

SARS-CoV-2 Spike protein (S1, D614G, His Tag)

Catalog # PVGS1956

Product Information

Primary Accession Species	P0DTC2 SARS-CoV-2
Sequence	Gln14-Arg685
Purity	> 90% as analyzed by SDS-PAGE
Endotoxin Level	
Biological Activity	SARS-CoV-2 Spike protein (S1, D614G, His Tag) can bind with human ACE2 (Cat.No.: Z03484) in functional ELISA assay.
Expression System	293 Cells
Theoretical Molecular Weight	79 kDa
Formulation	Supplied as a solution in PBS pH 7.2.
Storage & Stability	Upon receiving, this product remains stable for up to 6 months at -20°C or below. Avoid repeated freeze-thaw cycles.

Additional Information

Gene ID	43740568
Other Names	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}
Target Background	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. The spike protein mutation D614G became dominant during the globally pandemic. The D614G mutation enhances its replication ability in upper respiratory tract, transmission ability and viral loads in the patients' lung epithelial cells. It may also affect vaccine efficacy and antibody therapy.

Protein Information

Name	S {ECO:0000255 HAMAP-Rule:MF_04099}
Function	[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:32142651, PubMed:32155444, PubMed:33607086). When S2/S2' has been cleaved, binding to the receptor triggers direct fusion at the cell membrane (PubMed:34561887). 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:32075877, PubMed:32221306, PubMed:34903715, PubMed:36779763). Alternatively, may use NRP1/NRP2 (PubMed:33082294, PubMed:33082293) and integrin as entry receptors (PubMed:35150743). 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:33082293). The stalk domain of S contains three hinges, giving the head unexpected orientational freedom (PubMed:32817270).
Cellular Location	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}

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