In recent years, dermatologists in India have alerted the medical community to the widespread occurrence of recalcitrant extensive dermatophytosis across the country. Clinically, extensive dermatophytosis is characterized by tinea cruris, tinea corporis, or both, of the glabrous skin (1). The spread of this condition is thought to be a consequence of ill-advised use of over-the-counter corticosteroid-antifungal combinations, resulting in the emergence of terbinafine-resistant Trichophyton strains in India (2). T. mentagrophytes and T. interdigitale were suspected, but the correct identity of the etiologic agent of this outbreak was debated (3,4). Genomic data showed that India terbinafine-resistant isolates form a distinct clade from T. interdigitale and T. mentagrophytes (5). Recently, the clinical, mycological, and molecular features of 2 highly resistant T. interdigitale isolates from patients in Nepal and India, harboring mutations in the squalene epoxidase (SQLE) gene, have been analyzed and identified as a new species named T. indotineae (6).

The taxonomy of the T. mentagrophytes/T. interdigitale complex, including T. indotineae, has been revised based on multigene phylogeny revealing that analysis of the high-mobility group gene clearly demarcates the species, as suggested elsewhere (5,7). T. indotineae appears to be the primary contributor to terbinafine resistance (8). Migration and travel enable the spread of nonautochthonous pathogens in nonendemic areas. Terbinafine-resistant T. indotineae isolates have been recently identified in Europe (9–11). We present a series of extensive dermatophytosis cases in France caused by terbinafine-resistant T. indotineae and document the worldwide spread of T. indotineae using results from internal transcribed spacer (ITS) sequence-based screening.

The Study
Ten patients in 4 hospitals in France with clinical manifestations of tinea cruris or tinea corporis were diagnosed with extensive dermatophytosis caused by terbinafine-resistant Trichophyton indotineae harboring Phe397Leu and Leu393Ser substitutions in the squalene epoxidase enzyme was diagnosed in France. Analysis of internal transcribed spacer sequences revealed the wide spread of this species in Asia and Europe. Detection of T. indotineae in animals suggests their possible role as reservoirs.

Extensive dermatophytosis caused by terbinafine-resistant Trichophyton indotineae harboring Phe397Leu and Leu393Ser substitutions in the squalene epoxidase enzyme was diagnosed in France. Analysis of internal transcribed spacer sequences revealed the wide spread of this species in Asia and Europe. Detection of T. indotineae in animals suggests their possible role as reservoirs.

Extensive Dermatophytosis Caused by Terbinafine-Resistant Trichophyton indotineae, France

Arnaud Jabet, Sophie Brun, Anne-Cécile Normand, Sébastien Imbert, Mohammad Akhoundi, Eric Dannaoui, Laetitia Audiffred, François Chasset, Arezki Izri, Liliane Laroche, Renaud Piarroux, Claude Bachmeyer, Christophe Hennequin, Alicia Moreno Sabater.
stay in refugee camps in Turkey and Bangladesh. For patients 1, 4, and 8, skin lesions had appeared after living in France for 3–4 years, suggesting that human-to-human transmission could have occurred in France. All patients declared no contact with animals.

Eight out of 10 patients received oral treatment with terbinafine. Clinical cure or treatment failure were difficult to evaluate because most patients could not be reached for follow-up. We observed clinical cure in 2/8 patients (patients 1 and 3) and clinical improvement in 2/8 patients (patients 2 and 9). However, we observed no clinical improvement or relapse in 4/8 patients (patients 4, 5, 7, and 10). Patient 4 received a second round of oral griseofulvin treatment during the 3 months after initial treatment, and patient 10 received a second round of oral terbinafine for 3 months followed by griseofulvin treatment over 6 months, without clear clinical improvement for either. Finally, we observed clinical improvement in patient 10 after 1 month of a 2-month treatment with itraconazole (400 mg/d). Indeed, limited effectiveness of 4 oral antifungals (fluconazole, griseofulvin, itraconazole, and terbinafine) has been described in India, but itraconazole has shown better efficacy than the others (13). Increasing terbinafine exposure through higher doses or longer durations has been proposed to overcome treatment failure and could protect azole-based antifungal drugs from developing resistance (14).

Seven isolates initially diagnosed as *T. mentagrophytes* by microscopy, mass spectrometry, and clinical localization were available for further characterization (Table 1). ITS region sequencing revealed that all isolates were *T. indotineae*. Three out of 6 isolates grew on a solid terbinafine-containing medium (Appendix Figure 2). We confirmed terbinafine resistance by the EUCAST (https://eucast.org) method (15) with MICs of 2 or >8 μg/mL (Table 1). Sequencing the SQLE gene revealed that 4 resistant isolates contained Leu393Ser,
Phe397Leu, or Phe397Leu/Ala448Thr substitutions. Three terbinafine-susceptible isolates exhibited no substitutions or an Ala448Thr substitution alone. We determined low MICs for itraconazole, voriconazole, and amorolfine for 6 isolates (Table 1).

Using an ITS sequence-based screening of sequences stored in GenBank and a review of literature up through March 2021 (Appendix Table 2), we investigated the epidemiologic characteristics of T. indotineae. Information was available about the origin of the infection for 526/537 sequences found. Human-to-human transmission was predominant because 98.8% of the sequences identified were of human origin; however, 6 sequences indicated an animal origin. Two sequences detected during a survey that included 760 calves came from Egypt, 1
detected in an infected dog came from India, and 3 came from Poland but with no specific animal host indicated. These results suggest that animals can be reservoirs of *T. indotineae* and zoonotic transmission must be considered.

