

Free-Living Amoebae Recovered from Human Stool Samples in *Strongyloides* Agar Culture

Richard S. Bradbury*

MRC Keneba, Keneba Village, Kiang West, The Gambia

Our laboratory in The Gambia, West Africa, performs Koga agar (1.5% bacteriological agar, 0.5% sodium chloride, 0.5% meat extract, 0.1% bacteriological peptone) culture for larvae of strongyle nematodes on human fecal samples for which parasitological investigation has been requested. We have recovered free-living amoebae (FLA) from human fecal specimens on two occasions over a period of 9 months (representing 130 individual fecal cultures) on this agar (Fig. 1). The amoebae in both cases were identified as *Hartmannella* species based upon the morphology of trophozoites and cysts in agar culture (Fig. 2) and their inability to enflagellate in distilled water after 8 h of incubation at 37°C. No other types of FLA have been recovered. Due to resource constraints, sequencing of the partial 18S rRNA gene and internal transcribed spacer (ITS) region could not be performed, and this identification must remain presumptive. Specimens were collected into sterile containers, and Koga culture plates were sterile and sealed with Parafilm prior to incubation at 30°C for 5 days; therefore, environmental contamination with FLA was very unlikely. Repeat specimens could not be obtained from subjects, and so it was not possible to determine if these findings represent transient passage of FLA or true intestinal colonization.

It has been postulated that FLA provide a vehicle for bacterial pathogens to gain entry into the human respiratory tract (1). The possibility that free-living amoeba may have a similar role in the introduction of bacterial pathogens into the human intestinal tract (and particularly in defense against stomach acid during

transit to the duodenum) should be further considered, given these findings. FLA are potential environmental reservoirs of several important pathogens of the human intestinal tract, including *Salmonella* spp., *Shigella* spp., *Campylobacter* spp., and enterohemorrhagic *Escherichia coli* (2–5). Specifically, due to the large infective dose of *Vibrio cholerae* (10^8 to 10^9 cells), it has been suggested that Acanthamoebae may act as both environmental hosts and intracellular multipliers of this organism, allowing it to grow sufficiently to be able to cause human infection (6).

Human respiratory tract (7, 8) and urinary tract (9) colonization with *Acanthamoeba* spp. may occur. Recovery of FLA, including *Hartmannella* spp. from the intestinal tracts of rodents (10), reptiles (11), and fish and birds (12), has been previously reported. Only one previous reference in the literature describes the recovery of FLA from the human intestinal tract (13).

The potential for transient passage or true colonization of the

Published ahead of print 13 November 2013

Editor: B. A. Forbes

Address correspondence to Richard S. Bradbury, r.bradbury@cqu.edu.au.

* Present address: School of Medical and Applied Sciences, Central Queensland University, Rockhampton, Queensland, Australia.

Copyright © 2014, American Society for Microbiology. All Rights Reserved.

