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## Datasheet for ABIN7121366 SARS-CoV-2 Spike S1 Protein (BA.2 - Omicron) (His tag)



Overview

Quantity:	100 µg
Target:	SARS-CoV-2 Spike S1
Protein Characteristics:	BA.2 - Omicron
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 S1 protein is labelled with His tag.

## Product Details

Purpose:	SARS-CoV-2 Spike S1, His Tag (BA.2/Omicron)
Sequence:	Val 16 - Arg 685
Specificity:	SARS-CoV-2 Spike S1 (BA.2/Omicron)
Characteristics:	SARS-CoV-2 Spike S1, His Tag (BA.2/Omicron) (S1N-C52Hv) is expressed from human 293
	cells (HEK293). It contains AA Val 16 - Arg 685 (Accession # QHD43416.1(T19I, LPP24-26del,
	A27S, G142D, V213G, G339D, S371F, S373P, S375F, T376A, D405N, R408S, K417N, N440K,
	S477N, T478K, E484A, Q493R, Q498R, N501Y, Y505H, D614G, H655Y, N679K, P681H). The
	spike mutations are identified on the SARS-CoV-2 Omicron variant (Pango lineage: BA.2, GISAID
	clade: GRA, Nextstrain clade: 21L).
Purity:	>90 % as determined by SDS-PAGE.
Endotoxin Level:	Less than 1.0 EU per $\mu$ g by the LAL method.

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Target Details	
Target:	SARS-CoV-2 Spike S1
Alternative Name:	SARS-CoV-2 Spike (SARS-CoV-2 Spike S1 Products)
Target Type:	Viral Protein
Background:	Synonyms: Spike,S1 protein,Spike glycoprotein Subunit1,S glycoprotein Subunit1,Spike protein S1,
	Description: 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.
Molecular Weight:	76.7 kDa
Application Details	
Application Notes:	This protein carries a polyhistidine tag at the C-terminus. The protein has a calculated MW of 76.7 kDa. The protein migrates as kDa under reducing (R) condition due to glycosylation.
Restrictions:	For Research Use only
Handling	
Format:	Lyophilized
Buffer:	PBS, pH 7.4
Storage:	-20 °C
Storage Comment:	-20°C