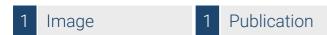


Datasheet for ABIN6952470

SARS-CoV-2 Spike S1 Protein (RBD) (His tag)





Overview

Overview						
Quantity:	100 μg					
Target:	SARS-CoV-2 Spike S1					
Protein Characteristics:	RBD					
Origin:	SARS Coronavirus-2 (SARS-CoV-2)					
Source:	HEK-293 Cells					
Purification tag / Conjugate:	This SARS-CoV-2 Spike S1 protein is labelled with His tag.					
Product Details						
Characteristics:	Recombinant SARS-CoV-2, S1 Subunit Protein (RBD) with C-terminal His tag, derived from the					
	transfected human HEK293 cells. AA 319-541,RBD					
Purification:	His-tag affinity purification by immobilized metal ion affinity chromatography (IMAC)					
Purity:	> 95 %					
Target Details						
Target:	SARS-CoV-2 Spike S1					
Abstract:	SARS-CoV-2 Spike S1 Products					
Target Type:	Viral Protein					
Gene ID:	43740568					
UniProt:	P0DTC2					

Application Details

Application Notes:	Recombinant protein product has a calculated molecular mass of 25 kDa. Due to the abur				
	glycosylation, it migrates as approximately 30 kDa protein bands in SDS-PAGE under DTT, beta-				
	mercaptoethanol reducing conditions. See deglycosylation analysis image below.				
Restrictions:	For Research Use only				

Handling

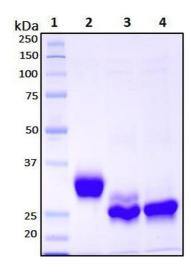
Format:	Liquid
Buffer:	Supplied as a 0.2 um filtered solution in PBS (pH 7.4)
Storage:	4 °C,-20 °C,-80 °C
Storage Comment:	Upon arrival, the protein may be stored for 2 weeks at 4 °C. For long term storage, it is recommended to store at -20 °C or -80 °C in appropriate aliquots. Avoid repeated freeze-thaw cycles.

Publications

Product cited in:

Bell, Powell, Rodriguez, Cochran, Kim: "Neutralizing antibodies targeting the SARS-CoV-2 receptor binding domain isolated from a naïve human antibody library." in: **Protein science : a publication of the Protein Society**, Vol. 30, Issue 4, pp. 716-727, (2021) (PubMed).

Images



SDS-PAGE

Image 1. Deglycosylation of purified recombinant proteins. Purified proteins were untreated (Lane 2) or treated with Protein Deglycosylation Kit under native (Lane 3) or reducing (Lane 4) conditions. Deglycosylation treatment resulted in a mobility shift of the protein to produce one major band at the expected size, thus indicating that the untreated recombinant protein (Lane 2) was glycosylated. Lane 1: Protein standard ladder (kDa) Lane 2: Untreated protein under reducing conditions Lane 3: Treated protein with deglycosylation enzymes under native conditions Lane 4: Treated protein with deglycosylation enzymes under

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reducing conditions.