**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/CO-CDPHE-2100156850/2020, complete genome**

GenBank: MW422256.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/MW422256.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/MW422256.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/MW422256.1" \l "goto1951422292_0)

LOCUS MW422256 29817 bp RNA linear VRL 30-DEC-2020

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/USA/CO-CDPHE-2100156850/2020, complete genome.

ACCESSION MW422256

VERSION MW422256.1

DBLINK BioProject: [PRJNA686984](https://www.ncbi.nlm.nih.gov/bioproject/PRJNA686984)

BioSample: [SAMN17187348](https://www.ncbi.nlm.nih.gov/biosample/SAMN17187348)

Sequence Read Archive: [SRR13332719](https://www.ncbi.nlm.nih.gov/sra/SRR13332719)

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 29817)

AUTHORS Bankers,L., Hetherington-Rauth,M.C., Ir,D., Ely,S., Matzinger,S.R.,

Totten,S.E. and Travanty,E.A.

TITLE Direct Submission

JOURNAL Submitted (30-DEC-2020) Microbiology, Colorado Department of Public

Health and Environment, 8100 E Lowry Blvd, Denver, CO 80230, USA

COMMENT ##Assembly-Data-START##

Assembly Method :: Arctic MinION v. JAN-2020

Sequencing Technology :: Oxford Nanopore

##Assembly-Data-END##

FEATURES Location/Qualifiers

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stem-loop 1"

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stem-loop 2"

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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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