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Frontiers Editorial Office, Frontiers Media SA, Switzerland

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Frontiers Production Office

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SPECIALTY SECTION

This article was submitted to Infectious Diseases: Epidemiology and Prevention,

a section of the journal Frontiers in Public Health

RECEIVED 23 February 2023 ACCEPTED 23 February 2023 PUBLISHED 03 March 2023

CITATION

Frontiers Production Office (2023) Erratum: Detection of SARS-CoV-2 infection clusters: The useful combination of spatiotemporal clustering and genomic analyses. Front. Public Health 11:1172435. doi: 10.3389/fpubh.2023.1172435

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Erratum: Detection of SARS-CoV-2 infection clusters: The useful combination of spatiotemporal clustering and genomic analyses

Frontiers Production Office*

Frontiers Media SA, Lausanne, Switzerland

KEYWORDS

SARS-CoV-2, COVID-19, epidemiology, spatiotemporal cluster, genomics, public health surveillance, superspreading, genetic similarities

An Erratum on

Detection of SARS-CoV-2 infection clusters: The useful combination of spatiotemporal clustering and genomic analyses

by Choi, Y., Ladoy, A., De Ridder, D., Jacot, D., Vuilleumier, S., Bertelli, C., Guessous, I., Pillonel, T., Joost, S., and Greub, G. (2022). Front. Public Health 10:1016169. doi: 10.3389/fpubh.2022.1016169

An omission to the funding section of the original article was made in error. The following sentence has been added: "Open access funding was provided by the University of Lausanne."

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