

SUPPORTING INFECTIOUS DISEASE RESEARCH

Product Information Sheet for NR-56695

SARS-Related Coronavirus 2, Isolate hCoV-19/France/PAC-IHUMI-6070/2022 (Lineage AY.4 plus omicron; XD Variant)

Catalog No. NR-56695

For research use only. Not for use in humans.

Contributor:

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Manufacturer:

BEI Resources

Product Description:

<u>Virus Classification</u>: Coronaviridae, Betacoronavirus <u>Species</u>: Severe acute respiratory syndrome-related coronavirus 2

Strain/Isolate: hCoV-19/France/PAC-IHUMI-6070/2022

Original Source: Severe acute respiratory syndrome-related coronavirus 2 (SARS-CoV-2), isolate hCoV-19/France/PAC-IHUMI-6070/2022 was isolated from a human on February 9, 2022, in France.¹ This isolate has been passaged one time in Vero cells prior to deposition.

Note: Genome sequence information is provided on the Certificate of Analysis and includes an analysis of all sequence variations observed for each lot.

Comments: Under the nomenclature system introduced by GISAID (Global Initiative on Sharing All Influenza Data), hCoV-19/France/PAC-IHUMI-SARS-CoV-2, isolate 6070/2022 is assigned lineage AY.4 plus omicron (Pango v.4.0.6 PLEARN-v1.8), Delta (AY.4-like) (Scorpio) and GISAID clade GKA using Phylogenetic Assignment of Named Global Outbreak lineages (PANGO) tool. 1,2,3 The complete genome of the clinical isolate of SARS-CoV-2, hCoV-19/France/PAC-IHUMI-6070/2022 has sequenced (GISAID: EPI ISL 10528736).1,2 The complete genome of the Vero cell-passaged isolate of SARS-CoV-2, hCoV-19/France/PAC-IHUMI-6070/2022 sequenced (GISAID: EPI ISL 10640045).1,2 The following mutations are present in this Vero cell-passaged isolate: Spike A27S, Spike D614G, Spike D796Y, Spike E156G, Spike E484A, Spike F157del, Spike G142D, Spike G339D, Spike G446S, Spike G496S, Spike H655Y, Spike ins214EPE, Spike K417N, Spike L212I, Spike L981F, Spike N211del, Spike N440K, Spike N501Y, Spike N679K, Spike N764K, Spike N856K, Spike N969K, Spike P681H, Spike Q493R, Spike Q498R, Spike Q954H, Spike R158del, Spike S371L, Spike S373P, Spike S375F, Spike S477N, Spike T19R, Spike T95I, Spike T478K, Spike T547K, Spike Y505H, Membrane (M) I82T, Nucleocapsid (N) D63G, N D377Y, N G215C, N R203M, NS3 D155Y, NS3 S92L, NS7a T120I, NS7a V82A, NS7b T40I, NSP2 (Non-structural protein 2) E172D, NSP3 (Non-structural protein 3) A488S, NSP3 A1711V, NSP3 P1469S, NSP4 (Non-structural protein 4) I57V, NSP4 T492I, NSP4 V167L, NSP6 (Nonstructural protein 6) T77A, NSP7 (Non-structural protein 7) V66I, NSP12 (Non-structural protein 12) G671S, NSP12 P323F, NSP13 (Non-structural protein 13) P77L, NSP14 (Non-structural protein 14) A394V.^{1,2} It was labeled as an XD variant by the World Health Organization (WHO).^{1,4}

In December 2019, an outbreak of a respiratory illness (COVID-19) began in Wuhan, Hubei Province, China. The outbreak is associated with a seafood market and although environmental samples from the market are positive for the novel coronavirus, an association with a particular animal has not been determined.⁵ SARS-CoV-2 has been isolated from patients from several countries and the sequences of some of these isolates have been deposited with GISAID.

Material Provided:

Each vial contains approximately 0.1 mL of spin-clarified cell lysate and supernatant from *Homo sapiens* lung adenocarcinoma epithelial cells (Calu-3) infected with SARS-CoV-2, isolate hCoV-19/France/PAC-IHUMI-6070/2022.

<u>Note</u>: If homogeneity is required for your intended use, please purify prior to initiating work.

Packaging/Storage:

NR-56695 was packaged aseptically in screw-capped plastic cryovials. The product is provided frozen and should be stored at -60°C or colder immediately upon arrival. For long-term storage, the vapor phase of a liquid nitrogen freezer is recommended. Freeze-thaw cycles should be avoided.

Growth Conditions:

<u>Host</u>: *Homo sapiens* lung adenocarcinoma epithelial cells (Calu-3; ATCC[®] HTB-55™)

Growth Medium: Eagle's Minimum Essential Medium containing Earle's Balanced Salt Solution, non-essential amino acids, 2 mM L-glutamine, 1 mM sodium pyruvate and 1500 milligrams per liter of sodium bicarbonate supplemented with 2% fetal bovine serum, or equivalent

<u>Infection</u>: Cells should be 70% to 90% confluent <u>Incubation</u>: 2 to 4 days at 37°C and 5% CO₂
<u>Cytopathic Effect</u>: Cell rounding and sloughing

Citation:

Acknowledgment for publications should read "The following reagent was obtained through BEI Resources, NIAID, NIH: SARS-Related Coronavirus 2, Isolate hCoV-19/France/PAC-IHUMI-6070/2022 (Lineage AY.4 plus omicron; XD Variant), NR-56695, contributed by Dr. Bernard La Scola."

Biosafety Level: 3

Appropriate safety procedures should always be used with this material. Laboratory safety is discussed in the following publication: U.S. Department of Health and Human Services, Public Health Service, Centers for Disease Control and Prevention, and National Institutes of Health. Biosafety in Microbiological and Biomedical Laboratories. 6th ed. Washington, DC: U.S. Government Printing Office, 2020; see www.cdc.gov/biosafety/publications/bmbl5/index.htm.

Disclaimers:

You are authorized to use this product for research use only.

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Use Restrictions:

SARS-CoV-2 materials provided by BEI Resources under the EUSLA are made available for any legitimate purpose, including commercial purposes as long as they are to rapidly prevent, detect, prepare for, and respond to, the spread or transmission of the 2019 SARS-CoV-2. Any further transfer of the original material or any unmodified progeny must be done under the terms of the EUSLA, documented as described above and you must notify BEI Resources of each subsequent transfer. Any new materials made by you that are not the original material or unmodified progeny are excluded from this requirement and you are free to share and commercialize those as your materials.

References:

- 1. GISAID
- Rambaut, A., et al. "A Dynamic Nomenclature Proposal for SARS-CoV-2 Lineages to Assist Genomic Epidemiology." <u>Nat. Microbiol.</u> 5 (2020): 1403-1407. PubMed: 32669681.
- Mercatelli, D. and F. M. Giorgi. "Geographic and Genomic Distribution of SARS-CoV-2 Mutations." <u>Front. Microbiol.</u> (2020): doi.org/10.3389/fmicb.2020.01800. PubMed: 32793182.
- 4. WHO
- Gralinski, L. E. and V. D. Menachery. "Return of the Coronavirus: 2019-nCoV." <u>Viruses</u> 12 (2020): 135. PubMed: 31991541.

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