

ER to Golgi vesicle-mediated transport, GO:0006888

- retrograde vesicle-mediated transport, GO:0006890
- Golgi organization, GO:0007030
- IRE1-mediated unfolded protein response, GO:0036498
- azurophil granule lumen, GO:0035578
- secretory granule membrane, GO:0030667
- exocytosis, GO:0006887
- COPII vesicle coating, GO:0048208
- trans-Golgi network membrane, GO:0032588
- endoplasmic reticulum-Golgi intermediate compartment membrane, GO:0033116
- Cul3-RING ubiquitin ligase complex, GO:0031463
- SNARE complex, GO:0031201
- syntaxin binding, GO:0019905
- platelet degranulation, GO:0002576
- transport vesicle, GO:0030133
- lipid particle, GO:0005811
- calcium-dependent protein binding, GO:0048306
- ficolin-1-rich granule membrane, GO:0101003
- vesicle fusion, GO:0006906
- SNAP receptor activity, GO:0005484
- ER to Golgi transport vesicle membrane, GO:0012507
- SNARE binding, GO:0000149
- dendritic shaft, GO:0043198
- ER to Golgi transport vesicle, GO:0030134
- pigmentation, GO:0043473
- hemostasis, GO:0007599
- COPI-coated vesicle membrane, GO:0030663
- endoplasmic reticulum exit site, GO:0070971
- vesicle docking involved in exocytosis, GO:0006904
- autophagosome maturation, GO:0097352
- COPII vesicle coat, GO:0030127
- calcium-dependent cysteine-type endopeptidase activity, GO:0004198
- COPI vesicle coat, GO:0030126
- vesicle docking, GO:0048278
- protein targeting to plasma membrane, GO:0072661
- intra-Golgi vesicle-mediated transport, GO:0006891
- platelet alpha granule lumen, GO:0031093
- COPI-coated vesicle, GO:0030137
- Golgi to plasma membrane protein transport, GO:0043001
- Golgi ribbon formation, GO:0090161
- response to calcium ion, GO:0051592
- WASH complex, GO:0071203
- neural crest cell development, GO:0014032
- cargo loading into COPII-coated vesicle, GO:0090110
- phospholipase activity, GO:0004620
- Golgi vesicle transport, GO:0048193
- nuclear envelope reassembly, GO:0031468
- serine-type peptidase activity, GO:0008236
- negative regulation of autophagosome assembly, GO:1902902
- regulation of protein localization to plasma membrane, GO:1903076
- neural crest formation, GO:0014029
- apical protein localization, GO:0045176
- endoplasmic reticulum organization, GO:0007029
- axolemma, GO:0030673
- fibrinolysis, GO:0042730
- GPI anchor binding, GO:0034235
- chloride channel inhibitor activity, GO:0019869
- regulation of complement activation, GO:0030449
- negative regulation of cytokine production involved in inflammatory response, GO:1900016
- synaptic transmission, GO:0035249
- TRAPP complex, GO:0030008
- barbed-end actin filament capping, GO:0051016
- negative regulation of proteolysis, GO:0045861
- Golgi lumen, GO:0005796
- costamere, GO:0043034
- negative regulation of fibrinolysis, GO:0051918
- signal peptide processing, GO:0006465
- angiotensin maturation, GO:0002003
- common-partner SMAD protein phosphorylation, GO:0007182
- activation of signaling protein activity involved in unfolded protein response, GO:0006987
- actin filament capping, GO:0051693
- zymogen activation, GO:0031638
- Sec61 translocon complex, GO:0005784
- T cell mediated cytotoxicity, GO:0001913
- compact myelin, GO:0043218
- negative regulation of endopeptidase activity, GO:0010951
- central nervous system projection neuron axonogenesis, GO:0021952
- ventricular cardiac muscle cell action potential, GO:0086005
- regulation of lipoprotein lipase activity, GO:0051004