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© 2024 Biswas, Mallick, Maity, Bhowmik, Mitra, Saha, Roy, Chakrabarti, Paul and Chakrabarti. 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. Corrigendum: Genomic surveillance and phylodynamic analyses reveal the emergence of novel mutations and co-mutation patterns within SARS-CoV-2 variants prevalent in India

Nupur Biswas^{1‡}, Priyanka Mallick^{1,2‡}, Sujay Krishna Maity^{2,3‡}, Debaleena Bhowmik^{1,2‡}, Arpita Ghosh Mitra⁴, Soumen Saha⁴, Aviral Roy⁴, Partha Chakrabarti^{2,3}, Sandip Paul^{1,2*†} and Saikat Chakrabarti^{1,2*}

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In the published article, there was an error in affiliation 2. Instead of "Academy of Scientific and Innovative Research (AcSIR), Ghaziabad, India," it should be "Academy of Scientific and Innovative Research (AcSIR), Ghaziabad- 201002, India."

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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