

Virus detail

Virus name: hCoV-19/Canada/NB-NML-30086/2021

Accession ID: EPI\_ISL\_1588284

Type: betacoronavirus

Clade GRY

**Pango Lineage** B.1.1.7 (version: 2021-04-28)

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

T716I, Spike V70del, Spike Y144del, N D3L, N G204R, N R203K, N S235F, NS8 K68stop, NS8 Q27stop, NS8 R52I, NS8 Y73C, NSP3 A890D, NSP3 I1412T, NSP3 M1441I, NSP3 T183I, NSP6 F108del, NSP6 G107del, NSP6

S106del, NSP12 P227L, NSP12 P323L

Variant VUI202012/01 GRY (B.1.1.7)

Passage details/history: Original

Sample information

Collection date: 2021-03-29

Location: North America / Canada / New Brunswick

Host: Human

Additional location information:

Gender: Male
Patient age: 70 - 79
Patient status: Symptomatic
Specimen source: Nasopharyngeal swab

Additional host information:

Sampling strategy: Targeted surveillance (outbreak)

Outbreak:
Last vaccinated:
Treatment:

Sequencing technology: Nanopore MinION

Assembly method: ARTIC Network Protocol v1

Coverage: 1732.1X

Comment: NS8\_K68stop results in 44.6% truncation of the protein sequence. NS8\_Q27stop results in 78.5% truncation of the

protein sequence. Gap of 19 nucleotides when compared to the reference sequence.

Institute information

Originating lab: NB-Hôpital Georges L. Dumont

Address: 330 Université Avenue Moncton NB Canada E1C 2Z3

Sample ID given by the originating laboratory:

Submitting lab: National Microbiology Laboratory (NML)

Address: 1015 Arlington St, Winnipeg, MB Canada R3E 3R2

Sample ID given by the submitting laboratory:

Authors: Anna Majer, Shari Tyson, Grace Seo, Philip Mabon, Elsie Grudeski, Rhiannon Huzarewich, Russell Mandes,

Anneliese Landgraff, Jennifer Tanner, Natalie Knox, Morag Graham, Gary Van Domselaar, Richard Garceau, Guillaume Desnoyers, Nathalie Bastien, Yan Li, Timothy Booth, Darian Hole, Madison Chapel, Kirsten Biggar,

CanCOGeN's metadata curation team, Public Health Agency of Canada CanCOGeN team

Submitter information

Submitter: Majer, Anna Submission Date: 2021-04-14

Address: 1015 Arlington Street R3E 3R2 Winnipeg

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