**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/ITA/ABR-IZSGC-TE30964/2021 ORF1ab polyprotein (ORF1ab), ORF1a polyprotein (ORF1ab), surface glycoprotein (S), ORF3a protein (ORF3a), envelope protein (E), membrane glycoprotein (M), ORF6 ...**

GenBank: MW642249.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/MW642249.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/MW642249.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/MW642249.1?expand-gaps=on" \l "goto1991545385_0)

LOCUS MW642249 29862 bp RNA linear VRL 20-FEB-2021

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/ITA/ABR-IZSGC-TE30964/2021 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 MW642249

VERSION MW642249.1

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 29862)

AUTHORS Jurisic,L., Delli Compagni,E., Mangone,I. and Lorusso,A.

TITLE Genome Sequence of three SARS-CoV-2 P.1 Strains Identified from

Patients Returning from Brazil to Italy

JOURNAL Unpublished

REFERENCE 2 (bases 1 to 29862)

AUTHORS Jurisic,L., Delli Compagni,E., Mangone,I. and Lorusso,A.

TITLE Direct Submission

JOURNAL Submitted (20-FEB-2021) Virology Unit, Istituto Zooprofilattico

Sperimentale dell'Abruzzo e Molise 'G.Caporale', Campo Boario,

Teramo 64100, Italy

COMMENT ##Assembly-Data-START##

Assembly Method :: iVar v.

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Sequencing Technology :: Illumina

##Assembly-Data-END##

FEATURES Location/Qualifiers

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

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