高 病 原 性 Cryptococcus gattiiの感染防衛に寄与する 樹状細胞ワクチン、口頭	上金大 浦金大亀澁宮野城保郎, 幸秀克和義書樹陽, 誠弘明, 京縣縣 建秀克和義縣, 一种	第63回日本感染症学会 東日本地方会学術集会	2014年10月	国内
真菌の薬剤耐性の現状と課 題、口頭	名木 稔, 田石野山 梅山越野山 大宮崎 養料, 宮崎	第63回日本感染症学会東日本地方会学術集会	2014年10月	国内
肺アスペルギローマとの鑑別が 困 難 で あっ た Pseudallescheria boydiiによる肺菌球症の1 例、口頭	本福今宮泉大柳宮早田河川田村崎川野原﨑田代野奈雄圭泰公秀克義隆野人の東京、東京、東京、東京、東京、東京、東京、東京、東京、東京、東京、東京、東京、東	第62回日本化学療法学 会西日本支部総会 57回日本感染症学会 日本地方会学術集定学会 第84回日本地方会学術集 西日本地方会学術集	2014年10月	国内

黒色菌糸症の1例、口頭	多田明子, 山藤本四井山藤口井山崎 瀬川井山崎 大田田明子, 東河井山崎 大田田明子, 東京 大田明子, 東京 大田明子, 東京 大田明子, 東京 大田明子, 東京 大田明子, 東京 大田明子, 東京 大田明子, 東京 大田明子, 東京 大田明子, 東京 大田明子, 東京 大田明寺, 東京 大田明寺, 東京 大田明寺, 東京 大田明寺, 東京 大田明寺, 東京 大田明寺, 大田明 大田明寺, 大田明 大田, 大田, 大田明 大田, 大田 大田 大田 大田 大田 大田 大田 大田 大田 大田 大 大 大田 大 大 大 大 大 大 大 大 大 大 大 大 大 十 大 大 十 大 十	第263回日本皮膚科学 会岡山地方会	2014年9月	国内
高病原性クリプトコックス症に対する樹状細胞ワクチンの効果、口頭	上大 清金浦水奈川大澁宮金野久郎水子井口良本野谷﨑城圭陽,公幸一裕拓 秀和義雄吾。一一次,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	第25回日本生体防御学 会学術総会	2014年7月	国内
カンジダ属の抗真菌薬耐性、口頭	田大名浦金梅山宮公秀 稔誠弘, 一明, 新弘, 年隆智継	第35回関東医真菌懇話会	2014年6月	国内

ミカファンギン耐性Candida glabrata株のin vitro性状解 析、口頭	田大名浦金梅山荒皿宮公秀、龍誠弘、大名浦金梅山荒皿宮の野木、井幸山越、北谷・・・・・・・・・・・・・・・・・・・・・・・・・・・・・・・・・・・・	第35回関東医真菌懇話会	2014年6月	国内
腹膜透析中に発症した Cryptococcus laurentiiによ る腹膜炎の一例、口頭	金稲狩本水名上山田梅川金大宮井幸垣谷大谷木野越辺山原城野﨑誠弘司芳郎、三真稔吾智一隆明樹明継、,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,,	第35回関東医真菌懇話会	2014年6月	国内
マウスモデルでの肺炎球菌蛋白・糖脂質併用ワクチンの感染防御効果の解析、口頭	金城雄樹, 金子幸弘, 梅山 隆, 川上和魏, 大石和義德, 宮崎義継	第88回日本感染症学会 学術講演会·第62回日 本化学療法学会総会合 同学会	2014年6月	国内

症例から学ぶ感染症セミナームーコル症の真菌同定検査、口頭	梅山 隆, 大野秀明, 田辺公一, 山越 名味 宮﨑義継	第88回日本感染症学会 学術講演会·第62回日 本化学療法学会総会合 同学会	2014年6月	国内
病 原 糸 状 菌 Aspergillus fumigatusのPolo-likeキナーゼ遺伝子破壊株の菌糸成長・分生子形成・抗真菌薬感受性への影響、ポスター	梅山 隆, 山田 超, 山田 名子城野、 金子城野、 宝崎 、 大宮崎 、 大宮崎	第88回日本感染症学会 学術講演会·第62回日 本化学療法学会総会合 同学会	2014年6月	国内
カンジダ属の抗真菌薬感受性の変貌、口頭	田大名浦金梅山知亀宮公秀 稔誠弘, 神, 神, 林, 神,	第88回日本感染症学会 学術講演会·第62回日 本化学療法学会総会合 同学会	2014年6月	国内

