Enteric Opportunistic Infection and the Impact of Antiretroviral Therapy among HIV/AIDS Patients from Tehran, Iran

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Abstract

Background: Opportunistic parasites have been identified as human pathogens, especially in immunodeficient patients. Microsporidian and coccidian infections cause chronic diarrhea as common clinical manifestation in HIV positive patients. In this study, the frequency of opportunistic infections, including microsporidian and coccidian infections, was evaluated in HIV/AIDS patients from Tehran and phylogenic analysis was performed for E. bieneusi isolates from these patients.

Methods: One hundred and two stool samples were collected from confirmed HIV/AIDS patients, referred to Consult Center of Behavior Diseases, West Health Center, Iran University of Medical Sciences in Tehran, Iran. The samples were transferred to Research Center of Pediatric Infectious Diseases, Institute of Immunology and Infectious Diseases, Iran University of Medical Sciences from Jan 2016 to Dec 2016. After conventional formalin-ether concentration, aniline blue staining method and acid-fast staining technique were used for detection of microsporidian spores and Cyclospora oocysts. DNA was extracted and nested PCR was performed.

Results: Two (1.96%) cases were found to be positive for intestinal microsporidia infection using aniline blue staining method and were confirmed as E. bieneusi by nested PCR. One patient was found with Cyclospora cayetanensis infection by acid-fast staining method and PCR. Giardia lamblia and Blastocystis hominis were detected as non-opportunistic parasites in 1/102 (0.98%) and 2/102 (1.96%) of the HIV positive patients, respectively.

Conclusion: With respect to the use of antiretroviral therapy (ART) in HIV positive patients, we found a low frequency of infection.

Keywords: Opportunistic infection; HIV; Coccidian; Microsporidian

Introduction

Opportunistic parasites including microsporidian and coccidian infections have been identified as human pathogens, especially in immunodeficient individuals like organ transplant recipients (1), cancer patients (2), and HIV/AIDS patients (3). However, there are some reports of microsporidal infection in immunocompetent people also, especially those like travelers, chi-
children, and healthy human population from all over the world (4, 5).

The most common microsporidial infections are *Enterocytozoon bieneusi* and *Encephalitozoon intestinalis* in humans (6). *E. bieneusi* is a zoonotic pathogen that infects animals and causes infection in humans (7, 8). *E. bieneusi* infection causes chronic diarrhea and biliary illness and it is has been reported in individuals infected with HIV (4). Prevalence rate of *E. bieneusi* has been reported between 2.5% - 51% for HIV-seropositive adult patients with diarrhea (9-11) and 4.6% for patients without diarrhea (12).

Diagnosis of microsporidia is based on the identification of spores by staining methods like chromotrope 2R, aniline blue and calcofluor white (13, 14), and various molecular methods (3, 12, 15).

*Cyclospora cayetanensis* infection has recently emerged as one of the opportunistic infections with worldwide distribution. A number of outbreaks have been reported in the United States and Canada (16). The oocyst of *C. cayetanensis* has been identified in the feces of immunocompetent people who travel to developing countries or consume contaminated fruits or salads, and in patients with AIDS. In Pune, India, *Cyclospora* was reported in 0.7%-3.3% of HIV positive individuals with diarrhea (17).

Oocysts of *Cyclospora* can be distinguished by direct examination of wet smear. They can also be observed by acid-fast staining method and diagnosis of oocysts sporulation after preservation in dichromate potassium (2.5%) can be made (16, 18).

This study aimed to evaluate the frequency of opportunistic infections, including microsporidian and coccidian infections, among HIV/AIDS patients in Tehran.

**Methods**

In this cross-sectional study, 102 stool samples were collected from confirmed HIV/AIDS patients, with or without diarrhea, referred to Consult Center of Behavior Diseases, West Health Center, Iran University of Medical Sciences, Tehran, Iran. The stool samples were transferred to Research Center of Pediatric Infectious Diseases, Institute of Immunology and Infectious Diseases, Iran University of Medical Sciences, from Jan 2016 to Dec 2016. Wet mount smear was prepared from concentrated samples and all samples were observed under light microscope. Acid-fast staining was carried out for all samples and slides were observed and evaluated for detecting *Cyclospora* and another coccidian like *cryptosporidium* oocysts.

**Ethical issues**

This study was approved by the Ethics Committee of Iran University of Medical Sciences in accordance with the Helsinki Declaration and guidelines and all human participation has been obtained in accordance with informed consent.

**Cyclospora**

The stool was evaluated by direct examination of wet smear and conventional formalin-ether concentration technique was performed, smear was prepared and fixed with methanol. Next, the slide was stained with modified acid-fast staining method (18). The stool was washed in Phosphate-buffered saline (PBS) and preserved in dichromate potassium (2.5%) for following up oocyst sporulation, and finally, autofluorescence of oocyst was observed with immunofluorescence microscope for further confirmation.

