

# 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

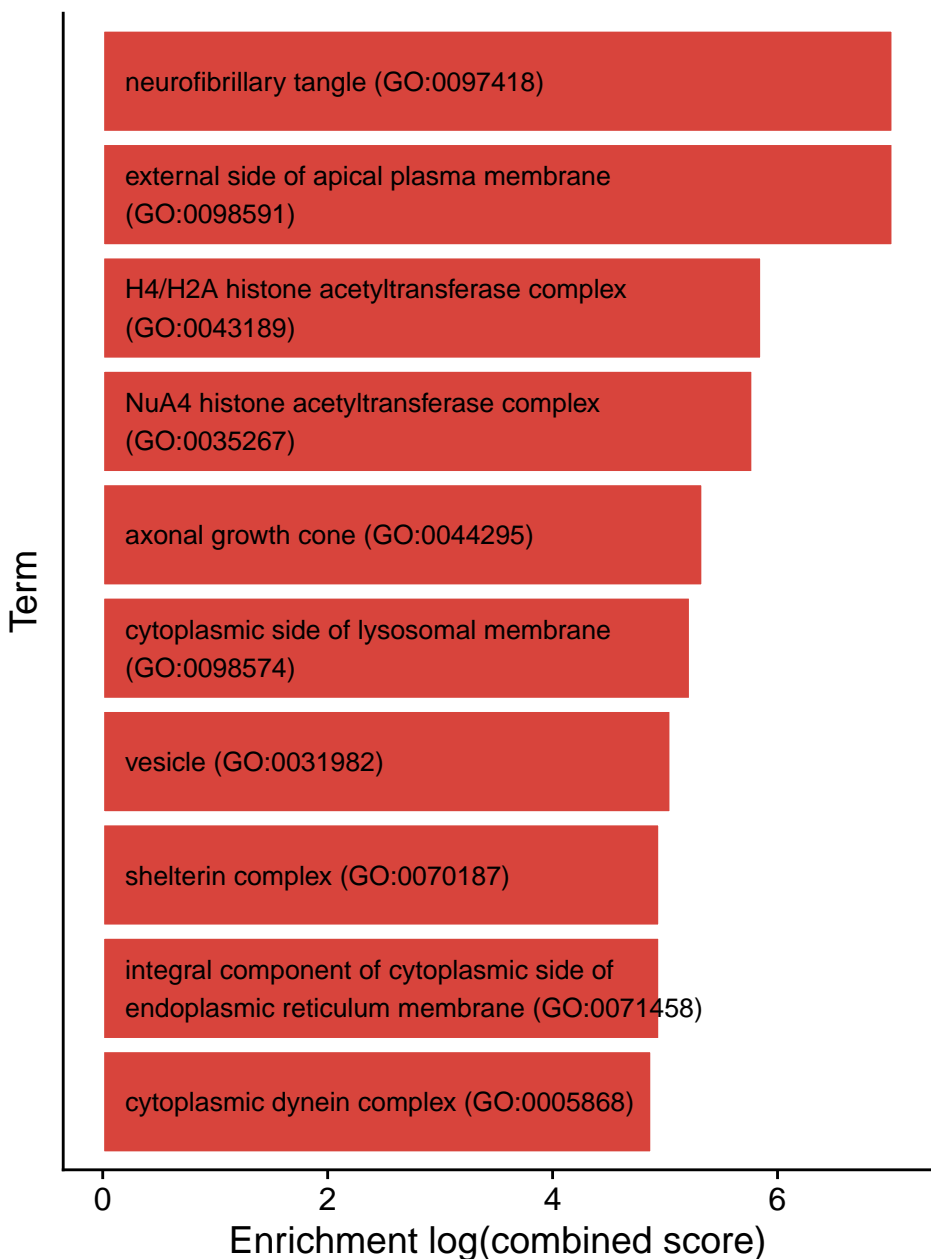
2

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)