

# GO\_Biological\_Process\_2021

Term

endothelial cell apoptotic process  
(GO:0072577)

transmembrane receptor protein tyrosine  
phosphatase signaling pathway (GO:0007185)

response to granulocyte macrophage  
colony-stimulating factor (GO:0097012)

chondrocyte development (GO:0002063)

cellular response to granulocyte macrophage  
colony-stimulating factor stimulus  
(GO:0097011)

regulation of transforming growth factor  
beta2 production (GO:0032909)

pharyngeal arch artery morphogenesis  
(GO:0061626)

regulation of Cdc42 protein signal  
transduction (GO:0032489)

negative regulation of systemic arterial  
blood pressure (GO:0003085)

epithelial cell apoptotic process  
(GO:1904019)

cellular response to leptin stimulus  
(GO:0044320)

0

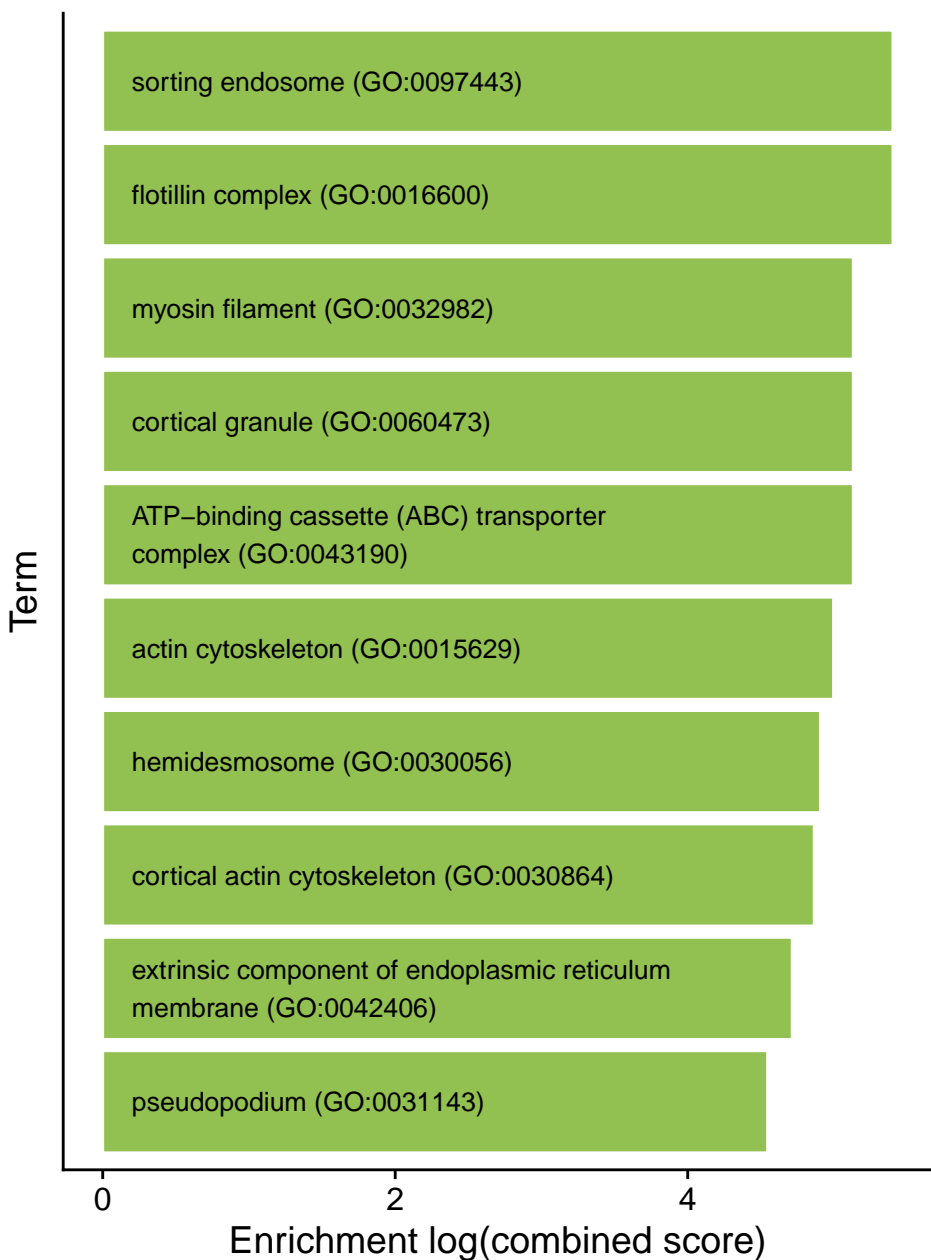
2

4

6

Enrichment log(combined score)

# GO\_Cellular\_Component\_2021



# GO\_Molecular\_Function\_2021

Term

death receptor binding (GO:0005123)

tumor necrosis factor binding (GO:0043120)

phosphatidylcholine floppase activity  
(GO:0090554)

oxo-acid-lyase activity (GO:0016833)  
oxidoreductase activity, acting on a sulfur

group of donors, disulfide as acceptor  
(GO:0016671)

interleukin-1 receptor activity (GO:0004908)

apolipoprotein A-I binding (GO:0034186)

transforming growth factor beta receptor  
binding (GO:0005160)

solute:cation symporter activity  
(GO:0015294)

primary miRNA binding (GO:0070878)  
glucuronosyl-N-acetylgalactosaminyl-proteoglycan

4-beta-N-acetylgalactosaminyltransferase  
activity (GO:0047238)

dihydropyrimidinase activity (GO:0004157)

cyclic nucleotide-dependent protein kinase  
activity (GO:0004690)

alpha-adrenergic receptor activity  
(GO:0004936)

BMP receptor activity (GO:0098821)

BH3 domain binding (GO:0051434)

BH domain binding (GO:0051400)

0

2

4

Enrichment log(combined score)