

Virus detail	
<b>Virus name:</b>	hCoV-19/Malaysia/IMR_125063/2021
<b>Accession ID:</b>	EPI_ISL_1585271
<b>Type:</b>	betacoronavirus
<b>Clade</b>	G
<b>Pango Lineage</b>	B.1.524 (version: 2021-04-01)
<b>AA Substitutions</b>	Spike A701V, Spike D614G, N S33I, N S194L, NS3 L53F, NS8 E110stop, NSP3 I483L, NSP3 P822S, NSP3 T1198I, NSP4 T28I, NSP5 T24A, NSP12 P323L, NSP13 L428F
<b>Variant</b>	
<b>Passage details/history:</b>	Original
Sample information	
<b>Collection date:</b>	2021-04-02
<b>Location:</b>	Asia / Malaysia
<b>Host:</b>	Human
<b>Additional location information:</b>	
<b>Gender:</b>	Male
<b>Patient age:</b>	40
<b>Patient status:</b>	Live
<b>Specimen source:</b>	
<b>Additional host information:</b>	
<b>Sampling strategy:</b>	
<b>Outbreak:</b>	
<b>Last vaccinated:</b>	
<b>Treatment:</b>	
<b>Sequencing technology:</b>	Illumina NextSeq 500
<b>Assembly method:</b>	
<b>Coverage:</b>	
<b>Comment:</b>	NS8_E110stop results in 9.9% truncation of the protein sequence.
Institute information	
<b>Originating lab:</b>	Labuan Hospital
<b>Address:</b>	81006 Labuan, WP Labuan
<b>Sample ID given by the originating laboratory:</b>	
<b>Submitting lab:</b>	Institute for Medical Research, Infectious Disease Research Centre, National Institutes of Health, Ministry of Health Malaysia
<b>Address:</b>	Jalan Setia Murni U13/52, Bandar Setia Alam, 40170 Selangor, Malaysia
<b>Sample ID given by the submitting laboratory:</b>	
<b>Authors:</b>	Suppiah J, Kamel K, Mohd Zawawi Z, Ramly N, Robert F, Thayan R
Submitter information	
<b>Submitter:</b>	Suppiah, Jeyanthi
<b>Submission Date:</b>	2021-04-14
<b>Address:</b>	

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