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The first report of onychomycosis caused by *Cryptococcus friedmannii* (*Naganishia friedmannii*) a basidiomycetous yeast



Masoome Ekhtiari^a, Shirin Farahyar^{a,*}, Mehraban Falahati^a, Elham Razmjou^a, Mahtab Ashrafi-Khozani^a, Zeinab Ghasemi^b, Ziba Abbasi-Nejat^{c,d}

^a Department of Medical Parasitology and Mycology, School of Medicine, Iran University of Medical Sciences, Tehran, Iran

^b Department of Medical Parasitology and Mycology, School of Public Health, Tehran University of Medical Sciences, Tehran, Iran

^c International Campus, Iran University of Medical Sciences, Tehran, Iran

^d Deputy of Research and Technology, Golestan University of Medical Sciences, Gorgan, Iran

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ABSTRACT

Yeasts are common etiologic agents of onychomycosis. This study reported a case of onychomycosis due to *Cryptococcus friedmannii* (*Naganishia friedmannii*). This yeast was isolated of the right great toenail of 57year-old man. Microscopic examination of nail scrapings showed budding cells with thin capsule. Sequence analyzes of the internal transcribed spacer regions was closely related to *Cryptococcus friedmannii*. The results of susceptibility testing showed the *Cryptococcus friedmannii* to be sensitive to fluconazole, itraconazole and amphotericin B.

1. Introduction

Onychomycosis is a common nail infection due to yeasts and filamentous fungi [1,2]. The uncommon fungi such as *Trichosporon mucoides* [3] *Aspergillus clavatus* [4] and *Cryptococcus* spp. (basidiomycetous yeasts) [5] were reported as emerging pathogens of onychomycosis.

The basidiomycetous yeasts are important in economic, agricultural and medical problems [6]. Cryptococcus neoformans (Cr. neoformans) was an important agent of cryptococcal infections in immunocompromised patients and Cr. *gatti* is the major pathogen in the genus as well [7,8]. However the reports of infections due to non-*neoformans* have been increased [9]. The species of Cryptococcus genus have been identified from different environmental sources such as air, water, soil, wood, and pigeon excreta. Cr. friedmannii was reported as a new species of yeast from Antarctic in 1985 [10]. The other studies reported that Cr. friedmannii isolated from Beringian and Icelandic soils which were more vegetated. This kind of yeast was reported from soil of Tabriz in Iran as well [11,12]. Cr. friedmannii and some of the yeast species were isolated from the Atacama Desert, with high daily temperature variations. Cr. friedmannii had ability to survive at moderate to cold temperatures [13]. Cr. friedmannii has not been reported as an agent of onychomycosis.

2. Case

A 57-year-old man was admitted to our department in October 2015 (at day 0) with distal subungual hyperkeratosis clinical type of onychomycosis on the first right toenail (Fig. 1). The patient did not have disease like diabetes, psoriasis, immunodeficiencies or any chronic disease. However, four months ago (at day - 4 months), the patient had a history of ungueal trauma and small traumatic lesion was emerged on toenail. Three direct microscopic examinations (at day 0) of nail scrapings, with 20% potassium hydroxide revealed single or budding yeast cells. Three nail specimens were cultured on sabourauds dextrose agar plates with chloramphenicol (at day 0), and incubated at 20-25 °C, which were produced creamy and smooth colonies after a few days (Fig. 2). The other fungi were not isolated. Microscopic examinations (at day 7) of the colonies with Chinese ink were shown single or budding cells with a thin capsule (Fig. 3). Also, lactophenol cotton blue wet mount slide of colonies showed budding cells (Fig. 4). For molecular identification, the yeast was cultured on yeast extract peptone dextrose agar and incubated at 20-25 °C.

Genomic DNA was extracted using Qiagen tissue kit (Germany) (at day 11). The ITS1-5.8S-ITS2 region was amplified with ITS1 (TCC GTA GGT GAA CCT GCG G) and ITS4 (TCC TCC GCT TAT TGA TAT GC) universal primers by the following profile: 98 °C (5 min), 40 cycles of 98 °C (30 s), annealing temperature 56 °C (30 s), and 72 °C (30 s), followed by a final extension of 72 °C (5 min). Amplification of the

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^{*} Correspondence to: Department of Medical Mycology and Parasitology, School of Medicine, Iran University of Medical Sciences, Shahid Hemmat hwy, Tehran 14155 Iran. *E-mail address:* farahyar.sh@iums.ac.ir (S. Farahyar).

