

Catalog Number: 230-30161

Recombinant SARS-CoV-2 S1 Subunit, Full Length

Source

Species SARS-CoV-2
 Accession Number QHD43416
 Expressed Region Val16 - Gln690

• Synonyms Spike protein, S Protein, S1 Subunit.

Preparation

• Expression System Human embryonic kidney 293 (HEK293) cells

Tag C-terminal His-tag

Purification
 His-tag affinity purification by immobilized metal ion affinity chromatography (IMAC)

• Purity >80%

Purity determined By SDS-PAGE under reducing conditions and visualized by Coomassie blue staining

Molecular Weight Recombinant protein product has a calculated molecular mass of ~75 kDa. Due to the abundant

glycosylation, it migrates as approximately ~120 kDa major protein band in SDS-PAGE under DTT,

beta-mercaptoethanol reducing conditions. See deglycosylation analysis image below.

Protein Specifications

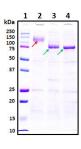
Format Liquid

• **Formulation** Supplied as a 0.2 μm filtered solution in PBS (pH 7.4)

Concentration Lot specific (see the label on the vial), determined by BCA protein assay.

SDS-PAGE Image Deglycosylation analysis

Deglycosylation analysis of purified recombinant proteins. Purified proteins were untreated (*Lane 2, right*) or treated with deglycosylation under native (*Lane 3*) or reducing (*Lane 4*) conditions. Deglycosylation treatment resulted in a mobility shift of the protein to produce one major band at the expected size (~75 kDa), thus indicating that the untreated recombinant protein (*Lane 2,* ~120 kDa) was glycosylated. **Lane 1**: Protein standard ladder (kDa). **Lane 2**: Untreated protein (~120 kDa, *red arrow*) under reducing conditions. **Lane 3**: Treated protein (~75 kDa, *green arrow*) with deglycosylation enzymes under native conditions. **Lane 4**: Treated protein (~75 kDa, *green arrow*) with deglycosylation enzymes under reducing conditions.



Shipping

The product is shipped with ice packs.

Storage/Stability

Upon arrival, the protein may be stored for 2 weeks at 4 $^{\circ}$ C. For long term storage, it is recommended to store at -20 $^{\circ}$ C or -80 $^{\circ}$ C in appropriate aliquots. Avoid repeated freeze-thaw cycles.

References

- F Wu, et al. A new coronavirus associated with human respiratory disease in China. Nature. 579, 265–269 (2020).
- N Dong, et al. Genomic and protein structure modelling analysis depicts the origin and infectivity of 2019-nCoV, a new coronavirus which caused a pneumonia outbreak in Wuhan, China. bioRxiv (2020).
- M Hoffmann, et al. SARS-CoV-2 Cell Entry Depends on ACE2 and TMPRSS2 and Is Blocked by a Clinically Proven Protease Inhibitor. Cell. 181, 1–10 (2020).
- W Li et al. Angiotensin-converting enzyme 2 is a functional receptor for the SARS coronavirus. Nature. 426, 450–454 (2003).

This product is furnished for LABORATORY RESEARCH USE ONLY.

Not for diagnostic or therapeutic use.







