

SARS-Related Coronavirus 2, Isolate hCoV-19/USA/WI-UW-4340/2021 (Lineage B.1.621)

Catalog No. NR-55698

For research use only. Not for use in humans.

Contributor:

Dr. Peter Halfmann, Ph.D., Associate Scientist, Influenza Research Institute, and Dr. Yoshihiro Kawaoka, Ph.D., Professor of Veterinary Virology, Influenza Research Institute, University of Wisconsin-Madison, Madison, Wisconsin, USA

Manufacturer:

BEI Resources

Product Description:

Virus Classification: *Coronaviridae*, *Betacoronavirus*

Species: Severe acute respiratory syndrome-related coronavirus 2

Strain/Isolate: hCoV-19/USA/WI-UW-4340/2021

Original Source: Severe acute respiratory syndrome-related coronavirus 2 (SARS-CoV-2), isolate hCoV-19/USA/WI-UW-4340/2021 was isolated from a human nasopharyngeal swab on April 22, 2021, in Madison, Wisconsin, USA.¹

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), SARS-CoV-2, isolate hCoV-19/USA/WI-UW-4340/2021 is assigned lineage B.1.621 (Pango v.4.1.1 PLEARN-v1.12), Mu (B.1.621-like) (Scorpio) and GISAID clade GH using the Phylogenetic Assignment of Named Global Outbreak lineages (PANGO) tool.^{1,2,3} The complete genome of SARS-CoV-2, isolate hCoV-19/USA/WI-UW-4340/2021 has been sequenced (GISAID: EPI_ISL_1827552).^{1,2} The following mutations are present in the clinical isolate: Spike D614G, Spike D950N, Spike E484K, Spike N501Y, Spike N1074K, Spike P681H, Spike R346K, Spike T95I, Spike Y145N, N (Nucleocapsid) T205I, NS3 Q57H, NS3 V256I, NS7b M1I, NS8 P38S, NS8 S67F, NS8 T11K, NSP3 (Non-structural protein 3) T237A, NSP3 T720I, NSP4 (Non-structural protein 4) T492I, NSP6 (Non-structural protein 6) Q160R, NSP12 (Non-structural protein 12) P323L, NSP13 (Non-structural protein 13) P419S.^{1,2} It was designated a variant of interest, Mu GH (B.1.621+B.1.621.1), and first detected in Colombia 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/USA/WI-UW-4340/2021.

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

Packaging/Storage:

NR-55698 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:

Host: *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

Infection: Cells should be 70% to 90% confluent

Incubation: 2 to 5 days at 37°C and 5% CO₂

Cytopathic Effect: 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/USA/WI-UW-4340/2021 (Lineage B.1.621), NR-55698, contributed by Peter Halfmann and Yoshihiro Kawaoka."

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:

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

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