

Description of Additional Supplementary Files

Supplementary Data 1.

GISAID acknowledgment table listing all SARS-CoV-2 genomes used in the study.

Supplementary Data 2.

BEAST v1.10.5 XML files used to conduct continent-level and country-level Bayesian phylogeographic analyses.

Sequences are redacted out of the XMLs, per GISAID's Database Access Agreement.