

CORRECTION

Open Access



# Correction: Whole-genome sequencing of non-typeable *Haemophilus influenzae* isolated from a tertiary care hospital in Surabaya, Indonesia

Made Ananda Krisna<sup>1,2\*</sup>, Lindawati Alimsardjono<sup>3</sup>, Korrie Salsabila<sup>1,6</sup>, Naritha Vermasari<sup>3</sup>,  
Wa Ode Dwi Daningrat<sup>1,4</sup>, Kuntaman Kuntaman<sup>3</sup>, Odile Barbara Harrison<sup>5</sup>,  
Martin Christopher James Maiden<sup>2</sup> and Dodi Safari<sup>1\*</sup>

**Correction: BMC Infect Dis 24, 1097 (2024)**  
<https://doi.org/10.1186/s12879-024-09826-8>

Following publication of the original article [1], it was flagged to us that an additional statement needs to be included in the Acknowledgements section (marked in bold below):

“The archived specimen in this study was obtained from the capacity-building project funded by the Grant or Cooperative Agreement Number NU2GGH001852-03, funded by the Centers for Disease Control and

Prevention. Its contents are solely the responsibility of the authors and do not necessarily represent the official views of the Centers for Disease Control and Prevention or the Department of Health and Human Services. **The genomic analyses in this study was funded by a Wellcome Trust Biomedical Resource Grant to MJCM (grant number 218205/Z/19/Z).** The funders had no role in the study design, data collection and analysis, decision to publish, or preparation of the manuscript.”

The original article has been corrected.

Published online: 21 November 2024

The original article can be found online at <https://doi.org/10.1186/s12879-024-09826-8>.

\*Correspondence:

Made Ananda Krisna  
made.krisna@biology.ox.ac.uk  
Dodi Safari  
dodi004@brin.go.id

<sup>1</sup> Eijkman Research Centre for Molecular Biology, National Research and Innovation Agency, Cibinong, West Java, Indonesia

<sup>2</sup> Department of Biology, University of Oxford, Oxford, UK

<sup>3</sup> Department of Clinical Microbiology, Dr. Soetomo Academic General Hospital, Surabaya, Indonesia

<sup>4</sup> Centre for Genomic Pathogen Surveillance, Nufeld Department of Clinical Medicine, University of Oxford, Oxford, UK

<sup>5</sup> Nufeld Department of Population Health, University of Oxford, Oxford, UK

<sup>6</sup> Graduate School of Medical and Pharmaceutical Sciences, Chiba University, Chiba, Japan

## Reference

1. Krisna, et al. Whole-genome sequencing of non-typeable *Haemophilus influenzae* isolated from a tertiary care hospital in Surabaya, Indonesia. *BMC Infect Dis.* 2024;24:1097. <https://doi.org/10.1186/s12879-024-09826-8>.



© The Author(s) 2024. **Open Access** This article is licensed under a Creative Commons Attribution 4.0 International License, which permits use, sharing, adaptation, distribution and reproduction in any medium or format, as long as you give appropriate credit to the original author(s) and the source, provide a link to the Creative Commons licence, and indicate if changes were made. The images or other third party material in this article are included in the article's Creative Commons licence, unless indicated otherwise in a credit line to the material. If material is not included in the article's Creative Commons licence and your intended use is not permitted by statutory regulation or exceeds the permitted use, you will need to obtain permission directly from the copyright holder. To view a copy of this licence, visit <http://creativecommons.org/licenses/by/4.0/>.