SUPPLEMENTARY FIGURE. Phylogenetic relationships* among the internal transcribed spacer regions of rDNA of *Trichophyton indotineae* isolates from patient A† and patient B§,¶ and sequences of two *Trichophyton interdigitale* isolates** and sequences deposited at the National Center for Biotechnology Information††

* Generated in CLC Genomics Workbench (version 23.0.2) using the Neighbor-Joining algorithm with a bootstrap analysis of 1,000 replicates. *T. quinckeanum* (KU257460) was used as an outgroup (distantly related comparison group).
† M20518: Patient A
§ M22558: Patient B
¶ The sequences of isolates from patients A and B were identical and grouped with the reference strain of *T. indotineae*.
** M22570 and M22691
†† *T. mentagrophytes XIV* (MK312950), *T. indotineae* (MH791418), and *T. interdigitale II* (MK447596)