**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/CA-CDPH-8000000114/2022, complete genome**

GenBank: ON618010.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/ON618010.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/ON618010.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/ON618010.1" \l "goto2245759458_0)

LOCUS ON618010 29752 bp RNA linear VRL 26-MAY-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/USA/CA-CDPH-8000000114/2022, complete genome.

ACCESSION ON618010

VERSION ON618010.1

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

BioSample: [SAMN28165494](https://www.ncbi.nlm.nih.gov/biosample/SAMN28165494)

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 29752)

AUTHORS Smith,E.

TITLE Direct Submission

JOURNAL Submitted (26-MAY-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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stem-loop 1"

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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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