

Certificate of Analysis for NR-53257

Spike Glycoprotein (Stabilized) from SARS-Related Coronavirus 2, Wuhan-Hu-1 with C-Terminal Histidine Tag, Recombinant from HEK293 Cells

Catalog No. NR-53257

This reagent is the property of the U.S. Government.

Product Description:

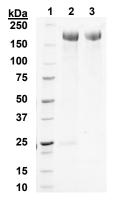
A recombinant form of the spike (S) glycoprotein from severe acute respiratory syndrome-related coronavirus 2 (SARS-CoV-2), Wuhan-Hu-1 (GenPept: YP 009724390) was produced in human embryonic kidney HEK293 cells, purified by immobilized metal affinity chromatography and dialyzed into buffer. NR-53257 contains a cleaved N-terminal mu-phosphatase signal sequence and 1198 residues (ectodomain) of the SARS-CoV-2 spike glycoprotein; the recombinant protein is stabilized by substitution at the furin S1/S2 cleavage site (RRAR to GSAS; residues 682 to 685) and stabilizing mutations (K986P and V987P, wild type numbering) and includes a C-terminal tobacco etch virus (TEV)-cleavage site, glycine-serine linker, T4 foldon trimerization domain and octa-histidine tag.

Lot: 70036058 Manufacturing Date: 19MAY2020

TEST	SPECIFICATIONS	RESULTS
Appearance	Clear solution, no particles present	Clear solution, no particles present
Purity		
SDS-PAGE analysis	Protein band of interest represents > 90% of total staining intensity	Dominant band of ~ 185 kDa represents > 95% of total staining intensity (Figure 1) ¹
SEC-HPLC	Report results	Peak observed at expected retention time; No aggregate or degradation observed (Figure 2)
Protein Concentration (A ₂₈₀)	Report results	0.77 mg per mL
Final Product		
Amount per vial	Report results	250 μg
Volume per vial	Report results	324 µL
Negative Stain Electron Microscopy	Report results	Well-folded sample with some background protein present (Figure 3)
Endotoxin	Report results	2.4 EU per mg

¹Test was performed prior to freeze/thaw. The recombinant protein migrated to a slightly larger size than was expected, likely caused by glycosylation common in recombinant spike proteins derived from coronaviruses. For more information, please see Chakraborti, S., et al. "The SARS Coronavirus S Glycoprotein Receptor Binding Domain: Fine Mapping and Functional Characterization." Virol. J. 2 (2005): 73. PubMed: 16122388.

Figure 1: SDS-PAGE Analysis



Lane 1: Bio-Rad Precision Plus Unstained

MW standard (5 μL)

Lane 2: NR-53257 (reduced; 1.5 μg) Lane 3: NR-53257 (non-reduced; 1.5 μg)

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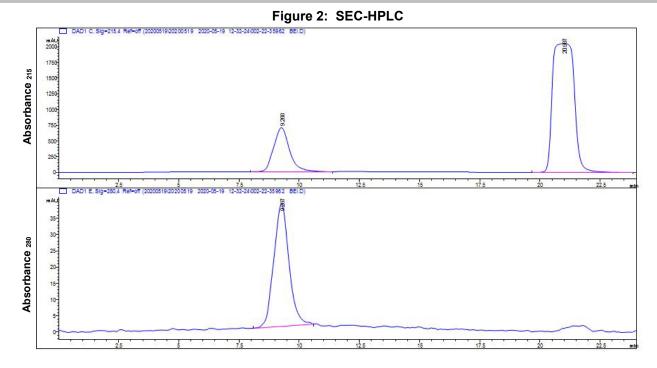
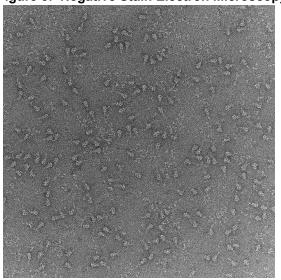


Figure 3: Negative Stain Electron Microscopy



/Heather Couch/ Heather Couch

05 JUN 2020

Program Manager or designee, ATCC Federal Solutions

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