

# GO\_Biological\_Process\_2021

Term

regulation of chloride transport  
(GO:2001225)

cerebellar Purkinje cell differentiation  
(GO:0021702)

neuron projection maintenance (GO:1990535)

regulation of anion channel activity  
(GO:0010359)

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

central nervous system neuron development  
(GO:0021954)

glucose catabolic process to pyruvate  
(GO:0061718)

canonical glycolysis (GO:0061621)

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

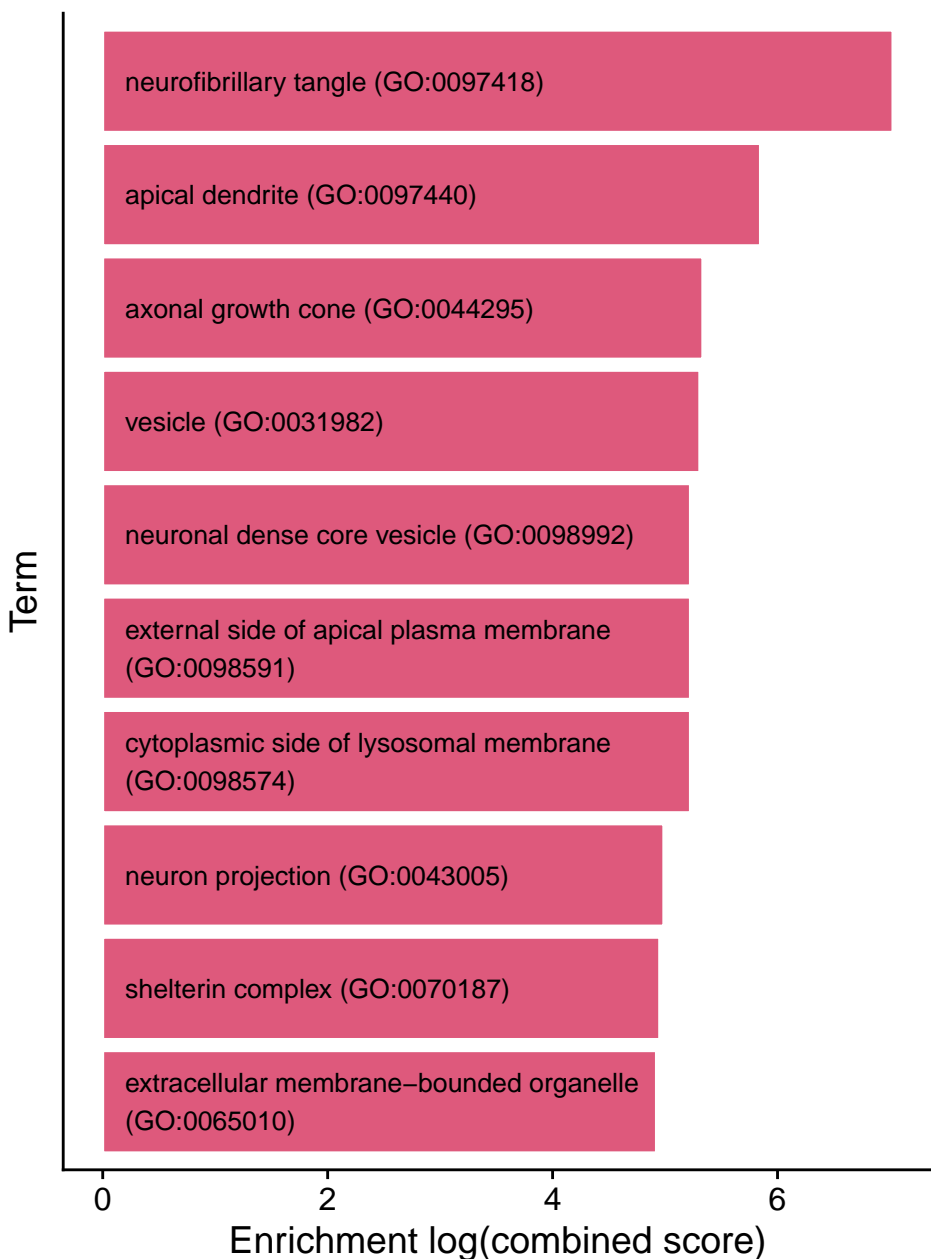
2

4

6

Enrichment log(combined score)

# GO\_Cellular\_Component\_2021



# GO\_Molecular\_Function\_2021

Term

ganglioside binding (GO:0035594)

adenylate cyclase binding (GO:0008179)

aspartic-type endopeptidase inhibitor  
activity (GO:0019828)

disordered domain specific binding  
(GO:0097718)

cuprous ion binding (GO:1903136)

tau protein binding (GO:0048156)

syntaxin-1 binding (GO:0017075)

titin binding (GO:0031432)

protein phosphatase activator activity  
(GO:0072542)

protein kinase A catalytic subunit binding  
(GO:0034236)

GDP-dissociation inhibitor activity  
(GO:0005092)

0

2

4

6

Enrichment log(combined score)