**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/CO-CDPHE-2103024458/2022, complete genome**

GenBank: ON123183.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/ON123183.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/ON123183.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/ON123183.1" \l "goto2214658102_0)

LOCUS ON123183 29752 bp RNA linear VRL 31-MAR-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/USA/CO-CDPHE-2103024458/2022, complete genome.

ACCESSION ON123183

VERSION ON123183.1

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

BioSample: [SAMN27101664](https://www.ncbi.nlm.nih.gov/biosample/SAMN27101664)

Sequence Read Archive: [SRR18537544](https://www.ncbi.nlm.nih.gov/sra/SRR18537544)

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 29752)

AUTHORS Bankers,L., Bogale,H.N., Hetherington-Rauth,M.C., Ir,D.,

Matzinger,S.R., Rossheim,A. and Smith,A.

TITLE Direct Submission

JOURNAL Submitted (31-MAR-2022) Department of Microbiology, CDPHE

SARS-CoV-2 submissions, 8100 E. Lowry Blvd., Denver, CO 80230, USA

COMMENT ##Assembly-Data-START##

Assembly Method :: BWA v. 0.7.17-r1188; iVar v. 1.3.1

Sequencing Technology :: Illumina

##Assembly-Data-END##

FEATURES Location/Qualifiers

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[stem\_loop](https://www.ncbi.nlm.nih.gov/nuccore/ON123183.1?from=13422&to=13476) 13422..13476

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SFYEDFLEYHDVRVVLDFI"

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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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