## Product Information

| Product SKU: | CAB20394 | Gene ID: | 918758 | Size: | 20uL, 100uL |
| :--- | :--- | :--- | :--- | :--- | :--- |
| Clone No: | - | Host Species: | Rabbit | Reactivity: | HCoV-229E |

## Additional Information

| Observed MW: | 160 kDa | Conjugate: | Unconjugated |
| :--- | :--- | :--- | :--- |
| Calculated MW: | 129 kDa | Isotype: | lgG |

## Immunogen Information

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\begin{array}{ll}\text { Background: } & \begin{array}{l}\text { S1 region attaches the virion to the cell membrane by interacting with host ANPEP/aminopeptidase N, } \\
\text { initiating the infection. Binding to the receptor probably induces conformational changes in the S }\end{array}
$$ <br>
glycoprotein unmasking the fusion peptide of S2 region and activating membranes fusion. S2 region <br>
belongs to the class I viral fusion protein. Under the current model, the protein has at least 3 <br>
conformational states: pre-fusion native state, pre-hairpin intermediate state, and post-fusion hairpin <br>
state. During viral and target cell membrane fusion, the coiled coil regions (heptad repeats regions <br>

assume a trimer-of-hairpins structure, positioning the fusion peptide in close proximity to the C-terminal\end{array}\right\}\)| region of the ectodomain. The formation of this structure appears to drive apposition and subsequent |
| :--- | :--- |
| fusion of viral and target cell membranes. |

