Fungal Melanonychia: Ungual Phaeohyphomycosis caused by *Botryosphaeria dothidea*

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The phaeoid fungus *Neoscytalidium dimidiatum* and a dermatophyte *Trichophyton rubrum* are the most frequently isolated agents causing fungal melanonychia, followed by the phaeoid fungi of the genera *Alternaria* and *Exophiala* (1, 2). Phaeohyphomycosis is an umbrella term describing infections in humans and other animals characterized primarily by the development of dark-coloured hyphae that are caused by phaeoid fungi. The term “phaeohyphomycosis” encompasses broad mycotic infections, regardless of the site of the lesion; the pattern of tissue response, granuloma, or abscess; or the taxonomy of the actiological agents (3). We describe here a classic example of ungual phaeohyphomycosis.

**CASE REPORT**

An 82-year-old retired Japanese farmer with chronic heart failure and hemiparesis noticed a black-pigmented area on his right thumbnail that had been present for 6 months. His internist (KS) considered the possibility of malignant melanoma. Upon presentation in February 2016, the nail was thickened and rough with a greyish-white surface, and black pigmentation was observed from its centre to the proximal side. Hutchinson’s sign was not observed (Fig. 1a). The discoloured area accounted for 91% of the entire nail plate (Image J version 1.48; National Institutes of Health). The opacity ratio decreased from 91% to 32% during that time.

Direct microscopy revealed branching brown septate hyphae (Fig. 2a), and haematoxylin and eosin and periodic acid-Schiff staining revealed black septate hyphae in the nail plate. The thick cell wall appeared to be double contoured (Fig. 2b). Plate culture on potato dextrose agar after 7 days at 30°C showed a greyish-white woolly colony with coal-black pigmentation on the reverse (Fig. 2c, d). Slide culture showed pigmented broad and unpigmented narrow branching hyphae and intercalary and acropetal chlamydoconidia measuring up to 40 μm (Fig. 2e). The sequence of the internal transcribed spacer 1 region of the ribosomal RNA gene from the nail and isolate had 99% homology to the *Botryosphaeria dothidea* type strain CBS 115476 (accession: KF766151) (4). Based on the morphological characteristics and gene analysis results, we diagnosed the patient with ungual phaeohyphomycosis due to *B. dothidea* (Moug. ex Fr.) Ces. & De Not. The minimum inhibitory concentrations for the isolate were as follows: amphotericin B, 0.25 μg/ml; efinaconazole (EFCZ), 1.0 μg/ml; fluconazole, > 64 μg/ml; 5-fluorocytosine, 2 μg/ml; itraconazole, > 16 μg/ml; miconafungin, 0.25 μg/ml; miconazole, 0.25 μg/ml; terbinafine, 0.5 μg/ml; and voriconazole, 0.03 μg/ml. Nail opacity was reduced to 74% and 32% of the nail surface after 1 and 4 months, respectively, of topical application of 10% EFCZ solution (Fig. 1c). The discoloration disappeared with negative conversion of the fungi after 7 months. The patient showed no recurrence of fungal infection during a 3-month follow-up, as of December 2016.

**Fig. 1**. Clinical characteristics of the thumb nail. (a) Clinical image at presentation. Melanonychia without Hutchinson’s sign. (b) Dermoscopic image showing a black homogenous area and coarse granules proximally. (c) Findings after 4 months of topical application of 10% efinaconazole solution. The opacity ratio decreased from 91% to 32% during that time.

**Fig. 2**. Mycologic characteristics. (a) Direct microscopic examination revealed branching black hyphae (KOH preparation, original magnification ×400). (b) Septate black hyphae in the nail plate (periodic acid-Schiff staining ×400). (c, d) Plate culture showed a greyish-white woolly colony with a coal-black pigmentation on the reverse. (e) Slide culture showed pigmented broad and unpigmented narrow hyphae and chlamydoconidia. (Lactophenol Cotton Blue staining ×400).
DISCUSSION

*Botryosphaeria dothidea* was first described as *Sphaeria dothidea* Moug. in 1823. The type species of the genus (family Botryosphaeriaceae, order Botryosphaerales, class Dothideomycetes, phylum Ascomycota) is *B. dothidea*. Fungi in the order Botryosphaerales are the most widespread and important plant-pathogens associated with twig, branch, and stem cankers; tip and branch dieback; fruit rot; blue stain; and plant death (5). To date there have been no reports of *B. dothidea* infection in humans or lower animals, while *N. dimidiatum* is the most frequent non-dermatophyte and non-*Candida* pathogen causing onychomycosis in tropical and subtropical regions (6). A survey in Brazil showed that the typical symptom associated with onychomycosis caused by *N. dimidiatum* was a lateral-distal and lateral-subungual lesion in the big toenail, and melanonychia accounted for 66.6% (20/30) of all cases. It developed in 27.2% (3/11) of Caucasians and 89.4% (17/19) of non-Caucasian individuals (7). The sequence of the internal transcribed spacer 1 region of the ribosomal RNA gene from the isolate and the nail in our case had 95% homology with the *N. dimidiatum* type strain CBS 499.66 (acccession: FM211432) (4). Our case occurred on the Kyushu island of Japan, which is located in the temperate region at 32° north latitude with a mean temperature of 17.2 ± 0.5°C. The annual mean temperature in the Northern hemisphere has increased by 0.85°C during the past 30 years (1983–2012) (8). The incidence of *B. dothidea* infections might therefore increase gradually even in temperate regions.

Although pigment cells exist in both the nailbed and the nail matrix, melanoma derived from the nailbed has an amelanotic tendency. It is important to differentiate melanonychia arising in the proximal part of the nail from malignant melanoma even if Hutchinson’s sign is not noted (9). The characteristic dermoscopic images of fungal melanonychia are homogeneous pigmented coarse granular lesions with fine black granules of a diameter of 0.1 mm or less (9), which were useful for ruling out a diagnosis of malignant melanoma. The clinical appearances of longitudinal melanonychia were similar to those caused by *Phialophora* species (10) and were different from those caused by *T. rubrum* presenting bundle-forming longitudinal lines with a brownish or yellow element (11).

Because onychomycosis caused by *N. dimidiatum* is refractory and resistant to itraconazole and terbinafine (6), the conventional therapy was expected to be ineffective against *B. dothidea* infection. In this case, topical application of 10% EFCZ solution was effective. EFCZ is a novel triazole antifungal agent that has good human nail permeability and retention properties (12, 13). Because of its broad spectrum of antifungal activities (14), EFCZ is a promising option in the treatment of non-dermatophyte onychomycosis including ungual phaeohyphomycosis (15).

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The authors declare no conflicts of interest.

REFERENCES