**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/BHR/22970088641/2022, complete genome**

GenBank: ON103153.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/ON103153.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/ON103153.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/ON103153.1" \l "goto2213992133_0)

LOCUS ON103153 29752 bp RNA linear VRL 30-MAR-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/BHR/22970088641/2022, complete genome.

ACCESSION ON103153

VERSION ON103153.1

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 29752)

AUTHORS Marhoon,A., AlHujairi,Z., AlTooq,M., Khamees,F., AlAnsari,W.,

AlAbbas,Z. and Almoamen,G.

TITLE Direct Submission

JOURNAL Submitted (30-MAR-2022) Communicable Disease Laboratory, Public

Health Directorate, 1124, Manama 12, Bahrain

COMMENT ##Assembly-Data-START##

Assembly Method :: ABIOMIX v. 1.0

Sequencing Technology :: Illumina

##Assembly-Data-END##

FEATURES Location/Qualifiers

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stem-loop 1"

[stem\_loop](https://www.ncbi.nlm.nih.gov/nuccore/ON103153.1?from=13422&to=13476) 13422..13476

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stem-loop 2"

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[CDS](https://www.ncbi.nlm.nih.gov/nuccore/ON103153.1?from=27681&to=27812) 27681..27812

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/codon\_start=1

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SFYEDFLEYHDVRVVLDFI"

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[gene](https://www.ncbi.nlm.nih.gov/nuccore/ON103153.1?from=29474&to=29590) 29474..29590

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/note="Coronavirus 3' UTR pseudoknot stem-loop 1"

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/note="Coronavirus 3' UTR pseudoknot stem-loop 2"

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

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