

This is an Accepted Manuscript for Epidemiology & Infection. Subject to change during the editing and production process.

DOI: 10.1017/S0950268825000159

1 ***Dientamoeba fragilis* Cases Identified by Molecular Detection, Utah, United**
2 **States, 2014-2024**

3
4 Anna Jones MD, MPH, ^a Marc Roger Couturier PhD,^{b,c} , Andrew T. Pavia,^d Daniel T.
5 Leung MD, MSc ^e

6
7 ^a Department of Pediatrics, University of Utah School of Medicine, Salt Lake City, UT

8 ^b Department of Pathology, University of Utah, Salt Lake City, UT

9 ^c Institute for Clinical and Experimental Pathology, ARUP Laboratories, Salt Lake City,
10 Utah

11 ^d Division of Pediatric Infectious Disease, Department of Pediatrics, University of Utah,
12 School of Medicine, Salt Lake City, UT

13 ^e Division of Infectious Disease, Department of Internal Medicine, University of Utah
14 School of Medicine, Salt Lake City, UT

15

16 **Corresponding author:**

17 Anna Jones, MD, MPH

18 University of Utah, Department of Pediatrics

19 Email: anna.jones@hsc.utah.edu

20 Phone: +1 (919) 928 3730

21

22 **Alternate Corresponding author:**

23 Daniel Leung, MD, MSc, FIDSA, FASTMH

24 Email: Daniel.Leung@utah.edu

25

26

27

This is an Open Access article, distributed under the terms of the Creative Commons Attribution-NonCommercial-NoDerivatives licence (<http://creativecommons.org/licenses/by-nc-nd/4.0/>), which permits non-commercial re-use, distribution, and reproduction in any medium, provided the original work is unaltered and is properly cited. The written permission of Cambridge University Press must be obtained for commercial re-use or in order to create a derivative work.

28 **Summary**

29 *Dientamoeba fragilis* (*D. fragilis*) is an intestinal protozoan parasite with uncertain
30 pathogenic potential. In the United States, data on *D. fragilis* in the era of molecular
31 detection are limited. The aim of this retrospective chart review was to evaluate the
32 epidemiology and clinical characteristics of *D. fragilis* cases identified using polymerase
33 chain reaction assays between 2016 and 2024 at our academic medical center located
34 in Utah. We identified 28 unique cases with varying gastrointestinal symptomatology
35 including diarrhea, abdominal pain, nausea, vomiting, and bloating. Approximately half
36 (52%) of patients with follow-up data demonstrated improvement in symptoms following
37 initial treatment for *D. fragilis*. The overall prevalence of *D. fragilis* was low among those
38 tested (0.6% positivity). Additional research, including case control studies, are needed
39 to better describe the etiologic role of *D. fragilis*.

40

Accepted Manuscript

41 *Dientamoeba fragilis* (*D. fragilis*) is an intestinal protozoan with unclear pathogenic
42 potential [1–3]. *Dientamoeba fragilis* is commonly reported in association with
43 gastrointestinal symptoms but has also been commonly detected in asymptomatic
44 persons [2,4,5]. *Dientamoeba fragilis* is frequently detected with other organisms,
45 complicating efforts to understand its pathogenicity [5,6]. The life cycle and transmission
46 of *D. fragilis* are not completely understood and multiple hypotheses exist to explain the
47 protozoan's presence in human gastrointestinal tracts given the fragile nature of the
48 trophozoite stage [7,8]. It has appropriately been called “a neglected protozoan” [2,4].
49 The reported prevalence of *D. fragilis* varies depending on geographic location, study
50 population, and diagnostic methods [2–4]. Additionally, the clinical presentation ranges
51 from asymptomatic carriage to diarrhea, abdominal pain, and peripheral eosinophilia [4–
52 6]. With the increasing availability of molecular diagnostic methods, the identification of
53 *D. fragilis* has been facilitated by use of both single- and multi-plex polymerase chain
54 reaction (PCR) assays, which have a significantly higher sensitivity than microscopy [3].
55 The majority of recent clinical and epidemiologic studies characterizing *D. fragilis* have
56 been conducted in Europe [3,4], with the most recent study in the United States (US)
57 being a microscopy-based study published over a decade ago [9]. At the time of this
58 writing, only one FDA cleared PCR assay is available from Genetic Signatures, and this
59 product has been used in Australia and Europe with excellent performance [10]. Our
60 primary objective was to describe the epidemiologic and clinical characteristics of PCR-
61 diagnosed *D. fragilis* patients by performing a retrospective chart review at our
62 academic medical center located in the US.

