

Detection of *Trichophyton indotineae* in Austria: a case report

I. Indikova¹, L. Teodorowicz¹, K. Loydl¹, A. Stary¹

Outpatients Centre for Diagnosis of Infectious Venero-dermatological Diseases, Vienna, Austria

Objectives and aim: *Trichophyton (T.) indotineae* (Fig. 1) is a newly described species of dermatophyte. This fungal pathogen is widespread in India and is responsible for chronic or recurrent widespread superficial infections (Fig. 2-3). It is often associated with resistance to terbinafine, that results from a point mutation in the gene encoding squalene epoxidase.

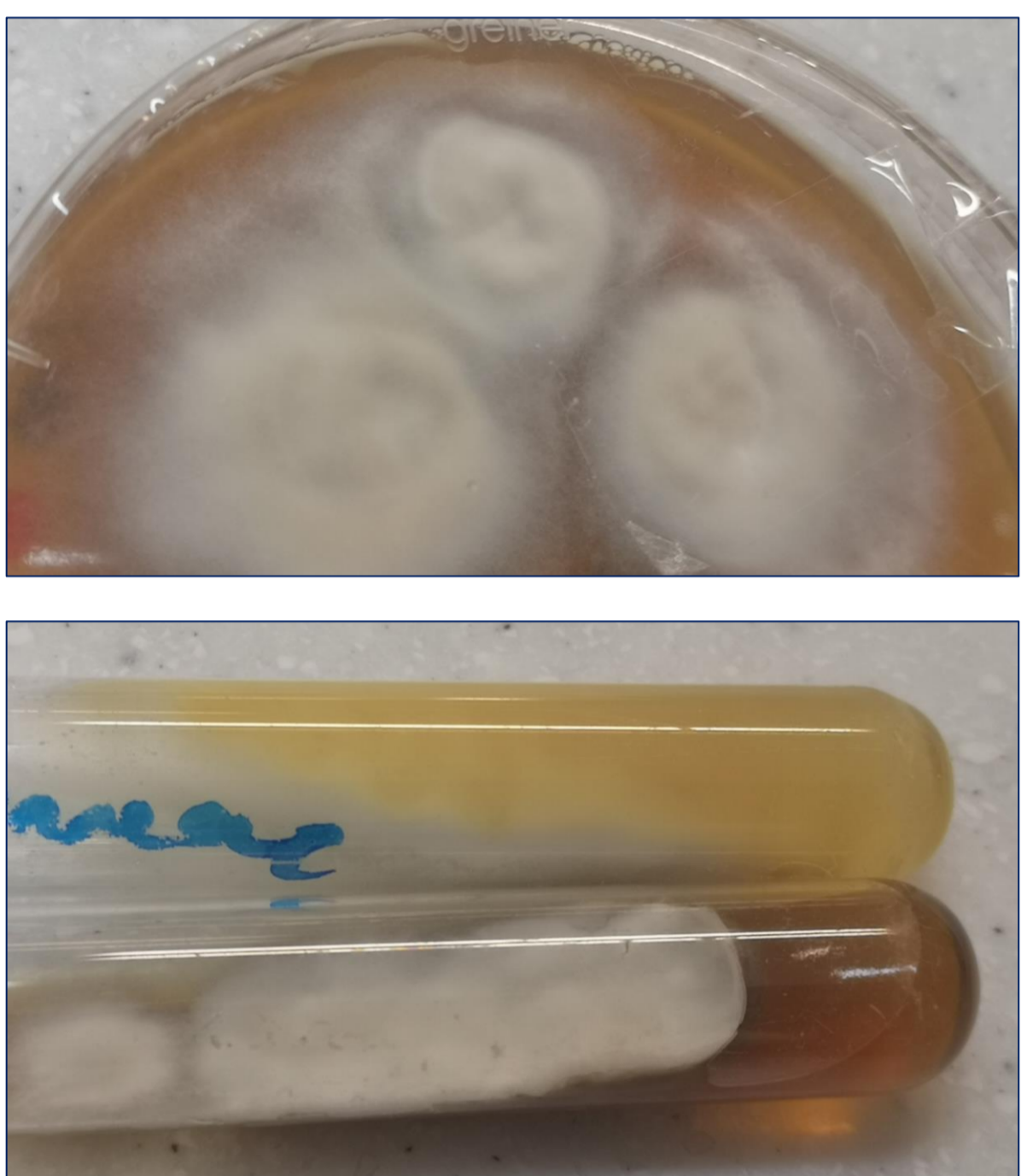


Fig 1: Culture of *T. indotineae*



Fig 2: Spreading of infections caused by *T. indotineae* (Jabet et al., 2023)



Fig 3: Extensive dermatomycosis by *T. indotineae* (Uhrlaß et al., 2022)

Patients and Methods:

- A female patient (32 years old), origin in India, with extensive tinea corporis (Fig. 4), referred to the Outpatient's centre (Juni 2022) for fungal diagnosis.
- Cultivation of skin scrapings on Sabourand agar at 28° for 21 days
- Initial identification based on both the colony morphology and microscopic observation.
- Final identification of the *Trichophyton* species by sequencing of the entire ITS region of the rDNA and a subsequent database search (courtesy of Labor Mölbis, Leipzig, Germany).
- Antifungal susceptibility by the Ezy MIC™ (Himedia).

Fig 4: Tinea corporis



Results:

- Diagnosis by culture: member of the *T. mentagrophytes/interdigitale* complex, suspicious for *T. indotineae*.
- Sequencing: clear and 100% match with the species *Trichophyton indotineae* = *T. mentagrophytes* ITS genotype VIII India.
- Antifungal susceptibility testing: undiminished susceptibility to terbinafine.

Conclusion:

The results proved the identity of the dermatophyte *Trichophyton indotineae* = *T. mentagrophytes* ITS genotype VIII India and confirmed the spread of the fungus across Europe. It was one of the first *T. indotineae* isolates in Austria.

Update:

In early 2023, we identified another case of dermatomycosis caused by *T. indotineae*. As before, the identity of the strain was confirmed by sequencing (Mölbis Laboratory, Leipzig, Germany). This time the strain showed complete resistance to terbinafine.

Literature:

• Uhrlaß, et al. J. Fungi 8 2022; 7:757.

• Nenoff, et al. Indian J Dermatol, Venereol Leprol. 2022; 88, 5: 586-589.

• Jabet, et al. Med Mycol 2023, 33 (3): 101383