**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/GA-GPHL-1184/2022, complete genome**

GenBank: OM907131.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/OM907131.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/OM907131.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/OM907131.1" \l "goto2203191499_0)

LOCUS OM907131 29728 bp RNA linear VRL 03-MAR-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/USA/GA-GPHL-1184/2022, complete genome.

ACCESSION OM907131

VERSION OM907131.1

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 29728)

AUTHORS Parrott,T., Reeves,S., Edwards,J., Fields,A., Smith,T. and

Talekar,S.

TITLE Direct Submission

JOURNAL Submitted (03-MAR-2022) Bacteriology, GPHL, 1749 Clairmont Road,

Decatur, GA 30033, USA

COMMENT ##Assembly-Data-START##

Assembly Method :: Illumina Dragen v. 3.5.8

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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