specific granule membrane, GO:0035579

electron carrier activity, GO:0009055 metalloendopeptidase activity, GO:0004222 regulation of ion transmembrane transport, GO:0034765 anchored component of membrane, GO:0031225 voltage-gated ion channel activity, GO:0005244 endodermal cell differentiation, GO:0035987 carbohydrate transport, GO:0008643 ceramide biosynthetic process, GO:0046513 integrin complex, GO:0008305 heterotypic cell-cell adhesion, GO:0034113 regulation of insulin secretion, GO:0050796 platelet alpha granule membrane, GO:0031092 phagocytosis, GO:0006911 NAD metabolic process, GO:0019674 nitric oxide mediated signal transduction, GO:0007263 natural killer cell activation, GO:0030101 superoxide metabolic process, GO:0006801 protein maturation, GO:0051604 unsaturated fatty acid biosynthetic process, GO:0006636 lipid storage, GO:0019915 cation channel activity, GO:0005261 synaptic transmission, GO:0007271 aldehyde dehydrogenase (NAD) activity, GO:0004029 aldo-keto reductase (NADP) activity, GO:0004033 positive chemotaxis, GO:0050918 cell adhesion mediated by integrin, GO:0033627 N-acylphosphatidylethanolamine-specific phospholipase D activity, GO:0070290 peptide transport, GO:0015833 filopodium membrane, GO:0031527 attachment of GPI anchor to protein, GO:0016255 negative regulation of vascular permeability, GO:0043116 regulation of potassium ion transmembrane transport, GO:1901379 negative regulation of T cell receptor signaling pathway, GO:0050860 glucose transmembrane transporter activity, GO:0005355 ethanol catabolic process, GO:0006068 aldehyde dehydrogenase [NAD(P)+] activity, GO:0004030 positive regulation of monocyte chemotaxis, GO:0090026