**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/BGD/Laila\_872/2022 ORF1ab polyprotein (ORF1ab), ORF1a polyprotein (ORF1ab), surface glycoprotein (S), ORF3a protein (ORF3a), envelope protein (E), membrane glycoprotein (M), ORF6 protein ...**

GenBank: OM570235.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/OM570235.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/OM570235.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/OM570235.1" \l "goto2190482944_0)

LOCUS OM570235 29762 bp RNA linear VRL 08-FEB-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/BGD/Laila\_872/2022 ORF1ab polyprotein (ORF1ab),

ORF1a polyprotein (ORF1ab), surface glycoprotein (S), ORF3a protein

(ORF3a), envelope protein (E), membrane glycoprotein (M), ORF6

protein (ORF6), ORF7a protein (ORF7a), ORF7b (ORF7b), ORF8 protein

(ORF8), nucleocapsid phosphoprotein (N), and ORF10 protein (ORF10)

genes, complete cds.

ACCESSION OM570235

VERSION OM570235.1

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 29762)

AUTHORS Banu,L.A.

TITLE Whole genome sequencing of SARRS-CoV-2 at Genomic Research

Laboratory, BSMMU

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 29762)

AUTHORS Banu,L.A., Ahmed,M., Hossain,M. and Hassan,Z.

TITLE Direct Submission

JOURNAL Submitted (07-FEB-2022) Genome research Center, Anatomy, BSMMU,

Bangabandhu Sheikh Mujib Medical University, Bangabandhu Sheikh

Mujib Medical University (BSMMU) Shahbag, Dhaka, Dhaka 1000,

Bangladesh

COMMENT ##Assembly-Data-START##

Assembly Method :: DRAGEN COVID Lineage 3.5.3 v.

05.021.510.3.5.88-85-g1aac5c73 and Bio-IT

Processor Version 0x04261818

Sequencing Technology :: Illumina

##Assembly-Data-END##

FEATURES Location/Qualifiers

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[CDS](https://www.ncbi.nlm.nih.gov/nuccore/OM570235.1?from=27699&to=27830) 27699..27830

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SFYEDFLEYHDVRVVLDFI"

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

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