**Severe acute respiratory syndrome coronavirus 2 isolate SARS-CoV-2/human/ITA/ABR-IZSGC-TE30968/2021, complete genome**

GenBank: MW642250.1

[FASTA](https://www.ncbi.nlm.nih.gov/nuccore/MW642250.1?report=fasta) [Graphics](https://www.ncbi.nlm.nih.gov/nuccore/MW642250.1?report=graph)

[Go to:](https://www.ncbi.nlm.nih.gov/nuccore/MW642250.1" \l "goto1991545398_0)

LOCUS MW642250 29859 bp RNA linear VRL 20-FEB-2021

DEFINITION Severe acute respiratory syndrome coronavirus 2 isolate

SARS-CoV-2/human/ITA/ABR-IZSGC-TE30968/2021, complete genome.

ACCESSION MW642250

VERSION MW642250.1

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 29859)

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JOURNAL Unpublished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 29859)

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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stem-loop 1"

[stem\_loop](https://www.ncbi.nlm.nih.gov/nuccore/MW642250.1?from=13479&to=13533) 13479..13533

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/product="ORF3a protein"

/protein\_id="[QRX39426.1](https://www.ncbi.nlm.nih.gov/protein/1991545402)"

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

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