**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/OH-GD-122821-315/2021, complete genome**

GenBank: OM084605.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/OM084605.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/OM084605.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/OM084605.1" \l "goto2172004250_0)

LOCUS OM084605 29771 bp RNA linear VRL 31-DEC-2021

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/USA/OH-GD-122821-315/2021, complete genome.

ACCESSION OM084605

VERSION OM084605.1

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 29771)

AUTHORS Diagnostics,G.

TITLE Direct Submission

JOURNAL Submitted (31-DEC-2021) Laboratory, Gravity Diagnostics, LLC, 812

Russell St, Covington, KY 41011, USA

COMMENT ##Assembly-Data-START##

Assembly Method :: DRAGEN COVID Lineage v. 3.5.4

Sequencing Technology :: Illumina

##Assembly-Data-END##

FEATURES Location/Qualifiers

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

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