**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/FL-BPHL-3478/2022, complete genome**

GenBank: OM970683.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/OM970683.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/OM970683.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/OM970683.1" \l "goto2206926674_0)

LOCUS OM970683 29786 bp RNA linear VRL 11-MAR-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/USA/FL-BPHL-3478/2022, complete genome.

ACCESSION OM970683

VERSION OM970683.1

DBLINK BioProject: [PRJNA630714](https://www.ncbi.nlm.nih.gov/bioproject/PRJNA630714)

BioSample: [SAMN26577219](https://www.ncbi.nlm.nih.gov/biosample/SAMN26577219)

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 29786)

AUTHORS Schmedes,S., Tarigopula,N., Splatt,T. and Blanton,J.

TITLE Direct Submission

JOURNAL Submitted (11-MAR-2022) BPHL, FLDOH, 1217 N Pearl St, Jacksonville,

FL 32202, USA

COMMENT ##Assembly-Data-START##

Assembly Method :: bwa v. v0.7.17; ivar v. v1.3.1

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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