Detection of giardia lamblia genotypes in sewage and in stool samples in Israel

Abidelfatah M Nasser1*, Neta-Lee Benisti1,2, Marina Taran-Benshoshan1,2, Valeria Kravitz1 and Yeshayahu Nitzan2

1Water Quality Research Laboratory, Ministry of Health, Israel
2Faculty of Life Sciences, Bar-Illan University, Ramat Gan, Israel

Abstract

Giardia lamblia is a protozoan parasite which causes a severe disease called Giardiasis. This study was conducted to determine the prevalence of Giardia cysts in raw wastewater and its prevalence in the study community. Furthermore, the prevalence of assemblages A and B in sewage was compared with their prevalence in stool samples tested positive for Giardia. All wastewater samples were found positive for Giardia at a concentration range of 10-12225 cysts/L. Positive stool samples contained Giardia at an average concentration of 1.4x10^5 cysts/gr. Giardia assemblage A alone was detected in 38.2% of wastewater samples, whereas assemblage B was not detected separately. However, 61.8% of the samples were found to contain both assemblages. In stool samples, the majority 27 out of 50 (54%) were found to contain assemblage B, 34.6% contained assemblage A and only 11.5% contained a mix of both assemblages. The results of the study indicate that Giardia is highly prevalent in wastewater in Israel reflecting its prevalence in the community. In addition, assemblage A is highly prevalent in wastewater, whereas assemblage B is more prevalent in stool samples, suggesting milder and/or carrier state of infection for assemblage A.

Introduction

Giardia intestinalis (also known as G. lamblia and G. duodenalis) is a waterborne protozoan parasite [1]. Giardia causes a self-limited clinical illness (giardiasis), typically characterized by diarrhea, often associated with dehydration, bloating, weight loss, abdominal pain, nausea, malabsorption and vomiting; asymptomatic infection also occurs frequently [2]. The transmission of the protozoan parasites occurs mainly through the fecal-oral route, person to person or by exposure to contaminated water and food [3–5]. The infectious dose is low; ingestion of 10 cysts has been reported to cause infection in humans (Rendtorff 1954) [6]. Infected persons have been reported to shed 10^8–10^9 cysts per day and excrete cysts for months [6].

The prevalence and concentration of Giardia cysts in raw wastewater depend on several factors such as sanitation conditions, socioeconomic status, levels of annual precipitation and on the sensitivity of the detection methods. Giardia cysts have been reported to be prevalent at higher levels in raw wastewater samples analyzed in various parts of the world. Giardia cysts were detected in almost 100% of the analyzed raw wastewater samples in various countries at a concentration range of 2 to 13,600 cysts/L and no substantial differences were observed in the concentration of Giardia cysts in raw wastewater from various countries [7].

A considerable amount of data has shown that Giardia should be considered as a species complex that comprises at least eight distinct genetic groups, referred to assemblages A to H [8–10]. To date, only assemblages A and B have been associated with human infections. Genotypes C to H are known as pathogens of animals. However, genetic traits that influence the virulence and other aspects of the infection are unknown and efforts to correlate the parasite genetic make-up and the clinical symptoms in the host have generated controversial results [11,12]. This study was performed to determine the prevalence and concentration of Giardia assemblages in Giardia positive stool in comparison with assemblages present in wastewater.

Citation: Nasser AM, Benisti NL, Benshoshan MT, Kravitz V, Nitzan Y (2020) Detection of giardia lamblia genotypes in sewage and in stool samples in Israel. J Clin Microbiol Biochem Technol 6(1): 029-032. DOI: https://dx.doi.org/10.17352/jcmbt.000040
Material and Methods

Raw wastewater and stool samples

Forty-two raw wastewater samples of one liter were collected from two wastewater treatment plants (WWTP), plant A serves 240,000 and plant B serves 120,000 inhabitants. Grab samples were collected once a month in 2012 and 2013. Giardia cysts were concentrated from 0.2 L of raw wastewater samples by centrifugation for 15 minutes at 11,000 x g. Fifty coded clinical stool samples (1 ml), positive for G. intestinalis were obtained from Clalit Health Services Laboratories. Concentrated parasites from stool and wastewater samples were affinity purified and concentrated by Immunomagnetic Separation (IMS) using magnetic beads coated with anti-Giardia monoclonal antibodies (Dynal, Lake Success, N.Y.) as previously described in method[16:23:1 of the USEPA [13]. At this stage, 20% of the purified sample was used for immune-fluorescent staining and microscopic enumeration of Giardia cysts. In parallel, the remaining 80% of the concentrated sample (complex of immune magnetic beads and parasites) was stored at –80°C for later DNA extraction for Giardia detection by PCR and genotyping. The recovery efficiency of Giardia cysts from raw wastewater samples was 49±14.5%.

