



Correction to: Whole-genome sequence analysis of a novel orthobunyavirus isolated in Japan in the 1980s

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In figure 1, a half of virus names are disappeared from the horizontal axis of graphs. Authors would like to replace Fig. 1 with the revised version to get this fixed. The original article has been corrected.

The original article can be found online at <https://doi.org/10.1007/s00705-022-05639-9>.

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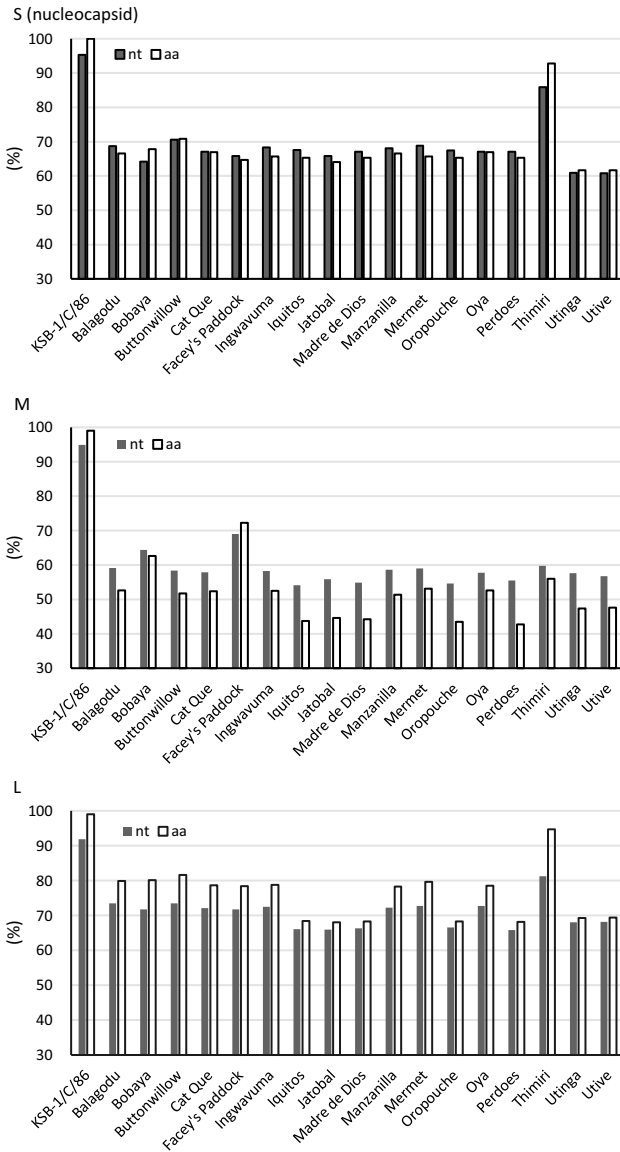


Fig. 1 Nucleotide (nt) and amino acid (aa) sequence comparisons of the coding regions of the S, M, and L RNA segments of KSB-1/C/85 and other orthobunyaviruses belonging to Simbu serogroup clade A. Histograms show percent nt (filled) and aa (open) sequence identity. The percent identity values can be found in Supplementary Table S2. The sequences used in the comparison are listed in Supplementary Table S3.

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