

# Spike protein RBD

Catalog # PVGS1571

## Product Information

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<b>Primary Accession Species</b>	<a href="#">P0DTC2</a> SARS-CoV-2
<b>Sequence</b>	Arg319-Ser591
<b>Purity</b>	> 95% as analyzed by SDS-PAGE
<b>Endotoxin Level</b>	
<b>Biological Activity</b>	SARS-CoV-2 Spike protein (RBD, His Tag, CHO-expressed) can bind with Human ACE2 (Cat. No.: Z03484) in functional ELISA assay.
<b>Expression System</b>	CHO
<b>Theoretical Molecular Weight</b>	30 kDa
<b>Formulation</b>	Supplied as a solution in PBS pH 7.4
<b>Storage &amp; Stability</b>	Upon receiving, this product remains stable for up to 6 months at -20°C or below. Avoid repeated freeze-thaw cycles.

## Additional Information

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<b>Gene ID</b>	43740568
<b>Other Names</b>	Spike glycoprotein {ECO:0000255 HAMAP-Rule:MF_04099}, S glycoprotein {ECO:0000255 HAMAP-Rule:MF_04099}, E2 {ECO:0000255 HAMAP-Rule:MF_04099}, Peplomer protein {ECO:0000255 HAMAP-Rule:MF_04099}, Spike protein S1 {ECO:0000255 HAMAP-Rule:MF_04099}, Spike protein S2 {ECO:0000255 HAMAP-Rule:MF_04099}, Spike protein S2' {ECO:0000255 HAMAP-Rule:MF_04099}, S {ECO:0000255 HAMAP-Rule:MF_04099}
<b>Target Background</b>	SARS-CoV-2 (Severe Acute Respiratory Syndrome Coronavirus 2) also known as 2019-nCoV (2019 Novel Coronavirus) is a virus that causes illnesses ranging from the common cold to severe diseases. SARS-CoV-2 Spike Protein is composed of S1 domain and S2 domain. S1 contains a receptor-binding domain (RBD) that can specifically bind to angiotensin-converting enzyme 2 (ACE2), the receptor on target cells. It is believed that SARS-CoV-2 Spike Protein (RBD) has potential value for the diagnosis of the virus.

## Protein Information

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<b>Name</b>	S {ECO:0000255 HAMAP-Rule:MF_04099}
<b>Function</b>	<p>[Spike protein S1]: Attaches the virion to the cell membrane by interacting with host receptor, initiating the infection. The major receptor is host ACE2 (PubMed:<a href="#">32142651</a>, PubMed:<a href="#">32155444</a>, PubMed:<a href="#">33607086</a>). When S2/S2' has been cleaved, binding to the receptor triggers direct fusion at the cell membrane (PubMed:<a href="#">34561887</a>). When S2/S2' has not been cleaved, binding to the receptor results in internalization of the virus by endocytosis using host TFRC and GRM2 and leading to fusion of the virion membrane with the host endosomal membrane (PubMed:<a href="#">32075877</a>, PubMed:<a href="#">32221306</a>, PubMed:<a href="#">34903715</a>, PubMed:<a href="#">36779763</a>). Alternatively, may use NRP1/NRP2 (PubMed:<a href="#">33082294</a>, PubMed:<a href="#">33082293</a>) and integrin as entry receptors (PubMed:<a href="#">35150743</a>). The use of NRP1/NRP2 receptors may explain the tropism of the virus in human olfactory epithelial cells, which express these molecules at high levels but ACE2 at low levels (PubMed:<a href="#">33082293</a>). The stalk domain of S contains three hinges, giving the head unexpected orientational freedom (PubMed:<a href="#">32817270</a>).</p>
<b>Cellular Location</b>	<p>Virion membrane {ECO:0000255 HAMAP-Rule:MF_04099, ECO:0000269 PubMed:32979942}; Single-pass type I membrane protein {ECO:0000255 HAMAP-Rule:MF_04099, ECO:0000269 PubMed:34504087}. Host endoplasmic reticulum-Golgi intermediate compartment membrane {ECO:0000255 HAMAP-Rule:MF_04099, ECO:0000269 PubMed:34504087}; Single-pass type I membrane protein {ECO:0000255 HAMAP-Rule:MF_04099}. Host cell membrane {ECO:0000255 HAMAP-Rule:MF_04099, ECO:0000269 PubMed:34504087}; Single-pass type I membrane protein {ECO:0000255 HAMAP-Rule:MF_04099}. Note=Accumulates in the endoplasmic reticulum-Golgi intermediate compartment, where it participates in virus particle assembly. Some S oligomers are transported to the host plasma membrane, where they may mediate cell-cell fusion (PubMed:34504087). An average of 26 +/-15 S trimers are found randomly distributed at the surface of the virion (PubMed:32979942) {ECO:0000255 HAMAP-Rule:MF_04099, ECO:0000269 PubMed:32979942, ECO:0000269 PubMed:34504087}</p>

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