**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/1014547886\_4-21-22/2022, complete genome**

GenBank: ON368355.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/ON368355.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/ON368355.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/ON368355.1" \l "goto2231875466_0)

LOCUS ON368355 29752 bp RNA linear VRL 28-APR-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/USA/1014547886\_4-21-22/2022, complete genome.

ACCESSION ON368355

VERSION ON368355.1

DBLINK BioProject: [PRJNA765540](https://www.ncbi.nlm.nih.gov/bioproject/PRJNA765540)

BioSample: [SAMN27935960](https://www.ncbi.nlm.nih.gov/biosample/SAMN27935960)

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 29752)

AUTHORS Harley,S.E., Laudadio,J., Henry,A., Morgan,C., Hendricks,S.,

Boileau,A., Williams,A., Stewart,T., Thurman,T.J., Bird,J.T.,

Byrum,S.D. and Kennedy,J.L.

TITLE SARS-CoV2 sequence detection

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 29752)

AUTHORS Harley,S.E., Laudadio,J., Henry,A., Morgan,C., Hendricks,S.,

Boileau,A., Williams,A., Stewart,T., Thurman,T.J., Bird,J.T.,

Byrum,S.D. and Kennedy,J.L.

TITLE Direct Submission

JOURNAL Submitted (28-APR-2022) Department of Pathology and Laboratory

Services, University of Arkansas for Medical Sciences, 4301 W.

Markham St., Little Rock, AR 72205, USA

COMMENT ##Assembly-Data-START##

Assembly Method :: Illumina DRAGEN COVID Lineage

Coverage :: NA

Sequencing Technology :: Illumina NextSeq 550Dx

##Assembly-Data-END##

FEATURES Location/Qualifiers

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

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