

# SARS-CoV-2 XBB.1.16.1 (Omicron) Spike S1+S2 trimer Recombinant Protein | SARS-CoV-2 recombinant protein

SARS-CoV-2 XBB.1.16.1 (Omicron) Spike S1+S2 trimer Protein (ECD, His Tag)(HPLC-verified)

|                               |                             |
|-------------------------------|-----------------------------|
| CATALOG #<br><b>AAA259236</b> | HOST<br><b>HEK293 Cells</b> |
|-------------------------------|-----------------------------|

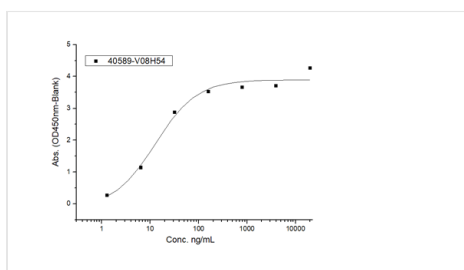
## PRODUCT SYNONYMS

SARS-CoV-2 XBB.1.16.1 (Omicron) Spike S1+S2 trimer; SARS-CoV-2 XBB.1.16.1 (Omicron) Spike S1+S2 trimer Protein (ECD; His Tag)(HPLC-verified); SARS-CoV-2 recombinant protein

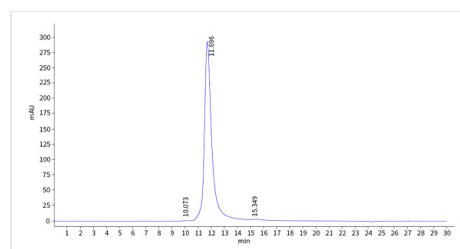
## PRODUCT SPECIFICATIONS

|                         |                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                              |
|-------------------------|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|
| Host                    | HEK293 Cells                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                 |
| Purity/Purification     | >= 90% as determined by SDS-PAGE.<br>= 95% as determined by SEC-HPLC.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                        |
| Form/Format             | Lyophilized from sterile 25mM sodium citrate, 200mM NaCl, 0.02% tween80, pH 6.0.<br>Normally 5%-8% trehalose, mannitol and 0.01% Tween80 are added as protectants before lyophilization.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                     |
| Species                 | SARS-CoV-2                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                   |
| Predicted N Terminal    | Val 16                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                       |
| Endotoxin               | <1.0 EU per ug protein as determined by the LAL method.                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                                      |
| Protein Construction    | A DNA sequence encoding the SARS-CoV-2 (XBB.1.16.1) Spike S1+S2 (YP_009724390.1, with mutations T19I, L24del, P25del, P26del, A27S, V83A, G142D, Y144del, H146Q, E180V, Q183E, V213E, G252V, G339H, R346T, L368I, S371F, S373P, S375F, T376A, D405N, R408S, K417N, N440K, V445P, G446S, N460K, S477N, T478R, E484A, F486P, F490S, Q498R, N501Y, Y505H, T547I, D614G, H655Y, N679K, P681H, N764K, D796Y, F817P, A892P, A899P, A942P, Q954H, N969K, K986P, V987P and furin cleavage site mutants) expressed with the bacteriophage T4 fibrin and a polyhistidine tag at the C-terminus. The mutations were identified in the SARS-CoV-2 variant (known as variant XBB.1.16.1). |
| Activity                | Immobilized Recombinant Human ACE2/Angiotensin-Converting Enzyme 2 Protein (Fc Tag) at 2ug/mL (100uL/well) can bind Recombinant SARS-CoV-2 XBB.1.16.1 Spike S1+S2 trimer Protein (ECD, His Tag) , the EC50 is 6-18 ng/mL.                                                                                                                                                                                                                                                                                                                                                                                                                                                    |
| Preparation and Storage | Samples are stable for up to twelve months from date of receipt at -20 degree C to -80 degree C. Store it under sterile conditions at -20 degree C to -80 degree C. It is recommended that the protein be aliquoted for optimal storage. Avoid repeated freeze-thaw cycles.                                                                                                                                                                                                                                                                                                                                                                                                  |

## IMAGES



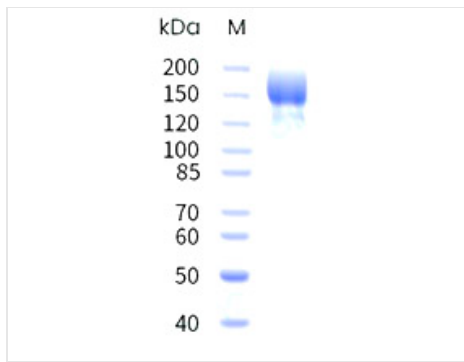
Application Data



Application Data

### Application Data

Immobilized Recombinant Human ACE2 / Angiotensin-Converting Enzyme 2 Protein (Fc Tag)  
(Cat: 10108-H05H) at 2ug/mL (100uL/well) can bind Recombinant SARS-CoV-2 XBB.1.16.1  
Spike S1+S2 trimer Protein (ECD, His Tag) (#AAA259236), the EC50 is 6-18 ng/mL.



Application Data

## ADDITIONAL INFORMATION

### Related Product Information for SARS-CoV-2 recombinant protein

The spike (S) glycoprotein of coronaviruses contains protrusions that will only bind to certain receptors on the host cell. Known receptors bind S1 are ACE2, angiotensin-converting enzyme 2; DPP4, dipeptidyl peptidase-4; APN, aminopeptidase N; CEACAM, carcinoembryonic antigen-related cell adhesion molecule 1; Sia, sialic acid; O-ac Sia, O-acetylated sialic acid. The spike is essential for both host specificity and viral infectivity. The term 'peplomer' is typically used to refer to a grouping of heterologous proteins on the virus surface that function together. The spike (S) glycoprotein of coronaviruses is known to be essential in the binding of the virus to the host cell at the advent of the infection process. It's been reported that SARS-CoV-2 (COVID-19 coronavirus, 2019-nCoV) 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. The main functions for the Spike protein are summarized as: Mediate receptor binding and membrane fusion; Defines the range of the hosts and specificity of the virus; Main component to bind with the neutralizing antibody; Key target for vaccine design; Can be transmitted between different hosts through gene recombination or mutation of the receptor binding domain (RBD), leading to a higher mortality rate.

### References

Shen S, et al. (2007) Expression, glycosylation, and modification of the spike (S) glycoprotein of SARS CoV. *Methods Mol Biol.* 379: 127-35.  
 Du L, et al. (2009) The spike protein of SARS-CoV--a target for vaccine and therapeutic development. *Nat Rev Microbiol.* 7 (3): 226-36.  
 Xiao X, et al. (2004) The SARS-CoV S glycoprotein. *Cell Mol Life Sci.* 61 (19-20): 2428-30.

## NCBI AND UNIPROT INFORMATION

### Molecular Weight

51.5 kDa

**For Research Use Only.** Not for use in diagnostic or therapeutic procedures. Small volumes of product may become entrapped in the seal of the vial during shipment and storage; if necessary, briefly centrifuge the vial on a tabletop centrifuge to dislodge any liquid in the cap.

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Online datasheet: <https://stage.aaabiotech.com/sars-cov-2-xbb-1-16-1-omicron-spike-s1-s2-trimer-recombinant-protein-hek293-cells/259236>