63

64 The University of Utah has used the Gastrointestinal (GI) Parasite Panel by PCR
65 developed by ARUP laboratories since October 2014. The panel includes
66 *Cryptosporidium hominis* and *parvum*, *Cyclospora* spp., *Giardia*, *Entamoeba histolytica*,
67 and *Dientamoeba fragilis* targets. The *D. fragilis* target is a conserved sequence within
68 the 18S rRNA gene. The analytical sensitivity is approximately 16,000 copies/ml of stool
69 (equal to approximately 200 copies/reaction). Analytical specificity was established for
70 each of the protozoal targets against each other and 42 additional viral, bacterial, and
71 parasitic organisms (including *Entamoeba* spp. and *Strongyloides*). *In silico* analysis
72 revealed no predicted cross-reactivity with other organisms, including all formally
73 sequenced protozoa. All specimens were frozen immediately after collection and
74 thawed only at the time of testing. This frozen stability was shown in validation to
75 preserve sensitivity consistent with testing fresh stool. ARUP laboratories recommend
76 use of the panel for individuals with chronic diarrhea and a travel history or other
77 relevant exposure history or those with a complicated clinic course; the decision to order
78 the test is ultimately left to the clinician [11].

79
80 Since the GI Parasite Panel by PCR became available, 4804 tests have been
81 performed on patients from University of Utah Health. The total positivity for any target
82 is 181 (3.8%). For our report, a case of *D. fragilis* was defined by a positive PCR test; a
83 patient with multiple positive PCR results was defined as one case if there was no
84 intervening negative result. We reviewed the charts of the *D. fragilis* cases to abstract
85 relevant demographic and clinical data. Study data were collected and managed using

86 REDCap electronic data capture tools hosted at University of Utah [12,13]. This study
87 was deemed exempt from full review by the University of Utah IRB (IRB_00101686).

88
89 Thirty-one samples were positive for *D. fragilis* (0.6% positivity). Of those 31, we
90 identified 28 unique cases of *D. fragilis*, detected between April 2016 and April 2024. At
91 least one case was identified each year, except for 2021. Apart from two cases, all
92 patients were diagnosed in the outpatient setting, with most patients evaluated and
93 treated in primary care clinics (Table 1). Several patients were diagnosed by
94 gastroenterology and infectious disease specialists. The two hospitalized patients had
95 underlying conditions, and their level of acuity was likely unrelated to the *D. fragilis*
96 infection. One hospitalized individual was a bone marrow transplant recipient with
97 concern for graft versus host disease as a possible etiology of their presentation and the
98 second was a patient with septic shock in the setting of a newly diagnosed HIV infection
99 and multiple co-infections.

100
101 At the time of data abstraction, 25 patients had addresses in urban Utah counties and 3
102 were from urban counties in nearby states. Median age was 33; the youngest patient
103 was 9 years-old and 17 (61%) patients were between the ages of 18 and 49 years
104 (Table 1). Seventeen (61%) were female. Eleven (39%) individuals reported history of
105 recent international travel. An additional two individuals (7%) had history of freshwater
106 exposure in the US. Most individuals presented with persistent gastrointestinal
107 symptoms, several with greater than 1 year of symptoms (Table 2) and most had
108 multiple gastrointestinal complaints (79%). Approximately 82% of patients reported

109 diarrhea. Abdominal pain (61%), nausea (46%), bloating (39%), and constipation (25%)
110 were also common.

