higher especially in young children who play in soil most of the time. The samples were taken from patients complaining diarrhoea from age 1 to 15 years. A total of 266 specimens were collected and processed at KCGRL. Samples from small kids were taken by the help of their care-takers. The filled and well-labeled containers were received and analyzed by microscopy on the following day; the samples were mixed with formalin and then stored under deep freeze for further processing.

### **Microscopic Examination**

Microscopy is inexpensive and rapid technique for the identification of *Entamoeba* parasite. It is the gold standard tool for the diagnosis of amoebiasis cases and a method of choice for many hospitals. Saline wet mount of fresh unpreserved stool samples were examined microscopically for demonstrating *Entamoeba* cysts and trophozoites (Parija and Prabhakar 1995). Similarly, iodine wet mounts were prepared by adding approximately 2 mg of stool to a drop of Lugol's iodine (diluted 1:5 with distilled water) on a glass microscope slide and placing a cover slip on the stool suspension. These wet mounts were microscopically examined initially by using a low-power (10×) objective and then using a high-power (40×) objective of a compound light microscope. Following the extraction of genomic DNA for protozoan parasite from all the stool specimens, the specimens showing the presence of

protozoan DNA were further re-checked for *Entamoeba* species using nested-PCR.

### **Extraction of *Entamoeba* genomic DNA**

gDNA was extracted from suspected faecal specimen using mdi Stool Genomic DNA (gDNA) Miniprep Kit according to the manufacturer's instructions. Approximately, 180-220mg faecal sample was taken in a 2ml microcentrifuge tube and then mixed with 1.2 ml lysis buffer MSL. Then the mixture was thoroughly homogenized and incubated at 70°C for 5 minutes. Followed by 15 seconds of vortexing and centrifugation at 10000 rpm for 5 minutes to pellet stool sample, 15 µl Proteinase k was pipetted into a new 1.5ml microcentrifuge tube. Now 200 µl supernatant from centrifuged sample was mixed with Proteinase k in new centrifuge tube and also added 200 µl Buffer SB. After short vortexing, this new mixture was incubated at 56°C in water bath for 15 minutes. Briefly centrifuge the mixture after incubation, add 200 µl Ethanol, mix and briefly centrifuge again. Then the mixture was passed into collection tube through spin column by spinning at 10000 rpm for 1 minute. Here, the spin column was washed in the new collection tube with 500 µl of buffer SW1 by centrifuging for 1 minute at ≥10000 rpm. This step was again repeated using 500ul of buffer SW2 but centrifugation was done for 3 minutes. Furthermore, the bound gDNA was eluted in the spin column placed in fresh 1.5ml microcentrifuge tube by adding 200 µl of buffer SE. After incubating for 2 minutes, it was spinned ≥10000 rpm for 1

minute. Hence obtained product possessed gDNA of the processed sample. DNA thus extracted was stored at  $-20^{\circ}\text{C}$  for PCR amplification.

### **Nested PCR amplification**

Nested PCR targeting genomic DNA was used to genetically characterize *E. histolytica*, *E. dispar* and *E. moshkovskii*. The PCR was carried out in a 25  $\mu\text{l}$  volume with the final mix containing 2.5  $\mu\text{l}$  PCR buffer (10 $\times$ ), 1.25 mM dNTPs (0.5  $\mu\text{l}$ ), 25 mM  $\text{MgCl}_2$  (2.5  $\mu\text{l}$ ), 10 pmole of each primer (0.15  $\mu\text{l}$ ), 2.5 U of Taq polymerase (0.3  $\mu\text{l}$ ), 2.5  $\mu\text{l}$  of DNA template and 16.4  $\mu\text{l}$  Nuclease free water.

Primary PCR for the detection of *Entamoeba* genus used forward primer E-1 (5'-TAA GAT GCA GAG CGA AA-3') and reverse primer E-2 (5'-GTA CAA AGG GCA GGG ACG TA-3'). For primary PCR, the master mix was heated to  $96^{\circ}\text{C}$  for 2 min, followed by 30 cycles of  $92^{\circ}\text{C}$  for 1 min (denaturing),  $56^{\circ}\text{C}$  for 1 minute (annealing),  $72^{\circ}\text{C}$  for 1 minute 30 s (extension) and a final extension at  $72^{\circ}\text{C}$  for 7 minutes.

