



# **Corrigendum: An Evolutionary Arms Race Between** *Burkholderia pseudomallei* **and Host Immune System: What Do We Know?**

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### **OPEN ACCESS**

#### Approved by:

Frontiers Editorial Office, Frontiers Media SA, Switzerland

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#### Specialty section:

This article was submitted to Evolutionary and Genomic Microbiology, a section of the journal Frontiers in Microbiology

Received: 26 October 2021 Accepted: 29 October 2021 Published: 22 November 2021

#### Citation:

Chomkatekaew C, Boonklang P, Sangphukieo A and Chewapreecha C (2021) Corrigendum: An Evolutionary Arms Race Between Burkholderia pseudomallei and Host Immune System: What Do We Know? Front. Microbiol. 12:801975. doi: 10.3389/fmicb.2021.801975 <sup>1</sup> Mahidol-Oxford Tropical Medicine Research Unit (MORU), Bangkok, Thailand, <sup>2</sup> Bioinformatics and Systems Biology Program, School of Bioresource and Technology, King Mongkut's University of Technology Thonburi, Bangkok, Thailand, <sup>3</sup> Wellcome Sanger Institute, Hinxton, United Kingdom

Keywords: melioidosis, evolution, burkholderia, host immune system, genetic variants

#### A Corrigendum on

## An Evolutionary Arms Race Between *Burkholderia pseudomallei* and Host Immune System: What Do We Know?

by Chomkatekaew, C., Boonklang, P., Sangphukieo, A., and Chewapreecha, C. (2020). Front. Microbiol. 11:612568. doi: 10.3389/fmicb.2020.612568

In the original article, there was a mistake in **Figure 1** as published. The annotation of core and accessory genes in the Figure 1 was misplaced. The corrected figure appears below.

The authors apologize for this error and state that this does not change the scientific conclusions of the article in any way. The original article has been updated.

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**FIGURE 1** Selection pressure acting on *B. pseudomallei* population. The histogram summarizes ranges of *dN/dS* calculated from predicted coding sequences from a collection of diverse *B. pseudomallei* population from northeast Thailand (Chewapreecha et al., 2019), and *B. thailandensis* genomes from the public database. *B. pseudomallei* and *B. thailandensis* have highly plastic genomes comprising of at least two chromosomes of ~7–8 Mb in size when combined. Using a pan-genome approach, all coding sequences could be categorized as "core" (present in all genomes) or "accessory" (variably present across studied genomes). Accessory genes display an elevated level of *dN/dS* which is signatures of positive selection or more relaxed purifying selection.