

## **Synonym**

Spike,S protein RBD,Spike glycoprotein Receptor-binding domain,S glycoprotein RBD,Spike protein RBD

#### Source

SARS-CoV-2 Spike RBD, His Tag (BA.2.12.1/Omicron) (SPD-C522q) is expressed from human 293 cells (HEK293). It contains AA Arg 319 - Lys 537 (Accession # QHD43416.1 (G339D, S371F, S373P, S375F, T376A, D405N, R408S, K417N, N440K, L452Q, S477N, T478K, E484A, Q493R, Q498R, N501Y, Y505H)). The spike mutations are identified on the SARS-CoV-2 Omicron variant (Pango lineage: BA.2.12.1; GISAID clade: GRA). Predicted N-terminus: Arg 319

# **Molecular Characterization**

This protein carries a polyhistidine tag at the C-terminus.

The protein has a calculated MW of 26.7 kDa. The protein migrates as 34-37 kDa when calibrated against <u>Star Ribbon Pre-stained Protein Marker</u> under reducing (R) condition (SDS-PAGE) due to glycosylation.

#### **Endotoxin**

Less than 1.0 EU per µg by the LAL method.

## **Purity**

>95% as determined by SDS-PAGE.

### **Formulation**

Lyophilized from  $0.22~\mu m$  filtered solution in PBS, pH7.4 with trehalose as protectant.

Contact us for customized product form or formulation.

### Reconstitution

Please see Certificate of Analysis for specific instructions.

For best performance, we strongly recommend you to follow the reconstitution protocol provided in the CoA.

### Storage

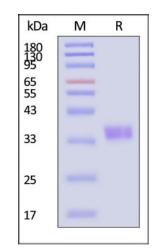
For long term storage, the product should be stored at lyophilized state at -20 $^{\circ}$ C or lower.

Please avoid repeated freeze-thaw cycles.

This product is stable after storage at:

- -20°C to -70°C for 12 months in lyophilized state;
- -70°C for 3 months under sterile conditions after reconstitution.

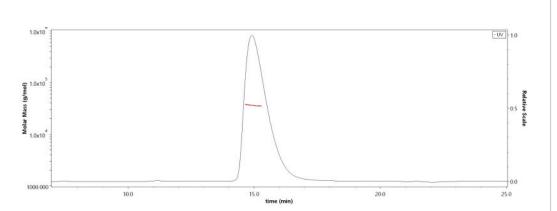
## **SDS-PAGE**



SARS-CoV-2 Spike RBD, His Tag (BA.2.12.1/Omicron) on SDS-PAGE under reducing (R) condition. The gel was stained with Coomassie Blue. The purity of the protein is greater than 95% (With <u>Star Ribbon Pre-stained Protein Marker</u>).

# **Bioactivity-ELISA**

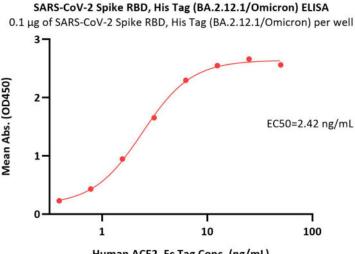
## SEC-MALS



The purity of SARS-CoV-2 Spike RBD, His Tag (BA.2.12.1/Omicron) (Cat. No. SPD-C522q) is more than 85% and the molecular weight of this protein is around 30-40 kDa verified by SEC-MALS.

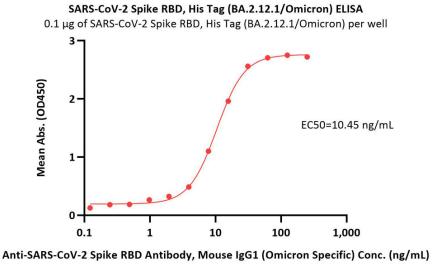
<u>Report</u>



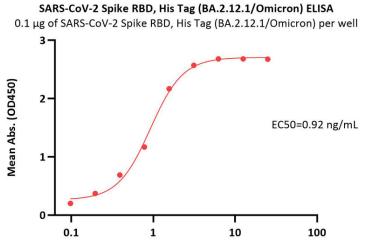


Mean Abs. (0D450) Human ACE2, Fc Tag Conc. (ng/mL)

Immobilized SARS-CoV-2 Spike RBD, His Tag (BA.2.12.1/Omicron) (Cat. No. SPD-C522q) at 1 μg/mL (100 μL/well) can bind Human ACE2, Fc Tag (Cat. No. AC2-H5257) with a linear range of 0.4-13 ng/mL (QC tested).



Immobilized SARS-CoV-2 Spike RBD, His Tag (BA.2.12.1/Omicron) (Cat. No. SPD-C522q) at 1 μg/mL (100 μL/well) can bind Anti-SARS-CoV-2 Spike RBD Antibody, Mouse IgG1 (Omicron Specific) (Cat. No. SPD-M305) with a linear range of 0.1-63 ng/mL (Routinely tested).



Anti-SARS-CoV-2 Spike RBD Broadly Neutralizing Antibody, Human IgG1 Conc. (ng/mL)

Immobilized SARS-CoV-2 Spike RBD, His Tag (BA.2.12.1/Omicron) (Cat. No. SPD-C522q) at 1 μg/mL (100 μL/well) can bind Anti-SARS-CoV-2 Spike RBD Broadly Neutralizing Antibody, Human IgG1 (Cat. No. SPD-M265) with a linear range of 0.1-6 ng/mL (Routinely tested).

## **Background**

It's been reported that Coronavirus can infect the human respiratory epithelial cells through interaction with the human ACE2 receptor. The spike protein is a large type I transmembrane protein containing two subunits, S1 and S2. S1 mainly contains a receptor binding domain (RBD), which is responsible for recognizing the cell surface receptor. S2 contains basic elements needed for the membrane fusion. The S protein plays key parts in the induction of neutralizing-antibody and T-cell responses, as well as protective immunity.

# **Clinical and Translational Updates**

