**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/WA-CDC-UW21121761830/2021 ORF1ab polyprotein (ORF1ab), ORF1a polyprotein (ORF1ab), surface glycoprotein (S), ORF3a protein (ORF3a), envelope protein (E), membrane glycoprotein (M), OR...**

GenBank: OM131858.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/OM131858.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/OM131858.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/OM131858.1" \l "goto2172789819_0)

LOCUS OM131858 29504 bp RNA linear VRL 05-JAN-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/USA/WA-CDC-UW21121761830/2021 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),

ORF8 protein (ORF8), nucleocapsid phosphoprotein (N), and ORF10

protein (ORF10) genes, complete cds.

ACCESSION OM131858

VERSION OM131858.1

DBLINK BioProject: [PRJNA738869](https://www.ncbi.nlm.nih.gov/bioproject/PRJNA738869)

BioSample: [SAMN24620169](https://www.ncbi.nlm.nih.gov/biosample/SAMN24620169)

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 29504)

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.,

Roychoudhury,P., Xie,H., Shrestha,L., Nguyen,T.V., Bakhash,S.M.,

Lin,M., Baker,N.R., Perez,R., Ellis,S., Breit,N., Livingston,R.J.,

Huang,M.-L., Jerome,K.R., Mathias,P., Greninger,A., Paden,C.R. and

MacCannell,D.

TITLE CDC Sars CoV2 Sequencing Baseline Constellation

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 29504)

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.,

Roychoudhury,P., Xie,H., Shrestha,L., Nguyen,T.V., Bakhash,S.M.,

Lin,M., Baker,N.R., Perez,R., Ellis,S., Breit,N., Livingston,R.J.,

Huang,M.-L., Jerome,K.R., Mathias,P., Greninger,A., Paden,C.R. and

MacCannell,D.

TITLE Direct Submission

JOURNAL Submitted (05-JAN-2022) Respiratory Viruses Branch, Division of

Viral Diseases, Centers for Disease Control and Prevention, 1600

Clifton Rd, Atlanta, GA 30329, USA

COMMENT ##Assembly-Data-START##

Assembly Method :: Taylor

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Sequencing Technology :: Illumina 1x100 Novaseq

##Assembly-Data-END##

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stem-loop 1"

[stem\_loop](https://www.ncbi.nlm.nih.gov/nuccore/OM131858.1?from=13275&to=13329) 13275..13329

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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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