**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/IND/17/2021, complete genome**

GenBank: ON052769.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/ON052769.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/ON052769.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/ON052769.1" \l "goto2211128971_0)

LOCUS ON052769 29736 bp RNA linear VRL 23-MAR-2022

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/IND/17/2021, complete genome.

ACCESSION ON052769

VERSION ON052769.1

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 29736)

AUTHORS Polipalli,S.K., Kumar,S., Lomash,A., Jindal,A., Kumar,S.,

Mohammed,F., Kapoor,S., Varughese,B., Dhakad,M.S., Sharma,A.,

Saxena,S., Manchanda,V., Siddiqui,O., Bothra,M., Aasif,M.K.,

Garg,S. and Suravajhala,P.N.

TITLE Emerging Variants of SARS-CoV2

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 29736)

AUTHORS Polipalli,S.K., Kumar,S., Lomash,A., Jindal,A., Kumar,S.,

Mohammed,F., Kapoor,S., Varughese,B., Dhakad,M.S., Sharma,A.,

Saxena,S., Manchanda,V., Siddiqui,O., Bothra,M., Aasif,M.K.,

Garg,S. and Suravajhala,P.N.

TITLE Direct Submission

JOURNAL Submitted (23-MAR-2022) Genome Sequencing Lab, MAMC & Lok Nayak

Hospital, JLN Marg, New Delhi, Delhi 110002, India

COMMENT ##Assembly-Data-START##

Assembly Method :: Nanopolish v. 0.10.1

Sequencing Technology :: ONT

##Assembly-Data-END##

FEATURES Location/Qualifiers

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stem-loop 1"

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SFYEDFLEYHDVRVVLDFI"

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LPQGTTLPKGFYAEGSRGGSQASSRSSSRSRNSSRNSTPGSSKRTSPARMAGNGGDAA

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

[stem\_loop](https://www.ncbi.nlm.nih.gov/nuccore/ON052769.1?from=29686&to=29726) 29686..29726

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

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