Similarly, the products of primary PCR were used for *Entamoeba* species-specific characterization. Amplification was achieved using primer sets EH-1 (5'-AAG CAT TGT TTC TAG ATC TGA G-3') and EH-2 (5'-AAG AGG TCT AAC CGA AAT TAG-3') to detect *E. histolytica* (439 bp); ED-1 (5'-TCT AAT TTC GAT TAG AAC TCT-3') and ED-2 (5'-TCC CTA CCTATT AGA CAT AGC-3') to detect *E. dispar* (174 bp); Mos-1 (5'-GAA ACC AAG AGT TTC ACA AC-3') and Mos-2 (5'-CAA

TAT AAG GCT TGG ATG AT-3') to detect *E. moshkovskii* (553 bp) (Khairnar and Parija, 2007).

For secondary PCR, the master mix was heated to  $96^{\circ}\text{C}$  for 2 min, followed by 30 cycles of  $92^{\circ}\text{C}$  for 1 min (denaturing),  $48^{\circ}\text{C}$  for 1 minute (annealing),  $72^{\circ}\text{C}$  for 1 minute (extension) and a final extension at  $72^{\circ}\text{C}$  for 7 minutes. For both amplifications, master mix was incubated in the thermal cycler (Palm cycler). The secondary PCR amplicons were subjected to electrophoresis in 2% agarose gel stained with ethidium bromide (EtBr) at 80V for 60 minute and visualized in UV illuminator.

## **RESULTS AND DISCUSSION**

*Entamoeba* species was detected under microscope on the basis of morphological characteristics of the cysts or trophozoites. The average cyst size ranges from 10–20  $\mu\text{m}$  in diameter. Qualitative data were estimated and presented as frequencies and percentage by using univariate analysis through SPSS statistical tool.

### **Prevalence of *Entamoeba* Infection via Microscopy and PCR**

A total of 266 stool samples collected were processed through microscopy and PCR methods separately and results thus obtained varied a lot. Out of all the stool samples processed, 47 (17.6%) samples were microscopically positive for *Entamoeba* cysts (Table 1). However, PCR results showed 52 (19.5%) positive cases of *Entamoeba* infection

which is higher than microscopy result (shown in Table 1).

Table 1. *Entamoeba* cysts analysis

Characteristics	Total (N)	Positive case(n)	%
Microscopy	266	47	17.6
PCR	266	52	19.5

Ngui *et al.* (2012) found that 48 (15.3%) children out of 313 children based on microscopy result suffered from amoebiasis. The parasitic infections are even higher in animals. In a study done by Huang *et al.* (2014) in dairy cattle in Vietnam, infection rate of protozoa alone was 81.3%.

#### Sex-wise Prevalence of *Entamoeba* Infection

Microscopy result showed *Entamoeba* infection was more in males

30(19.7%) compared to females 17 (14.9%), however, it was not statistically significant (data not shown). Similarly, PCR result showed *Entamoeba* infection was more in males 33(21.7%) compared to females 19 (16.6%), as shown in table 2. In contrast to this result, Ngui *et al.* (2012) found 75(17.6%) of the causative agents of illness were *E.histolytica/E.dispar* in microscopy in which infection was more prevalent in females (19.1%) compared to males (15.9%).

Table 2. *Entamoeba* infection by sex

Sex-wise	Total (N)	Microscopy			PCR	
		Positive Case(n)	%	95% CI	Positive Case(n)	%
Male	152	30	19.7		33	21.7
Female	114	17	14.9		19	16.6
Total	266	47	17.6		52	19.5

#### Age-wise Prevalence of *Entamoeba* Infection

Age-wise study of *Entamoeba* infection detected via microscopy and PCR methods showed following results as shown in Table 3.

Table 3. *Entamoeba* infection by Age-wise

Age-group (year)	Total(N)	PCR			
		Positive Case(n)	%	Positive Case(n)	%
1-5	59	8	13.6%	9	15.2%
5-10	132	21	15.9%	22	16.6%
10-15	75	18	24%	20	26.6%
Total	266	47	17.6%	52	19.5%

This table states that out of total samples processed through microscopy and

PCR, highest prevalence was found in age-group 10-15 years i.e. 18 (24%) and 20

(26.6%) respectively. But the age-group 1-5 years had least prevalence of 8 (13.6%) and 9 (15.2%) in microscopy and PCR respectively.

The PCR results were positive for only one species of *Entamoeba* i.e. *E. histolytica*. Other species i.e. *E. dispar* and *E. moshkovskii* were not detected in any of the samples. Thus the chances of mixed infection were also ignored.

Although, the microscopic observation showed 47 positive cases of *Entamoeba* infection out of 266 total stool specimens, it was not considered overall prevalence in this study population. The present study found an overall prevalence of *Entamoeba* infection as determined by PCR as 17.6% (52/266). Thus the molecular method has been found more effective in contrast to microscopy. This

study also suggests that relying only on microscopy cannot diagnose accurate cases of amoebiasis.

