

**specific granule membrane, GO:0035579**

- positive regulation of MAPK cascade, GO:0043410
- integrin-mediated signaling pathway, GO:0007229
- regulation of ion transmembrane transport, GO:0034765
- voltage-gated ion channel activity, GO:0005244
- integrin complex, GO:0008305
- metalloendopeptidase activity, GO:0004222
- ceramide biosynthetic process, GO:0046513
- transforming growth factor beta binding, GO:0050431
- phagocytosis, GO:0006911
- insulin-like growth factor I binding, GO:0031994
- excitatory postsynaptic potential, GO:0060079
- superoxide metabolic process, GO:0006801
- blood circulation, GO:0008015
- heterotypic cell-cell adhesion, GO:0034113
- protein maturation, GO:0051604
- integral component of nuclear inner membrane, GO:0005639
- cell adhesion mediated by integrin, GO:0033627
- substrate-specific transmembrane transporter activity, GO:0022891
- synaptic vesicle fusion to presynaptic active zone membrane, GO:0031629
- cation channel activity, GO:0005261
- aldo-keto reductase (NADP) activity, GO:0004033
- lamellipodium membrane, GO:0031258
- negative regulation of platelet aggregation, GO:0090331
- positive regulation of monocyte chemotaxis, GO:0090026
- synaptic vesicle docking, GO:0016081
- positive regulation of focal adhesion assembly, GO:0051894
- phosphorus-oxygen lyase activity, GO:0016849
- nitric oxide mediated signal transduction, GO:0007263
- endodermal cell differentiation, GO:0035987
- response to ATP, GO:0033198
- bleb assembly, GO:0032060
- phospholipase C-activating G-protein coupled receptor signaling pathway, GO:0007200
- aldehyde dehydrogenase (NAD) activity, GO:0004029
- cellular aldehyde metabolic process, GO:0006081
- lipid storage, GO:0019915
- NAD metabolic process, GO:0019674
- negative regulation of granulocyte differentiation, GO:0030853
- leukocyte migration involved in inflammatory response, GO:0002523
- unsaturated fatty acid biosynthetic process, GO:0006636
- positive regulation of bone resorption, GO:0045780
- filopodium membrane, GO:0031527
- positive chemotaxis, GO:0050918
- calcium-release channel activity, GO:0015278
- positive regulation of cell-cell adhesion, GO:0022409
- positive regulation of cellular extravasation, GO:0002693
- smooth muscle contraction, GO:0006939
- cell-cell adhesion via plasma-membrane adhesion molecules, GO:0098742
- bile acid transmembrane transporter activity, GO:0015125
- positive regulation of superoxide anion generation, GO:0032930
- acylglycerol acyl-chain remodeling, GO:0036155
- lymphocyte chemotaxis, GO:0048247
- C-X3-C chemokine binding, GO:0019960
- regulation of toll-like receptor signaling pathway, GO:0034121
- interleukin-1 beta secretion, GO:0050702
- calcium activated cation channel activity, GO:0005227
- ethanol catabolic process, GO:0006068
- 2-acylglycerol O-acyltransferase activity, GO:0003846
- positive regulation of tumor necrosis factor biosynthetic process, GO:0042535