DNA extraction was performed by (Roche Diagnostics GmbH, Mannheim, Germany) according to the manufacturer’s instructions, with some modifications. All primers used for *C. cayetanensis* were used from previously defined regions of the 18S ribosomal RNA gene in *C. cayetanensis*. PCR was performed in a 25 µL mixture containing the template (3 µL of DNA), 2.5 U of Taq DNA polymerase, 2.5 µL of 10x PCR buffer, 20 pmol of each primer, 100 µmol dNTPs, and 0.15 mmol MgCl₂.

Nested PCR was performed by initial denaturation of 5 min at 95 °C, followed by a cycling program consisting of 35 cycles of denaturation at 94
°C for 30 sec, annealing at 53 °C for 30 sec, and extension at 72 °C for 90 sec. A final extension at 72 °C for 10 min was followed. A 636-bp product was expected after this round of PCR. The second round was similar to the first one, with the exception that the annealing temperature was 60 °C. The inner primers were expected to produce a 294-bp product in case of the presence of C. cayetanensis DNA as the template.

Nested PCR for the detection of Cyclospora was performed using the outer primers F1E (5’-TACCCATGAAAAACAGTIT-3’) and R2B (5’-CAGGAGAAGCCACCTCTCTTCTT-3’). The inner primers used were F3E (5’-CCITCCTCGGCTCTCGCTCTTGTT-3’) and R4B (5’-CGTCTCTAAAACCCCTACTG-3’) (15).

**E. bieneusi**

Formalin-ether concentration method was performed and after 5 min of methanol fixation, aniline blue staining was done for the detection of microsporidial spores (13).

DNA extraction was performed by (Roche Diagnostics GmbH, Mannheim, Germany) according to the manufacturer’s instructions and nested PCR was performed using E. bieneusi specific primers, designed based on the small subunit (SSU) of rRNA gene.

The nested PCR was done using a set of primers that were specific for E. bieneusi ITS region as well as a portion of the flanking large and small subunit ribosomal RNA genes (400 bp). The outer primers were EBITS3 (5’-GTCGATAGGAGGAAGGAG-3’) and EBITS4 (5’-TTCCAGTTCTTTCCGCGCT-3’), and the inner primers were EBITS1 (5’-GCTCCTAATCTATGCTGCT-3’) and EBITS2.4 (5’-ATCGCGGACGGATCCAGTG-3’). Finally, these reactions produced a fragment of 389 bp (12).

The reaction mixture (25 µL) contained 0.15 mM MgCl2, 2.5 µL of 10x PCR buffer, 100 µM dNTPs, 20 pmol of each primer, and 2.5 U of Taq DNA polymerase. PCR was performed as follows: denaturation at 94 °C for 3 min, followed by 35 cycles of amplification (denaturation at 94 °C for 30 sec, annealing at 57 °C for 30 sec, and elongation at 72 °C for 40 sec), and a final extension at 72 °C for 10 min. Conditions for the secondary PCR were identical to that of the primary PCR, except that only 30 cycles were carried out at an annealing temperature of 55 °C. These reactions produced fragments of 435 and 389 bp, respectively (12).

Finally, PCR products were electrophoresed on 2% agarose gel. Negative and positive controls were included in all sets of PCRs.

### Sequencing

The second round of PCR was performed with inner primers and PCR products were purified using the High Pure PCR Product Purification Kit (Roche Diagnostic, Mannheim, Germany) according to the manufacturer's instructions and were used for direct sequencing using the dye termination method and an ABI 3730xl sequencer (19).

### Results

Out of 102 HIV positive patients, 70 (68.6%) were males and 32 (31.4%) were females including 9 patients with CD4 T-cell count< 200 cells /µl, 17 patients with CD4 200-500 cells/µl, and 76 patients with CD4 > 500 cells/µl. The mean age of patients was 31 yr (range of 19-48 yr). Among 102 stool samples, two (1.96%) cases were found to be positive for intestinal microsporidia infection using aniline blue staining method and were confirmed as E. bieneusi by nested PCR. One patient (0.98%) was found with C. cayetanensis infection by acid-fast staining method and PCR. G. lamblia and B. hominis were detected as non-opportunistic parasites in 1/102 (0.98%) and 2/102 (1.96%) of the HIV positive patients, respectively (Table 1).

