

GO_Biological_Process_2021

Term

axon choice point recognition (GO:0016198)

regulation of extracellular matrix
disassembly (GO:0010715)

astral microtubule organization (GO:0030953)

lipid phosphorylation (GO:0046834)

negative regulation of platelet-derived
growth factor receptor signaling pathway
(GO:0010642)

glycerolipid metabolic process (GO:0046486)

regulation of ventricular cardiac muscle
cell action potential (GO:0098911)

transport along microtubule (GO:0010970)

regulation of homotypic cell-cell adhesion
(GO:0034110)

aorta development (GO:0035904)

0

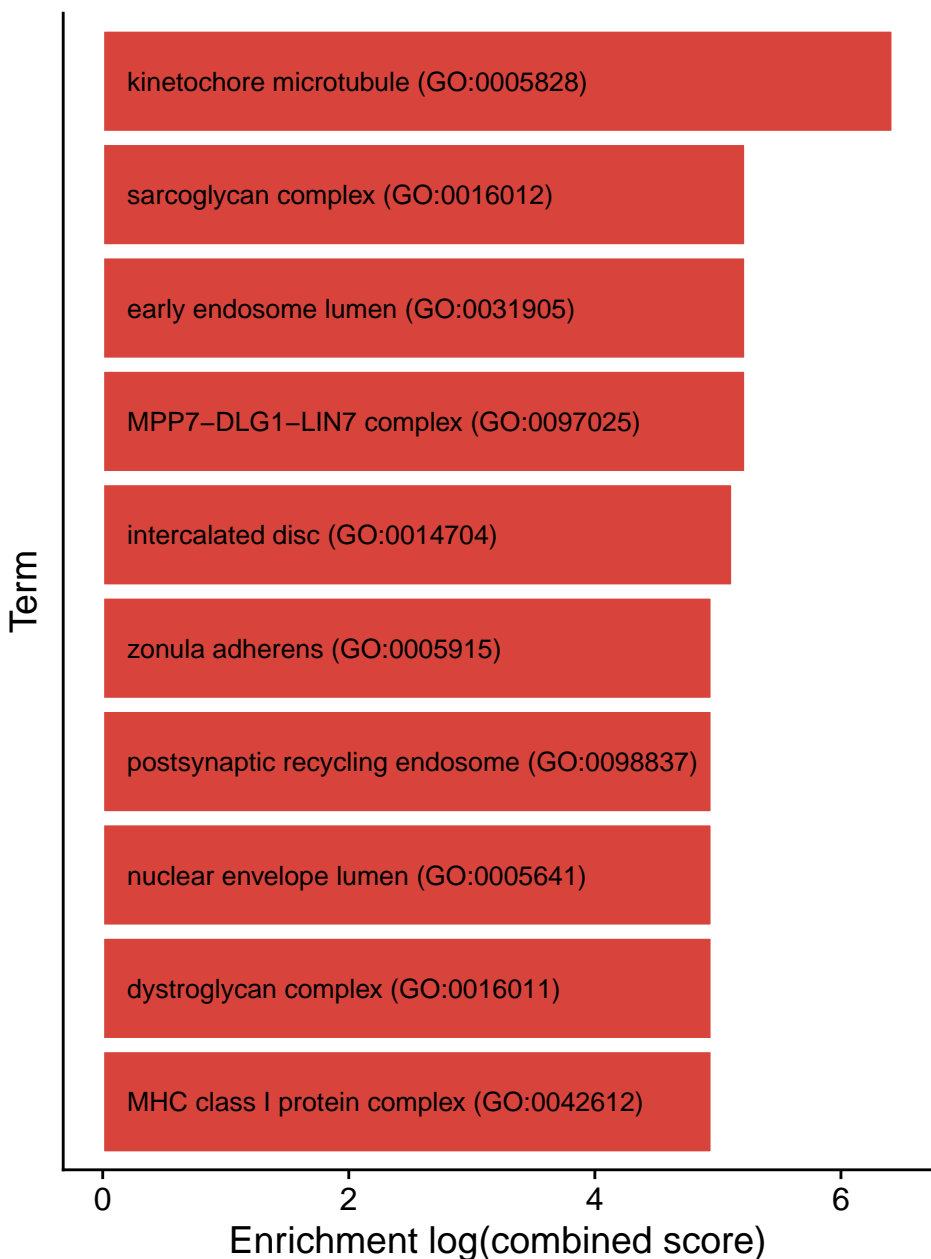
2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021



GO_Molecular_Function_2021

Term

diacylglycerol kinase activity (GO:0004143)

microtubule minus-end binding (GO:0051011)

histone acetyltransferase activity (H3-K23 specific) (GO:0043994)

transmembrane receptor protein tyrosine

phosphatase activity (GO:0005001)

transmembrane receptor protein phosphatase activity (GO:0019198)

protein phosphatase 2B binding (GO:0030346)

primary miRNA binding (GO:0070878)

1-phosphatidylinositol-4-phosphate 5-kinase activity (GO:0016308)

3',5'-cyclic-nucleotide phosphodiesterase activity (GO:0004114)

phosphatidic acid transfer activity (GO:1990050)

oxysterol binding (GO:0008142)

oxidoreductase activity, acting on metal

ions, oxygen as acceptor (GO:0016721)

group of donors, oxygen as acceptor

(GO:0016670)

nitric-oxide synthase binding (GO:0050998)

histone threonine kinase activity

(GO:0035184)

ferroxidase activity (GO:0004322)

ferric iron binding (GO:0008199)

3',5'-cyclic-GMP phosphodiesterase activity

(GO:0047555)

0

2

4

Enrichment log(combined score)