111
112 Enteric co-detections were not commonly identified. Twenty-five (89%) cases had
113 infectious diarrhea testing in addition to the GI Parasite Panel PCR (Table 3). One
114 patient was also positive for astrovirus (identified by comprehensive GI pathogen PCR
115 panel), and another individual was positive for *Blastocystis* (identified by stool ova and
116 parasite testing). A third patient was newly diagnosed with HIV and was also positive for
117 *Shigella* and EPEC (also identified by GI pathogen PCR panel). In the ten patients with
118 ova and parasite (O&P) examination results, none were positive for *D. fragilis*. In the ten
119 patients with CBC results, one (10%) demonstrated eosinophilia; this was the
120 aforementioned patient with recently diagnosed HIV and *Shigella* and EPEC co-
121 detections. An additional patient was evaluated due to history of persistent eosinophilia
122 and ultimately was diagnosed with systemic mastocytosis, a likely contributor to the
123 eosinophilia.

124
125 All individuals were treated for *D. fragilis*. The majority were prescribed metronidazole
126 (89%) as initial treatment. One individual was prescribed paromomycin, another
127 individual was prescribed tinidazole due to history of multiple rounds of metronidazole
128 for *Blastocystis* treatment, and a third was treated for concomitant chlamydia infection
129 with doxycycline. In the 25 cases with follow-up data available, symptoms improved in
130 13 (52%) after one round of treatment. Seven (26%) patients were retested due to
131 persistent symptoms following treatment; only two remained positive for *D. fragilis* tests

132 upon retesting (Supplemental Table 1). Four (15%) received additional rounds of
133 treatment with either metronidazole or doxycycline; none of those who received
134 additional rounds of treatment experienced resolution of symptoms.

135
136 In this single-center retrospective study of PCR-positive *D. fragilis* cases over a 10-year
137 period of PCR testing availability, we found an overall test positivity rate of 0.6%. Prior
138 prevalence estimates vary considerably based on geographic region, population
139 studied, and diagnostic method employed [2–4]. Our positivity rate was higher than a
140 2010 study of intestinal infections in the Rocky Mountain region, which found a 0.04%
141 prevalence of *D. fragilis* identified using microscopy [14] and notably lower than the
142 reported prevalence of *D. fragilis* identified using PCR in symptomatic individuals in
143 European countries and Australia [2,5,15]. Due to the limited availability of *D. fragilis*
144 PCR in the US, the clinical presentation and treatment outcomes of patients with *D.*
145 *fragilis* in the US is not well known.

146
147 Testing was requested only on symptomatic individuals; without a control group we
148 cannot clearly attribute *D. fragilis* as the cause of the symptoms. Additional viral or
149 bacterial testing was documented on most (89%) patients. Most patients (89%) had *D.*
150 *fragilis* identified as a single organism. However, three had a co-detection documented
151 and we identified alternative diagnoses through chart review in two (irritable bowel
152 syndrome and systemic mastocytosis). The scarcity of co-detections and alternative
153 diagnoses is a strength of our case series as these have limited the ability to
154 understand the pathogenicity of *D. fragilis* [6,16].

155

156 The range of gastrointestinal symptoms of the patients in our study was similar
157 compared to other studies [2,16,17]. Interestingly, only 10% had eosinophilia, which
158 differs from prior reports [2,5,16,18], though only approximately one third of patients had
159 CBC results for evaluation. Additionally, among the one-third of cases which also had an
160 O&P examination performed, none were positive for *D. fragilis*. This is not unexpected
161 given the high sensitivity of PCR and challenging nature of direct microscopy [19].

162

163 This study may have limited generalizability due to the single center of data collection.
164 Additionally, all patients in our review were tested due to the presence of
165 gastrointestinal symptoms, limiting our ability to draw conclusions about the etiologic
166 role of *D. fragilis*. It is possible that other underlying causes, such as IBS, may
167 contribute to symptomatology seen in patients in whom *Dientamoeba* is detected. The
168 lack of follow-up data in this retrospective study limits our assessment of treatment
169 efficacy.

170

171 We found that among patients from the Intermountain West who were tested using a
172 multi parasite PCR assay, the prevalence of *D. fragilis* was low. Case control studies in
173 the US could help determine the prevalence among asymptomatic persons and better
174 describe the etiologic role of *D. fragilis*. The reasons for the low prevalence in this
175 sample of US patients compared to the prevalence in Europe requires further study.

176

177

178 **ACKNOWLEDGEMENTS:**

179

180 ***Data Availability:*** Deidentified data that support the findings of this study are available
181 from the corresponding author upon reasonable request.

