Virus detail

Virus name: hCoV-19/USA/OH-ODH-SC3032044/2023
Accession ID: EPI_ISL_18138566
Type: betacoronavirus
Clade: GRA
Pango Lineage: BA.2.86 (consensus call)

AA Substitutions:

Variant: VUM GRA (BA.2.86) first detected in Denmark/Israel/USA

Passage details/history: Original

Sample information

Collection date: 2023-07-29
Location: North America / USA / Ohio / Cuyahoga
Host: Human

Additional location information:
- Gender: unknown
- Patient age: 26
- Patient status: unknown

Specimen source:

Additional host information:

Sampling strategy:

Outbreak:

Last vaccinated:

Treatment:

Sequencing technology: Illumina MiSeq

Assembly method:

Coverage:

Comment: Insertion of 12 nucleotides when compared to the reference WIV04 sequence. Gap of 68 nucleotides when compared to the reference WIV04 sequence.

Institute information

Originating lab: Pathology and Laboratory Medicine Institute, Cleveland Clinic, Ohio, USA
Address: 9500 Euclid Avenue, Cleveland, Ohio 44195, USA
Sample ID given by the originating laboratory: 2023032044

Submitting lab: Ohio Department of Health Laboratory
Address: 8995 East Main Street, Reynoldsburg, Ohio 43068, USA
Sample ID given by the submitting laboratory: 2023032044

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Submitter information

Submitter: Chill, Samantha Sevilla
Submission Date: 2023-08-26
Address:

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Database Access Agreement in respect of such data in the same manner as if they were data relating to influenza viruses.