Spike trimer translocates to ERGIC

Acencio, ML., Stephan, R.
Introduction

Reactome is open-source, open access, manually curated and peer-reviewed pathway database. Pathway annotations are authored by expert biologists, in collaboration with Reactome editorial staff and cross-referenced to many bioinformatics databases. A system of evidence tracking ensures that all assertions are backed up by the primary literature. Reactome is used by clinicians, geneticists, genomics researchers, and molecular biologists to interpret the results of high-throughput experimental studies, by bioinformaticians seeking to develop novel algorithms for mining knowledge from genomic studies, and by systems biologists building predictive models of normal and disease variant pathways.

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Literature references


Reactome database release: 77

This document contains 1 reaction (see Table of Contents)
Spike trimer translocates to ERGIC

**Stable identifier:** R-HSA-9694780

**Type:** omitted

**Compartments:** endoplasmic reticulum membrane, endoplasmic reticulum-Golgi intermediate compartment

**Diseases:** COVID-19

**Inferred from:** Spike trimer translocates to ERGIC (Homo sapiens)

This COVID-19 event has been created by a combination of computational inference (see https://reactome.org/documentation/inferred-events) from SARS-CoV-1 data and manual curation, as described in the summation for the overall SARS-CoV-2 infection pathway.

Trimmed palmitoylated Spike protein trimers become associated with the ERGIC (ER-Golgi Intermediate Compartment) (Fung & Liu 2019)

**Literature references**


**Editions**

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