

SARS-CoV-2 S protein RBD-SD1 (HEK293, mFc)

Cat. No.:	HY-P7432
Synonyms:	2019-nCoV RBD Protein; 2019-nCoV Spike RBD Protein; S protein RBD; 2019-nCoV S protein RBD
Species:	Virus
Source:	HEK293
Accession:	QHD43416.1 (R319-P589)
Gene ID:	43740568
Molecular Weight:	60-80 kDa

PROPERTIES

AA Sequence	R V Q P T E S I V R F P N I T N L C P F G E V F N A T R F A S V Y A W N R K R I S N C V A D Y S V L Y N S A S F S T F K C Y G V S P T K L N D L C F T N V Y A D S F V I R G D E V R Q I A P G Q T G K I A D Y N Y K L P D D F T G C V I A W N S N N L D S K V G G N Y N Y L Y R L F R K S N L K P F E R D I S T E I Y Q A G S T P C N G V E G F N C Y F P L Q S Y G F Q P T N G V G Y Q P Y R V V V L S F E L L H A P A T V C G P K K S T N L V K N K C V N F N F N G L T G T G V L T E S N K K F L P F Q Q F G R D I A D T T D A V R D P Q T L E I L D I T P
Appearance	Solution.
Formulation	Supplied as a 0.2 µm filter solution of PBS, pH 7.4.
Endotoxin Level	<1 EU/µg, determined by LAL method.
Reconstitution	N/A
Storage & Stability	Stored at -80°C for 1 year. It is stable at -20°C for 3 months after opening. It is recommended to freeze aliquots at -80°C for extended storage. Avoid repeated freeze-thaw cycles.
Shipping	Shipping with dry ice.

DESCRIPTION

Background	The transmembrane CoV S-protein spike trimer is composed of interwoven protomers that include an N-terminal receptor-binding S1 subunit and a C-terminal S2 subunit that contains the fusion elements. The S1 subunit is subdivided into the N-terminal domain (NTD) followed by the receptor-binding domain (RBD) and two structurally conserved subdomains (SD1 and SD2).
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REFERENCES

- [1]. Renhong Yan, et al. Structural basis for the recognition of SARS-CoV-2 by full-length human ACE2. Science. 2020 Mar 27; 367(6485): 1444–1448.
- [2]. Rory Henderson, et al. Controlling the SARS-CoV-2 Spike Glycoprotein Conformation. bioRxiv. 2020 May 18;2020.05.18.102087.
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Caution: Product has not been fully validated for medical applications. For research use only.

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