A study conducted on children in Kathmandu, Nepal reported 10% (50/500) of the causative agents of diarrhoea were *E. histolytica* (Pokharel *et al.*, 2007) which is consistent with this study. A recent study in India also has more or less similar results of 19% positive *Entamoeba* infection based on microscopic examination (Nath *et al.*, 2015). In a study conducted by Tuzemen and Dogan (2015), among 354 stool samples, 84 (23.7%) were found *E. histolytica/E. dispar* positive by direct microscopy and 9 (2.5%) by PCR methods. Although there is some similarity with the microscopy result with the current study but PCR result has a big difference.

Table 4. Molecular technique of detecting other *Entamoeba* species

Age group (year)	Total (N)	PCR			%	
		Positive case (n)	<i>E. histolytica</i>	<i>E. dispar</i>		<i>E. moshkovskii</i>
1-5	62	9	9	0	0	15.2%
5-10	137	22	22	0	0	16.6%
10-15	81	20	20	0	0	26.6%
Total	280	52	52	0	0	19.5%

This molecular technique showed that *E. histolytica* (100%) was the only agent among *Entamoeba* species causing illness in the study population. *E. dispar/E. moshkovskii* were not detected in any of the samples taken from symptomatic cases (shown in Table 4). From this finding, it can also be predicted that the causative agents of diarrhoea were either *E. histolytica* or bacteria or virus or some other agents but not *E. dispar/E.*

*moshkovskii*. Since, molecular studies targeting the *Entamoeba* species have not yet been done in Nepal; it is very difficult to understand the prevalence due to *E. dispar/E. moshkovskii* in Nepal. *E. dispar/E. moshkovskii* although are non-pathogenic, they have already been identified in neighboring countries (India and Bangladesh) (Khairnarnar, Parija 2007; Ali *et al.*, 2003) and thus it can be

speculated that they may exist in Nepal even though not identified in this study.

Although human cases infected with *E. moshkovskii* have been reported sporadically from different parts of the world including India (Khairnar and Parija 2007) Bangladesh (Ali *et al.*, 2003) and Australia (Fotedar *et al.*, 2007), they have been attributed to asymptomatic infection most frequently. The global prevalence of *E. dispar/E. histolytica* is quite uncommon with this study but it cannot be ignored that a single study with limited number of subject's i.e. only children may not be sufficient to assess the overall prevalence of *Entamoeba* species over certain geographic location. Therefore, future investigation which includes the clinical impact of *E. moshkovskii* and other *Entamoeba* species is imperative for a better understanding of a true pathogenic potential of *E. moshkovskii*.

Sex-wise study shows higher prevalence in males in comparison to females both in microscopy and PCR. This could be attributed to the fact that male members have freedom to participate in outside activities in southern plains of Nepal while the female children have family restriction and need permission of their parents to move out of their house. This situation play a bigger role in transmitting the amoebiasis in male children than female children as one would eat contaminated food or drink water while they are outside their houses. Furthermore, open defecation and eating without washing the hands has further impact of being infected.

Age-wise study shows increasing prevalence of amoebiasis with increasing age of children. The possible reason for this may be more or less the same as sex-wise prevalence. As a child grows older, he can go outside their house to play with their friends, swim in the nearby rivers or lakes which are already polluted by open defaecation. Moreover, there is no provision of clean drinking water in most of the schools of Nepal as well. Contaminated water or food may contain *Entamoeba* cysts and can easily transmit the infection (Chakraborty, 2001).

The participants of this study were from the southern plains of Nepal where there is no easy access to clean drinking water and living condition is poor. Given the faeco-oral route of amoebiasis, habits related to eating, defaecation, personal hygiene, cleanliness and level of education may have an impact on the prevalence rates (Nguí *et al.* 2012). Because of such unhygienic practices, the chances of parasitic infections have persisted (Sherchand SB 1997 and Maharjan *et al.*, 2007). The majority of the people in Nepal still live in the rural areas where literacy rate is quite lower. Unavailability of medical facilities and social awareness in such regions may contribute to high prevalence of *Entamoeba* infection.

## CONCLUSION

In conclusion, PCR is a more effective method for the identification of *Entamoeba* infection than microscopy. In the southern plains of Nepal, *E. histolytica*

is the major cause of amoebiasis and male children are more prone to *Entamoeba* infection. Besides, older children are more vulnerable to amoebiasis than young children.

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