

ER to Golgi vesicle-mediated transport, GO:0006888

- protein transporter activity, GO:0008565
- IRE1-mediated unfolded protein response, GO:0036498
- secretory granule membrane, GO:0030667
- COPII vesicle coating, GO:0048208
- synaptic vesicle, GO:0008021
- endoplasmic reticulum-Golgi intermediate compartment, GO:0005793
- transport vesicle, GO:0030133
- cis-Golgi network, GO:0005801
- Golgi cisterna membrane, GO:0032580
- endoplasmic reticulum-Golgi intermediate compartment membrane, GO:0033116
- Cul3-RING ubiquitin ligase complex, GO:0031463
- ficolin-1-rich granule membrane, GO:0101003
- Rab guanyl-nucleotide exchange factor activity, GO:0017112
- ER to Golgi transport vesicle membrane, GO:0012507
- antigen processing and presentation of peptide antigen via MHC class I, GO:0002474
- phospholipid binding, GO:0005543
- Ras guanyl-nucleotide exchange factor activity, GO:0005088
- endoplasmic reticulum unfolded protein response, GO:0030968
- protein secretion, GO:0009306
- response to calcium ion, GO:0051592
- mitotic cytokinesis, GO:0000281
- protein N-linked glycosylation via asparagine, GO:0018279
- dendritic shaft, GO:0043198
- integral component of Golgi membrane, GO:0030173
- ATP-dependent microtubule motor activity, GO:0008574
- Golgi stack, GO:0005795
- COPI-coated vesicle, GO:0030137
- anchored component of membrane, GO:0031225
- pigmentation, GO:0043473
- ARF guanyl-nucleotide exchange factor activity, GO:0005086
- microtubule-based process, GO:0007017
- Golgi transport complex, GO:0017119
- intrinsic component of the cytoplasmic side of the plasma membrane, GO:0031235
- dynein intermediate chain binding, GO:0045505
- endoplasmic reticulum exit site, GO:0070971
- axolemma, GO:0030673
- establishment of Golgi localization, GO:0051683
- dynein light intermediate chain binding, GO:0051959
- M band, GO:0031430
- COPI-coated vesicle membrane, GO:0030663
- positive regulation of ubiquitin protein ligase activity, GO:1904668
- COPI vesicle coat, GO:0030126
- heart looping, GO:0001947
- TRAPP complex, GO:0030008
- protein retention in ER lumen, GO:0006621
- nuclear migration, GO:0007097
- Golgi to plasma membrane protein transport, GO:0043001
- perinuclear endoplasmic reticulum, GO:0097038
- regulation of complement activation, GO:0030449
- lipid particle organization, GO:0034389
- melanosome transport, GO:0032402
- COPII vesicle coat, GO:0030127
- blood coagulation, GO:0007597
- protein localization to Golgi apparatus, GO:0034067
- acute-phase response, GO:0006953
- lipoprotein transport, GO:0042953
- ankyrin binding, GO:0030506
- positive regulation of cytokine-mediated signaling pathway, GO:0001961
- response to alkaloid, GO:0043279
- signal peptide processing, GO:0006465
- anchored component of external side of plasma membrane, GO:0031362