doi:10.1128/JCM.02738-13

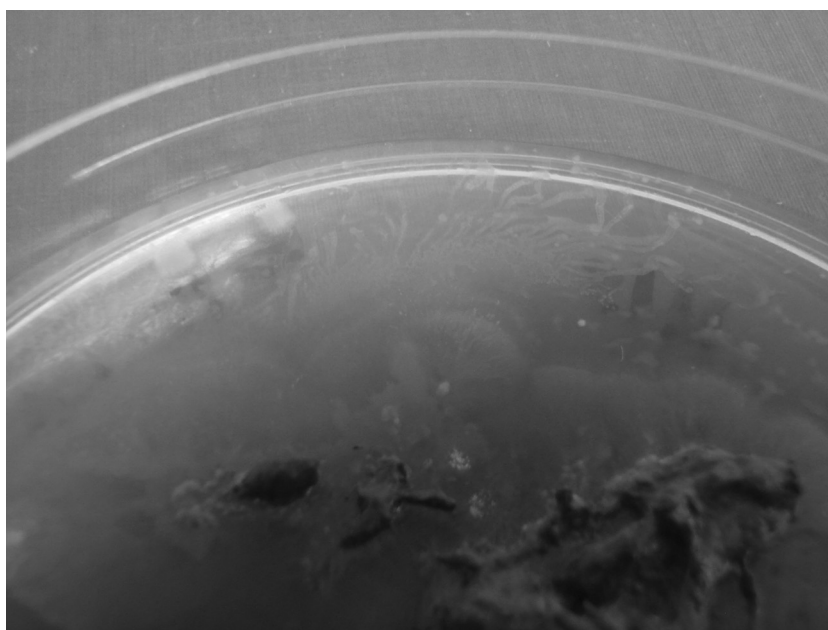


FIG 1 Tracks of presumptive *Hartmannella* species on the surface of a Koga agar culture of human feces after 5 days of incubation at 30°C.

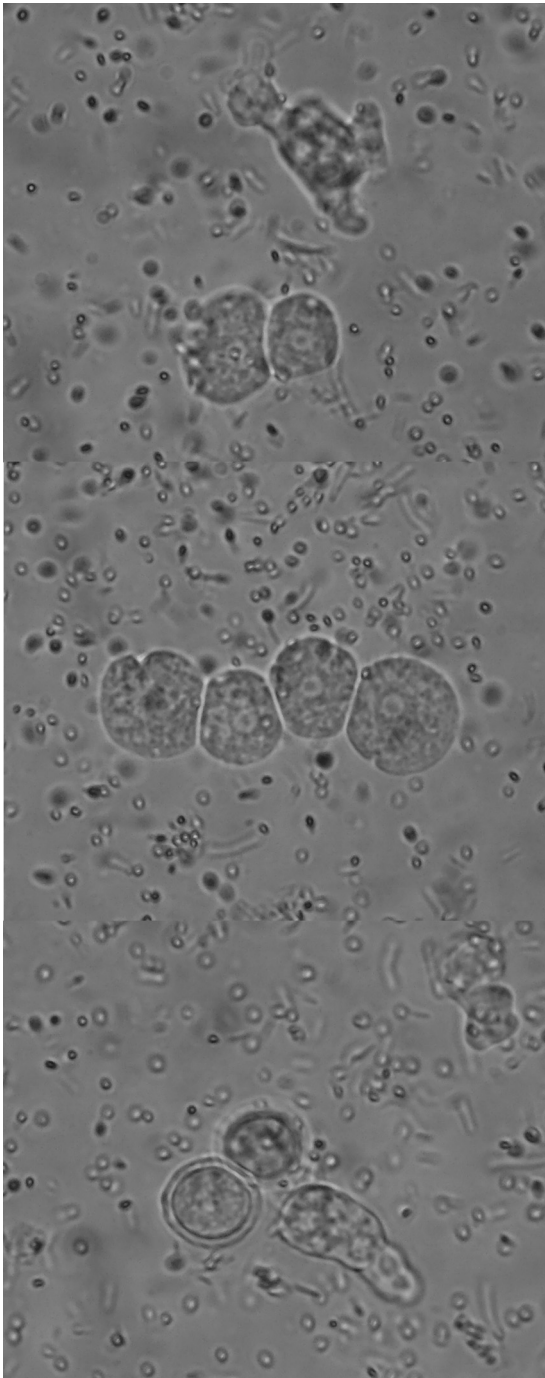


FIG 2 Characteristic morphology of presumptive *Hartmannella* cysts and trophozoites on agar culture.

human intestinal tract with FLA may also provide a novel potential portal of entry into the human body for FLA causing primary amoebic meningitis (PAM) and granulomatous amoebic encephalitis (GAE). These findings may also require a review of safety practices when dealing with such cultures, including the use of biosafety cabinets during manipulation of Koga agar cultures to avoid cultures potentially containing *Naegleria fowleri* being inappropriately manipulated on a benchtop in the laboratory.

In summary, we report the recovery of FLA in Koga agar culture of feces from two separate individuals in West Africa. It is unknown if this represents transient passage or true intestinal colonization. These findings demonstrate an important novel mechanism of entry for bacterial intestinal pathogens and possibly pathogenic FLA themselves into the human body. They also raise the need for awareness of the potential for FLA to be recovered from Koga agar culture in order to ensure the health and safety of laboratory staff performing such cultures.

