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Approved by:
Frontiers Editorial Office,
Frontiers Media SA, Switzerland

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RECEIVED 01 June 2023

ACCEPTED 01 June 2023

PUBLISHED 14 June 2023

CITATION

Frontiers Production Office (2023)
Erratum: Microbiological diagnostic
performance of metagenomic
next-generation sequencing compared
with conventional culture for patients
with community-acquired pneumonia.
Front. Cell. Infect. Microbiol. 13:1233180.
doi: 10.3389/fcimb.2023.1233180

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Erratum: Microbiological diagnostic performance of metagenomic next-generation sequencing compared with conventional culture for patients with community- acquired pneumonia

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KEYWORDS

metagenomic next-generation sequencing, culture, community-acquired pneumonia,
conventional microbiological test, pathogen detection

An Erratum on

[Microbiological diagnostic performance of metagenomic next-generation
sequencing compared with conventional culture for patients with
community-acquired pneumonia](#)

by Lin T, Tu X, Zhao J, Huang L, Dai X, Chen X, Xu Y, Li W, Wang Y, Lou J, Wu S and Zhang H
(2023) *Front. Cell. Infect. Microbiol.* 13:1136588. doi: 10.3389/fcimb.2023.1136588

Due to a production error, the Data Availability Statement that was originally published was incorrect.

The Data Availability Statement should be:

“The original contributions presented in the study are publicly available. This data can be found here: NCBI SRA under project ID: PRJNA917446.”

The publisher apologizes for this mistake.

The original version of this article has been updated.