SUPPORTING INFECTIOUS DISEASE RESEARCH

Recombinant Vesicular Stomatitis Virus Expressing SARS-CoV-2 Spike (S) with Enhanced Green Fluorescent Protein (eGFP)

Catalog No. NR-55284

Product Description:

rVSV-SARS-CoV-2 S was generated by modification of a Vesicular Stomatitis Virus (VSV) antigenome to replace its native glycoprotein (G) with the full-length, wild-type severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) Wuhan-Hu-1 Spike (S) gene (GenBank: <u>MN908947.3</u>). The antigenome also encodes for enhanced green fluorescent protein (eGFP). NR-55284 lot 70043361 was produced by infecting *Cercopithecus aethiops* kidney cells (Vero; ATCC[®] CCL-81TM) in Dulbecco's Modified Eagle's Medium (ATCC[®] 30-2002TM) supplemented with 2% fetal bovine serum (ATCC[®] 30-2020TM) and 1% Penicillin/Streptomycin solution (ATCC[®] 30-2300TM) for 3 days at 37°C with 5% CO₂. Cell lysate and supernatant were clarified by centrifuging at 1240 × g for 15 minutes at 4°C.

Passage History:

V(7)/V(1) (Prior to deposit at BEI Resources/BEI Resources); V = Vero cells

Lot: 70043361

Manufacturing Date: 09APR2021

TEST	SPECIFICATIONS	RESULTS
Identification by Infectivity in Vero Cells	Syncytia and detachment	Syncytia and detachment
Identification by eGFP Expression in Vero Cells	Fluorescence observed	Fluorescence observed (Figure 1)
Genotypic Analysis by Next-Generation Sequencing	≥ 98% identity with Vesicular Stomatitis Indiana virus (GenBank: J02428.1) VSV G sequence absent SARS-CoV-2 S sequence confirmed	99.3% identity with Vesicular Stomatitis Indiana virus (GenBank: J02428.1) VSV G sequence absent SARS-CoV-2 S sequence confirmed ¹
Identification by eGFP Expression in Calu-3 Cells	Fluorescence observed	Fluorescence observed
Titer by TCID₅0 Assay in Vero Cells by Cytopathic Effectand eGFP Expression²(5 days at 37°C and 5% CO₂)	Report results	8.9 × 10⁵ TCID₅₀ per mL
Sterility (21-day incubation) Harpo's HTYE broth, 37°C and 26°C, aerobic ³ Trypticase Soy broth, 37°C and 26°C, aerobic Sabouraud broth, 37°C and 26°C, aerobic Sheep blood agar, 37°C, aerobic Sheep blood agar, 37°C, anaerobic Thioglycollate broth, 37°C, anaerobic DMEM with 10% FBS, 37°C, aerobic	No growth No growth No growth No growth No growth No growth No growth	No growth No growth No growth No growth No growth No growth No growth
Mycoplasma Contamination Agar and broth culture (14-day incubation at 37°C) DNA detection by PCR of extracted Test Article nucleic acid	None detected None detected	None detected None detected

¹rVSV-SARS-CoV-2 S is reported to have numerous point mutations within the SARS-CoV-2 S gene along with a 21 to 24 base pair deletion at the 3' end. The sequence obtained for NR-55284 is consistent with the depositor's sequence. For more information, please refer to Dieterle, M. E., et al. "A Replication-Competent Vesicular Stomatitis Virus for Studies of SARS-CoV-2 Spike-Mediated Cell Entry and Its Inhibition." <u>Cell Host Microbe</u> 23 (2020): 486-496. PubMed: 32738193.

²The 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. ³Atlas, Ronald M. <u>Handbook of Microbiological Media</u>. 3rd ed. Ed. Lawrence C. Parks. Boca Raton: CRC Press, 2004, p. 798. bieii resources

Certificate of Analysis for NR-55284

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Figure 1: Identification by Infectivity in Vero Cells by GFP Expression



/Heather Couch/ Heather Couch

Program Manager or designee, ATCC Federal Solutions

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