**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/MD-MDH-0957/2021, complete genome**

GenBank: MW644498.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/MW644498.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/MW644498.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/MW644498.1" \l "goto1992174343_0)

LOCUS MW644498 29785 bp RNA linear VRL 21-FEB-2021

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/USA/MD-MDH-0957/2021, complete genome.

ACCESSION MW644498

VERSION MW644498.1

DBLINK BioProject: [PRJNA639505](https://www.ncbi.nlm.nih.gov/bioproject/PRJNA639505)

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 29785)

AUTHORS Keller,E., Loomis,J. and Moore,T.L.

TITLE Direct Submission

JOURNAL Submitted (19-FEB-2021) CSL, Maryland Department of Health, 1770

Ashland Ave, Baltimore, MD 21205, USA

COMMENT ##Assembly-Data-START##

Assembly Method :: CLC Genomics v. 20

Sequencing Technology :: Illumina

##Assembly-Data-END##

FEATURES Location/Qualifiers

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stem-loop 1"

[stem\_loop](https://www.ncbi.nlm.nih.gov/nuccore/MW644498.1?from=13441&to=13495) 13441..13495

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/product="ORF3a protein"

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SFYEDFLEYHDVRVVLDFI"

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/gene="N"

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LPQGTTLPKGFYAEGSRGGSQASSRSSSRSRNSSRNSTPGSSKRTSPARMAGNCGDAA

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/note="Coronavirus 3' UTR pseudoknot stem-loop 2"

[stem\_loop](https://www.ncbi.nlm.nih.gov/nuccore/MW644498.1?from=29660&to=29700) 29660..29700

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

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