

GO_Biological_Process_2021

Term

membrane raft assembly (GO:0001765)

cAMP biosynthetic process (GO:0006171)

positive regulation of vesicle fusion
(GO:0031340)

cAMP metabolic process (GO:0046058)

membrane raft organization (GO:0031579)

skeletal muscle tissue regeneration
(GO:0043403)

negative regulation of phospholipase
activity (GO:0010519)

negative regulation of B cell apoptotic
process (GO:0002903)

activation of protein kinase A activity
(GO:0034199)

response to forskolin (GO:1904321)

regulation of vesicle fusion (GO:0031338)

cellular response to forskolin (GO:1904322)

bone cell development (GO:0098751)

0

2

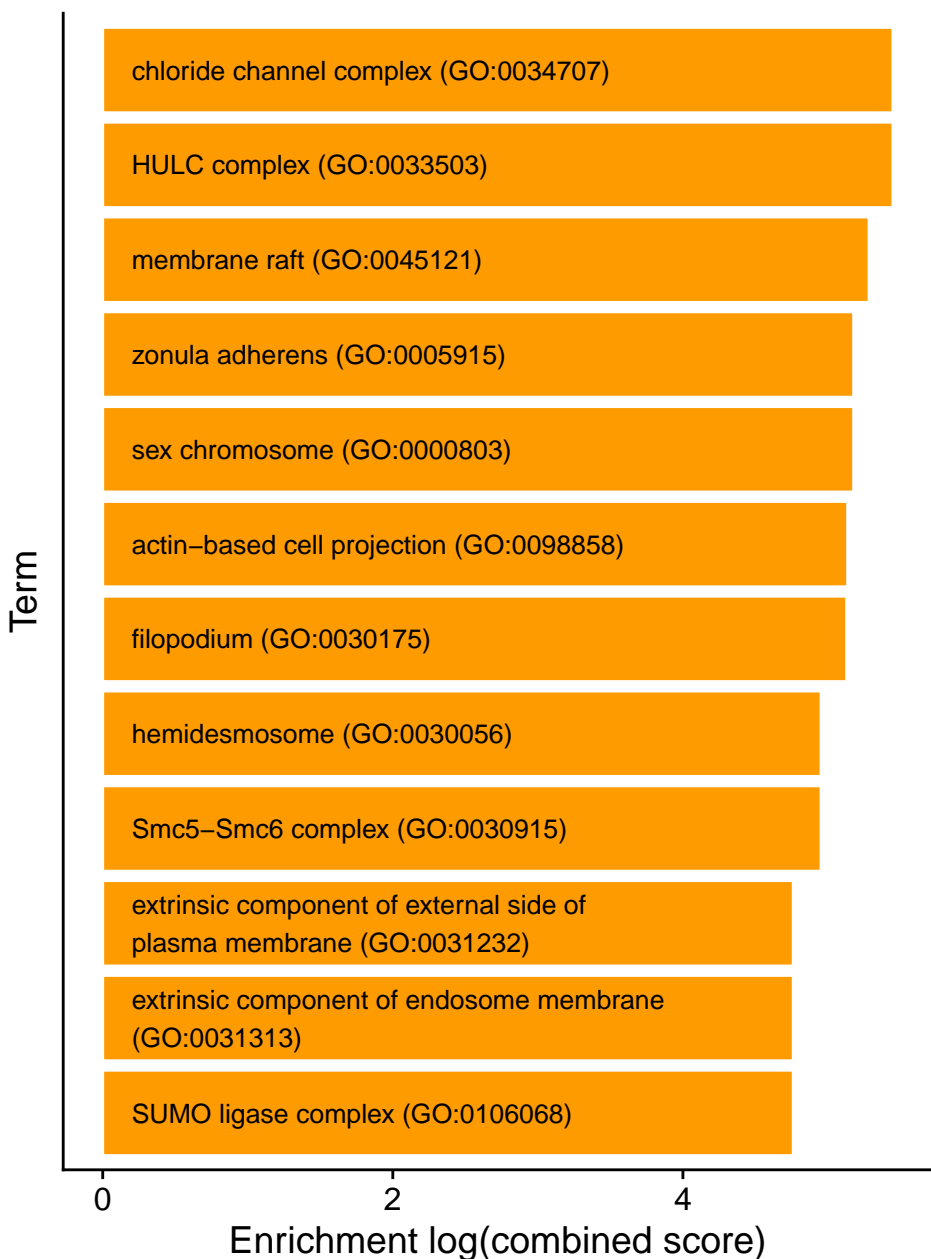
4

6

8

Enrichment log(combined score)

GO_Cellular_Component_2021



GO_Molecular_Function_2021

Term

phospholipase inhibitor activity
(GO:0004859)

adenylate cyclase binding (GO:0008179)

voltage-gated calcium channel activity
involved in cardiac muscle cell action
potential (GO:0086007)

nerve growth factor binding (GO:0048406)

hydroxymethyl-, formyl- and related
transferase activity (GO:0016742)

fibroblast growth factor-activated receptor
activity (GO:0005007)

cytoskeleton-nuclear membrane anchor
activity (GO:0140444)

calmodulin-dependent protein phosphatase
activity (GO:0033192)

cadherin binding involved in cell-cell
adhesion (GO:0098641)

protein phosphatase 2B binding (GO:0030346)

non-membrane spanning protein tyrosine
phosphatase activity (GO:0004726)

dopamine receptor binding (GO:0050780)

alpha-N-acetylneuraminate
alpha-2,8-sialyltransferase activity
(GO:0003828)

0

2

4

6

Enrichment log(combined score)