We obtained geographic information for all 537 sequences and temporal information for 486 sequences. Our study revealed that *T. indotineae* was present in India, Australia, Iran, and Oman during 2004–2013 (Figure 1, panel A). After 2014, a substantial increase in reported cases was observed, related to the outbreak in India. Since 2019, the number of reported *T. indotineae* cases has increased in Europe, confirming its spread. Currently, 76% of the known sequences have been identified in India, 12.8% in the Middle East, 9.6% in Europe, and 1.1% in other countries (Figure 1, panel B). Cases reported in Europe are supposed to have been introduced by migrants or travelers from India, Bangladesh, Pakistan, Bahrain, Libya, Saudi Arabia, or Thailand, suggesting the presence of *T. indotineae* in those countries. The cases imported from Bangladesh that were reported in France, together with those reported in Germany (9), suggest that *T. indotineae* transmission could be endemic in Bangladesh. Results obtained in this study were limited to data obtained from sources available in GenBank; *T. indotineae* distribution is probably greater than what is documented here.

The epidemiology of terbinafine-resistant *T. indotineae* isolates was difficult to assess because studies recording available molecular analysis and in vitro antifungal susceptibility testing were scarce. In India, 71.3% (n = 279) of reported isolates were resistant to terbinafine, but in Iran, 71.8% (n = 32) were susceptible. Of the isolates from this study, 50% (n = 29) from Germany and 57.1% (n = 7) from France were resistant. Isolates from India showed 11 different single or combined missense mutations of the SQLE gene with a large range of terbinafine MICs (Figure 2). Of note, Phe397Leu and Leu393Phe substitutions, associated with terbinafine resistance, were predominant in isolates from India and Germany, probably related to population movements between the 2 countries. Leu393Ser substitution was predominant in France in the isolates from Bangladesh.

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**Figure 2.** *Trichophyton indotineae* susceptibility to terbinafine reported from 4 countries. A) Relationships between terbinafine MIC and codon changes reported in isolates from different countries. Grey line shows terbinafine susceptibility threshold of 0.2 μg/mL. Available MICs were determined using the Clinical Laboratory and Standards Institute (https://clsi.org) or EUCAST (https://eucast.org) methods. Data show mean values. B) Prevalence of substitution points in the gene encoding SQLE. Sources shown in the Appendix (https://wwwnc.cdc.gov/EID/article/28/1/21-0883-App1.pdf). SQLE, squalene epoxidase enzyme.
Conclusions
The medical community and organizations receiving migrants and travelers should be aware that extensive dermatophytosis linked to terbinafine-resistant *T. indotineae* has reached France. Efficient systems to promptly identify terbinafine-resistant *T. indotineae* isolates must be implemented to halt the progression of this pathogen throughout Europe.

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About the Author
Dr. Jabet was a final-year resident at La Pitié-Salpêtrière Hospital (now a medical mycologist at La Pitié-Salpêtrière Hospital and Saint Antoine Hospital) in Paris, France. His primary research interests are dermatophytosis infections and emerging fungal pathogens. Dr. Brun is a medical mycologist working at Avicenne Hospital in Bobigny, France, and is particularly interested in dermatophytosis and fungal infections observed in migrants.

References

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Extensive Dermatophytosis Caused by Terbinafine-Resistant *Trichophyton indotineae*, France

Appendix

**Appendix Table.** PCR primers used in this study and annealing temperatures*

<table>
<thead>
<tr>
<th>DNA portion</th>
<th>Primer</th>
<th>Sequence</th>
<th>Annealing temperature</th>
</tr>
</thead>
<tbody>
<tr>
<td>ITS</td>
<td>ITS1</td>
<td>TCCGTAGGTGAACCTGCGG</td>
<td>55°C</td>
</tr>
<tr>
<td></td>
<td>ITS3</td>
<td>GCATCGATGAAGAACGCAGC</td>
<td></td>
</tr>
<tr>
<td></td>
<td>ITS4c</td>
<td>TCCCTCCGTTATTGATATGC</td>
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<tr>
<td>Squalene epoxidase</td>
<td>Erg1-2-F</td>
<td>CCAGACTGATGGCAAGCAAGA</td>
<td>60°C (used only for sequencing)</td>
</tr>
<tr>
<td></td>
<td>Erg1-2_R</td>
<td>ATAAGCTCCAGGCCCAGAA</td>
<td></td>
</tr>
<tr>
<td></td>
<td>TrSQLE-F1</td>
<td>ATGGTTGTAGAGGCTCTCCC</td>
<td></td>
</tr>
<tr>
<td></td>
<td>TrSQLE-R1</td>
<td>CTAGCTTTGAAGTGTCGAAA</td>
<td></td>
</tr>
</tbody>
</table>

* Primers were previously described by Yamada et al. (1). ITS, internal transcribed spacer.

**Appendix Figure 1.** Clinical presentation of extensive dermatophytosis diagnosed in France. A–B) The various lesions consisted in multiple and large erythematous or brown, round to oval macular lesions on the abdomen, legs, arms and inguinal folds, slightly scaly, with a raised inflammatory edge. Patients reported peripheral spreading but no central clearing. C) An atypical clinical presentation was observed in one patient (patient 9) with large and brown annular macular lesions and no inflammatory edge.
Appendix Figure 2. Terbinafine susceptibility screening using the solid plate method. Photos show antifungal susceptibility to terbinafine determined using the growth method involving Sabouraud-chloramphenicol-cycloheximide medium (SCC) with or without terbinafine at 0.2 μg/mL (lower and upper photos respectively). A–F) Patients 3, 5, 6, 7, 9, 10, respectively. Terbinafine susceptibility using this method could be determined for 6 isolates (loss of viability for isolate 4). Isolates from patients 5, 7, 10 were able to grow in terbinafine-containing solid medium.

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Additional Sources

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