**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/MA-MGB-04990/2022, complete genome**

GenBank: ON337887.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/ON337887.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/ON337887.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/ON337887.1" \l "goto2230382159_0)

LOCUS ON337887 29810 bp RNA linear VRL 25-APR-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/USA/MA-MGB-04990/2022, complete genome.

ACCESSION ON337887

VERSION ON337887.1

DBLINK BioProject: [PRJNA759255](https://www.ncbi.nlm.nih.gov/bioproject/PRJNA759255)

BioSample: [SAMN27743079](https://www.ncbi.nlm.nih.gov/biosample/SAMN27743079)

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 29810)

AUTHORS Uddin,R., Adams,G., Shaw,B., Park,D., Loreth,C., Branda,J.,

Turbett,S., Rosenberg,E. and Lemieux,J.E.

TITLE Direct Submission

JOURNAL Submitted (25-APR-2022) Department of Infectious Disease,

Massachusetts General Hospital, 55 Fruit St, Boston, MA 02114, USA

COMMENT ##Assembly-Data-START##

Assembly Method :: Broad viral-ngs v. v2.1.33

Sequencing Technology :: Illumina

##Assembly-Data-END##

FEATURES Location/Qualifiers

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