

# Screening Libraries

**Proteins** 

## ors • Sci

## **Product** Data Sheet

## MCE RedChemExpress

### HCoV-HKU1 Spike Protein (Q0ZME7, sf9, His)

Cat. No.: HY-P77386

Synonyms: Human coronAvirus HKU1 (isolate N5) (HCoV-HKU1) Spike Protein (S1+S2, His)

Species: Virus

**Source:** Sf9 insect cells

Accession: Q0ZME7 (M1-P1295)

Gene ID: /

Molecular Weight: Approximately 144.4 kDa.

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Appearance	Lyophilized powder.
Formulation	Lyophilized from a 0.2 μm filtered solution of 20 mM PB, 300 mM NaCl, 10% Glycerol, pH 7.5. Normally 5 % - 8 % trehalose, mannitol and 0.01% Tween 80 are added as protectants before lyophilization.
Endotoxin Level	<1 EU/µg, determined by LAL method.
Reconsititution	It is not recommended to reconstitute to a concentration less than 100 $\mu g/mL$ in ddH $_2$ O.
Storage & Stability	Stored at -20°C for 2 years. After reconstitution, it is stable at 4°C for 1 week or -20°C for longer (with carrier protein). It is recommended to freeze aliquots at -20°C or -80°C for extended storage.
Shipping	Room temperature in continental US; may vary elsewhere.

#### **DESCRIPTION**

Background

The HCoV-HKU1 Spike/S1 Protein plays a pivotal role in the viral infection process by attaching the virion to the cell membrane through interaction with host receptors, thereby initiating infection. Functioning as a class I viral fusion protein, it mediates the fusion of the virion with cellular membranes. The protein undergoes at least three conformational states: a pre-fusion native state, a pre-hairpin intermediate state, and a post-fusion hairpin state. During the fusion process between viral and target cell membranes, the coiled coil regions (heptad repeats) adopt a trimer-of-hairpins structure, positioning the fusion peptide in close proximity to the C-terminal region of the ectodomain. This structural rearrangement is crucial for driving 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.

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