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SUPPORTING INFECTIOUS DISEASE RESEARCH

Spike Glycoprotein Receptor Binding Domain (RBD) from SARS-Related Coronavirus 2, G476S Variant with C-Terminal Histidine Tag, Recombinant from HEK293 Cells

Catalog No. NR-55401 ACROBiosystems Catalog No. SPD-C52H4

For research use only. Not for use in humans.

Contributor and Manufacturer:

ACROBiosystems, Newark, Delaware, USA

Product Description:

A recombinant form of the spike (S) glycoprotein receptor binding domain (RBD) from severe acute respiratory syndrome-related coronavirus 2 (SARS-CoV-2), G476S variant was produced by transient transfection in human embryonic kidney HEK293 cells and purified by affinity chromatography.¹ NR-55401 lacks the signal sequence, contains 223 residues of the SARS-CoV-2 S glycoprotein RBD (amino acid residues R319 to F541) and features a C-terminal poly-histidine tag. NR-55401 is from a variant of SARS-CoV-2 which contains the G476S mutation in the S glycoprotein as compared to the SARS-CoV-2 reference sequence (GenPept: QHD43416).^{1,2} The predicted protein sequence is shown in Figure 1.1 NR-55401 has a theoretical molecular weight of 27,000 daltons. The crystal structure for the wild-type S glycoprotein from SARS-CoV-2 has been solved at 2.8 Å resolution (PDB: 6VXX) and that of SARS-CoV-2 spike RBD with ACE2 has been solved at 2.45 Å resolution (PDB: 6M0J).^{3,4}

Representative SDS-PAGE, ELISA, Surface Plasmon Resonance (SPR) and Bio-Layer Interferometry (BLI) analysis results are shown in Figures 2 to 6.¹

The S glycoprotein mediates viral binding to the host angiotensin converting enzyme 2 (ACE2). This protein forms a trimer, and when bound to a host receptor allows fusion of the viral and cellular membranes.⁵ The G476S mutation is widespread and was identified in multiple viral strains. The G476S mutation was reported to decrease the affinity of binding to ACE2.⁶

Material Provided:

Each vial contains approximately 100 μ g of purified recombinant protein lyophilized in phosphate-buffered saline, pH 7.4 and 10% trehalose.

Packaging/Storage:

NR-55401 was packaged aseptically in glass vials. The product is provided lyophilized and should be placed in a closed, dry environment with desiccants and stored at -20°C or colder immediately upon arrival. A frost-free freezer should be avoided, since changes in moisture and temperature may affect protein stability.

Functional Activity:

The biological activity of NR-55401 was measured by its binding ability in a functional ELISA (Figure 3), in which immobilized human ACE2 protein (Fc tag) (ACROBiosystems AC2-H5257) at 1 μ g per mL (100 μ L per well) can bind NR-55401; the linear range is 2 to 16 ng per mL.¹ The biological activity of NR-55401 was measured in a functional ELISA (Figure 4), in which anti-SARS-CoV-2, human IgG (AcroBiosystems SAD-S35) can bind NR-55401. The linear range is 2 to 8 ng per mL.¹ The biological activity of NR-55401 was also measured by its binding ability using biosensor analysis, in which human ACE2 protein (Fc tag) (ACROBiosystems AC2-H5257) can bind NR-55401; the affinity constant is 5.03 nM by Biacore T200 (Figure 5) and 5.49 nM by ForteBio Octet Red96e (Figure 6).¹

Reconstitution:

NR-55401 should be reconstituted with 167 μ L sterile deionized water to a stock solution of 600 μ g per mL. Add water at room temperature with occasional gentle mixing. Carrier protein [e.g., 0.1% (w/v) bovine serum albumin] must be included in the reconstitution buffer if the final protein concentration is lower than recommended or NR-55401 is aliquoted to less than 10 μ g per vial. <u>Note</u>: Avoid vigorous shaking or vortexing.