DNA extraction from giardia cysts

DNA was extracted from Giardia cysts bound to the immune magnetic beads without dissociation of cysts using the QiAamp DNA mini kit (QIAGEN, Valencia, Calif.) according to the manufacturer’s instructions with minor modifications. Briefly, 180 μL ATL buffer, from the kit, was added into 1.5-mL tubes containing the beads and subjected to ten freeze–thaw cycles in liquid nitrogen. To break the cysts wall, samples were incubated in 4 U μL–1 protease (Sigma-Aldrich, St. Louis, MO. USA) for 10 minutes at 56°C. DNA extraction from the cysts was then performed following the manufacturer–recommended protocol. After purification, a 200 μl DNA sample was stored at –80°C for later analysis.

PCR amplification

Extracted Giardia DNA from stool or wastewater samples was subjected to PCR amplification of β-Giardin gene fragment of 759 and/or a 384 bp fragment as described by Caccio, et al. 2002 [14]. These fragments were utilized to differentiate assemblage A and assemblage B as determined by PCR–RFLP. PCR was conducted as previously described by Caccio, et al. 2002 [14] and Bovine Serum Albumin (BSA) was added to the PCR mixture at a concentration of 400ng/μL to prevent inhibition of the DNA–polymerase activity. The PCR products were visualized by 1.5% agarose gel electrophoresis and staining with ethidium bromide under UV irradiation.

RFLP - Restriction Fragment Length Polymorphism analysis

The PCR products were subjected to digestion by restriction enzyme HaeIII (FastDigest HaeIII, Thermo Scientific) as previously described [15]. This restriction enzyme produces specific fragments which enable the classification of the detected species as assemblages A and B. Restriction enzyme HaeIII produces the following fragment sizes (bp) 202, 201, 150, 126, 70, 34, 4 with Giardia duodenalis Assemblage A and 202, 176, 150, 117, 84, 34, 24 Giardia duodenalis Assemblage B (Figure 1). The results of some samples were a mixture of the fragments for assemblage A and B, as shown in Figure 1.

Results

Forty two raw wastewater samples from two treatment plants were analyzed and all (100%) samples were found positive for Giardia cysts. The concentration of Giardia cysts in raw wastewater ranged between 10 to 12,225 cysts/L and the average concentration was 3685 cysts/L (Table 1).

In stool samples positive for Giardia, assemblage B was dominant and was detected in 27 out of 50 (53.85%) samples (Figure 2). Assemblage A was less prevalent than assemblage B in Giardia positive stool samples and was detected in 17 out of 50 (34.62%) samples. Assemblage A and B mixture was detected in 6 out of 50 (12%) of the stool samples.

Figure 3 presents the results of the relationship between the Giardia assemblage present in the stool samples and the level of Giardia cysts detected. The majority (71.4%) of stool samples of assemblage B contained Giardia at a concentration range of 10 to 10^4 cysts/gr, whereas, 66.4% of stool samples of either assemblage A or of the mixture contained Giardia at a concentration range of 10^4 to 10^6 cysts/gr. The results show that stool samples of assemblage B contain one to two orders of magnitude Giardia cysts greater than samples of assemblage A.

The results of genotyping of Giardia lamblia cysts present in raw wastewater have shown that in 16 out of 42 (38.2%) of the raw wastewater samples assemblage A was the only genotype detected. Whereas, assemblage B alone was not detected in any one of the raw wastewater samples. A mixture of assemblages A and B was detected in 26 out of 42 (62%) of the raw wastewater samples. The results demonstrate that assemblage A of Giardia is more prevalent than assemblage B.

Table 1: Prevalence and concentration of Giardia cysts in raw wastewater.

<table>
<thead>
<tr>
<th>% positives</th>
<th>WWTP-A</th>
<th>WWTP-B</th>
</tr>
</thead>
<tbody>
<tr>
<td>Avg. concentration cyst/L</td>
<td>3584</td>
<td>3786</td>
</tr>
<tr>
<td>Mean concentration cyst/L</td>
<td>2315</td>
<td>3160</td>
</tr>
<tr>
<td>Concentration range cyst/L</td>
<td>10-11,875</td>
<td>60-12,225</td>
</tr>
</tbody>
</table>
in raw wastewater samples which may reflect its circulation in the community.

Discussion

All tested wastewater samples were found positive for Giardia cysts at a concentration range of 10 to 12,225 cysts/L and the mean concentration was 3160 cysts/L. The results are within the range reported previously and reviewed by Nasser, et al. 2012 [7]. The concentration range reported by Nasser, et al. 2012 [7] was up to 100,000 cysts/L with a prevalence of 50 to 100%. The variation in the prevalence and levels of Giardia cysts detected in raw wastewater may be the result of sanitation conditions, wastewater quality and the sensitivity of the detection methods [7].