182

183 ***Financial support:*** This work was supported by the National Institutes of Health
184 (R01AI135114, K24166087 to Daniel T. Leung and R38 HL143605 to Anna Jones
185 through Utah Stimulating Access to Research in Residency (StARR)). Additionally, the
186 research reported in this publication was supported by the National Center for
187 Advancing Translational Sciences of the National Institutes of Health under Award
188 Number UM1TR004409 NCATS/NIH

189

190 ***Conflicts of interest:*** Dr. Pavia has served as a consultant to Sanofi, GSK and Haleon,
191 unrelated to the current work. No other authors have conflicts of interest.

192

Accepted Manuscript

193 **References**

- 194 1. **Shasha D, et al.** The clinical significance of *Dientamoeba fragilis* and *Blastocystis* in
195 human stool—retrospective cohort study. *Clinical Microbiology and Infection* 2024;
196 **30**: 130–136.
- 197 2. **Stark D, et al.** *Dientamoeba fragilis*, the Neglected Trichomonad of the Human
198 Bowel. *Clinical Microbiology Reviews* 2016; **29**: 553–580.
- 199 3. **Gestel RS van, Kusters JG, Monkelbaan JF.** A clinical guideline on *Dientamoeba*
200 *fragilis* infections. *Parasitology* Cambridge University Press, 2019; **146**: 1131–1139.
- 201 4. **Garcia LS.** *Dientamoeba fragilis*, One of the Neglected Intestinal Protozoa. *Journal of*
202 *Clinical Microbiology* 2016; **54**: 2243–2250.
- 203 5. **Calderaro A, et al.** Prevalence of Intestinal Parasitoses in a Non-Endemic Setting
204 during a 10-Year Period (2011–2020): A Focus on *Dientamoeba fragilis*.
205 *Microorganisms* 2022; **10**: 426.
- 206 6. **Venturini E, et al.** Epidemiology and clinical features of intestinal protozoan
207 infections detected by Real-time PCR in non-native children within an Italian tertiary
208 care children’s hospital: A cross-sectional study. *Travel Medicine and Infectious*
209 *Disease* 2021; **43**: 102107.
- 210 7. **Hall LM, et al.** Observations on the transmission of *Dientamoeba fragilis* and the cyst
211 life cycle stage. *Parasitology* 2024; **151**: 337–345.
- 212 8. **CDC - DPDx - *Dientamoeba fragilis* Infection.**
213 2019(<https://www.cdc.gov/dpdx/dientamoeba/index.html>). Accessed 8 November
214 2024.
- 215 9. **Chang AH, et al.** Decreasing Intestinal Parasites in Recent Northern California
216 Refugees. *The American Journal of Tropical Medicine and Hygiene* The American
217 Society of Tropical Medicine and Hygiene, 2013; **88**: 191–197.
- 218 10. **Gough R, Ellis J, Stark D.** Comparison and Recommendations for Use of
219 *Dientamoeba fragilis* Real-Time PCR Assays. *Journal of Clinical Microbiology*
220 American Society for Microbiology, 2019; **57**: 10.1128/jcm.01466-18.
- 221 11. **Infectious Diarrhea Testing Algorithm | Choose the Right Test.**
222 (https://arupconsult.com/algorithm/infectious-diarrhea-testing-algorithm?_ga=2.251007558.1152959944.1731085974-533400701.1728511603&_gl=1*1kk5czk*_ga*NTMzNDAwNzAxLjE3Mjg1MTE2MDM.*_ga_Z8H49DQE4D*MTczMTA5MDc0OS40LjAuMTczMTA5MDc0OS4wLjAuMA..)
223 algorithm?
224 533400701.1728511603&_gl=1*1kk5czk*_ga*NTMzNDAwNzAxLjE3Mjg1MTE2MD
225 M.*_ga_Z8H49DQE4D*MTczMTA5MDc0OS40LjAuMTczMTA5MDc0OS4wLjAuMA..) .
226 . Accessed 8 November 2024.