Microsporidian spores with aniline blue staining were oval and measured 1-1.5 µm in size (Fig. 1). The positive samples showed positive results for nested PCR for E. bieneusi also. A band of 389 bp corresponding to E. bieneusi was observed in positive cases (Fig. 2). These two positive E. bieneusi patients had chronic diarrhea as main clinical manifestation and showed a CD4 count of less than 200 cells/ µl.
Table 1: Enteric opportunistic and non-opportunistic parasites among HIV/AIDS patients

| Parasites               | CD4 > 500 cells/µl N=76 | CD4 200–500 cells/µl N=17 | CD4 < 200 cells/µl N=9 | Total N (%) |
|-------------------------|--------------------------|---------------------------|------------------------|-------------|
| E. bieneusi             | 2                        | 0                         | 0                      | 2(1.96%)    |
| Cyclospora cayetanensis | 1                        | 1                         | 0                      | 1(0.98%)    |
| Giardia lamblia         | 1                        | 1                         | 0                      | 1(0.98%)    |
| Blastocystis hominis    | 1                        | 1                         | 2                      | 2(1.96%)    |
| Total                   | 2(1.96%)                 | 1(0.98%)                  | 3(2.9%)                | 6/102(5.9)  |

The result of sequencing of two E. bieneusi positive patients in this study based on SSU rRNA gene ITS region demonstrated 99% identity with E. bieneusi isolates HNSC9, 178, HNZM19 and Chan L46, and 98% identity with isolates HNZM54, 356, and 601. The sequences were deposited in GenBank with accession numbers MF163429 and MF163430 (Fig. 3).

C. cayetanensis oocysts were found in a HIV positive patient with watery diarrhea and CD4 count less than 200 cells/µl. The oocysts were almost 8-10 µm in diameter and partial acid-fast positive with wrinkled edge in some oocyst cases (Fig. 4). The result of preservation in dichromate potassium (2.5%) and oocysts sporulation showed that oocysts had two sporocysts and each sporocyst contained two sporozoites.

For further confirmation, autofluorescence of oocyst was observed under immunofluorescence microscope.

C. cayetanensis was also confirmed by nested PCR. A band of 294 bp corresponding to Cyclospora was observed in positive cases (Fig. 5). The result of sequencing and blast showed 99% identity with C. cayetanensis isolates H6 (1-4), PC1, and HMCCPR2. The result of 18S ribosomal rRNA sequences of C. cayetanensis described in this paper was deposited in GenBank with the accession number KY769936.

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Fig. 3: Phylogenetic tree based on *E. bieneusi* nucleotide sequences from two individuals with *E. bieneusi* infection and those corresponding to different *E. bieneusi* genotypes taken from the GenBank database. Bootstrap values ≥70 achieved after 1000 replicates are shown at the nodes.

*G. lamblia* and *B. hominis* were detected as non-opportunistic parasites in 1/102 (0.98%) and 2/102 (1.96%) HIV positive patients with CD4 cell count more than 200 cells/µl in this study.

**Discussion**

Opportunistic infections are found frequently in HIV positive patients, especially when the CD4 T-cell count is less than 200 cells/µl (3). Microsporidiosis in HIV positive patients have been reported from Southeast Asia (India, Thailand), the Middle-East (Turkey), Europe, Africa (Tunisia, Mali, Uganda, Senegal, Zimbabwe), and Latin America (Brazil, Peru) (20).

Diagnosis of *E. bieneusi* infection is based on staining methods and visualization by light microscopy and electron microscopy, and PCR methods. The comparison of results obtained is difficult because of the differences between these methods (21). Prevalence of *E. bieneusi* in HIV positive patients with diarrhea in developed countries like Europe, Australia, and North America has been reported to be between 2%-
78% (21-23), however, this rate was reported from 1.4%-4.3% in HIV positive patients without diarrhea (24).

In this study, 102 HIV positive patients were evaluated to assess the occurrence of protozoa opportunistic infection and two cases were shown to be E. bieneusi positive by staining and nested PCR. Both of these patients had chronic diarrhea. Sequencing confirmed the isolates as E. bieneusi, with 99% identity to E. bieneusi isolates HNSC9, 178, HNZM19, and Chan L46; and 98% identity with isolates HNZM54, 356, and 601.

Prevalence of enteric E. bieneusi in HIV positive patients has been reported in various countries, like Russia (1.2%) (25), Thailand (5.6%) (26), Vietnam (7.1%) (27), Nigeria (2.6%) (28), Switzerland (12.7%) in patients with diarrhea and 0.4% in patients without diarrhea (29), Holland (7.7%) (30), UK (8.3%) (31), Peru (3.9%) (32) using fecal samples by PCR method.

In a study, from 356 HIV positive patients, eight (2.2%) cases were identified as E. bieneusi by staining method and PCR in Shiraz (3). In a study conducted in North India, E. bieneusi was reported in 2.5% of the HIV-infected individuals. Furthermore, both C. cayetanensis and B. hominis were detected in 3.3% of the patients (9).

