



Retraction Note: Origin and cross-species transmission of bat coronaviruses in China

Retraction to: *Nature Communications*
<https://doi.org/10.1038/s41467-020-17687-3>,
published online 25 August 2020

<https://doi.org/10.1038/s41467-024-55454-w>

Published online: 19 December 2024



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Following publication, errors in the sequence data included in the analyses in the article were identified. Out of 1246 sequences analysed, 41 coronavirus (CoV) sequences (12 alpha-CoVs and 29 beta-CoVs) obtained from bats sampled in the border region of northern Lao PDR (Luang Namtha province) were erroneously included in the analyses. These sequences will remain available on GenBank under the following accession codes: MN312304-MN312315, MN312608-MN312631, and MN312666-MN312670. Additionally, 27 duplicated sequences were erroneously included in the analyses.

A revised article reports the analyses with the Laotian and duplicate sequences removed. It has undergone peer review independently from the original article's review process and has been published¹.

All Authors agree with this retraction.

Reference

1. Latinne, A. et al. Origin and cross-species transmission of bat coronaviruses in China. *Nat. Commun.* <https://doi.org/10.1038/s41467-024-55384-7> (2024).

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