SARS-CoV-2 S1 (RBD Mutant (L452R, T478K)), His-tag

Cat. no. P2020-060



Product Information

Protein: SARS-CoV-2 S1 (RBD) (L452R, T478K), His-tag (~ 27.6 kDa)

Uniprot#: PODTC2

Sequence: MRVQPTESIVRFPNITNLCPFGEVFNATRFASVYAWNRKRISNCVADYSVLYNSASFSTF

KCYGVSPTKLNDLCFTNVYADSFVIRGDEVRQIAPGQTGKIADYNYKLPDDFTGCVIAWN SNNLDSKVGGNYNYRYRLFRKSNLKPFERDISTEIYQAGSKPCNGVEGFNCYFPLQSYGF

QPTNGVGYQPYRVVVLSFELLHAPATVCGPKKSTNLVKNKCVNF

Methionine at pos. 1 present due to cloning constraints, C-terminal His-tag not shown in

sequence.

X indicates mutation sites.

Source: Recombinantly expressed in HEK293 cells.

Tag(s): His-tag, C-terminal

Purification: Purified by affinity chromatography and subsequent buffer exchange.

Formulation: PBS; pH 7.4

Liquid, stored and shipped at -80 °C.

Purity: > 75 % (will be determined by densitometry of Coomassie stained gel, example next page)

Concentration: Will be determined by BCA-Assay.

Long-term storage: No recommendations.

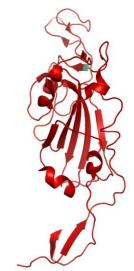
Comment: Protein migrates at higher molecular weight during SDS-PAGE due to posttranslational

modifications.

Background Information:

The spike (S) glycoprotein of coronaviruses is essential for binding of the virus to the host cell at the beginning of the infection process. The target protein is also a major immunogen and a possible target for entry inhibitors.

The SARS-CoV-2 spike (S) protein is a large type I transmembrane protein composed of two subunits, S1 and S2. The S1 subunit contains a receptor-binding domain (RBD) responsible for binding to the host cell receptor angiotensin-converting enzyme 2 (ACE2). Several mutants of the spike protein are known. A new SARS-CoV-2 lineage called B.1.617, exhibits 13 mutations. Compared to the previously circulating variants, the mutations L452R and T478K of the SARS-CoV-2 Spike S1 (RBD) may cause a stronger affinity of the spike protein to hACE2 and also conferring an increasing ability to evade the hosts' immune system.



Structural model of the receptor binding domain (RBD) of the spike protein. The location of the mutated part is indicated by the green circle.

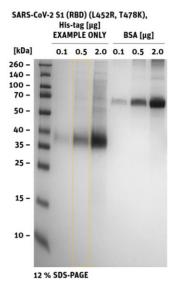
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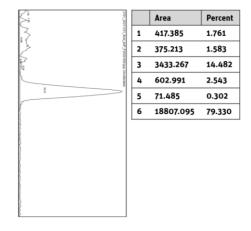


Product Information

Quality Information (provided for each lot):



SDS-PAGE/Coll.Coomassie



Histogram (of marked lane in gel picture)