

ER to Golgi vesicle-mediated transport, GO:0006888

- vesicle, GO:0031982
- secretory granule membrane, GO:0030667
- IRE1-mediated unfolded protein response, GO:0036498
- trans-Golgi network membrane, GO:0032588
- COPII vesicle coating, GO:0048208
- exocytosis, GO:0006887
- endoplasmic reticulum-Golgi intermediate compartment membrane, GO:0033116
- ficolin-1-rich granule membrane, GO:0101003
- Golgi cisterna membrane, GO:0032580
- ER to Golgi transport vesicle membrane, GO:0012507
- transport vesicle, GO:0030133
- platelet degranulation, GO:0002576
- lipid particle, GO:0005811
- protein N-linked glycosylation via asparagine, GO:0018279
- protein oligomerization, GO:0051259
- SNARE complex, GO:0031201
- integral component of Golgi membrane, GO:0030173
- SNAP receptor activity, GO:0005484
- endoplasmic reticulum organization, GO:0007029
- hemostasis, GO:0007599
- cis-Golgi network, GO:0005801
- syntaxin binding, GO:0019905
- autophagosome maturation, GO:0097352
- intra-Golgi vesicle-mediated transport, GO:0006891
- vesicle docking involved in exocytosis, GO:0006904
- SNARE binding, GO:0000149
- transport vesicle membrane, GO:0030658
- protein targeting to plasma membrane, GO:0072661
- vesicle fusion, GO:0006906
- dendritic shaft, GO:0043198
- dynein light intermediate chain binding, GO:0051959
- vesicle docking, GO:0048278
- COPI-coated vesicle membrane, GO:0030663
- regulation of complement activation, GO:0030449
- Golgi to plasma membrane protein transport, GO:0043001
- response to calcium ion, GO:0051592
- COPI vesicle coat, GO:0030126
- TRAPP complex, GO:0030008
- platelet alpha granule lumen, GO:0031093
- protein localization to pre-autophagosomal structure, GO:0034497
- calcium-dependent cysteine-type endopeptidase activity, GO:0004198
- axolemma, GO:0030673
- COPII vesicle coat, GO:0030127
- dynein intermediate chain binding, GO:0045505
- serine-type peptidase activity, GO:0008236
- mammary gland alveolus development, GO:0060749
- synaptic transmission, GO:0035249
- negative regulation of blood coagulation, GO:0030195
- Golgi ribbon formation, GO:0090161
- Golgi vesicle transport, GO:0048193
- Golgi-associated vesicle, GO:0005798
- barbed-end actin filament capping, GO:0051016
- costamere, GO:0043034
- anchored component of membrane, GO:0031225
- actin filament capping, GO:0051693
- nucleophagy, GO:0044804
- pre-autophagosomal structure membrane, GO:0034045
- copper ion binding, GO:0005507
- atrial cardiac muscle cell action potential, GO:0086014
- intrinsic component of the cytoplasmic side of the plasma membrane, GO:0031235
- voluntary musculoskeletal movement, GO:0050882
- muscle fiber development, GO:0048747
- regulation of lipoprotein lipase activity, GO:0051004
- auditory receptor cell stereocilium organization, GO:0060088
- regulation of AMPA receptor activity, GO:2000311
- phospholipase activity, GO:0004620
- signal peptide processing, GO:0006465
- Golgi cis cisterna, GO:0000137
- rough endoplasmic reticulum, GO:0005791
- perinuclear endoplasmic reticulum, GO:0097038
- labyrinthine layer blood vessel development, GO:0060716
- response to cholesterol, GO:0070723
- T cell mediated cytotoxicity, GO:0001913