

specific granule membrane, GO:0035579

- tertiary granule membrane, GO:0070821
- regulation of ion transmembrane transport, GO:0034765
- voltage-gated ion channel activity, GO:0005244
- anchored component of membrane, GO:0031225
- lipid transport, GO:0006869
- metalloendopeptidase activity, GO:0004222
- clathrin-coated vesicle membrane, GO:0030665
- phospholipid transport, GO:0015914
- integrin complex, GO:0008305
- carbohydrate transport, GO:0008643
- glucose transmembrane transport, GO:1904659
- phospholipid translocation, GO:0045332
- ceramide biosynthetic process, GO:0046513
- heterotypic cell-cell adhesion, GO:0034113
- positive regulation of membrane protein ectodomain proteolysis, GO:0051044
- low-density lipoprotein receptor activity, GO:0005041
- phospholipase C-activating G-protein coupled receptor signaling pathway, GO:0007200
- ethanol catabolic process, GO:0006068
- phospholipid-translocating ATPase activity, GO:0004012
- substrate-specific transmembrane transporter activity, GO:0022891
- regulation of neurotransmitter secretion, GO:0046928
- lipoprotein transport, GO:0042953
- glucose transmembrane transporter activity, GO:0005355
- platelet alpha granule membrane, GO:0031092
- cell-cell adhesion via plasma-membrane adhesion molecules, GO:0098742
- melanosome organization, GO:0032438
- specific granule, GO:0042581
- triglyceride biosynthetic process, GO:0019432
- sperm-egg recognition, GO:0035036
- intestinal cholesterol absorption, GO:0030299
- azurophil granule, GO:0042582
- regulation of potassium ion transmembrane transport, GO:1901379
- diacylglycerol metabolic process, GO:0046339
- attachment of GPI anchor to protein, GO:0016255
- interleukin-1 beta secretion, GO:0050702
- regulation of anion transmembrane transport, GO:1903959
- sodium-independent organic anion transport, GO:0043252
- filopodium membrane, GO:0031527
- regulation of proteolysis, GO:0030162