

GO_Biological_Process_2021

Term

negative regulation of smooth muscle
contraction (GO:0045986)

negative regulation of response to reactive
oxygen species (GO:1901032)

regulation of vascular associated smooth
muscle contraction (GO:0003056)

regulation of DNA topoisomerase
(ATP-hydrolyzing) activity (GO:2000371)

positive regulation of transcription from
RNA polymerase II promoter in response to
hypoxia (GO:0061419)

positive regulation of metalloproteinase
activity (GO:1905050)

positive regulation of DNA topoisomerase
(ATP-hydrolyzing) activity (GO:2000373)

regulation of hydrogen peroxide-induced cell
death (GO:1903205)

Rap protein signal transduction (GO:0032486)

response to manganese ion (GO:0010042)

presynaptic active zone organization
(GO:1990709)

0

2

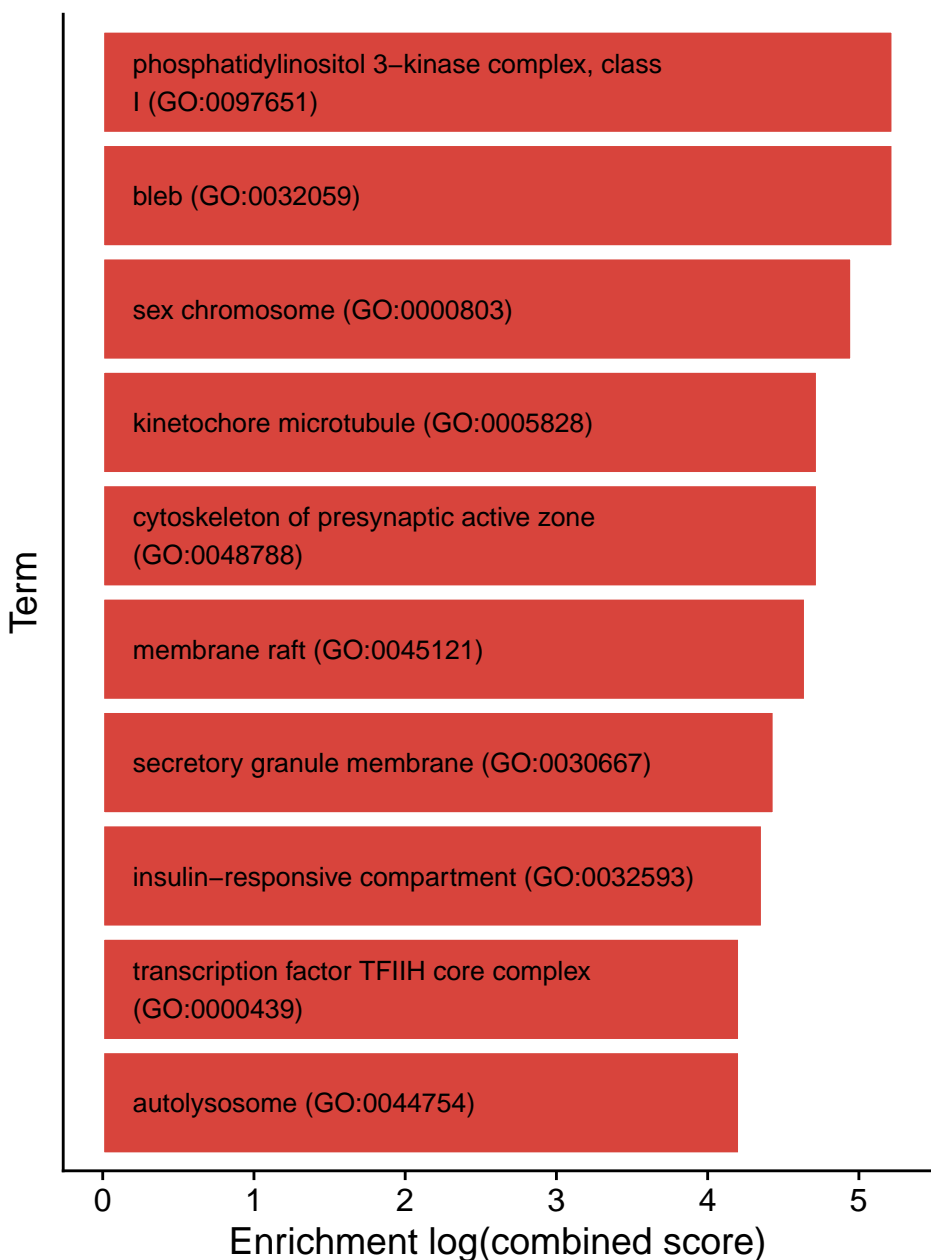
4

6

8

Enrichment log(combined score)

GO_Cellular_Component_2021



GO_Molecular_Function_2021

Term

tau-protein kinase activity (GO:0050321)

3',5'-cyclic-AMP phosphodiesterase activity
(GO:0004115)

histone demethylase activity (GO:0032452)

tumor necrosis factor binding (GO:0043120)

syndecan binding (GO:0045545)

solute:proton antiporter activity
(GO:0015299)

histone demethylase activity (H3-K27
specific) (GO:0071558)

histone demethylase activity
(H3-trimethyl-K4 specific) (GO:0034647)

ganglioside binding (GO:0035594)

fructose-2,6-bisphosphate 2-phosphatase
activity (GO:0004331)

ErbB-3 class receptor binding (GO:0043125)

ATPase inhibitor activity (GO:0042030)

0

2

4

6

Enrichment log(combined score)