

GO_Biological_Process_2021

Term

astral microtubule organization (GO:0030953)

regulation of basement membrane assembly
involved in embryonic body morphogenesis
(GO:1904259)

protein retention in Golgi apparatus
(GO:0045053)

positive regulation of basement membrane
assembly involved in embryonic body
morphogenesis (GO:1904261)

membrane depolarization during SA node cell
action potential (GO:0086046)

beta-catenin destruction complex assembly
(GO:1904885)

positive regulation of sodium ion
transmembrane transport (GO:1902307)

microtubule anchoring (GO:0034453)

positive regulation of protein localization
to centrosome (GO:1904781)

SA node cell action potential (GO:0086015)

0

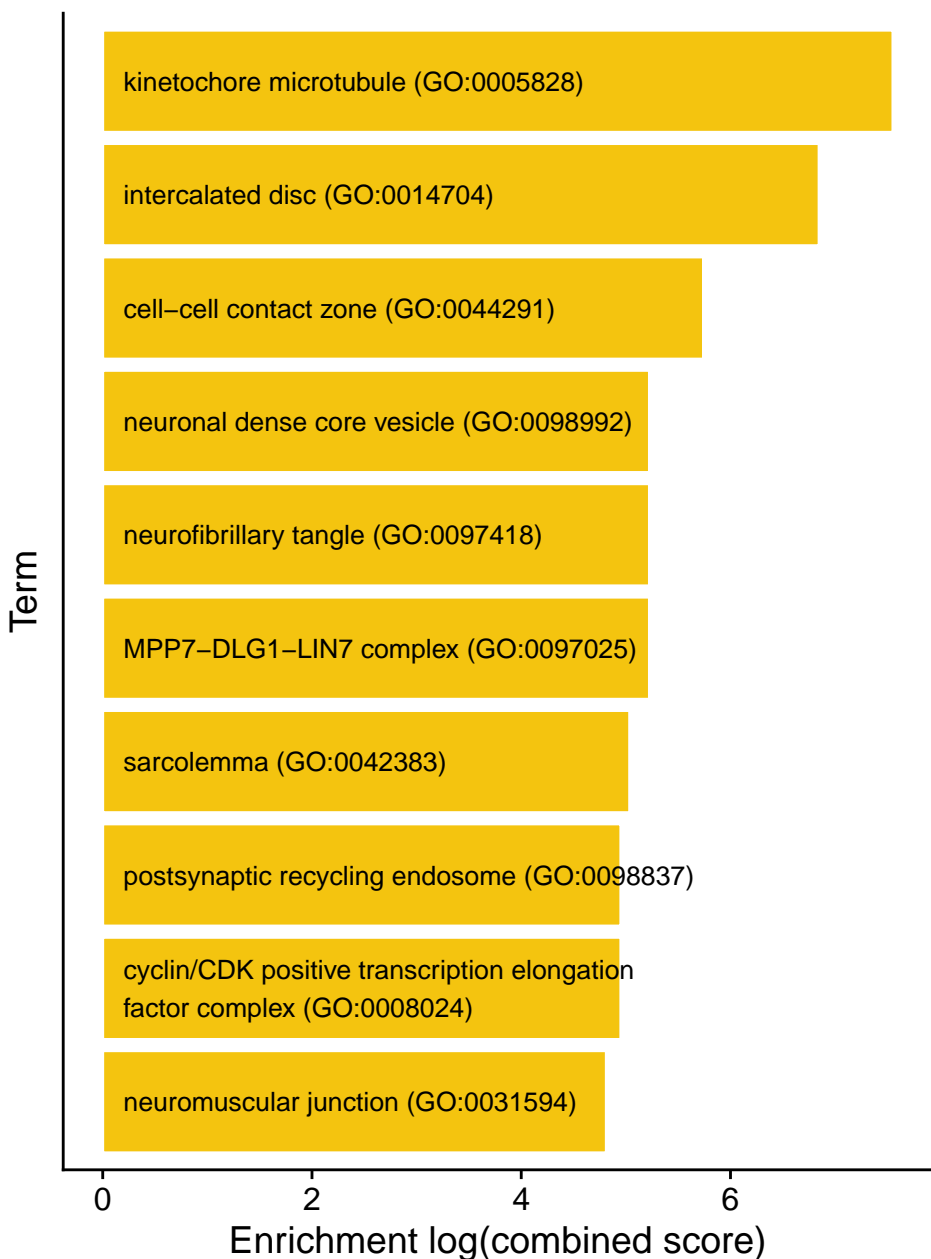
2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021



GO_Molecular_Function_2021

Term

microtubule plus-end binding (GO:0051010)

guanylate kinase activity (GO:0004385)

dynactin binding (GO:0034452)

GTPase binding (GO:0051020)

voltage-gated calcium channel activity
involved in cardiac muscle cell action
potential (GO:0086007)

kainate selective glutamate receptor
activity (GO:0015277)

TPR domain binding (GO:0030911)

C-X-C chemokine binding (GO:0019958)

MHC class II protein complex binding
(GO:0023026)

DNA polymerase binding (GO:0070182)

0

2

4

6

Enrichment log(combined score)