

## Genetic Diversity and Acquired Drug Resistance Mutations Detected by Deep Sequencing in Virologic Failures Among Antiretroviral Treatment Experienced Human Immunodeficiency Virus-1 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.

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