

Virus detail	
Virus name:	hCoV-19/Denmark/DCGC-50693/2021
Accession ID:	EPI_ISL_1127084
Type:	betacoronavirus
Clade	GV
Pango Lineage	B.1.177.82 (version: 2021-04-14)
AA Substitutions	Spike A222V, Spike A262S, Spike D614G, Spike P272L, M H125Y, N A220V, N T366I, NS7a S36F, NSP12 P323L, NSP14 M49I
Variant	
Passage details/history:	Original
Sample information	
Collection date:	2021-02-15
Location:	Europe / Denmark / Hovedstaden
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:	Oxford Nanopore
Assembly method:	ARTIC-ncov2019 v1.1.0
Coverage:	
Comment:	
Institute information	
Originating lab:	Department of Virus and Microbiological Special Diagnostics, Statens Serum Institut, Copenhagen, Denmark
Address:	Artillerivej 5, 2300 Copenhagen S, Denmark
Sample ID given by the originating laboratory:	
Submitting lab:	Aalborg University
Address:	Fredrik Bajers Vej 5 9100 Aalborg Denmark
Sample ID given by the submitting laboratory:	
Authors:	Danish Covid-19 Genome Consortium
Submitter information	
Submitter:	Michaelsen, Thomas Yssing
Submission Date:	2021-03-03
Address:	Fredrik Bajers Vej 5 9100 Aalborg Denmark

Important note: In the GISAID EpiFlu™ Database Access Agreement, you have accepted certain terms and conditions for viewing and using data regarding influenza viruses. To the extent the Database contains data relating to non-influenza viruses, the viewing and use of these data is subject to the same terms and conditions, and by viewing or using such data you agree to be bound by the terms of the GISAID EpiFlu™ Database Access Agreement in respect of such data in the same manner as if they were data relating to influenza viruses.