Proteins

Inhibitors



Product Data Sheet

SARS-CoV-2 S glycoprotein (HEK293, His, Flag)

Cat. No.: HY-P72035

Synonyms: E2 Peplomer protein

Species: Virus Source: HEK293

Accession: P0DTC2 (V16-R685)

Gene ID: 43740568

Molecular Weight: Approximately 120 kDa

PROPERTIES

AA Sequence				
		AYTNSFTRG	VYYPDKVFRS	SVLHSTQDLF
	LPFFSNVTWF H	AIHVSGTNG	TKRFDNPVLP	FNDGVYFAST
	EKSNIIRGWI F	GTTLDSKTQ	SLLIVNNATN	VVIKVCEFQF
	C N D P F L G V Y Y H	KNNKSWMES	EFRVYSSANN	CTFEYVSQPF
	L M D L E G K Q G N F	KNLREFVFK	NIDGYFKIYS	KHTPINLVRD
	L P Q G F S A L E P L	VDLPIGINI	TRFQTLLALH	RSYLTPGDSS
	SGWTAGAAAY	VGYLQPRTF	LLKYNENGTI	TDAVDCALDP
	L S E T K C T L K S F	TVEKGIYQT	SNFRVQPTES	IVRFPNITNL
	C P F G E V F N A T R	FASVYAWNR	KRISNCVADY	SVLYNSASFS
	T F K C Y G V S P T K	LNDLCFTNV	YADSFVIRGD	EVRQIAPGQT
	G K I A D Y N Y K L P	DDFTGCVIA	WNSNNLDSKV	GGNYNYLYRL
	FRKSNLKPFE R	DISTEIYQA	GSTPCNGVEG	FNCYFPLQSY
	G F Q P T N G V G Y Q	PYRVVVLSF	ELLHAPATVC	GPKKSTNLVK
	NKCVNFNFNG L	TGTGVLTES	NKKFLPFQQF	GRDIADTTDA
	VRDPQTLEIL D	ITPCSFGGV	SVITPGTNTS	NQVAVLYQDV
	N C T E V P V A I H A	DQLTPTWRV	YSTGSNVFQT	RAGCLIGAEH
	V N N S Y E C D I P I	GAGICASYQ	TQTNSPRRAR	
Biological Activity	ivity Measured by its binding ability in a functional ELISA. Immobilized SARS-CoV-2-S at 2 μ g/mL can bind human ACE2, the EC ₅₀ of SARS-CoV-2-S protein is 56.64 - 103.6 ng/mL.			
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Appearance	Lyophilized powder.			
Formulation	Lyophilized from a 0.2 μm solution of PBS, 6% Trehalose, pH 7.4.			
Endotoxin Level	<1 EU/μg, determined by LAL method.			
Reconsititution	It is not recommended to reconstitute to a concentration less than 100 μ g/mL in ddH ₂ O.			
	20 Pg/ 2			
Storage & Stability	Stored at -20°C for 2 years. After reconstitution, it is stable at 4°C for 1 week or -20°C for longer (with carrier protein). It is recommended to freeze aliquots at -20°C or -80°C for extended storage.			
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Shipping

Room temperature in continental US;may vary elsewhere.

DESCRIPTION

Background

The SARS-CoV-2 S glycoprotein plays a crucial role in infection by attaching the virion to the host cell membrane through interaction with the primary receptor, host ACE2. Upon cleavage of S2/S2', binding to the ACE2 receptor initiates either direct fusion at the cell membrane or internalization of the virus via endocytosis, leading to fusion of the virion membrane with the host endosomal membrane. Additionally, the glycoprotein may utilize NRP1/NRP2 and integrin as alternative entry receptors, possibly explaining the virus's tropism in human olfactory epithelial cells. The stalk domain of S exhibits three hinges, providing unexpected orientational freedom to the head. Acting as a class I viral fusion protein, the glycoprotein undergoes an extensive and irreversible conformational change during virus entry, triggered by host TMPRSS2 or CSTL, leading to fusion of the viral envelope with the cellular cytoplasmic membrane and release of viral genomic RNA into the host cell cytoplasm. The glycoprotein exhibits at least three conformational states: pre-fusion native, pre-hairpin intermediate, and post-fusion hairpin, with the coiled coil regions adopting a trimer-of-hairpins structure during fusion, facilitating the apposition and subsequent fusion of viral and target cell membranes.

Caution: Product has not been fully validated for medical applications. For research use only.

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