

Virus detail

Virus name: hCoV-19/England/2/2020
Accession ID: EPI_ISL_610245.1
Type: betacoronavirus

Clade: S

Pango Lineage: B (Pango v.4.0.6 PLEARN-v1.8)

AA Substitutions: Spike A684del, Spike N679del, Spike P681del, Spike R682del, Spike R683del, Spike R685del, Spike S680del,

Spike S686del, Spike V687I, NS8 L84S, NSP14 I150T

Variant:

Passage details/history: Vero E6

Sample information

Collection date: 2020-01-29

Location: Europe / United Kingdom / England

Host: Human

Additional location information:

Gender: unknown
Patient age: unknown
Patient status: unknown

Specimen source:

Additional host information:

Sampling strategy: Outbreak:

Last vaccinated:

Treatment:

Sequencing technology: Illumina

Assembly method: Geneious v. 10.2.3

Coverage:

Comment: Gap of 24 nucleotides found at refpos 23598. %UniqueMutations 0.08%.

Institute information

Originating lab: Virology, National Institute for Biological Standards and Control
Address: Blanche Lane, South Mimms EN6 3QG, United Kingdom

Sample ID given by the originating laboratory:

Submitting lab:Virology, National Institute for Biological Standards and ControlAddress:Blanche Lane, South Mimms EN6 3QG, United Kingdom

Sample ID given by the submitting laboratory:

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Submitter information

Submitter: Chew, Yi Hong
Submission Date: 2020-09-30

Address:

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