

GO_Biological_Process_2021

Term

Golgi transport vesicle coating (GO:0048200)
 COPI-coated vesicle budding (GO:0035964)
 COPI coating of Golgi vesicle (GO:0048205)
 positive regulation of mRNA metabolic
 regulation of proteolysis involved in
 process (GO:1903313)
 cellular protein catabolic process
 mitochondrial RNA metabolic process
 (GO:1903050)
 spontaneous conformational changes to
 generate catalytic conformation (GO:0000393)
 tRNA transport (GO:0051029)
 RNA import into mitochondrion (GO:0035927)
 regulation of miRNA catabolic process
 (GO:2000635)
 cell proliferation involved in
 regulation of adenylate cyclase-activating G
 heart morphogenesis (GO:2000136)
 protein-coupled receptor signaling pathway
 protein localization to microtubule plus-end
 (GO:0106070)
 protein localization to microtubule end
 (GO:1905225)
 (GO:1905725)
 positive regulation (GO:0009249) of
 (GO:2009249)
 positive regulation of endothelial response to
 heparin (GO:0061419) via fibronectin
 positive regulation of cell proliferation
 (GO:1904906)
 positive regulation by histone H3 (GO:20000138)
 replication (GO:0001489) mRNA at the site of
 negative regulation of oxidative
 transcription (GO:0071033)
 stress-induced neuron intrinsic apoptotic
 negative regulation of macrophage cytokine
 signaling pathway (GO:1903377)
 negative regulation of leukocyte adhesion to
 vascular endothelial cells (GO:1904995)
 negative regulation of cell cycle (GO:1903208)
 mitochondrial tRNA processing
 (GO:0001007)
 (GO:0000095) of animal organ identity
 (GO:0048496)
 immature B cell differentiation (GO:0002327)
 extracellular exosome biogenesis
 (GO:0007344)
 GTP-AMP transmembrane import across
 plasma membrane (GO:0140059)
 differentiation
 (GO:0060370)

0

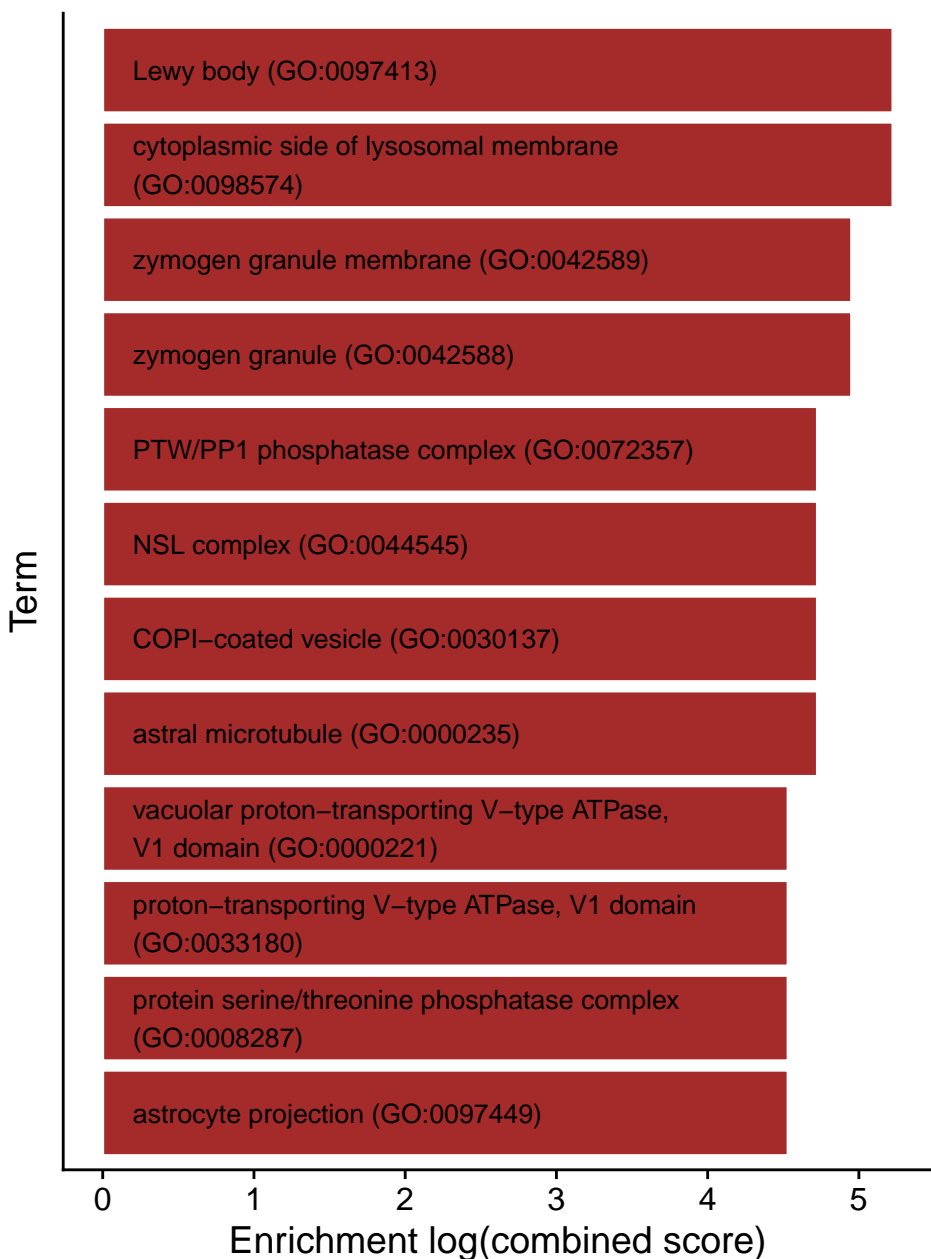
2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021



GO_Molecular_Function_2021

Term

alpha-2A adrenergic receptor binding
(GO:0031694)

chitinase activity (GO:0004568)

C3HC4-type RING finger domain binding
(GO:0055131)

phosphatidic acid transfer activity
(GO:1990050)

protein kinase B binding (GO:0043422)

poly(G) binding (GO:0034046)

calcium-dependent protein kinase activity
(GO:0010857)

volume-sensitive anion channel activity
(GO:0005225)

double-stranded telomeric DNA binding
(GO:0003691)

armadillo repeat domain binding (GO:0070016)

0

1

2

3

4

5

Enrichment log(combined score)