**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/CA-CDPH-7000007788/2022, complete genome**

GenBank: OP769083.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/OP769083.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/OP769083.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/OP769083.1" \l "goto2325136267_0)

LOCUS OP769083 29811 bp RNA linear VRL 03-NOV-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/USA/CA-CDPH-7000007788/2022, complete genome.

ACCESSION OP769083

VERSION OP769083.1

DBLINK BioProject: [PRJNA750736](https://www.ncbi.nlm.nih.gov/bioproject/PRJNA750736)

BioSample: [SAMN31556089](https://www.ncbi.nlm.nih.gov/biosample/SAMN31556089)

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 29811)

AUTHORS Smith,E.

TITLE Direct Submission

JOURNAL Submitted (03-NOV-2022) IDLB VRDL/COVIDNet, California Department

of Public Health, 850 Marina Bay Blvd, Richmond, CA 94804, USA

COMMENT ##Assembly-Data-START##

Assembly Method :: iVar v. 1.3.1

Sequencing Technology :: Illumina

##Assembly-Data-END##

FEATURES Location/Qualifiers

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/product="ORF3a protein"

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/gene="ORF8"

[CDS](https://www.ncbi.nlm.nih.gov/nuccore/OP769083.1?from=27862&to=28227) 27862..28227

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SFYEDFLEYHDVRVVLDFI"

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/note="Coronavirus 3' UTR pseudoknot stem-loop 2"

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/note="Coronavirus 3' stem-loop II-like motif (s2m)"

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