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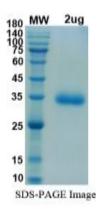
Recombinant SARS-CoV-2 (2019-nCoV) S Protein RBD(N501Y) His Tag (UK variant)

| Catalog No. | CSI99106 | Quantity: | 100 µg |
|-----------------------|--|----------------------------|---|
| Alternate Names: | UK variant (B.1.1.7) Spike glycoprotein, Spike S1 subunit receptor binding domain | | |
| Description: | The spike (S) glycoprotein of coronaviruses contains protrusions that will only bind to certain receptors on the host cell. Known receptors that bind S1 are ACE2, angiotensin- converting enzyme 2; DPP4, dipeptidyl peptidase-4; APN, aminopeptidase N; CEACAM, carcinoembryonic antigen-related cell adhesion molecule 1; Sia, sialic acid; O-ac Sia, O- acetylated sialic acid. The spike is essential for both host specificity and viral infectivity. The spike (S) glycoprotein of coronaviruses is known to be essential in the binding of the virus to the host cell at the advent of the infection process. The spike protein is a large type I transmembrane protein containing two subunits, S1 and S2. S1 mainly contains a receptor binding domain (RBD), which is responsible for recognizing the cell surface receptor. S2 mediates fusion of the virus in the induction of neutralizing-antibody and T-cell responses, as well as protective immunity. The main functions for the Spike protein are summarized as: Mediate receptor binding and membrane fusion; Defines the range of the hosts and specificity of the virus; Main component to bind with the neutralizing antibody; Key target for vaccine design; Can be transmitted between different hosts through gene recombination or mutation of the receptor binding domain (RBD), leading to a higher mortality rate. | | |
| UniProt ID: | P0DTC2 | | |
| Accession Number: | YP_009724390.1(N501Y) | | |
| Protein Construction: | A DNA sequence encoding t with a C-terminal His tag. | he SARS-CoV-2 (2019-nC | CoV) Spike S1 RBD was expressed |
| Source: | Mammalian cells | | |
| Formulation: | Lyophilized from sterile-filtered PBS, pH 7.5 | | |
| Molecular Weight: | 35 kDa | | |
| Purity: | > 90 % as determined by SDS-PAGE. | | |
| Endotoxin Level: | < 1.0 EU per µg protein as d | etermined by the LAL met | hod. |
| Reconstitution: | Centrifuge vial prior to ope mg/mL and gently pipette the DO NOT VORTEX . Allow se | e solution up and down the | |
| Storage & Stability: | Stable for up to 1 year from for 1 week at 2-8°C, or store Avoid repeated freeze-tha | working aliquots at -20°C | -80°C. After reconstitution, stable to -80°C. |



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