

Erratum: Genomic and Epidemiological Analysis of SARS-CoV-2 Viruses in Sri Lanka

Frontiers Production Office*

Frontiers Media SA, Lausanne, Switzerland

Keywords: SARS-CoV-2, genomic sequencing, pangolin lineages, B.1.411, B.1.1.7, molecular epidemiology

OPEN ACCESS An Erratum on

Approved by:

Frontiers Editorial Office, Frontiers Media SA, Switzerland

*Correspondence:

Frontiers Production Office production.office@frontiersin.org

Specialty section:

This article was submitted to Infectious Agents and Disease, a section of the journal Frontiers in Microbiology

> Received: 17 March 2022 Accepted: 17 March 2022 Published: 11 April 2022

Citation:

Frontiers Production Office (2022) Erratum: Genomic and Epidemiological Analysis of SARS-CoV-2 Viruses in Sri Lanka. Front. Microbiol. 13:898684. doi: 10.3389/fmicb.2022.898684 Genomic and Epidemiological Analysis of SARS-CoV-2 Viruses in Sri Lanka

by Jeewandara, C., Jayathilaka, D., Ranasinghe, D., Hsu, N. S., Ariyaratne, D., Jayadas, T. T., Panambara Arachchige, D. M., Lindsey, B. B., Gomes, L., Parker, M. D., Wijewickrama, A., Karunaratne, M., Ogg, G. S., de Silva, T. I., and Malavige, G. N. (2021). Front. Microbiol. 12:722838. doi: 10.3389/fmicb.2021.722838

Due to a production error, affiliations 4 and 5 were incorrectly published as:

⁴MRC Weatherall Institute of Molecular Medicine, National Institute of Infectious Diseases, Angoda, Sri Lanka

⁵University of Oxford, Oxford, United Kingdom

The correct affiliations 4 and 5 are:

⁴ National Institute of Infectious Diseases, Angoda, Sri Lanka

⁵ MRC Weatherall Institute of Molecular Medicine, University of Oxford, Oxford, United Kingdom

The publisher apologizes for this mistake. The original version of this article has been updated.

Copyright © 2022 Frontiers Production Office. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.