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高 病 原 性 Cryptococcus gattii由来莢膜多糖の免疫細 胞に及ぼす影響、口頭	浦金田梅山金大杉宮湖子辺山越城野田﨑誠弘一條智樹明。	第88回日本感染症学会 学術講演会·第62回日 本化学療法学会総会合 同学会	2014年6月	国内
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Research Article

Association between *Giardia duodenalis* and Coinfection with Other Diarrhea-Causing Pathogens in India

Avik K. Mukherjee,¹ Punam Chowdhury,¹ Krishnan Rajendran,² Tomoyoshi Nozaki,³ and Sandipan Ganguly¹

- ¹ Division of Parasitology, National Institute of Cholera and Enteric Diseases, P-33 CIT Road, Scheme XM, Beliaghata, Kolkata, West Bengal 700010, India
- ² Division of Data Management, National Institute of Cholera and Enteric Diseases, P-33 CIT Road, Scheme XM, Beliaghata, Kolkata, West Bengal 700010, India

Correspondence should be addressed to Sandipan Ganguly; sandipanganguly@gmail.com

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Giardia duodenalis, is often seen as an opportunistic pathogen and one of the major food and waterborne parasites. Some insights of Giardia infestation in a diarrhoea-prone population were investigated in the present study. Our primary goal was to understand the interaction of this parasite with other pathogens during infection and to determine some important factors regulating the diarrhoeal disease spectrum of a population. Giardia showed a steady rate of occurrence throughout the entire study period with a nonsignificant association with rainfall (P > 0.05). Interestingly coinfecting pathogens like Vibrio cholerae and rotavirus played a significant ($P \le 0.001$) role in the occurrence of this parasite. Moreover, the age distribution of the diarrhoeal cases was very much dependent on the coinfection rate of Giardia infection. As per our findings, Giardia infection rate seems to play a vital role in regulation of the whole diarrhoeal disease spectrum in this endemic region.

1. Introduction

Giardia duodenalis is present worldwide but is more prevalent in developing countries where the lack of sanitation and hygiene awareness is a matter of concern [1, 2]. Considering its high endemicity in some countries, research on Giardia is of low priority as the infection it causes is self-limiting, a situation that enhances its propagation. Giardiasis is caused by the protozoan parasite Giardia duodenalis [3] which is usually transmitted through ingesting contaminated food and water. A wide variety of pathogens can cause diarrhea, but G. duodenalis impacts the economic growth of a country by affecting the Disability Adjusted Life Year (DALY) rates [4]. Giardiasis has much lower mortality rates associated with it than do other diarrheagenic pathogens such as Vibrio cholerae or Shigella [5]; nevertheless, it may still play an important role in regulating the spectrum of diarrheal diseases in diarrhea-prone regions. The study described herein was designed to survey the prevalence of G. duodenalis among diarrheal patients within Kolkata, India. Kolkata is a densely populated city with a variable socioeconomic and climatic background and is frequently affected by outbreaks of diarrheal disease; hence that is why the area was chosen for disease transmission studies [6].

Fecal samples were tested from patients attending the Infectious Diseases and Beliaghata General (IDBG) Hospital in Kolkata city throughout a period of 56 months. These patients only complained of diarrhea. A systemic sampling procedure [7] allowed us to collect enough data to demarcate the catchment areas for diarrhea within the city and to interpret the epidemiological aspects of *Giardia* infestation in an urban region of this developing country.

2. Methods

2.1. Ethics Statement. This study received ethical clearance from the National Institute of Cholera and Enteric Diseases (NICED) ethical committee, the host institute.

³ Department of Parasitology, National Institute of Infectious Diseases, 1-23-1 Toyama, Shinjuku-ku, Tokyo 162-8640, Japan

2 BioMed Research International

2.2. Study Design. The study was performed through collaboration between NICED and IDBG Hospital, Kolkata. IDBG is located within the city of Kolkata and is the largest infectious diseases hospital in India. IDBG treats around 25000 cases of diarrhea every year and most of these patients are residents of the city [6]. Thus, the prevalence of diarrheal diseases in the city can be estimated by surveying IDBG patients. Every fifth patient visiting IDBG who complained of only diarrheal symptoms on two randomly selected days per week was enrolled in the study. The study ran from November 2007 to June 2012. A single fecal sample was sent to the laboratory for analysis by trained healthcare professionals who also obtained the patient's background history via a systematically designed questionnaire. Patient consent for the study was obtained at the same time. The system remained unbiased with regard to sex, age, or other physical factors with nearly proportional distribution of male and female subjects and age ranging from 0 to 60 years in the majority of cases.

