

Virus detail	
Virus name:	hCoV-19/USA/NY-NYUMC267/2020
Accession ID:	EPI_ISL_428781
Type:	betacoronavirus
Passage details/history:	Original
Sample information	
Collection date:	2020-04-06
Location:	North America / USA / New York / Brooklyn
Host:	Human
Additional location information:	
Gender:	Male
Patient age:	75
Patient status:	
Specimen source:	
Additional host information:	
Outbreak:	
Last vaccinated:	
Treatment:	
Sequencing technology:	Targeted capture, Illumina Novaseq
Assembly method:	bwa/0.7.17, mapped to NC_045512v2, bases below 20x coverage masked
Coverage:	19,620x
Comment:	
Institute information	
Originating lab:	NYU Langone Health
Address:	150 55th St, Brooklyn, NY 11220
Sample ID given by the sample provider:	
Submitting lab:	Departments of Pathology and Medicine, New York University School of Medicine
Address:	New York, NY 10016
Sample ID given by the submitting laboratory:	
Authors:	Maria Aguero-Rosenfeld, Brendan Belovarac, Margaret Black, Ludovic Boytard, John Cadley, Paolo Cotzia, John Chen, Dacia Dimartino, Xiaojun Feng, Tatyana Gindin, Emily Guzman, Adriana Heguy, Megan Hogan, Emily Huang, George Jour, Andrew Lytle, Christian Marier, Matthew T. Maurano, Mark J. Mulligan, Peter Meyn, Iman Osman, Jared Pinnell, Vanessa Raabe, Sitharam Ramaswami, Amy Rapkiewicz, Marie Samanovic-Golden, Antonio Serrano, Guomiao Shen, Matija Snuderl, Theodore Vougiouklakis, Nick Vulpescu, Gael Westby, Paul Zappile, Yutong Zhang
Submitter information	
Submitter:	Maurano, Matthew Thomas
Submission Date:	2020-04-22
Address:	NYU - Science Building Room 801, 435 East 30th Street 10016 New York

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.