Storage of Reconstituted Protein:

Reconstituted NR-55401 should be stored at -70°C or colder immediately and used within 3 months. Avoid repeated freeze-thaw cycles.

Citation:

Acknowledgment for publications should read "The following reagent was obtained through BEI Resources, NIAID, NIH: Spike Glycoprotein Receptor Binding Domain (RBD) from SARS-Related Coronavirus 2, G476S Variant with C-Terminal Histidine Tag, Recombinant from HEK293 Cells, NR-55401."

Biosafety Level: 1

Appropriate safety procedures should always be used with this material. Laboratory safety is discussed in the following publication: U.S. Department of Health and Human Services, Public Health Service, Centers for Disease Control and Prevention, and National Institutes of Health. <u>Biosafety in Microbiological and Biomedical Laboratories</u>. 6th ed. Washington, DC: U.S. Government Printing Office, 2020; see www.cdc.gov/biosafety/publications/bmbl5/index.htm.

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References:

1. Chen, J., Personal Communication.

- Wu, F., et al. "A New Coronavirus Associated with Human Respiratory Disease in China." <u>Nature</u> 579 (2020): 265-269. PubMed: 32015508.
- Walls, A. C., et al. "Structure, Function, and Antigenicity of the SARS-CoV-2 Spike Glycoprotein." <u>Cell</u> 181 (2020): 281-292. PubMed: 32155444.
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- Chakraborty, S. "Evolutionary and Structural Analysis Elucidates Mutations on SARS-CoV-2 Spike Protein with Altered Human ACE2 Binding Affinity." <u>Biochem.</u> <u>Biophys. Res. Commun.</u> 534 (2021): 374-380. PubMed: 33602511.

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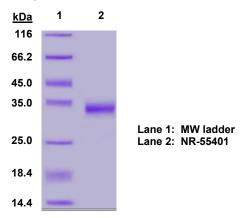
Figure 1: Predicted Protein Sequence

51 YNSASFSTFK CYGVSPTKLN DLCFTNVYAD SFVIRGDEVR QIAPGQTO
101 ADYNYKLPDD FTGCVIAWNS NNLDSKVGGN YNYLYRLFRK SNLKPFEF
151 STEIYQASST PCNGVEGFNC YFPLQSYGFQ PTNGVGYQPY RVVVLSFE
201 HAPATVCGPK KSTNLVKNKC VNFGGGSGGG SHHHHHHHHH H

RBD – **Residues 1 to 223** (represents amino acid residues 319 to 541) G476S mutation – **Residue 158**

Poly-histidine tag - Residues 232 to 241

Figure 2: Representative SDS-PAGE



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Product Information Sheet for NR-55401

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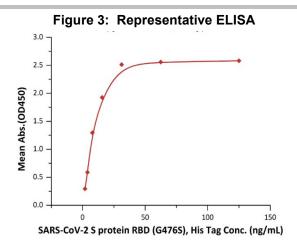


Figure 5: Representative SPR Analysis

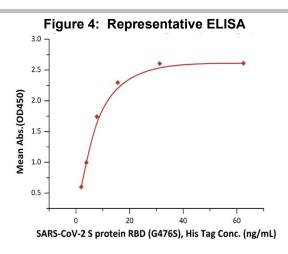
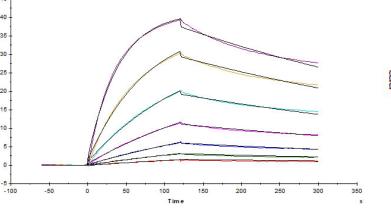
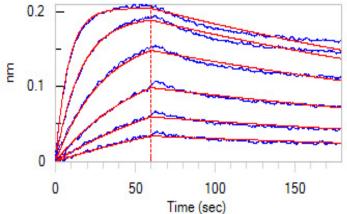


Figure 6: Representative BLI Analysis





RU 45

> 40 35 30

25 Response 20

15 10

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