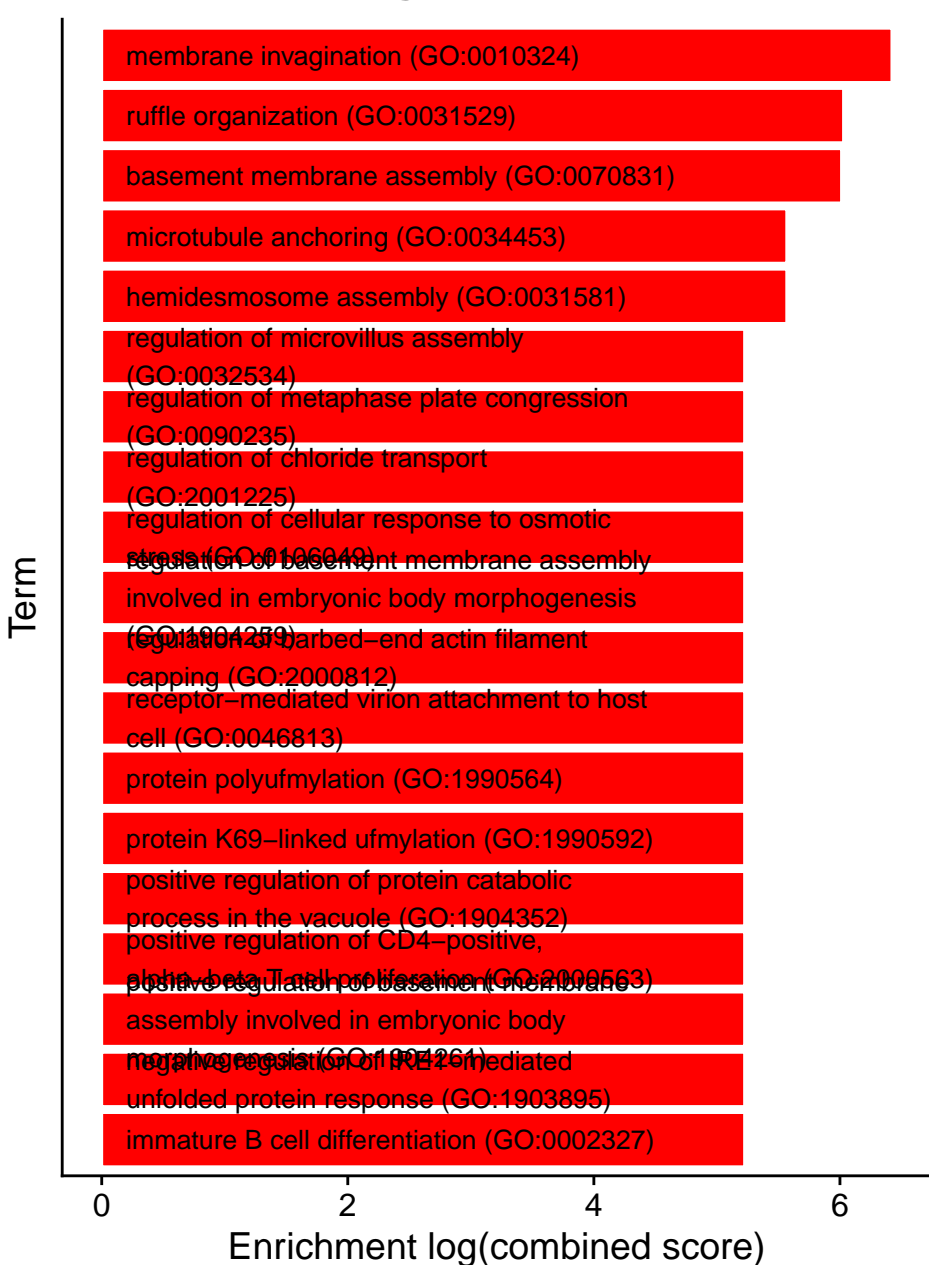


GO_Biological_Process_2021



GO_Cellular_Component_2021

Term

hemidesmosome (GO:0030056)

phosphatidylinositol 3-kinase complex, class I (GO:0097651)

endoplasmic reticulum tubular network membrane (GO:0098826)

Cul4B-RING E3 ubiquitin ligase complex (GO:0031465)

basement membrane (GO:0005604)

uniplex complex (GO:1990246)

dystroglycan complex (GO:0016011)

pinosome (GO:0044352)

macropinosome (GO:0044354)

endoplasmic reticulum tubular network (GO:0071782)

0

2

4

6

Enrichment log(combined score)

GO_Molecular_Function_2021

Term

phosphatidylserine flippase activity
(GO:0140346)

phosphatidylcholine flippase activity
(GO:0140345)

protein carboxyl O-methyltransferase
activity (GO:0051998)

MHC class II protein binding (GO:0042289)

K48-linked polyubiquitin
modification-dependent protein binding
(GO:0036435)

coreceptor activity involved in Wnt
signaling pathway, planar cell polarity
pathway (GO:1904929)

leucine binding (GO:0070728)

hydrolase activity, acting on acid
anhydrides, in phosphorus-containing
anhydrides (GO:0016818)

aminophospholipid flippase activity
(GO:0015247)

1-phosphatidylinositol-4-phosphate 3-kinase
activity (GO:0035005)

0

1

2

3

4

5

Enrichment log(combined score)