

### **Synonym**

Spike, Sprotein, Spike glycoprotein, Sglycoprotein

#### Source

SARS-CoV-2 Spike Trimer Protein, His Tag (XBB.2.3/Omicron) (SPN-C524t) is expressed from human 293 cells (HEK293). It contains AA Val 16 - Pro 1213 (Accession # QHD43416.1 (T19I, LPP24-26del, A27S, V83A, G142D, Y144del, H146Q, Q183E, V213E, D253G, G339H, R346T, L368I, S371F, S373P, S375F, T376A, D405N, R408S, K417N, N440K, V445P, G446S, N460K, S477N, T478K, E484A, F486P, F490S, Q498R, N501Y, Y505H, P521S, D614G, H655Y, N679K, P681H, N764K, D796Y, Q954H, N969K, R683A, R685A, F817P, A892P, A899P, A942P, K986P, V987P)). The spike mutations are identified on the SARS-CoV-2 Omicron variant (Pango lineage: XBB.2.3). The recombinant protein is expressed from human 293 cells (HEK293) with T4 fibritin trimerization motif and a polyhistidine tag at the C-terminus. Proline substitutions (F817P, A892P, A899P, A942P, K986P, V987P) and alanine substitutions (R683A and R685A) are introduced to stabilize the trimeric prefusion state of SARS-CoV-2 S protein and abolish the furin cleavage site, respectively.

Predicted N-terminus: Val 16

#### **Molecular Characterization**

This protein carries a polyhistidine tag at the C-terminus

The protein has a calculated MW of 137.8 kDa. The protein migrates as >150 kDa when calibrated against Star Ribbon Pre-stained Protein Marker under reducing (R) condition (SDS-PAGE).

## Endotoxin

Less than 1.0 EU per µg by the LAL method.

# **Purity**

>90% as determined by SDS-PAGE.

#### **Formulation**

Lyophilized from 0.22 µm filtered solution in PBS 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**

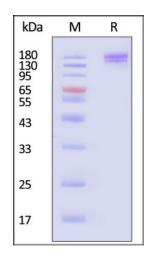
For long term storage, the product should be stored at lyophilized state at -20°C or lower.

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.

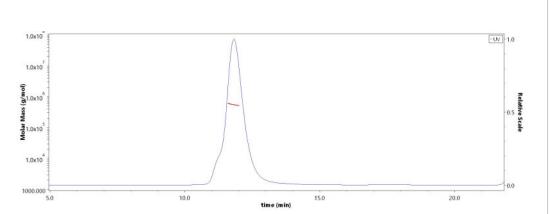
# SDS-PAGE



SARS-CoV-2 Spike Trimer Protein, His Tag (XBB.2.3/Omicron) on SDS-PAGE under reducing (R) condition. The gel was stained with Coomassie Blue. The purity of the protein is greater than 90% (With <u>Star Ribbon Pre-stained Protein Marker</u>).

**Bioactivity-ELISA** 

# SEC-MALS

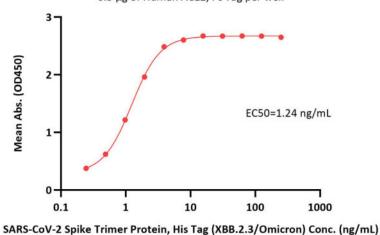


The purity of SARS-CoV-2 Spike Trimer Protein, His Tag (XBB.2.3/Omicron) (Cat. No. SPN-C524t) is more than 85% and the molecular weight of this protein is around 525-575 kDa verified by SEC-MALS.

Report



SARS-CoV-2 Spike Trimer Protein, His Tag (XBB.2.3/Omicron) ELISA 0.5 µg of Human ACE2, Fc Tag per well



Mean Abs. (OD450)

EC20=2.05 ng/mL

0.1

SARS-CoV-2 Spike Trimer Protein, His Tag (XBB.2.3/Omicron) ELISA

0.1 µg of SARS-CoV-2 Spike Trimer Protein, His Tag (XBB.2.3/Omicron) per well

Immobilized Human ACE2, Fc Tag (Cat. No. AC2-H5257) at 5  $\mu g/mL$  (100

μL/well) can bind SARS-CoV-2 Spike Trimer Protein, His Tag (XBB.2.3/Omicron) (Cat. No. SPN-C524t) with a linear range of 0.2-4 ng/mL (QC tested).

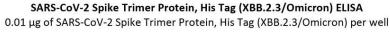
Immobilized SARS-CoV-2 Spike Trimer Protein, His Tag (XBB.2.3/Omicron) (Cat. No. SPN-C524t) at 1  $\mu$ g/mL (100  $\mu$ L/well) can bind Human ACE2, Fc Tag (Cat. No. AC2-H5257) with a linear range of 0.2-6 ng/mL (Routinely tested).

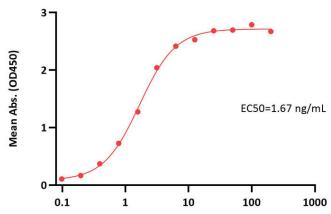
10

Human ACE2, Fc Tag Conc. (ng/mL)

100

1000





Anti-SARS-CoV-2 Spike RBD Antibody, Chimeric mAb, Human IgG1 (AM130) Conc. (ng/mL)

Immobilized SARS-CoV-2 Spike Trimer Protein, His Tag (XBB.2.3/Omicron) (Cat. No. SPN-C524t) at 0.1  $\mu$ g/mL (100  $\mu$ L/well) can bind Anti-SARS-CoV-2 Spike RBD Antibody, Chimeric mAb, Human IgG1 (AM130) (Cat. No. S1N-M13A1) with a linear range of 0.1-3 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**

Please contact us via <u>TechSupport@acrobiosystems.com</u> if you have any question on this product.