A systematic review and meta-analysis had reported the overall prevalence of microsporidia infection in immunocompromised patients to be 8.18% in Iran. In addition, prevalence of microsporidia infection was 15.4% in immunocompromised patients with chronic diarrhea, 4.1% in patients without diarrhea, and 12.9% in patients with CD4 less than 200 cells/µL (33).

Some studies carried out in developed countries demonstrated that the prevalence of E. bieneusi in HIV positive patients is decreasing with the use of highly active antiretroviral therapy (HAART) (29, 34).

Prevalence of intestinal parasites in pre-ART and patients with ART treatment was 39% and 17.6%, respectively. All positive cases related to Cryptosporidium spp were found in the pre-ART group and significantly related to CD4 <200 cells/µL (35).

Prevalence of intestinal parasites in HIV/AIDS positive patients was compared between pre-HAART and HAART groups in Brazil. Frequency of Isospora belli, Cryptosporidium sp. and G. lamblia were (4.8% and 1%), (8.1% and 0) and (7.9% and 1%) in pre-HAART and HAART treatment respectively that may due to effect of HAART treatment on intestinal parasites (36).

Effect of antiretroviral therapy on opportunistic
infection like cryptosporidiosis and microsporidiosis in HIV positive patients were evaluated. The result confirmed combination antiretroviral therapy is able to improve the course of cryptosporidiosis and microsporidiosis in HIV-1 positive patients (37). HAART treatment in HIV positive patients in Australia has led to a decrease in microsporidial infections from 11% in 1995 to 0% in 2004 (38).

Contrary to this, prevalence of intestinal parasites remained high in pre- ART in comparison with on-ART patients 84.6% and 82.3% respectively. Opportunistic infection like C. parvum, Microsporidium spp. and I. belli were associated with CD4 <200cells/ µL in on-ART (39).

HIV/AIDS patients referred to Consult Center of Behavior Diseases, West Health Center, Iran University of Medical Sciences in Tehran, were used for the analysis. Two cases of E. bieneusi among 102 HIV positive patients were found with chronic diarrhea as main clinical manifestation. Furthermore, low prevalence of E. bieneusi in this study was interpreted by administration of ART treatment in all patients and use sulfamethoxazole-trimethoprim treatment during the course of AIDS for patients with CD4 T-cell count less than 200 cells/ µl.

C. cayetanensis is a coccidian protozoan observed in many different countries. There are some reports about presence of C. cayetanensis in HIV positive patients from Argentina (40), Turkey (41), and Iran (18). Prevalence of C. cayetanensis has been reported to be 3% in Cuba (42), 11% in Haiti (43), between 0.7%-3.3% in India (9, 17), 2.2% in Thailand (44), and 1% in Tanzania (45).

In this study, one C. cayetanensis positive sample was detected in 102 stool samples collected from HIV positive patients. The sequencing of C. cayetanensis and blast alignment showed 99% identity with C. cayetanensis isolates H6 (1-4), PC1, and HMCCPR2.

No Cryptosporidium spp. was found from 102 stool samples collected from HIV positive patients by modified acid-fast staining method and G. lamblia and B. hominis as non-opportunistic parasites were detected in 1/102 (0.98%), and 2/102 (1.96%) HIV positive patients with CD4 T-cell count more than 200 cells/µl.

Some studies have reported a prevalence range from (1.5%-9.4%) of enteric coccidian, G. lamblia with prevalence range (3.1%-7.3%) and B. hominis (4.4%) as most common non-opportunistic parasites in HIV/AIDS patients in Iran (46, 47).

In this study, E. bieneusi and C. cayetanensis cases were detected from HIV positive patients as opportunistic infection. Phylogenetic mapping indicated that these Iranian E. bieneusi isolates were most closely related to E. bieneusi isolates HNSC9, 178, HNZM19, and Chan L46; and C. cayetanensis isolates showed identity with C. cayetanensis isolates H6 (1-4), PC1, and HMCCPR2.

Conclusion

In this study, with the use of ART treatment in HIV positive patients and administration of sulfamethoxazole-trimethoprim during the course of AIDS for patients with CD4 T-cell counts less than 200 cells/ µl, frequency of microsporidian and coccidion infection was low.

Ethical considerations

This study was approved by the Ethics Committee of Iran University of Medical Sciences code number (IR.IUMS.REC1394-01-131-25831). Ethical issues (Including plagiarism, informed consent, misconduct, data fabrication and/or falsification, double publication and/or submission, redundancy, etc.) have been completely observed by the authors.

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Conflict of interest

The authors declare that there is no conflict of interests.
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