

Certificate of Analysis for NR-54001

SARS-Related Coronavirus 2, Isolate USA-WA1/2020, Recombinant Infectious Clone (icSARS-CoV-2-WT)

Catalog No. NR-54001

Product Description:

NR-54001 is a recombinant infectious cDNA clone (icSARS-CoV-2-WT) of severe acute respiratory syndrome-related coronavirus 2 (SARS-CoV-2), isolate USA-WA1/2020. Note: NR-54001 label lists this clone as wildtype (WT); however, NR-54001 is not WT and includes a T15102A silent mutation. NR-54001 lot 70041548 was produced by infecting human lung adenocarcinoma cells (Calu-3; ATCC® HTB-55™) with the deposited material in Eagle's Minimum Essential Medium (ATCC® 30-2003™) supplemented with 2% fetal bovine serum (ATCC® 30-2020™) for 5 days at 37°C with 5% CO₂.

Passage History:

V(1)/C(1) (Prior to deposit at BEI Resources/BEI Resources); V = Vero; C = Calu-3 cells

Lot: 70041548 Manufacturing Date: 18JAN2021

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TEST	SPECIFICATIONS	RESULTS Cell rounding and detachment		
Identification by Infectivity in Calu-3 Cells	Cell rounding and detachment			
Next-Generation Sequencing (NGS) of Complete Genome Using Illumina® iSeq™ 100 Platform (Refer to Appendix I for NGS information)	ne ≥ 98% identity with synthetic construct clone icSARS-CoV-2-WT (GenBank: MT461669.1) 99.96% identity with synth construct clone icSARS-CoV-2-WT (GenBank: MT461669.1)			
Titer by TCID₅ Assay in Calu-3 Cells by Cytopathic Effect¹ (5 days at 37°C and 5% CO₂)	Report results	2.8 × 10 ⁴ TCID ₅₀ per mL		
Sterility (21-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, 37°C, 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.

/Heather Couch/

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Program Manager or designee, ATCC Federal Solutions

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APPENDIX I: NGS Information for NR-54001 lot 70041548

Sequence analysis using SBC v2.0 pipeline resulted in the discovery of eleven SNPs when compared to the reference sequence GenBank MT461669.1 (see Table I below). Additionally, both the reference sequence MT461669.1 and NR-54001 lot 70041548 contained three SNPs when compared to GenBank MN908947 (SARS-CoV-2, isolate Wuhan-Hu-1, complete genome) (see Table II below). Quality scores over 60 indicate it is improbable that the variant call is incorrect.

Table I: Variants with different nucleotides between NR-54001 lot 70041548 and reference sequence MT461669.1

Position in NR-54001_ 70041548 Sequence	Position in MN908947 Wuhan- Hu-1 Sequence	Position in MT461669 Reference Sequence	Reported MN908947 Wuhan- Hu-1 Sequence	Reported MT461669 Reference Sequence	Identified Alternative Base	Quality	Variant Type	Length of Variant	Frequency of Variant
8771	8782	8782	С	Т	С	45868	SNP	1	1.000000
14566	14577	14577	Т	Т	Α	8630	SNP	1	0.153876
16053	16064	16064	Α	Α	С	1267	SNP	1	0.050161
19723	19734	19734	Т	Т	G	5928	SNP	1	0.146205
20211	20222	20222	Α	Α	С	7076	SNP	1	0.152614
22103	22114	22114	Т	Т	С	389	SNP	1	0.058224
22290	22301	22301	Α	Α	С	3408	SNP	1	0.104343
24938	24949	24949	Α	Α	G	1701	SNP	1	0.050839
26256	26267	26267	Α	А	Т	15437	SNP	1	0.324007
28242	28253	28253	С	С	Т	6909	SNP	1	0.152501
29828	29839	29839	Α	Α	G	203	SNP	1	0.053691

Table II: Variants with different nucleotides between NR-54001 lot 70041548 and GenBank MN908947 (SARS-CoV-2, isolate Wuhan-Hu-1, complete genome)

Position in NR-54001_ 70041548 Sequence	Position in MN908947 Wuhan- Hu-1 Sequence	Position in MT461669 Reference Sequence	Reported MN908947 Wuhan- Hu-1 Sequence	Reported MT461669 Reference Sequence	Identified Alternative Base	Quality	Variant Type	Length of Variant	Frequency of Variant
15091	15102	15102	Т	Α	Α	N/A	SNP	1	1.000000
18059	18060	18060	С	T	Т	N/A	SNP	1	1.000000
28133	28144	28144	T	С	С	N/A	SNP	1	1.000000

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