

GO_Biological_Process_2021

Term

platelet activating factor metabolic process

(GO:0046469)

vesicle fusion with Golgi apparatus

(GO:0048280)

ubiquitin-dependent glycoprotein ERAD

pathway (GO:0097466)

regulation of heart growth (GO:0060420)

gamma-interferon receptor signaling

pathway involved in phagocytosis

regulation of basement membrane assembly

(GO:1905149)

involved in embryonic body morphogenesis

reelin-mediated signaling pathway

(GO:1904259)

protein degradation involved in

positive regulation of histone H3 K27

methylation (GO:0061087)

positive regulation of Golgi to plasma

positive regulation of basement membrane

membrane protein transport (GO:0042998)

assembly involved in embryonic body

morphogenesis (GO:1904864)

phosphatase (GO:0098166)

negative regulation of IRE1-mediated

unfolded protein response (GO:1903895)

mRNA cleavage involved in gene silencing by

miRNA (GO:0035278)

microtubule nucleation by microtubule

organizing center (GO:0051418)

meiotic mismatch repair (GO:0000710)

mannose trimming involved in glycoprotein

ERAD pathway (GO:1904382)

diphosphonucleoside polyphosphate metabolic

process (GO:0071543)

diadenosine pentaphosphate metabolic process

(GO:1901906)

diadenosine pentaphosphate catabolic process

(GO:1901907)

diadenosine hexaphosphate metabolic process

(GO:1901908)

diadenosine hexaphosphate catabolic process

(GO:1901909)

caveolin-mediated endocytosis (GO:0072584)

caveola assembly (GO:0070836)

adenosine 5'-(hexahydrogen pentaphosphate)

metabolic process (GO:1901910)

adenosine 5'-(hexahydrogen pentaphosphate)

catabolic process (GO:1901911)

0

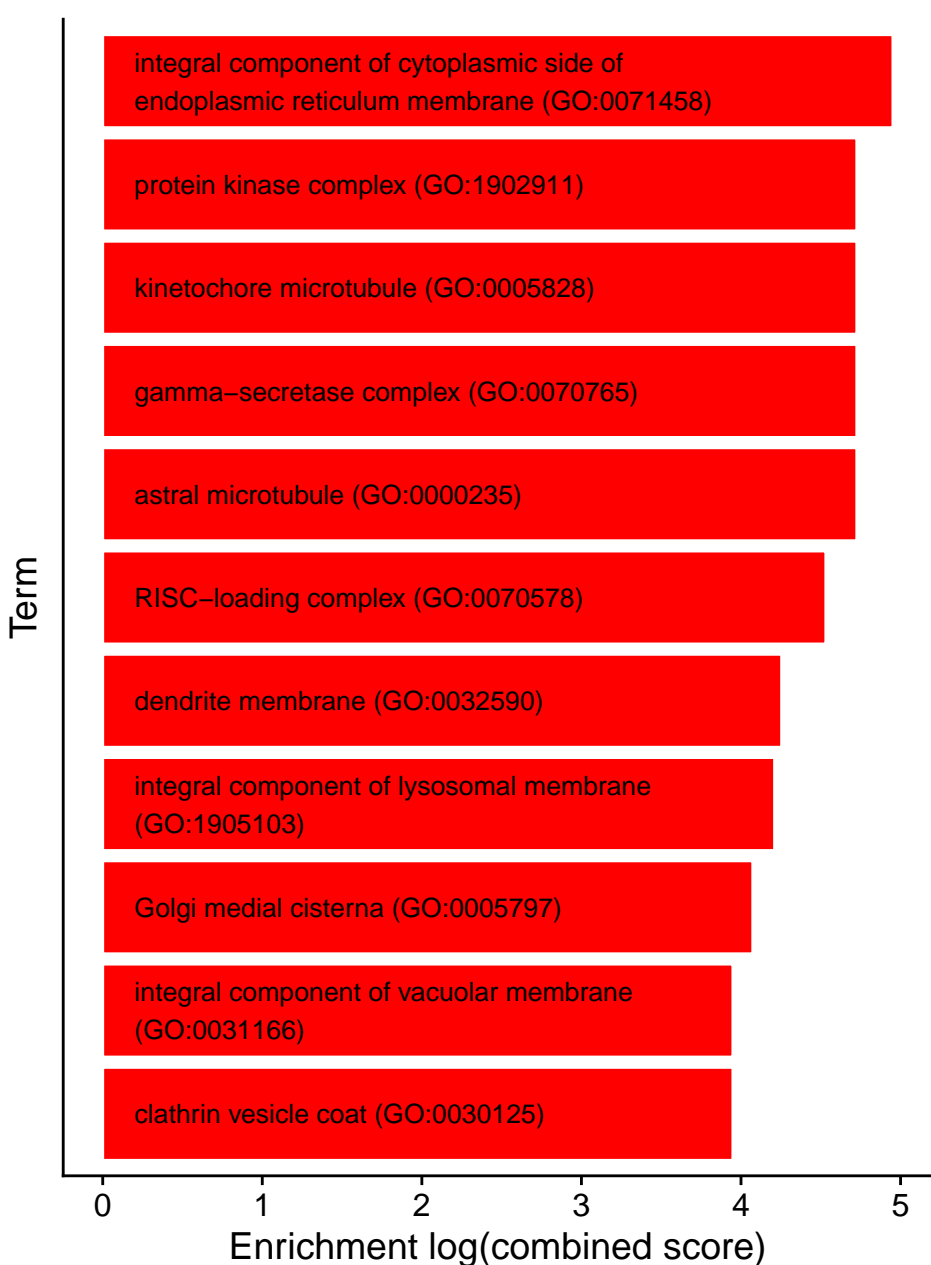
2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021



GO_Molecular_Function_2021

Term

1-acylglycerol-3-phosphate O-acyltransferase
activity (GO:0003841)

protein serine/threonine/tyrosine kinase
activity (GO:0004712)

MAP kinase kinase activity (GO:0004708)

lysophosphatidic acid acyltransferase
activity (GO:0042171)

tumor necrosis factor binding (GO:0043120)

RNA polymerase II C-terminal domain
phosphoserine binding (GO:1990269)

protein-arginine omega-N
monomethyltransferase activity (GO:0035241)

NAADP-sensitive calcium-release channel
activity (GO:0072345)

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

endopolyphosphatase activity (GO:0000298)

diphosphoinositol-polyphosphate
diphosphatase activity (GO:0008486)

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

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

0

2

4

Enrichment log(combined score)