**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/UT-UPHL-220107941042/2021, complete genome**

GenBank: OM295527.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/OM295527.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/OM295527.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/OM295527.1" \l "goto2180102971_0)

LOCUS OM295527 29790 bp RNA linear VRL 18-JAN-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/USA/UT-UPHL-220107941042/2021, complete genome.

ACCESSION OM295527

VERSION OM295527.1

DBLINK BioProject: [PRJNA614995](https://www.ncbi.nlm.nih.gov/bioproject/PRJNA614995)

BioSample: [SAMN25044703](https://www.ncbi.nlm.nih.gov/biosample/SAMN25044703)

Sequence Read Archive: [SRR17646820](https://www.ncbi.nlm.nih.gov/sra/SRR17646820)

KEYWORDS purposeofsampling:baselinesurveillance.

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 29790)

AUTHORS Linares-Perdomo,O.J.

TITLE Direct Submission

JOURNAL Submitted (18-JAN-2022) Utah Public Health Laboratory, Utah Public

Health Laboratory Infectious Disease submission group, 4431 S 2700

W, Salt Lake City, UT 84129, USA

COMMENT ##Assembly-Data-START##

Assembly Method :: Illumina DRAGEN COVID pipeline v. 1.0.1

Sequencing Technology :: Illumina

##Assembly-Data-END##

FEATURES Location/Qualifiers

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/gene="ORF8"

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/gene="ORF8"

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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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