

Genomic RNA from SARS-Related Coronavirus 2, Isolate USA-CA1/2020

Catalog No. NR-52504

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Product Description:

Genomic RNA was extracted from a preparation of cell lysate and supernatant from *Cercopithecus aethiops* kidney epithelial cells (Vero E6; ATCC® CRL-1586™) infected with severe acute respiratory syndrome-related coronavirus 2 (SARS-CoV-2), isolate USA-CA1/2020 (BEI Resources NR-52382 lot 70034877), using QIAamp® Viral RNA Mini Kit (Qiagen 52904). The viral genomic RNA is in a background of cellular nucleic acid and carrier RNA.

Lot: 70035255

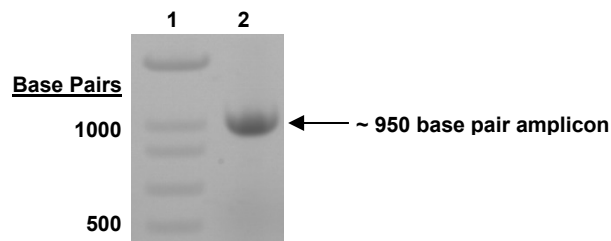
Manufacturing Date: 02JUN2020

TEST	SPECIFICATIONS	RESULTS
Sequencing of Species-Specific Region (~ 870 nucleotides)	≥ 98% identity with SARS-CoV-2, isolate USA-CA1/2020 (GenBank: MN994467.1)	100% identity with SARS-CoV-2, isolate USA-CA1/2020 (GenBank: MN994467.1)
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-CA1/2020 (GenBank: MN994467.1)	99.9% identity with SARS-CoV-2, isolate USA-CA1/2020 (GenBank: MN994467.1)
Functional Activity by RT-PCR Amplification¹	~ 950 base pair amplicon	~ 950 base pair amplicon (Figure 1)
Genome Copy Number Using BioRad QX200 Droplet Digital PCR (ddPCR™) System (Post vial; 6 replicates)	Report results	4.18 × 10 ⁷ genome equivalents/mL
Virus Inactivation (14 Days, 2 Passages) 10% of total yield inoculated on Vero E6 cells and evaluated for cytopathic effect ² Passage 1 Passage 2	No viable virus detected No viable virus detected	No viable virus detected No viable virus detected
Virus Inactivation (14 Days, 2 Passages) 100% of total yield inoculated on Vero E6 cells and evaluated for cytopathic effect ² Passage 1 Passage 2	No viable virus detected No viable virus detected	No viable virus detected No viable virus detected

¹Amplified using iTaq™ Universal SYBR Green One-step Kit (Bio-Rad 172-5151) with 5 µL of NR-52504 in a 50 µL reaction

²Supernatant was tested by qPCR to confirm absence of CPE and no evidence of replicative RNA was detected. Samples from both passages were tested by qPCR at the end of day 14 of the passages.

Figure 1: Functional Activity of NR-52504 by RT-PCR Amplification



Lane 1: Invitrogen™ TrackIt™ 1 Kb Plus DNA Ladder
Lane 2: PCR product from 1 µL of NR-52504

/Heather Couch/

Heather Couch

18 NOV 2020

Program Manager or designee, ATCC Federal Solutions

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APPENDIX I: NGS Information for NR-52504 lot 70035255

Sequence analysis resulted in the discovery of one SNP and two deletions when compared to GenBank MN994467.1 (see Table below). Quality scores over 60 indicate it is improbable that the variant call is incorrect.

Position in NR-52504_70035255 Sequence	Position in MN994467.1	Reported MN994467.1 Sequence	Identified Alternative Base	Quality	Variant Type	Length of Variant	Frequency of Variant
495	507	ATGGTCATGTT ATGGTTG	ATG	228	Indel	15	0.8965517
21772	21799	T	A	225	SNP	1	1.0000000
23555	23582	TATCAGACTCA GACTAAT	TAT	222	Indel	15	0.6134454