**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/NY-PV58389/2022, complete genome**

GenBank: ON619375.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/ON619375.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/ON619375.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/ON619375.1" \l "goto2245777284_0)

LOCUS ON619375 29633 bp RNA linear VRL 30-DEC-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/USA/NY-PV58389/2022, complete genome.

ACCESSION ON619375

VERSION ON619375.1

DBLINK BioProject: [PRJNA623586](https://www.ncbi.nlm.nih.gov/bioproject/PRJNA623586)

KEYWORDS .

SOURCE Severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2)

ORGANISM [Severe acute respiratory syndrome coronavirus 2](https://www.ncbi.nlm.nih.gov/Taxonomy/Browser/wwwtax.cgi?id=2697049)

Viruses; Riboviria; Orthornavirae; Pisuviricota; Pisoniviricetes;

Nidovirales; Cornidovirineae; Coronaviridae; Orthocoronavirinae;

Betacoronavirus; Sarbecovirus.

REFERENCE 1 (bases 1 to 29633)

AUTHORS Direct Submission.

TITLE Analysis of pandemic Covid sequences from New York in 2022

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 29633)

CONSRTM NIAID Centers of Excellence for Influenza Research and Response

(CEIRR);NIAID Centers of Excellence for Influenza Research and

Surveillance (CEIRS)

TITLE Direct Submission

JOURNAL Submitted (26-MAY-2022) CEIRS Data Processing and Coordinating

Center, Center for Research on Influenza Pathogenesis (CRIP), New

York, NY 10029-6574, USA

COMMENT This submission was made by the CEIRR Influenza Data Processing and

Communication Center (iDPCC) on behalf of the Center for Research

on Influenza Pathogenesis (CRIP). This work was supported by

National Institute of Allergy and Infectious Diseases, National

Institutes of Health, grant HHSN272201400008C.

##Assembly-Data-START##

Assembly Method :: Minimap2/Pilon via Bakel Lab SARS-CoV-2

Amplicon Set v3, 1.5kb and 2kb, forward

and reverse v. 2.21-r1071p1.24

Sequencing Technology :: Illumina

##Assembly-Data-END##

FEATURES Location/Qualifiers

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