

SUPPORTING INFECTIOUS DISEASE RESEARCH

# **Product Information Sheet for NR-55277**

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

# Catalog No. NR-55277

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

# For research use only. Not for use in humans.

### Contributor:

Florian Krammer, Ph.D. and Fatima Amanat, Department of Microbiology, Icahn School of Medicine at Mount Sinai, New York, New York, USA, supported partially under government contract HHSN272201400008C, NIAID CEIRS program

#### Manufacturer:

**BEI Resources** 

# **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), Alpha variant [also referred to as United Kingdom (UK) variant; B.1.1.7 lineage] was produced in human embryonic kidney HEK293 cells and purified by affinity chromatography. NR-55277 lacks the signal sequence and contains 223 residues of the SARS-CoV-2 S glycoprotein RBD and features a C-terminal hexa-histidine tag. 1.2.3.4 NR-55277 is an Alpha variant of SARS-CoV-2, which includes a N501Y mutation in the S glycoprotein RBD as compared to the SARS-CoV-2 reference sequence (GenPept: QHD43416). 1.5.6 The predicted protein sequence is shown in Figure 1. NR-55277 has a theoretical molecular weight of 25,970 daltons. The crystal structure for trimeric S glycoprotein from SARS-CoV-2, B.1.1.7 lineage has been solved at 3.22 Å resolution (PDB: 7LWS).6

Note: For a detailed protocol and list of related items, see <a href="https://labs.icahn.mssm.edu/krammerlab/covid-19/">https://labs.icahn.mssm.edu/krammerlab/covid-19/</a>

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.<sup>7</sup> The Alpha variant of SARS-CoV-2 includes multiple mutations that were first identified in the United Kingdom, and the most studied is N501Y.<sup>8</sup> Structural modeling and mouse studies indicate N501Y increases S glycoprotein binding to ACE2, resulting in increased SARS-CoV-2 virulence.<sup>9,10</sup>

# **Material Provided:**

Each vial contains approximately 0.1 mL of NR-55277 in phosphate buffered saline (PBS), pH 7.4. The concentration, expressed as mg per mL, is shown on the Certificate of Analysis.

Note: The long-term stability of this preparation is not known at this time. It is recommended that users confirm the activity of the product if not used within three months of receipt.

## Packaging/Storage:

NR-55277 was packaged aseptically in cryovials. The product is provided on dry ice and should be stored at -60°C or colder immediately upon arrival. Freeze-thaw cycles should be avoided.

## 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, Alpha Variant with C-Terminal Histidine Tag, Recombinant from HEK293 Cells, NR-55277."

# 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/bmbl5/index.htm.

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## References:

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# Figure 1 – Predicted Protein Sequence

- 1 RVQPTESIVR FPNITNLCPF GEVFNATRFA SVYAWNRKRI SNCVADYSVL
- 51 YNSASFSTFK CYGVSPTKLN DLCFTNVYAD SFVIRGDEVR QIAPGQTGKI
- 101 ADYNYKLPDD FTGCVIAWNS NNLDSKVGGN YNYLYRLFRK SNLKPFERDI
- 151 STEIYQAGST PCNGVEGFNC YFPLQSYGFQ PTYGVGYQPY RVVVLSFELL
- 201 HAPATVCGPK KSTNLVKNKC VNFHHHHHH

RBD – **Residues 1 to 223** (represents amino acid residues 319 to 541) N501Y Mutation – <u>Residue 183</u> Hexa-histidine tag – <u>Residues 224 to 229</u>

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Tel: 800-359-7370

Fax: 703-365-2898