## SARS-CoV-2 S glycoprotein (HEK293, His, Flag)

| Cat. No.: | HY-P72035 |
| :--- | :--- |
| Synonyms: | E2 Peplomer protein |
| Species: | Virus |
| Source: | HEK293 |
| Accession: | P0DTC2 (V16-R685) |
| Gene ID: | 43740568 |
| Molecular Weight: | Approximately 120 kDa |

## PROPERTIES

## AA Sequence

Biological Activity

Appearance

Formulation

Endotoxin Level

Reconsititution

Storage \& Stability

| VNLTTRTQLP | PAYTNS F TRG | $V Y Y P D K V F R S$ | SVLHSTQDLF |
| :---: | :---: | :---: | :---: |
| LPFFSNVTWF | HA I HVSGTNG | T K R F D N P V L P | F NDGVYFAST |
| EKSNIIRGW I | FGTTLDSKTQ | SLLIVNNATN | VVIKVCEFQF |
| CNDPFLGVYY | HKNNKSWMES | EFRVYSSANN | CTFEYVSQPF |
| LMDLEGKQ ${ }^{\text {L }}$ | FKNLREFVFK | N I D G Y F K I Y S | K H T P I NLVRD |
| LPQGFSALEP | LVDLPIGINI | T R F Q T L A L H | RSYLTPGDSS |
| S GWTAGAAAY | Y V G Y L Q PRTF | L L K Y N E G T I | T DAVDCALDP |
| LSETKCTLKS | F TVEKGIYQT | SNFRVQPTES | I VRFPNITNL |
| CPFGEVFNAT | R F A S V Y W N R | KRISNCVADY | SVLYNSASFS |
| T F K C Y G S P T | KLNDLCFTNV | YADSFVIRGD | $E \vee R Q I A P G Q T$ |
| GKIADYNYKL | PDDFTGCVIA | W N S N L D K V | G G N Y N Y L Y R L |
| FRKSNLKPFE | RDISTEIYQA | G S T P CNGVEG | F NCYFPLQS Y |
| G F Q P T N V G Y | Q P Y R V V V L F | ELLHAPATVC | G P K K S N L V K |
| NKCVNFNFNG | LTGTGVLTES | NKKFLPFQQ F | GRD I A D T DA |
| VRDPQTLEIL | D I TPCSFGGV | SVITPGTNTS | NQ V $\mathrm{A} V \mathrm{~L} Y$ Q D V |
| NCTEVPVA I H | A D L T P TWRV | Y S T G S V F Q T | RAGCLIGAEH |
| VNNSYECDIP | IGAG I CASYQ | TQ T N S PRRAR |  |

Measured by its binding ability in a functional ELISA. Immobilized SARS-CoV-2-S at $2 \mu \mathrm{~g} / \mathrm{mL}$ can bind human ACE2 , the EC 50 of SARS-CoV-2-S protein is $56.64-103.6 \mathrm{ng} / \mathrm{mL}$.

Lyophilized powder.

Lyophilized from a $0.2 \mu \mathrm{~m}$ solution of PBS, $6 \%$ Trehalose, pH 7.4 .
$<1 \mathrm{EU} / \mu \mathrm{g}$, determined by LAL method.

It is not recommended to reconstitute to a concentration less than $100 \mu \mathrm{~g} / \mathrm{mL}$ in ddH2 $\mathrm{O}_{2}$.

Stored at $-20^{\circ} \mathrm{C}$ for 2 years. After reconstitution, it is stable at $4^{\circ} \mathrm{C}$ for 1 week or $-20^{\circ} \mathrm{C}$ for longer (with carrier protein). It is recommended to freeze aliquots at $-20^{\circ} \mathrm{C}$ or $-80^{\circ} \mathrm{C}$ for extended storage.

## DESCRIPTION

## Background

The SARS-CoV-2 S glycoprotein plays a crucial role in infection by attaching the virion to the host cell membrane through interaction with the primary receptor, host ACE2. Upon cleavage of S2/S2', binding to the ACE2 receptor initiates either direct fusion at the cell membrane or internalization of the virus via endocytosis, leading to fusion of the virion membrane with the host endosomal membrane. Additionally, the glycoprotein may utilize NRP1/NRP2 and integrin as alternative entry receptors, possibly explaining the virus's tropism in human olfactory epithelial cells. The stalk domain of S exhibits three hinges, providing unexpected orientational freedom to the head. Acting as a class I viral fusion protein, the glycoprotein undergoes an extensive and irreversible conformational change during virus entry, triggered by host TMPRSS2 or CSTL, leading to fusion of the viral envelope with the cellular cytoplasmic membrane and release of viral genomic RNA into the host cell cytoplasm. The glycoprotein exhibits at least three conformational states: pre-fusion native, pre-hairpin intermediate, and post-fusion hairpin, with the coiled coil regions adopting a trimer-of-hairpins structure during fusion, facilitating the apposition and subsequent fusion of viral and target cell membranes.

## Caution: Product has not been fully validated for medical applications. For research use only.

Tel: 609-228-6898 Fax: 609-228-5909 E-mail: tech@MedChemExpress.com
Address: 1 Deer Park Dr, Suite O, Monmouth Junction, NJ 08852, USA

