


**From:** GISAID Secretariat [REDACTED]@gisaid.org   
**Subject:** data packages made available for download  
**Date:** October 1, 2025 at 2:35 AM  
**To:** Trevor Bedford [REDACTED]@bedford.io

GS

Dear Mr. Bedford,

During a recent review of data packages made available for download in EpiCoV we noticed that your package ‘nextregion’ in the ‘Genomic epidemiology’ section is for the most part no longer being downloaded by users. To be precise, during the past seven months (Mar-Sep 2025) only 79 users have made a total of 88 downloads of your package ‘nextregion.’

Because GISAID’s resources are limited, we have made the determination to no longer make the ‘nextregion’ package available for download in EpiCoV.

Consequently, the data provision GISAID has generated for you on a weekly basis for the past five years to facilitate updates of the ‘nextregion’ package in the ‘Genomic epidemiology’ section, will no longer be produced.

Of course, GISAID will gladly work with you to keep up-to-date your global subsampled phylogenetic tree of the ‘Genomic epidemiology of novel coronavirus (nCoV)’ GISAID agreed to support in January 2020 (see your email update 26-Jan-2020).

Additionally, we draw your attention to an outstanding matter. We came across an email you sent on 2. January 2023 to our legal counsel, Mr. Michael Songer (see attached). You indicated at the time that you would follow up on his questions, once you are able following a medical leave. Our counsel, you will recall, had previously provided you with ‘detailed clarification on data use under the general DAA vs. web application data agreements’ you sought. We would appreciate if you provide us with your answers to our counsel’s questions, and we apologize in advance, if we have misplaced your reply.

Best wishes

GISAID Secretariat

email 26-Jan-2020 Bedford  
global tree Jan-2020.pdf



eMail reply Bedford 2-Jan-2023  
.pdf

