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Author Correction: Genome-wide analysis of SARS-CoV-2 virus strains circulating worldwide implicates heterogeneity

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The Acknowledgments section in the original version of this Article was incomplete.

“The authors of this manuscript would like to extend their thank to all who have contributed sequences to the GISAID database (<https://www.gisaid.org/>).”

now reads:

“The authors of this manuscript would like to extend their thanks to all who have contributed sequences (Acknowledgements File) to the GISAID database (<https://www.gisaid.org/>).”

The original Article and accompanying Supplementary Information files have been corrected.



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