late endosome membrane, GO:0031902

late endosome, GO:0005770 specific granule membrane, GO:0035579 azurophil granule membrane, GO:0035577 autophagosome, GO:0005776 tertiary granule membrane, GO:0070821 antigen processing and presentation, GO:0019882 T cell costimulation, GO:0031295 positive regulation of neuron projection development, GO:0010976 MHC class II protein complex, GO:0042613 antigen processing and presentation of peptide or polysaccharide antigen via MHC class II, GO:0002504 retrograde transport, GO:1990126 lysosome localization, GO:0032418 cation transport, GO:0006812 HOPS complex, GO:0030897 antiporter activity, GO:0015297 AP-3 adaptor complex, GO:0030123 response to zinc ion, GO:0010043 vacuolar transport, GO:0007034 chloride channel activity, GO:0005254 ESCRT III complex, GO:0000815 zinc II ion transport, GO:0006829 calcium-dependent phospholipid binding, GO:0005544 positive regulation of cellular protein catabolic process, GO:1903364 response to interferon-beta, GO:0035456 regulation of centrosome duplication, GO:0010824 MHC class II receptor activity, GO:0032395 zinc II ion transmembrane transport, GO:0071577 T-helper 1 type immune response, GO:0042088 phosphatidylcholine binding, GO:0031210 multivesicular body, GO:0005771 ESCRT I complex, GO:0000813 regulated exocytosis, GO:0045055 cholesterol transporter activity, GO:0017127 collagen fibril organization, GO:0030199 sterol transporter activity, GO:0015248 phospholipase inhibitor activity, GO:0004859 autophagosome membrane, GO:0000421 cadherin binding involved in cell-cell adhesion, GO:0098641 sterol transport, GO:0015918 regulation of anion transport, GO:0044070 establishment of protein localization to membrane, GO:0090150 CORVET complex, GO:0033263 negative regulation of macrophage derived foam cell differentiation, GO:0010745 specific granule, GO:0042581 negative regulation of serine-type endopeptidase activity, GO:1900004 Golgi-associated vesicle membrane, GO:0030660 cation transmembrane transport, GO:0098655 solute:proton antiporter activity, GO:0015299 regulation of muscle contraction, GO:0006937