**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/OH-UHTL-2178/2022, complete genome**

GenBank: ON256603.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/ON256603.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/ON256603.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/ON256603.1" \l "goto2225501568_0)

LOCUS ON256603 29850 bp RNA linear VRL 18-APR-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/USA/OH-UHTL-2178/2022, complete genome.

ACCESSION ON256603

VERSION ON256603.1

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 29850)

AUTHORS Sadri,N., Alouani,D. and Song,X.

TITLE Direct Submission

JOURNAL Submitted (18-APR-2022) UHTL, University Hospitals, 7100 Euclid

Ave, Ste 190, Cleveland, OH 44103, USA

COMMENT ##Assembly-Data-START##

Assembly Method :: Torrent Suite v. 5.10

Sequencing Technology :: IonTorrent

##Assembly-Data-END##

FEATURES Location/Qualifiers

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

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