

immune response influences the course of leprosy infection; it is challenging to understand the genetics of disease susceptibility and immunopathogenesis of leprosy (4,5).

With an inference of only a single case study, it is hard to say that *M. lepromatosis* lepromatous leprosy is a travel-related hazard for all US citizens. More surveillance data, such as patients' immunity toward the disease, their genetic susceptibility, and travel history, are needed to explore the travel-related hazard. In addition, evolutionary knowledge and how widely the disease is circulating in nonendemic regions will help in understanding the nature of the disease.

## References

1. Virk A, Pritt B, Patel R, Uhl JR, Bezalel SA, Gibson LE, et al. *Mycobacterium lepromatosis* lepromatous leprosy in US citizen who traveled to disease-endemic areas. *Emerg Infect Dis*. 2017;23:1864–6. <http://dx.doi.org/10.3201/eid2311.171104>

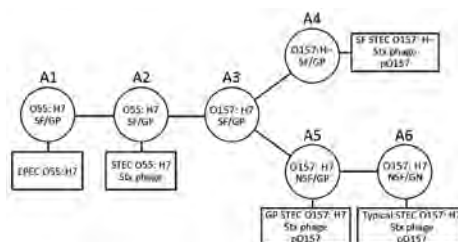
2. Singh P, Benjak A, Schuenemann VJ, Herbig A, Avanzi C, Busso P, et al. Insight into the evolution and origin of leprosy bacilli from the genome sequence of *Mycobacterium lepromatosis*. *Proc Natl Acad Sci U S A*. 2015;112:4459–64. <http://dx.doi.org/10.1073/pnas.1421504112>
3. Pinheiro RO, de Souza Salles J, Sarno EN, Sampaio EP. *Mycobacterium leprae*–host-cell interactions and genetic determinants in leprosy: an overview. *Future Microbiol*. 2011; 6:217–30. <http://dx.doi.org/10.2217/fmb.10.173>
4. Araújo S, Lobato J, Reis EM, Souza DOB, Gonçalves MA, Costa AV, et al. Unveiling healthy carriers and subclinical infections among household contacts of leprosy patients who play potential roles in the disease chain of transmission. *Mem Inst Oswaldo Cruz*. 2012;107(Suppl 1):55–9. <http://dx.doi.org/10.1590/S0074-02762012000900010>
5. Blake LA, West BC, Lary CH, Todd JR IV. Environmental nonhuman sources of leprosy. *Rev Infect Dis*. 1987;9:562–77. <http://dx.doi.org/10.1093/clinids/9.3.562>

Address for correspondence: Gaurav Sharma, Department of Chemistry, University of Miami, 1301 Memorial Dr, Coral Gables, FL 33146, USA; email: gxs183@miami.edu

## Corrections

### Vol. 24, No. 12

In Figure 1 of the article Genomic Characterization of  $\beta$ -Glucuronidase-Positive *Escherichia coli* O157:H7 Producing Stx2a (Y. Ogura et al.), Shiga toxin-producing *Escherichia coli* O157 was mislabeled several times, and the term STEC was incompletely defined. The corrected figure is shown, and the article has been corrected online ([https://wwwnc.cdc.gov/eid/article/24/12/18-0404\\_article](https://wwwnc.cdc.gov/eid/article/24/12/18-0404_article)).



### Vol. 25, No. 1

Table 3 misstated the number of animals tested during 2001–2004 and the first author's biographical sketch was incorrect in Multiple Introductions of Domestic Cat Feline Leukemia Virus in Endangered Florida Panthers (E.S. Chiu et al.). The article has been corrected online ([https://wwwnc.cdc.gov/eid/article/25/1/18-1347\\_article](https://wwwnc.cdc.gov/eid/article/25/1/18-1347_article)).

Author Sang-Ho Choi's name was listed incorrectly and author affiliations were unclear in Clinical and Radiologic Characteristics of Human Metapneumovirus Infections in Adults, South Korea (H.J. Koo et al.). The article has been corrected online ([https://wwwnc.cdc.gov/eid/article/25/1/18-1131\\_article](https://wwwnc.cdc.gov/eid/article/25/1/18-1131_article)).

Two locations in Figure 1 were shown incorrectly in Risk Factors for *Elizabethkingia* Acquisition and Clinical Characteristics of Patients, South Korea (M.H. Choi et al.), and the article has been corrected online ([https://wwwnc.cdc.gov/eid/article/25/1/17-1985\\_article](https://wwwnc.cdc.gov/eid/article/25/1/17-1985_article)).

