

Anti-SARS-CoV-2 Spike glycoprotein antibody (C617)

Purified Rabbit Polyclonal Antibody (Pab)

Catalog # AP22388I

Product Information

Application	E
Primary Accession	P0DTC2
Host	Rabbit
Clonality	Polyclonal
Isotype	Rabbit IgG
Clone Names	R00327-330
Calculated MW	141178

Additional Information

Gene ID	43740568
Target/Specificity	This Anti-SARS-CoV-2 Spike glycoprotein antibody is generated from rabbits immunized with a KLH conjugated synthetic peptide between 603-645 amino acids from the N-terminal region of human SARS-CoV-2 Spike glycoprotein.
Format	Purified polyclonal antibody supplied in PBS with 0.05% (V/V) Proclin 300. This antibody is purified through a protein A column, followed by peptide affinity purification.
Storage	Maintain refrigerated at 2-8°C for up to 2 weeks. For long term storage store at -20°C in small aliquots to prevent freeze-thaw cycles.
Precautions	Anti-SARS-CoV-2 Spike glycoprotein antibody (C617) is for research use only and not for use in diagnostic or therapeutic procedures.

Protein Information

Name	S {ECO:0000255 HAMAP-Rule:MF_04099}
Function	[Spike protein S1]: Attaches the virion to the cell membrane by interacting with host receptor, initiating the infection. The major receptor is host ACE2 (PubMed: 32142651 , PubMed: 33607086 , PubMed: 32155444). When S2/S2' has been cleaved, binding to the receptor triggers direct fusion at the cell membrane (PubMed: 34561887). When S2/s2' has not been cleaved, binding to the receptor results in internalization of the virus by endocytosis leading to fusion of the virion membrane with the host endosomal membrane (PubMed: 32221306 , PubMed: 32075877). Alternatively, may use NRP1/NRP2 (PubMed: 33082294 , PubMed: 33082293) and integrin as entry receptors (PubMed: 35150743). The use of NRP1/NRP2 receptors may explain the tropism of the virus in human olfactory epithelial cells, which express these molecules at high levels but ACE2 at low levels (PubMed: 33082293). The stalk

domain of S contains three hinges, giving the head unexpected orientational freedom (PubMed:[32817270](#)).

Cellular Location

Virion membrane {ECO:0000255 | HAMAP-Rule:MF_04099, ECO:0000269 | PubMed:32979942}; Single-pass type I membrane protein {ECO:0000255 | HAMAP-Rule:MF_04099, ECO:0000269 | PubMed:34504087}. Host endoplasmic reticulum-Golgi intermediate compartment membrane {ECO:0000255 | HAMAP-Rule:MF_04099, ECO:0000269 | PubMed:34504087}; Single-pass type I membrane protein {ECO:0000255 | HAMAP-Rule:MF_04099}. Host cell membrane {ECO:0000255 | HAMAP-Rule:MF_04099, ECO:0000269 | PubMed:34504087}; Single-pass type I membrane protein {ECO:0000255 | HAMAP-Rule:MF_04099}. Note=Accumulates in the endoplasmic reticulum-Golgi intermediate compartment, where it participates in virus particle assembly. Some S oligomers are transported to the host plasma membrane, where they may mediate cell-cell fusion (PubMed:34504087). An average of 26 +/-15 S trimers are found randomly distributed at the surface of the virion (PubMed:32979942) {ECO:0000255 | HAMAP-Rule:MF_04099, ECO:0000269 | PubMed:32979942, ECO:0000269 | PubMed:34504087}

Background

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. It's been reported that 2019-nCoV can infect the human respiratory epithelial cells through interaction with the human ACE2 receptor. 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 contains basic elements needed for the membrane fusion. The S protein plays key parts 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.