ACKNOWLEDGMENTS

This research is jointly funded by the United Kingdom Medical Research Council (MRC; grant MC-A760-5QX00) and the United Kingdom Department for International Development (DFID) under the MRC/DFID Concordat agreement.

REFERENCES

1. Khan NA. 2006. *Acanthamoeba*: biology and increasing importance in human health. *FEMS Microbiol. Rev.* 30:564–595. <http://dx.doi.org/10.1111/j.1574-6976.2006.00023>.
2. Wildschutte H, Lawrence JG. 2007. Differential *Salmonella* survival against communities of intestinal amoebae. *Microbiology* 153:1781–1789. <http://dx.doi.org/10.1099/mic.0.2006/003616-0>.
3. Bui XT, Winding A, Qvortrup K, Wolff A, Bang DD, Creuzenet C. 2012. Survival of *Campylobacter jejuni* in co-culture with *Acanthamoeba castellanii*: role of amoeba-mediated depletion of dissolved oxygen. *Environ. Microbiol.* 14:2034–2047. <http://dx.doi.org/10.1111/j.1462-2920.2011.02655.x>.
4. Jeong HJ, Jang ES, Han BI, Lee KH, Ock MS, Kong HH, Chung DI, Seol SY, Cho DT, Yu HS. 2007. *Acanthamoeba*: could it be an environmental host of *Shigella*? *Exp. Parasitol.* 115:181–186. <http://dx.doi.org/10.1016/j.exppara.2006.08.002>.
5. Chekabab SM, Daigle F, Charette SJ, Dozois CM, Harel J. 2012. Survival of enterohemorrhagic *Escherichia coli* in the presence of *Acanthamoeba castellanii* and its dependence on Pho regulon. *Microbiologyopen* 1:427–437. <http://dx.doi.org/10.1002/mbo3.40>.
6. Abd H, Shanan S, Saeed A, Sandström G. 2011. Survival of *Vibrio cholerae* inside *Acanthamoeba* and detection of both microorganisms from natural water samples may point out the amoeba as a protozoal host for *V. cholerae*. *J. Bacteriol. Parasitol.* S1-003. <http://dx.doi.org/10.4172/2155-9597.S1-003>.
7. Madrigal Sesma MJ, Santillana López I. 1989. Isolation of free-living amoebas from samples of respiratory origin. *Rev. Sanid. Hig. Publica (Madr.)* 63:63–72.
8. Rivera F, Medina F, Ramírez P, Alcocer J, Vilaclara G, Robles E. 1984. Pathogenic and free-living protozoa cultured from the nasopharyngeal and oral regions of dental patients. *Environ. Res.* 33:428–440. [http://dx.doi.org/10.1016/0013-9351\(84\)90040-9](http://dx.doi.org/10.1016/0013-9351(84)90040-9).
9. Santos LC, Oliveira MS, Lobo RD, Higashino HR, Costa SF, van der Heijden IM, Giudice MC, Silva AR, Levin AS. 2009. *Acanthamoeba* spp. in urine of critically ill patients. *Emerg. Infect. Dis.* 15:1144–1146.
10. Kerdraon G, Simitzis-Le Flohic AM, Couatarmanac'h A. 1986. Amphizoic stage of free amoebae, myth or reality (could oxygen be the limiting factor?). *Bull. Soc. Pathol. Exot. Filiales* 79:66–73.
11. Hassl A, Benyr G, Appelt S. 2000. Free-living amoebas as opportunistic intestinal parasites of reptiles. *Mitt. Osterr. Ges. Tropenmed. Parasitol.* 22:49–54.
12. Franke ED, Mackiewicz JS. 1982. Isolation of *Acanthamoeba* and *Naegleria* from the intestinal contents of freshwater fishes and their potential pathogenicity. *J. Parasitol.* 68:164–166. <http://dx.doi.org/10.2307/3281343>.
13. de Moura H, Salazar HC, Fernandes O, Lisboa DC, de Carvalho FG. 1985. Free-living amoeba in the human intestine. Evidences of parasitism. *Rev. Inst. Med. Trop. Sao Paulo* 27:150–156. <http://dx.doi.org/10.1590/S0036-46651985000300007>.