

Certificate of Analysis for NR-52281

SARS-Related Coronavirus 2, Isolate USA-WA1/2020

Catalog No. NR-52281

Product Description:

Severe acute respiratory syndrome-related coronavirus 2 (SARS-CoV-2), isolate USA-WA1/2020 was isolated from an oropharyngeal swab from a patient with a respiratory illness who had recently returned from travel to the affected region of China and developed clinical disease (COVID-19) in January 2020 in Washington, USA. Deposited and labeled as 2019 Novel Coronavirus (2019 nCoV) prior to the determination of the official name. NR-52281 lot 70036318 was produced by infecting *Cercopithecus aethiops* kidney cells (Vero E6; ATCC® CRL-1586™) with the deposited material in Eagle's Minimum Essential Medium (ATCC® 30-2003) supplemented with 2% fetal bovine serum (ATCC® 30-2020) for 6 days at 37°C with 5% CO₂.

Passage History:

V(3)/VE6(1) (Centers for Disease Control and Prevention/BEI Resources); V = Vero cells; VE6 = Vero E6 cells

Lot: 70036318 Manufacturing Date: 28MAY2020

TEST	SPECIFICATIONS	DECLII TO		
1531	SPECIFICATIONS	RESULTS		
Identification by Infectivity in Vero E6 Cells	Cell rounding and detachment	Cell rounding and detachment		
Next-Generation Sequencing (NGS) of Complete Genome Using Illumina [®] iSeq [™] 100 Platform (Refer to Appendix I for NGS information)	≥ 98% identity with SARS-CoV-2, isolate USA-WA1/2020 (GenBank: MN985325.1)	99.99% identity with SARS-CoV- 2, isolate USA-WA1/2020 (GenBank: MN985325.1)		
Sequencing of Species-Specific Region				
(~ 930 nucleotides) (~ 930 nucleotides)	≥ 98% identity with SARS-CoV-2, isolate USA-WA1/2020 (GenBank: MN985325.1) ≥ 98% identity with SARS-CoV-2, strain FDAARGOS_983 isolate USA-WA1/2020 (GenBank: MT246667.1)	100% identity with SARS-CoV-2, isolate USA-WA1/2020 (GenBank: MN985325.1) 100% identity with SARS-CoV-2, strain FDAARGOS_983 isolate USA-WA1/2020 (GenBank: MT246667.1)		
Titer by TCID ₅₀ Assay in Vero E6 Cells by Cytopathic Effect ¹ (5 days at 37°C and 5% CO ₂)	Report results	1.6 × 10 ⁶ TCID ₅₀ per mL		
Sterility (33-day incubation)				
Harpo's HTYE broth, 37°C and 26°C, aerobic ²	No growth	No growth		
Trypticase Soy broth, 37°C and 26°C, aerobic	No growth	No growth		
Sabouraud broth, 37°C and 26°C, aerobic	No growth	No growth		
Sheep blood agar, 37°C, aerobic	No growth	No growth		
Sheep blood agar, 37°C, anaerobic	No growth	No growth		
Thioglycollate broth, 37°C, anaerobic	No growth	No growth		
DMEM with 10% FBS, aerobic	No growth	No growth		
Mycoplasma Contamination				
Agar and broth culture (14-day incubation at 37°C)	None detected	None detected		
DNA detection by PCR of extracted Test Article nucleic acid	None detected	None detected		

¹The Tissue Culture Infectious Dose 50% (TCID₅₀) endpoint is the 50% infectious endpoint in cell culture. The TCID₅₀ is the dilution of virus that under the conditions of the assay can be expected to infect 50% of the culture vessels inoculated, just as a Lethal Dose 50% (LD₅₀) is expected to kill half of the animals exposed. A reciprocal of the dilution required to yield the TCID₅₀ provides a measure of the titer (or infectivity) of a virus preparation.
²Atlas, Ronald M. Handbook of Microbiological Media. 3rd ed. Ed. Lawrence C. Parks. Boca Raton: CRC Press, 2004, p. 798.

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21 AUG 2020

Program Manager or designee, ATCC Federal Solutions

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APPENDIX I: NGS Information for NR-52281 lot 70036318

Sequence analysis resulted in the discovery of two SNPs when compared to the reference sequence from GenBank MN985325.1. Additionally, both the reference sequence GenBank MN985325.1 and NR-52281_70036318 contained three SNPs when compared to GenBank MN908947 (SARS-CoV-2, isolate Wuhan-Hu-1, complete genome) (see Table below). Quality scores over 60 indicate it is improbable that the variant call is incorrect.

Position in NR- 52281_ 70036318 Sequence	Position in MN985325.1 Reference Sequence	Position in MN908947 Wuhan- Hu-1 Sequence	Reported MN908947 Wuhan- Hu-1 Sequence	Reported MN985325.1 Reference Sequence	Identified Alternative Base	Quality	Variant Type	Length of Variant	Frequency of Variant
8771	8782	8782	С	Т	Т	n/a	SNP	1	1.0000000
18049	18060	18060	С	Т	T	n/a	SNP	1	1.0000000
22471	22482	22482	O	С	Т	111	SNP	1	0.1718464
23596	23607	23607	G	G	T	142	SNP	1	0.1593870
28133	28144	28144	Т	С	С	n/a	SNP	1	1.0000000

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