**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/H.sapiens/USA/TB-90009100/2021, complete genome**

GenBank: OP929381.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/OP929381.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/OP929381.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/OP929381.1" \l "goto2370572313_0)

LOCUS OP929381 29852 bp RNA linear VRL 01-DEC-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/H.sapiens/USA/TB-90009100/2021, complete genome.

ACCESSION OP929381

VERSION OP929381.1

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 29852)

AUTHORS Sullivan,T.J.

TITLE Sars CoV2 Surveillance Western New Hampshire, Upper Valley

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 29852)

AUTHORS Sullivan,T.J.

TITLE Direct Submission

JOURNAL Submitted (30-NOV-2022) Department of Biomedical Data Science,

Geisel School of Medicine, Dartmouth College, 1 Medical Center Dr.,

Lebanon, NH 03766, 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"

[stem\_loop](https://www.ncbi.nlm.nih.gov/nuccore/OP929381.1?from=13483&to=13537) 13483..13537

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/product="ORF3a protein"

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LLVQ"

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/product="ORF8 protein"

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SFYEDFLEYHDVRVVLDFI"

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/gene="N"

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LPQGTTLPKGFYAEGSRGGSQASSRSSSRSRNSSRNSTPGSSKRTSPARMAGNGGDAA

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/note="Coronavirus 3' UTR pseudoknot stem-loop 1"

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

[stem\_loop](https://www.ncbi.nlm.nih.gov/nuccore/OP929381.1?from=29720&to=29760) 29720..29760

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

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