Assemblage B was the dominant G. duodenalis genotype detected in clinical stool samples and it was detected in 53.8% of the samples, whereas, assemblage A was detected in only 34.2%. The higher prevalence of assemblage B in stool samples can stem from differences of the symptoms caused by the two genotypes. Conflicting results have been reported concerning the correlation between disease and Giardia genotype in humans. A survey conducted in Ethiopia found a significant correlation between symptomatic infection and the presence of assemblage B [16]. A similar correlation was reported by Homan & Mank, 2001 [11], with assemblage B isolates associated with persistent diarrhea, whereas assemblage A infections were associated with intermittent diarrhea [11]. On the other hand, Sahagun, et al. 2008 [17], found a strong correlation between symptomatic infection and assemblage A2 in patients from Spain [17]. Interestingly, the proportion of asymptomatic to symptomatic infections with assemblage A was similar for three studies from Ethiopia, Bangladesh and Spain, 62%, 57% and 67%, respectively [16-18]. The key difference was that 100% of detected assemblage B infections reported from Ethiopia were associated with diarrhea, compared with 16% of infections resulting in diarrhea reported from Bangladesh and 42% from Spain [16-18]. It is expected that patients with severe diarrhea may seek medical help and stool samples may be submitted for diagnosis, whereas in cases of mild, intermittent and asymptomatic infection, the incidence may go undetected. It is worth noting that patients infected with assemblage B of G. duodenalis shed on average twice greater concentration of cysts compared with those infected with assemblage A which makes the microscopic detection easier. This can be one of the reasons for higher parentage of assemblage B detected in the stool samples.

Both Assemblages A and B were detected as mixture in 62% of the tested wastewater samples and the rest of the samples (38%) contained assemblage A only. None of the wastewater samples contained assemblage B alone. This indicates that assemblage A is dominant in the studied raw wastewater. Bertrand & Schwartzbrod [19] found that assemblages A and B were detected in urban wastewater with a predominance of assemblage A [19]. G. duodenalis assemblage A was detected in 92% and 100% of wastewater samples of domestic wastewater treatment plants, whereas assemblage B was detected in 42% and 92% of the same samples [19]. Similar results were reported by Caccio, et al. 2003 [15] who reported the presence of assemblage A in 16 out of 16 (100%) wastewater samples, while assemblage B was detected in only 50% of the samples [15]. Sulaiman, et al. 2004 [20] reported the presence of G. duodenalis assemblage A in 85% of raw urban wastewater samples, whereas assemblage B was detected in only 15% of the samples [20]. These results were supported by Robertson, et al. 2006 [21], who reported on the predominance of G. duodenalis assemblage A in raw domestic wastewater in Norway [21]. In contrast with the results observed with urban wastewater samples, the genotyping of G. duodenalis cysts from human feces classified a majority of samples as assemblage B [22]. This difference between environmental and clinical samples has also been observed in other studies [21,23]. It is possible to use the levels of G. duodenalis in raw wastewater to estimate the circulation of the parasite in the studied community. Since giardiasis is reportable disease in many countries, it is important to evaluate the real infection rate of giardiasis based on the data established
by wastewater monitoring. By examining wastewater samples Oda, et al. [24], estimated the prevalence of *Giardia* infection in inhabitants in the two study areas, in Japan to be 0.1-1.1 and 0.1-1.3% [24]. In 2010, 1956 cases of giardiasis were reported to the Department of Epidemiology of the Ministry of Health in Israel, which means. 27.9 cases of giardiasis per 100,000. However, according to the concentration of *Giardia* cysts in raw wastewater, the incidence was 1.6% of the population, indicating that reporting of giardiasis cases to the Department of Epidemiology underestimates the prevalence of giardiasis in the population [7]. Giardiasis reported in the USA is lower than that reported in Israel (average 6.2 per 100,000 as compared to 27.9), these figures are also an underestimation of the number of giardiasis cases, since the mean concentration in sewage was reported to be about 5000 cysts/L [20]. Under-reporting can stem from various reasons such as: asymptomatic infection of giardiasis, low levels of excreted cysts in stool samples and the low efficiency of the reporting program.

### Conclusions

The results presented demonstrate that *G. lamblia* is highly prevalent at high concentrations in raw wastewater in Israel. *G. lamblia* assemblage B is more prevalent, where 27 out of 50 (54%), than assemblage A in the infected stool samples and is shed at higher concentrations. *G. lamblia* assemblage A is more prevalent in raw wastewater than assemblage B. Reporting of *Giardia* incidence in the community to the Department of Epidemiology in Israel is lower than the true incidence (as estimated from the concentration in sewage) because infection can be mild or in a carrier status.

### References