- 227 12. **Harris PA, et al.** Research electronic data capture (REDCap)—A metadata-driven
228 methodology and workflow process for providing translational research informatics
229 support. *Journal of Biomedical Informatics* 2009; **42**: 377–381.
- 230 13. **Harris PA, et al.** The REDCap consortium: Building an international community of
231 software platform partners. *Journal of Biomedical Informatics* 2019; **95**: 103208.
- 232 14. **Church C, Neill A, Schotthoefer AM.** Intestinal Infections in Humans in the Rocky
233 Mountain Region, United States. *The Journal of Parasitology* [The American Society
234 of Parasitologists, Allen Press], 2010; **96**: 194–196.
- 235 15. **Stark D, et al.** Comparison of microscopy, two xenic culture techniques,
236 conventional and real-time PCR for the detection of *Dientamoeba fragilis* in clinical
237 stool samples. *European Journal of Clinical Microbiology & Infectious Diseases*
238 2010; **29**: 411–416.
- 239 16. **Miguel L, et al.** Clinical and Epidemiological Characteristics of Patients with
240 *Dientamoeba fragilis* Infection. *The American Journal of Tropical Medicine and*
241 *Hygiene* 2018; **99**: 1170–1173.
- 242 17. **Clemente L, et al.** *Dientamoeba fragilis* in the North-East of Italy: Prevalence study
243 and treatment. *Parasitology International* 2021; **80**: 102227.
- 244 18. **Garg P, et al.** A tale of two studies: is peripheral eosinophilia associated with
245 *Dientamoeba fragilis* detection in adult stool samples? *Pathology* 2024; **56**: 688–
246 695.
- 247 19. **Calderaro A, et al.** Evaluation of a real-time polymerase chain reaction assay for
248 the detection of *Dientamoeba fragilis*. *Diagnostic Microbiology and Infectious*
249 *Disease* 2010; **67**: 239–245.

250

251

253 **Table 1.** Demographic characteristics of cases.

	n (%)
Total cases	28
Female sex	17 (61)
Male sex	11 (39)
Median age	33
<5 years	0 (0)
5-17 years	5 (18)
18-49 years	17 (61)
50+ years	6 (21)
Encounter type	
Clinic	23 (82)
Hospital	2 (7)
Other	3 (11)
Insurance type	
Private	20 (71)
Other	8 (29)
Provider specialty	
Primary care	15(54)
Infectious disease	5 (18)
Gastroenterology	4 (14)
Other	4 (14)

History of international travel

Yes*	11 (39)
No	6 (21)
Unknown	11 (39)

Immunocompromised state

Yes	3 (11)
No	25 (89)

254 * Destinations visited: Columbia, Japan, Madagascar,
255 Malawi, Mexico (4), Pacific Islands, Peru (3),
256 Philippines, Puerto Rico, Singapore, Spain, Vietnam

Accepted Manuscript

257 **Table 2.** Reported symptoms.

	n (%)
Median length of symptoms in days (min, max)*	45 (3, 700)
Reported diarrhea	23 (82)
3 or more loose stools per day	9 (32)
Blood in stools	3 (11)
Abdominal pain	17 (61)
Nausea	13 (46)
Vomiting	6 (21)
Bloating	11 (39)
Constipation	7 (25)
Subjective fever	5 (18)
Objective fever	0 (0)
Weight loss	4 (14)
Anorexia	4 (14)
Fatigue	4 (14)
Anal pruritus	4 (14)
Multiple gastrointestinal complaints	22 (79)

258 * Missing in 4 cases

259

260

261

262

263 **Table 3.** Additional infectious diarrhea testing. Additional testing was performed on 25

264 (89%) cases.

	Testing ordered n (% of cases)	Test positivity n (% of tests ordered)
Comprehensive GI pathogen PCR panel	2 (7)	2 (100)
Stool viral PCR panel	5 (18)	0 (0)
Stool bacterial PCR panel	4 (14)	0 (0)
Stool culture	12 (43)	0 (0)
Stool Ova & Parasite	10 (36)	1 (10)
<i>C. difficile</i> toxin by EIA	14 (50)	0 (0)
<i>Campylobacter</i> antigen	8 (29)	0 (0)
<i>H. pylori</i> antigen	5 (18)	0 (0)
<i>Strongyloides</i> antibody	4 (14)	0 (0)
<i>Schistosoma</i> antibody	1 (4)	0 (0)
<i>Giardia</i> antigen	1 (4)	0 (0)
Pinworm	1 (4)	0 (0)

265

266

267

268