

## Spike Protein S1

Catalog # PVGS1588

## **Product Information**

Primary Accession PODTC2
Species SARS-CoV-2

**Sequence** Gln14-Arg685 (E484Q, L452R)

**Purity** ≥ 95% as analyzed by SDS-PAGE

**Endotoxin Level** 

**Biological Activity**This protein is validated to bind with human ACE2 in functional ELISA assay.

Expression System CHO

Theoretical Molecular Weight 75.7 kDa

**Formulation** Supplied as a solution in PBS, pH 7.4.

**Storage & Stability** Upon receiving, this product remains stable for up to 6 months at -20°C or

below. Please 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. Lineage B.1.617, also known as G/452.V3, was first identified in October 2020 in India. This variant has the double mutations E484Q and L452R in the spike proteins. Emerging research suggests the variant may be more transmissible than previously evolved ones. Whether the effectiveness of currently-deployed vaccines is affected remains

under investigation. Moreover, the sublineages B.1.617.2 has been redesignated as "variant of concern" (VOC-21APR-02) in May 2021, which

spreads more quickly than the original version of the virus.

## **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}

Please note: All products are 'FOR RESEARCH USE ONLY. NOT FOR USE IN DIAGNOSTIC OR THERAPEUTIC PROCEDURES'.