

Virus detail

hCoV-19/Italy/CAM-TIGEM-IZSM-COLLI-9211/2021 Virus name:

Accession ID: EPI ISL 1673660 betacoronavirus Type:

Clade

Pango Lineage B.1.1.7 (version: 2021-04-14)

Spike A570D, Spike D614G, Spike D1118H, Spike H69del, Spike N501Y, Spike P681H, Spike S982A, Spike **AA Substitutions** 

> T716I, Spike V70del, Spike Y144del, N D3L, N G204R, N R203K, N S235F, NS8 Q27stop, NS8 R52I, NS8 Y73C, NSP3 A890D, NSP3 I1412T, NSP3 T183I, NSP3 T749A, NSP6 F108L, NSP6 G107S, NSP8 Q24R, NSP12 P323L,

NSP12 V742L, NSP13 K460R

Variant VUI202012/01 GRY (B.1.1.7)

Passage details/history: Original

Sample information

Collection date: 2021-04-02

Europe / Italy / Campania Location:

**Host:** Human

Additional location

information:

Male Gender: 54 Patient age: Patient status: unknown

Specimen source: Oropharyngeal swab

**Additional host information:** 

Sampling strategy:

**Outbreak:** 

Last vaccinated:

Treatment:

MGI DNBSEQ-G400 Sequencing technology:

SARS-CoV-2\_Multi-PCR\_v1.0 Assembly method:

Coverage:

**Comment:** NS8\_Q27stop results in 78.5% truncation of the protein sequence. Gap of 10 nucleotides when compared to the

reference sequence.

Institute information

Originating lab: Istituto Zooprofilattico Sperimentale del Mezzogiorno Address: Via della Salute 2 - 80055 Portici-Napoli-Italy

Sample ID given by the

originating laboratory:

**TIGEM Submitting lab:** 

Via dei Campi Flegrei 34 Pozzuoli Napoli - Italy

Sample ID given by the submitting laboratory:

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Submitter information

**Submitter:** Grimaldi, Antonio **Submission Date:** 2021-04-22

Address:

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