

GO_Biological_Process_2021

Term

regulation of long-term neuronal synaptic
plasticity (GO:0048169)

regulation of chloride transport
(GO:2001225)

cerebellar Purkinje cell differentiation
(GO:0021702)

regulation of anion channel activity
(GO:0010359)

positive regulation of tau-protein kinase
activity (GO:1902949)

long-term synaptic depression (GO:0060292)

regulation of protein dephosphorylation
(GO:0035304)

positive regulation of protein
dephosphorylation (GO:0035307)

regulation of cyclic-nucleotide
phosphodiesterase activity (GO:0051342)

positive regulation of ryanodine-sensitive
calcium-release channel activity
(GO:0060316)

negative regulation of neuron projection
regeneration (GO:0070571)

0

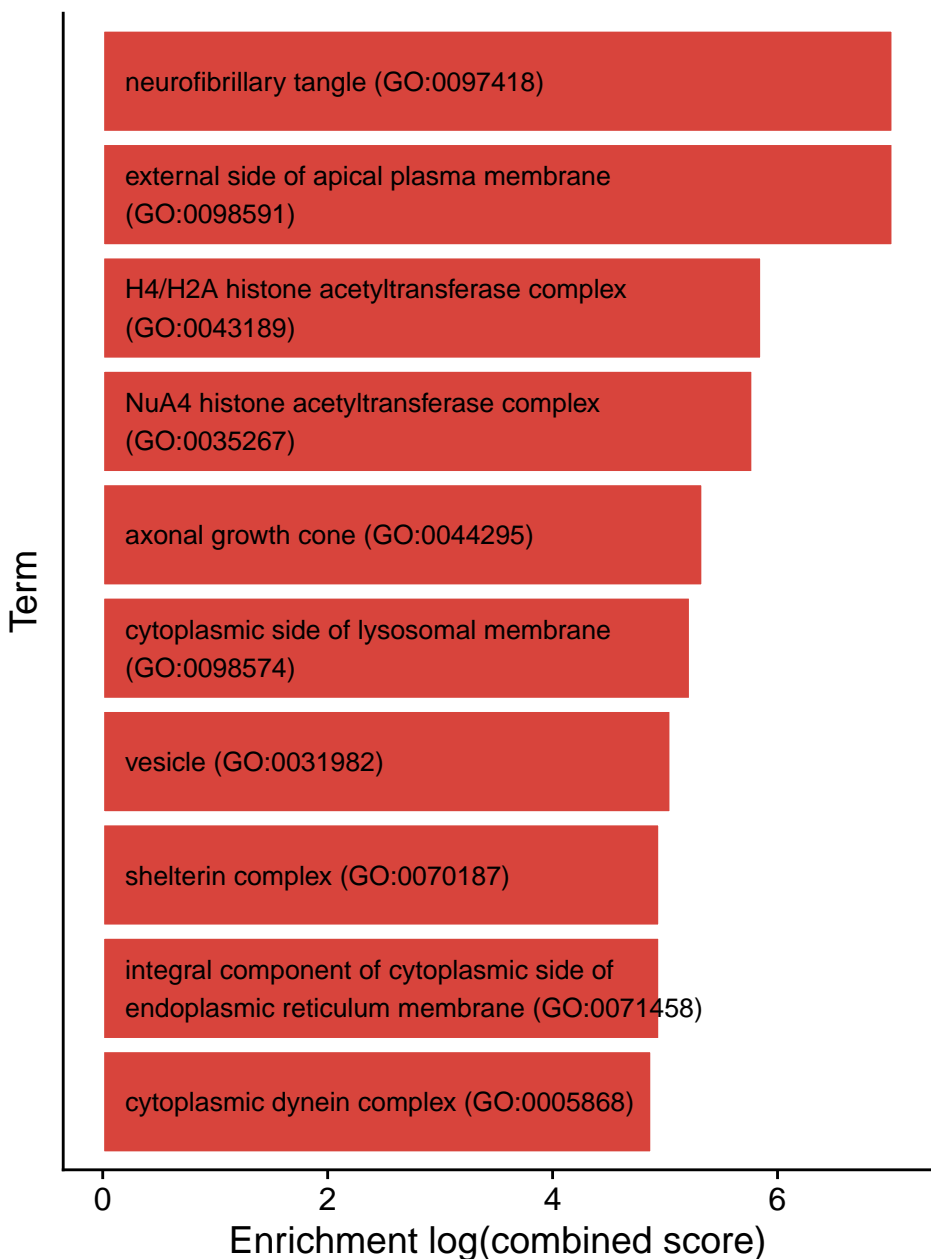
2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021



GO_Molecular_Function_2021

Term

ganglioside binding (GO:0035594)

adenylate cyclase binding (GO:0008179)

disordered domain specific binding
(GO:0097718)

aspartic-type endopeptidase inhibitor
activity (GO:0019828)

MHC class II protein complex binding
(GO:0023026)

phosphoserine residue binding (GO:0050815)

ionotropic glutamate receptor binding
(GO:0035255)

syntaxin-1 binding (GO:0017075)

titin binding (GO:0031432)

protein phosphatase activator activity
(GO:0072542)

protein kinase A catalytic subunit binding
(GO:0034236)

RNA 7-methylguanosine cap binding
(GO:0000340)

GDP-dissociation inhibitor activity
(GO:0005092)

0

2

4

6

Enrichment log(combined score)