**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/TX-CDC-QDX31492531/2021, complete genome**

GenBank: OL897126.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/OL897126.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/OL897126.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/OL897126.1" \l "goto2166950875_0)

LOCUS OL897126 29752 bp RNA linear VRL 17-DEC-2021

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/USA/TX-CDC-QDX31492531/2021, complete genome.

ACCESSION OL897126

VERSION OL897126.1

DBLINK BioProject: [PRJNA716985](https://www.ncbi.nlm.nih.gov/bioproject/PRJNA716985)

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 29752)

AUTHORS Howard,D., Batra,D., Cook,P.W., Caravas,J., Rambo-Martin,B.,

Sammons,S., Unoarumhi,Y., Schmerer,M., Lacek,K.A., Kendall,T.,

Caban Figueroa,V., Morrison,S., Gulvick,C., Sula,E.,

Rosenthal,S.H., Gerasimova,A., Kagan,R.M., Anderson,B., Hua,M.,

Liu,Y., Bernstein,L.E., Livingston,K.E., Perez,A., Shlyakhter,I.A.,

Rolando,R.V., Owen,R., Tanpaiboon,P., Lacbawan,F., Paden,C.R. and

MacCannell,D.

TITLE CDC Sars CoV2 Sequencing Baseline Constellation

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 29752)

AUTHORS Howard,D., Batra,D., Cook,P.W., Caravas,J., Rambo-Martin,B.,

Sammons,S., Unoarumhi,Y., Schmerer,M., Lacek,K.A., Kendall,T.,

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Rosenthal,S.H., Gerasimova,A., Kagan,R.M., Anderson,B., Hua,M.,

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TITLE Direct Submission

JOURNAL Submitted (17-DEC-2021) Respiratory Viruses Branch, Division of

Viral Diseases, Centers for Disease Control and Prevention, 1600

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FEATURES Location/Qualifiers

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

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