Genetic Diversity and Acquired Drug Resistance Mutations Detected by Deep Sequencing in Virologic Failures Among Antiretroviral Treatment Experienced Human Immunodeficiency Virus-I Patients in a Pastoralist Region of Ethiopia [Corrigendum]

Tachbele E, Kyobe S, Katabazi FA, et al. *Infect Drug Resist.* 2021;14:4833–4847.

The authors have advised affiliation 4 on page 4833 is incorrect. The correct affiliation should read “J. Craig Venter Institute, Rockville, MD, USA”.

Rembert Pieper was not affiliated with Janssen Biopharma, South San Francisco, CA, USA, when the work was conducted. It is their current affiliation and should have only been noted as an acknowledgment. The authors apologize for this error.

Publish your work in this journal

Infection and Drug Resistance is an international, peer-reviewed open-access journal that focuses on the optimal treatment of infection (bacterial, fungal and viral) and the development and institution of preventive strategies to minimize the development and spread of resistance. The journal is specifically concerned with the epidemiology of

antibiotic resistance and the mechanisms of resistance development and diffusion in both hospitals and the community. The manuscript management system is completely online and includes a very quick and fair peer-review system, which is all easy to use. Visit <http://www.dovepress.com/testimonials.php> to read real quotes from published authors.

Submit your manuscript here: <https://www.dovepress.com/infection-and-drug-resistance-journal>

<https://doi.org/10.2147/IDR.S352888>