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## SARS-CoV-2 Spike Protein (BA.2.76 - Omicron, Trimer) (His tag)



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#### Overview

Quantity:	50 μg
Target:	SARS-CoV-2 Spike
Protein Characteristics:	BA.2.76 - Omicron, Trimer
Origin:	SARS Coronavirus-2 (SARS-CoV-2), SARS CoV-2 Omicron
Source:	HEK-293 Cells
Protein Type:	Recombinant
Purification tag / Conjugate:	This SARS-CoV-2 Spike protein is labelled with His tag.

### **Product Details**

- Toddot Details	
Purpose:	SARS-CoV-2 Spike Trimer, His Tag (BA.2.76/Omicron) (MALS verified)
Sequence:	Val 16 - Pro 1213
Characteristics:	SARS-CoV-2 Spike Trimer, His Tag (BA.2.76/Omicron) (SPN-C522i) is expressed from human
	293 cells (HEK293). It contains AA Val 16 - Pro 1213 (Accession # QHD43416.1 (T19I, LPP24-
	26del, A27S, G142D, V213G, Y248N, G339D, R346T, S371F, S373P, S375F, T376A, D405N,
	R408S, K417N, N440K, S477N, T478K, E484A, Q493R, Q498R, N501Y, Y505H, 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: BA.2.76). 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.

#### Product Details

Storage Comment:

-20°C

Product Details	
Purity:	>90 % as determined by SDS-PAGE.
Endotoxin Level:	Less than 1.0 EU per µg by the LAL method.
Grade:	MALS verified
Target Details	
Target:	SARS-CoV-2 Spike
Abstract:	SARS-CoV-2 Spike Products
Background:	Synonyms:Spike,S protein,Spike glycoprotein,S glycoprotein,Description:It's been reported that SARS-CoV-2 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.
Molecular Weight:	138.0 kDa
Application Details	
Application Notes:	This protein carries a polyhistidine tag at the C-terminus. The protein has a calculated MW of 138.0 kDa. The protein migrates as kDa under reducing (R) condition due to glycosylation.
Restrictions:	For Research Use only
Handling	
Format:	Lyophilized
Buffer:	PBS
Storage:	-20 °C