**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/CA-CDC-STM-P019/2020 ORF1ab polyprotein (ORF1ab), ORF1a polyprotein (ORF1ab), surface glycoprotein (S), ORF3a protein (ORF3a), envelope protein (E), membrane glycoprotein (M), ORF6 pr...**

GenBank: MW430967.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/MW430967.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/MW430967.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/MW430967.1" \l "goto1951969947_0)

LOCUS MW430967 29812 bp RNA linear VRL 27-JUL-2021

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/USA/CA-CDC-STM-P019/2020 ORF1ab polyprotein

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

ORF3a protein (ORF3a), envelope protein (E), membrane glycoprotein

(M), ORF6 protein (ORF6), ORF7a protein (ORF7a), ORF7b (ORF7b), and

truncated ORF8 protein (ORF8) genes, complete cds; nucleocapsid

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

gene, complete cds.

ACCESSION MW430967

VERSION MW430967.1

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 29812)

AUTHORS Cook,P.W., Batra,D., de Feo,E., Antico,J., Tran,C., Tolentino,M.,

Wickline,S., Gietzen,K., Sickler,B., Liu,J., Allen,E., Febbo,P.,

Galloway,S., Washington,N.L., White,S., Levan,G., Barret,K.S.,

Cirulli,E., Bolze,A., Ascencio,A., Rivera-Garcia,C., Cho,R.,

Nguyen,J., Wang,S., Ramirez,J., Cassens,T., Sandoval,E.,

Isaksson,M., Lee,W., Becker,D., Laurent,M., Lu,J., Paden,C.R.,

Tong,S. and MacCannell,D.

TITLE CDC Sars CoV2 Sequencing Baseline Constellation

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 29812)

AUTHORS Cook,P.W., Batra,D., de Feo,E., Antico,J., Tran,C., Tolentino,M.,

Wickline,S., Gietzen,K., Sickler,B., Liu,J., Allen,E., Febbo,P.,

Galloway,S., Washington,N.L., White,S., Levan,G., Barret,K.S.,

Cirulli,E., Bolze,A., Ascencio,A., Rivera-Garcia,C., Cho,R.,

Nguyen,J., Wang,S., Ramirez,J., Cassens,T., Sandoval,E.,

Isaksson,M., Lee,W., Becker,D., Laurent,M., Lu,J., Paden,C.R.,

Tong,S. and MacCannell,D.

TITLE Direct Submission

JOURNAL Submitted (01-JAN-2021) Respiratory Viruses Branch, Centers for

Disease Control and Prevention, 1600 Clifton Rd, Atlanta, GA 30329,

USA

COMMENT ##Assembly-Data-START##

Assembly Method :: COVIDSEQ v. DEC-2020

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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