**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/TX-DSHS-18056/2022 ORF1ab polyprotein (ORF1ab), ORF1a polyprotein (ORF1ab), and surface glycoprotein (S) genes, complete cds; ORF3a protein (ORF3a) and membrane glycoprotein (M) genes...**

GenBank: ON254531.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/ON254531.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/ON254531.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/ON254531.1?expand-gaps=on" \l "goto2225445905_0)

LOCUS ON254531 29693 bp RNA linear VRL 18-APR-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/USA/TX-DSHS-18056/2022 ORF1ab polyprotein

(ORF1ab), ORF1a polyprotein (ORF1ab), and surface glycoprotein (S)

genes, complete cds; ORF3a protein (ORF3a) and membrane

glycoprotein (M) genes, partial cds; ORF6 protein (ORF6) and ORF7a

protein (ORF7a) genes, complete cds; ORF7b (ORF7b) and nucleocapsid

phosphoprotein (N) genes, partial cds; and ORF10 protein (ORF10)

gene, complete cds.

ACCESSION ON254531

VERSION ON254531.1

DBLINK BioProject: [PRJNA639066](https://www.ncbi.nlm.nih.gov/bioproject/PRJNA639066)

BioSample: [SAMN27590041](https://www.ncbi.nlm.nih.gov/biosample/SAMN27590041)

Sequence Read Archive: [SRR18761918](https://www.ncbi.nlm.nih.gov/sra/SRR18761918)

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 29693)

AUTHORS Pokharel,A., Bobier,K., Lu,J., Oh,B., Tuladhar,R., Pedrueza,M.,

Rahman,M., Long,K., Lee,R., Kubin,G. and Sun,Y.

TITLE Direct Submission

JOURNAL Submitted (18-APR-2022) Laboratory Services Section, Texas

Department of State Health Services-SARS-CoV-2 submission group,

1100 W. 49th St, Austin, TX 78756, USA

COMMENT ##Assembly-Data-START##

Assembly Method :: bwa/iVar v. 0.7.17-r1188/1.3.1

Sequencing Technology :: Illumina

##Assembly-Data-END##

FEATURES Location/Qualifiers

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

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