

Spike Glycoprotein Receptor Binding Domain (RBD) from SARS-Related Coronavirus 2, L452R Variant with C-Terminal Histidine Tag, Recombinant from HEK293 Cells

Catalog No. NR-55403
ACROBiosystems Catalog No. SPD-C52He

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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), L452R variant was produced by transient transfection in human embryonic kidney HEK293 cells and purified by affinity chromatography.¹ NR-55403 lacks the signal sequence, contains 219 residues of the SARS-CoV-2 S glycoprotein (amino acid residues R319 to K537) and features a C-terminal poly-histidine tag. NR-55403 is from a variant of SARS-CoV-2 which contains the L452R 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-55403 has a theoretical molecular weight of 26,600 daltons.

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.³ The L452R mutation has been shown to decrease sensitivity to neutralizing antibodies, increase viral infectivity and enhance viral replication capacity.^{4,5}

Material Provided:

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

Packaging/Storage:

NR-55403 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-55403 was measured by its binding ability in a functional ELISA (Figure 3), in which immobilized NR-55403 at 1 µg per mL (100 µL per well) can bind human ACE2 protein (Fc tag) (ACROBiosystems AC2-H5257); the linear range is 0.1 to 2 ng per mL.¹

The biological activity of NR-55403 was also measured by its binding ability using biosensor analysis. Human ACE2 protein (Fc tag) (ACROBiosystems AC2-H5257) or Anti-SARS-CoV-2 RBD Neutralizing Antibody, Human IgG1 (Cat. No. SAD-S35) can bind NR-55403 with the affinity constants 20.1 nM and 17.4 nM, respectively, using ForteBio Octet Red96e (Figures 4 and 5). NR-55403 can bind ACROBiosystems SAD-S35 with an affinity constant of 19.6 nM by Biacore 8K (Figure 6).¹

Reconstitution:

NR-55403 should be reconstituted with 167 µL sterile deionized water to a stock solution of 600 µg 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-55403 is aliquoted to less than 10 µg per vial. Note: Avoid vigorous shaking or vortexing.

Storage of Reconstituted Protein:

Reconstituted NR-55403 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, L452R Variant with C-Terminal Histidine Tag, Recombinant from HEK293 Cells, NR-55403.”

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. Biosafety in Microbiological and Biomedical Laboratories. 6th ed. Washington, DC: U.S. Government Printing Office, 2020; see www.cdc.gov/biosafety/publications/bmb15/index.htm.

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References:

1. Chen, J., Personal Communication.
2. Wu, F., et al. "A New Coronavirus Associated with Human Respiratory Disease in China." *Nature* 579 (2020): 265-269. PubMed: 32015508.
3. Hulswit, R. J. G., C. A. M. de Haan and B.-J. Bosch. "Coronavirus Spike Protein and Tropism Changes." *Adv. Virus Res.* 96 (2016): 29-57. PubMed: 27712627.
4. Motozono, C., et al. "SARS-CoV-2 Spike L452R Variant Evades Cellular Immunity and Increases Infectivity." *Cell Host Microbe* 29 (2021): 1124-1136.e11. PubMed: 34171266.
5. Li, Q., et al. "The Impact of Mutations in SARS-CoV-2 Spike on Viral Infectivity and Antigenicity." *Cell* 182 (2020): 1284-1294.e9. PubMed: 32730807.

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Figure 1: Predicted Protein Sequence

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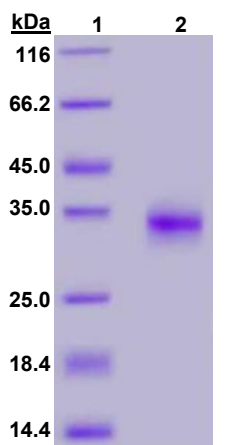
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101 ADYNYKLPDD FTGCVIAWNS NNLDSKVGGN YNYRYRLFRK SNLKPFERDI
151 STEIYQAGST PCNGVEGFNC YFPLQSYGFQ PTNGVGYQPY RVVVLSFELL
201 HAPATVCGPK KSTNLVKNKG GSGGGGSHHH HHHHHHH
    
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RBD domain – **Residues 1 to 219** (represents amino acid residues 319 to 537)

L452R mutation – **Residue 134**

Poly-histidine tag – **Residues 228 to 237**

Figure 2: Representative SDS-PAGE



Lane 1: MW ladder
Lane 2: NR-55403

Figure 3: Representative ELISA

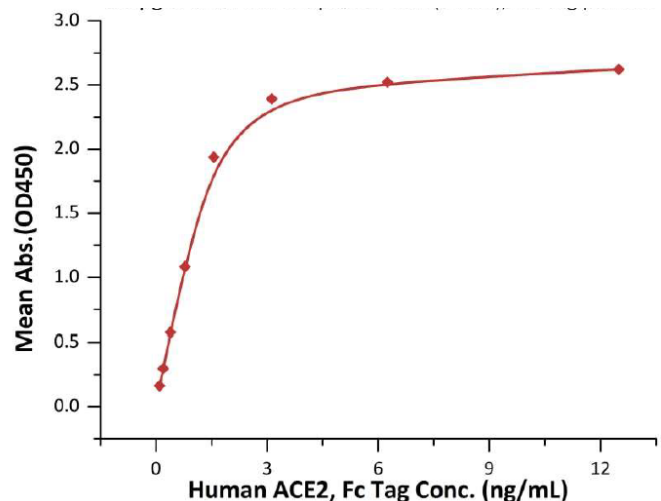


Figure 4: Representative Bioactivity-BLI

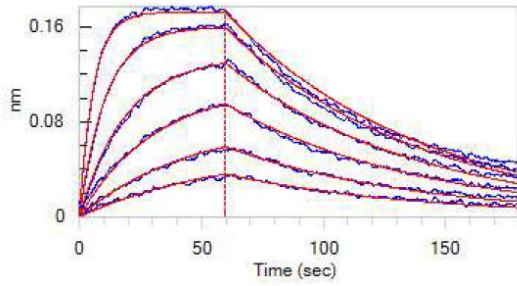


Figure 5: Representative Bioactivity-BLI

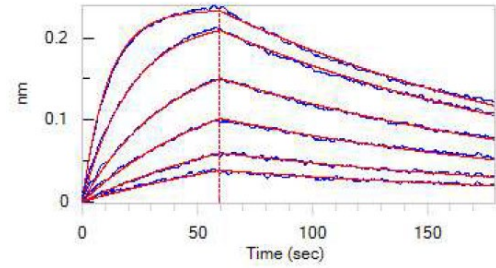


Figure 6: Representative Bioactivity-SPR

