

GO_Biological_Process_2021

Term

positive regulation of B cell
differentiation (GO:0045579)

regulation of triglyceride metabolic process
(GO:0090207)

regulation of glial cell migration
(GO:1903975)

positive regulation of CD8-positive,
alpha-beta T cell differentiation
(GO:0043378)

B cell homeostasis (GO:0001782)

negative regulation of myeloid leukocyte
mediated immunity (GO:0002887)

substrate-dependent cell migration, cell
extension (GO:0006930)

regulation of myosin-light-chain-phosphatase
activity (GO:0035507)

regulation of microglial cell migration
(GO:1904139)

regulation of CD8-positive, alpha-beta T
cell differentiation (GO:0043376)

positive regulation of microtubule
nucleation (GO:0090063)

epithelial cell differentiation involved in
kidney development (GO:0035850)

0

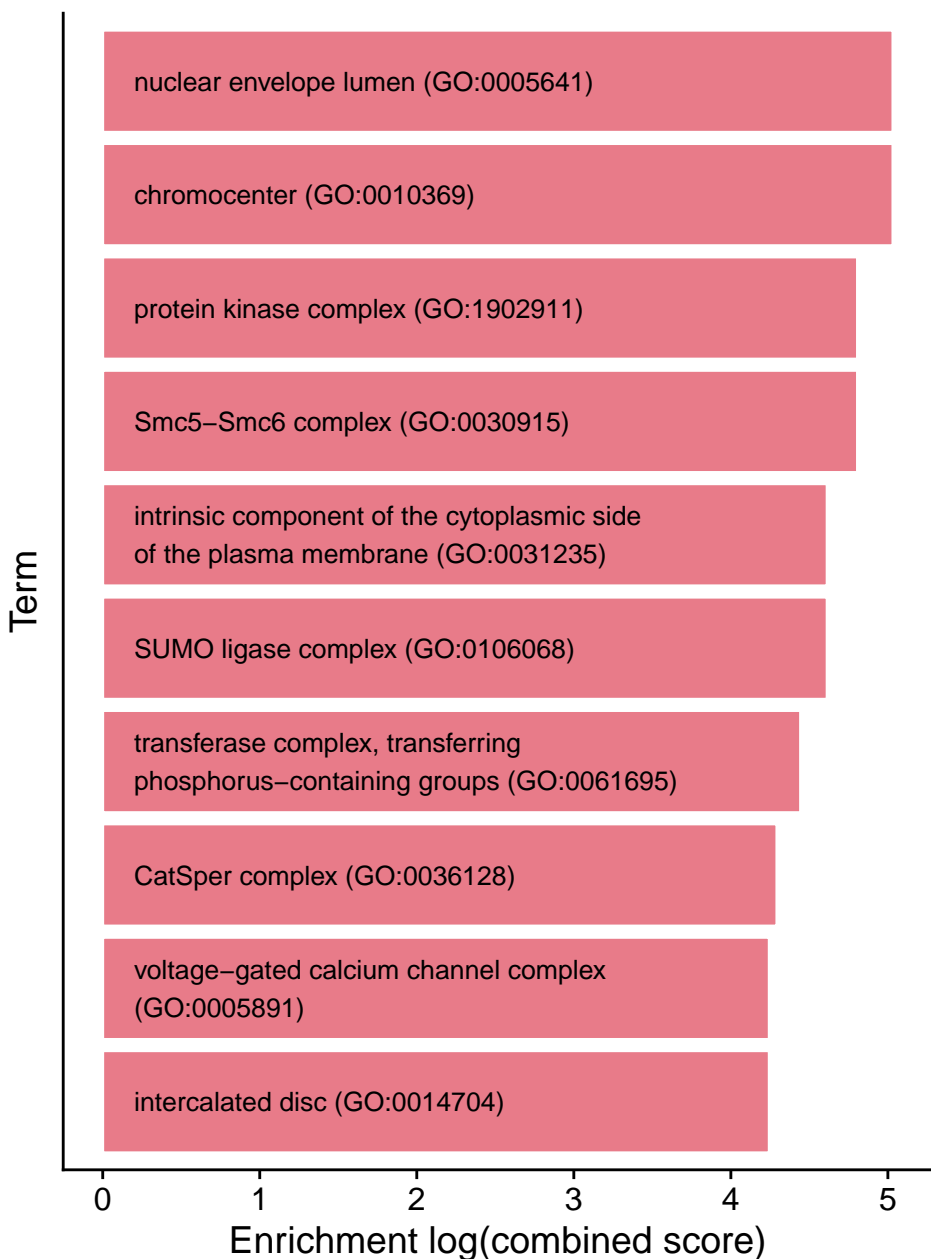
2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021



GO_Molecular_Function_2021

Term

phospholipase binding (GO:0043274)

nucleotide diphosphatase activity
(GO:0004551)

tumor necrosis factor binding (GO:0043120)

protein tyrosine kinase inhibitor activity
(GO:0030292)

phosphodiesterase I activity (GO:0004528)

endopolyphosphatase activity (GO:0000298)

diphosphoinositol–polyphosphate
diphosphatase activity (GO:0008486)

calmodulin–dependent protein phosphatase
activity (GO:0033192)

bis(5'–adenosyl)–pentaphosphatase activity
(GO:0034432)

bis(5'–adenosyl)–hexaphosphatase activity
(GO:0034431)

C–X3–C chemokine binding (GO:0019960)

C–X–C chemokine binding (GO:0019958)

0

2

4

Enrichment log(combined score)