

GO_Biological_Process_2021

Term

regulation of basement membrane assembly

involved in embryonic body morphogenesis

(GO:1904259)

reelin-mediated signaling pathway

(GO:0038026)

protein C-linked glycosylation via

tryptophan (GO:0018317)

protein C-linked glycosylation via

2'-alpha-mannosyl-L-tryptophan (GO:0018406)

positive regulation of macrophage

proliferation (GO:0120041)

positive regulation of blood-brain barrier

permeability (GO:1905605)

positive regulation of basement membrane

assembly involved in embryonic body

morphogenesis (GO:1904261)

negative regulation of posttranscriptional

gene silencing (GO:0060149)

negative regulation of interferon-alpha

production (GO:0032687)

negative regulation of gene silencing by RNA

(GO:0060967)

negative regulation of collateral sprouting

(GO:0048671)

melanin biosynthetic process (GO:0042438)

extracellular exosome biogenesis

(GO:0097734)

beta-catenin destruction complex assembly

(GO:1904885)

0

1

2

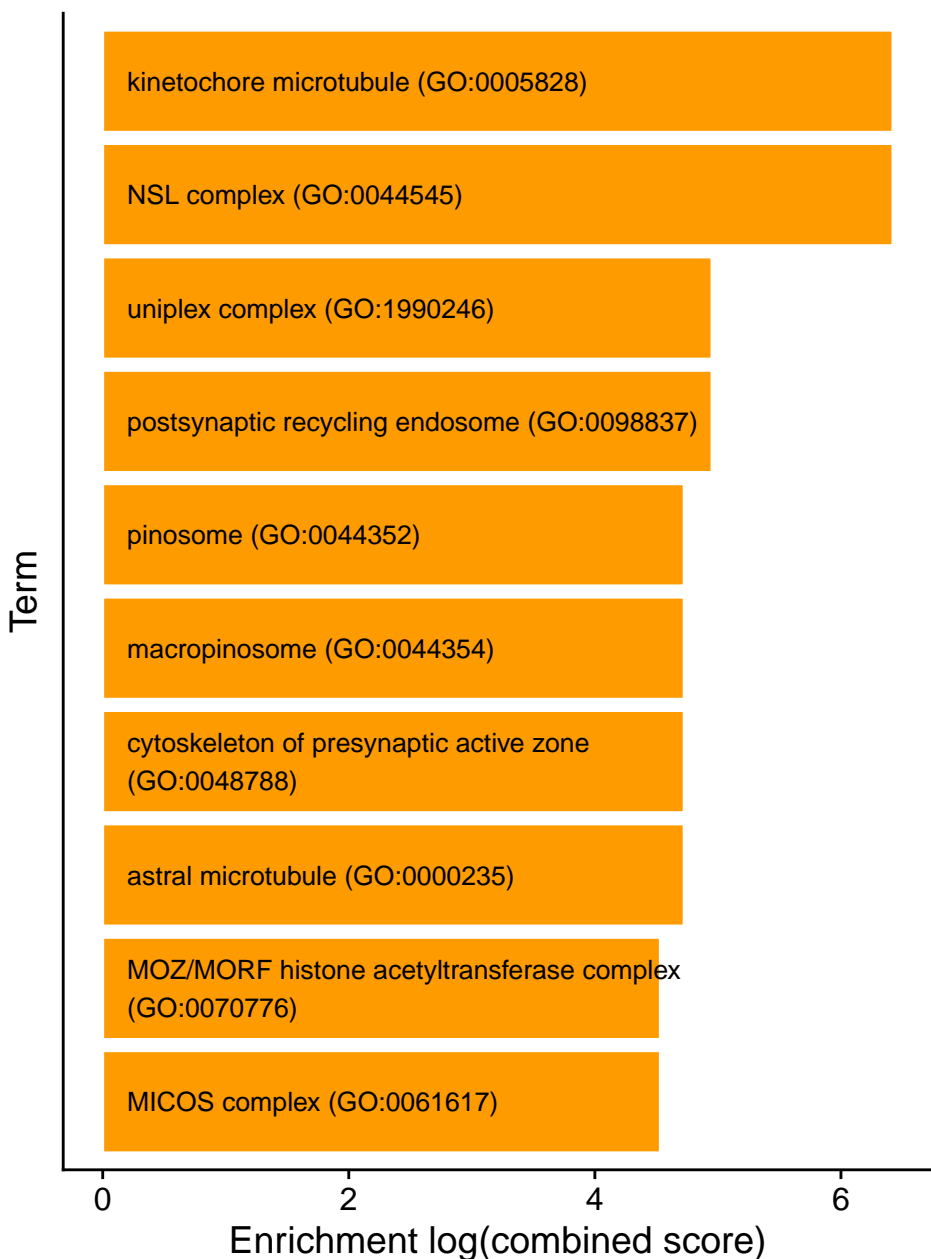
3

4

5

Enrichment log(combined score)

GO_Cellular_Component_2021



GO_Molecular_Function_2021

Term

microtubule plus-end binding (GO:0051010)

poly(G) binding (GO:0034046)

phosphotransferase activity, nitrogenous
group as acceptor (GO:0016775)

nucleoside kinase activity (GO:0019206)

histone acetyltransferase activity (H3-K23
specific) (GO:0043994)

creatine kinase activity (GO:0004111)

CoA carboxylase activity (GO:0016421)

histone acetyltransferase binding
(GO:0035035)

oxysterol binding (GO:0008142)

oxidoreductase activity, acting on a sulfur
group of donors, oxygen as acceptor
(GO:0016670)

leucine binding (GO:0070728)

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

JUN kinase binding (GO:0008432)

3',5'-cyclic-GMP phosphodiesterase activity
(GO:0047555)

0

2

4

6

Enrichment log(combined score)