**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/E8DAF5E01C0/2022, complete genome**

GenBank: OP364296.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/OP364296.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/OP364296.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/OP364296.1" \l "goto2294994662_0)

LOCUS OP364296 29811 bp RNA linear VRL 05-SEP-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/USA/E8DAF5E01C0/2022, complete genome.

ACCESSION OP364296

VERSION OP364296.1

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 Dean,D. and Noorai,R.E.

TITLE Direct Submission

JOURNAL Submitted (05-SEP-2022) Clemson University Genomics &

Bioinformatics Facility, Clemson University, 190 Collings St,

Clemson, SC 29634, USA

COMMENT ##Assembly-Data-START##

Assembly Method :: nf-core/viralrecon v. 2.4.1

Sequencing Technology :: Illumina

##Assembly-Data-END##

FEATURES Location/Qualifiers

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stem-loop 1"

[stem\_loop](https://www.ncbi.nlm.nih.gov/nuccore/OP364296.1?from=13481&to=13535) 13481..13535

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/note="Coronavirus frameshifting stimulation element

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SFYEDFLEYHDVRVVLDFI"

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/note="Coronavirus 3' UTR pseudoknot 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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