**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/USA/B017088/2021, complete genome**

GenBank: OP745925.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/OP745925.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/OP745925.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/OP745925.1" \l "goto2323451353_0)

LOCUS OP745925 29684 bp RNA linear VRL 31-OCT-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/USA/B017088/2021, complete genome.

ACCESSION OP745925

VERSION OP745925.1

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

BioSample: [SAMN31531542](https://www.ncbi.nlm.nih.gov/biosample/SAMN31531542)

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 29684)

AUTHORS Kandel,S., Bird,J.T., Byrum,S.D., Thurman,T.J., Kennedy,J.L. and

Ussery,D.W.

TITLE SARS-CoV2 sequence detection

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 29684)

AUTHORS Kandel,S., Bird,J.T., Byrum,S.D., Thurman,T.J., Kennedy,J.L. and

Ussery,D.W.

TITLE Direct Submission

JOURNAL Submitted (31-OCT-2022) Department of Biomedical Informatics,

University of Arkansas for Medical Sciences, 501 Jack Stephens

Drive, Little Rock, AR 72205, USA

COMMENT ##Assembly-Data-START##

Assembly Method :: ONTdeCIPHER

Coverage :: NA

Sequencing Technology :: Nanopore GridION

##Assembly-Data-END##

FEATURES Location/Qualifiers

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[stem\_loop](https://www.ncbi.nlm.nih.gov/nuccore/OP745925.1?from=13422&to=13476) 13422..13476

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SFYEDFLEYHDVRVVLDFI"

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[stem\_loop](https://www.ncbi.nlm.nih.gov/nuccore/OP745925.1?from=29644&to=29684) 29644..29684

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

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