2.3. Screening for G. duodenalis in Stool Samples. G. duodenalis was detected in stool samples by using three different procedures. Stool samples were divided into three aliquots immediately after reaching the laboratory. The first aliquot was used for microscopic analysis with iodine wet-mount and trichrome staining [8] after concentration using "Ridley's concentration technique" [9]. The second aliquot was used in an antigen capture enzyme-linked immunoabsorbent assay using a GIARDIAII kit (TechLAB, Blacksburg, VA, USA) as per the manufacturer's protocol. DNA was extracted directly from the third aliquot of each stool sample using a DNA Stool Minikit (Qiagen, USA), according to the manufacturer's protocol. PCR was performed using G. duodenalis-specific primers and the DNA extracted by the kit as template following previously published protocols [7, 10]. All of the G. duodenalis-positive cases were also investigated for coinfections with other common pathogens as described previously [7]. The bacterial and viral coinfection status of a sample was investigated with assistance from Drs. T. Ramamurthy, T. Krishnan, and M. C. Sarkar in their laboratories at NICED [6].

2.4. Statistics and GIS Mapping. Data were entered into the predesigned format of the *pro forma* in the SQL server that has an inbuilt entry validation checking facilitated program by trained data entry professionals. Data were randomly checked and matched for consistency and validity. Edited data were exported and analyzed using SPSS.19.0 and Epi-info 3.5.4 [11].

The inferential age group was explored for *G. duodenalis*-positive cases by multinomial logistic regression [12, 13]. The aim of this was to determine the age groups that were most likely to be infected with *G. duodenalis*. Five age groups were classified, that is, up to 5 years, >5–10 years, >10–20 years, >20–30 years, 30–40 years, and >40 years, and were coded as 1–6, respectively. The relationships between the risk-dependent variable and each of the categorical explanatory variables are shown in Table 1. Infections caused by *G. duodenalis* were classified "1" when the pathogen was present

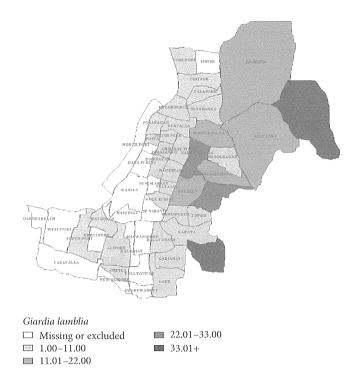


FIGURE 1: *Giardia duodenalis* distribution area. Map of the study region showing the catchment areas for the *Giardia duodenalis* cases according to our surveillance report (November 2007 to June 2008).

or "2" when absent. The extreme values of the classified age group were fixed as a reference category.

Associations between G. duodenalis infection and other variables such as rainfall or coinfection with other pathogens were tested using EpiInfo 3.5.4. Where the presence of G. duodenalis was considered an outcome variable, factors like rainfall, overall coinfection, and major coinfection were assigned as dependent variables. Where the P value was ≤ 0.05 , this was considered a valid association [14].

A choropleth map was constructed to display the data from the area where all the positive samples had originated within the city [15]. For this map, the different colors and patterns were combined to depict the different values of the attribute variable associated with each area. Each area is colored according to the category into which its corresponding attribute value had fallen. *G. duodenalis*-positive cases were embedded on the thematic map by the geographical information system (GIS) to visualize the infections. The boundary map shows that the prevalence of *G. duodenalis* was highest in Rajarhat and Tiljala (31.0%), followed by Narkeldanga and Tangra (22–33%), while the values for Dum Dum, Salt Lake, Beliaghata, Maniktala, and Entally regions ranged from 11 to 22 percent (Figure 1).

3. Results and Discussion

Single stool samples from 4039 diarrheal patients were examined throughout a 56-month period, and 413 (i.e., 10.2%) of them tested positive for *G. duodenalis*. All the data were categorized on a monthly basis to assess any