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Fig. 1. Distal subungual thickening and hyperkeratosis of patient.



Fig. 2. Creamy and smooth colonies of *Cr. friedmannii* on sabourauds dextrose agar plate with chloramphenicol.



Fig. 3. Budding cells with a thin capsule with Chinese ink wet mount (×400).



Fig. 4. Budding cells in lactophenol cotton blue wet mount slide of colonies (×400).

isolate with ITS1 and ITS4 primers yielded 500 bp fragment (at day 11).

The PCR products were sequenced by Macrogen (Korea). Sequence analyzes was compared with the reference sequences of GenBank database using BLAST (basic local alignment search tool) (http:// www.ncbi.nlm.nih.gov/BLAST). Sequence analyzes was 100% related to *Cr. friedmannii* (*Naganishia friedmannii*) with the accession number KM243311.1 (at day 23). The sequence of the ITS region was submitted to the GeneBank as the accession number KX268322 (Fig. 5).

The antifungal susceptibilities were conducted according to the Clinical and Laboratory Standard Institute method (document M27-S3). The susceptibility tests of *Cr. friedmannii* to fluconazole, Itraconazole and amphotericin B was determined at 72 h. The MICs results revealed this isolate was susceptible to these drugs with MIC values of 0.25, 0.125 and 0.25 μ g/ml for fluconazole, itraconazole and amphotericin B, respectively (at day 30). The patient was treated with oral itraconazole at the dosage of 200 mg daily. Three months later, the symptoms of disease complete remission of the nail. Direct microscopic examinations of nail scrapings with 20% potassium hydroxide had not revealed single or budding yeast cells and culture of the nail sample was negative.

3. Discussion

The yeast *Cr. friedmannii* was first reported as a new species of basidiomycetous yeast of Antarctic in 1985 [10]. The incidence of infection due to non- *Cr. neoformans* has increased. This increased may reveal enhanced immunocompromised patients with impaired cell-mediated immunity, organ transplantation, diabetic patients, azole prophylaxis, and etc [9]. Some of non- *Cr. neoformans* species, such as *Cr. laurentii*, *Cr. albidus*, *Cr. luteolus*, *Cr. uniguttulatus*, *Cr. curvatus* and *Cr. humicola* were reported as opportunistic pathogens over the last few years [9,14]. A case of *Cr. laurentii* keratitis was reported in diabetic patient with onychomycosis. Cultures of the nail and of the contact lens have shown *Cr. laurentii* and *Fusarium solani* [15]. Three species of *Cryptococcus* (*Cr. albidus*, *Cr. uniguttulatus* and *Cr. laurentii*) were isolated of onychomycosis in type 2 diabetes mellitus

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1 gattagaaag cggtctttag teetgeaaca gggeeateeg aagatgteet tagegaaata
61 ettattaege caagteaaae catgtegega gacagateea getattaett ttaagaegag
121 eegaettgae ateggeaaae gteeaaatee aageeaaaga aaggteaaat aaceaateta
181 aggttgaggg tttteatgae acteaaaeag geatgeteet eggaataeea aggagegeaa
241 ggtgegttea aagattegat gatteaetga attetgeaat teaeatteet tategeattt
301 egetgegtte tteategatg egagageeaa gagateegtt gttgaaagtt ttattttgtt
361 ataataagae taeatttgt acattattgt ttagtgtaag tggatgaeta ttgaetttag
421 gttttaeeet atgeettaga ateeteeaae aagtgeaeag gtgttatgga tatgatagaa
481 ageeegtgag caagetegae aggeatetgt atteattaat gateetteeg
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Fig. 5. The sequence of the ITS region was submitted to the GeneBank as the accession number KX268322.

patients [5]. Also, *Prothoteca* spp was reported as the first causative agent of onychomycosis in Brazil which was the 3rd case in the world [16]. The distribution of pathogens of onychomycosis is not uniform. It is related to several factors such as geographical region and climatic conditions. Furthermore, onychomycosis is related to the host ones such as age, occupation, chronic diseases, nail care and lifestyle. Deformity and pain in nails are caused by the fungi in onychomycosis. Thus, affected individuals may experience difficulties while walking and writing. Therefore, accurate diagnosis of the agents is important for suitable therapeutic approach. This study reports *Cr. friedmannii* is the first case of onychomycosis in the world.

Conflict of interest

There are none.

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