

# GO\_Biological\_Process\_2021

Term

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

negative regulation of amyloid precursor  
protein biosynthetic process (GO:0042985)

negative regulation of amyloid fibril  
formation (GO:1905907)

regulation of amyloid fibril formation  
(GO:1905906)

negative regulation of amyloid-beta  
formation (GO:1902430)

regulation of cellular component biogenesis  
(GO:0044087)

negative regulation of glycoprotein  
biosynthetic process (GO:0010561)

negative regulation of amyloid precursor  
protein catabolic process (GO:1902992)

regulation of tau-protein kinase activity  
(GO:1902947)

regulation of amyloid precursor protein  
biosynthetic process (GO:0042984)

positive regulation of phospholipid  
transport (GO:2001140)

0

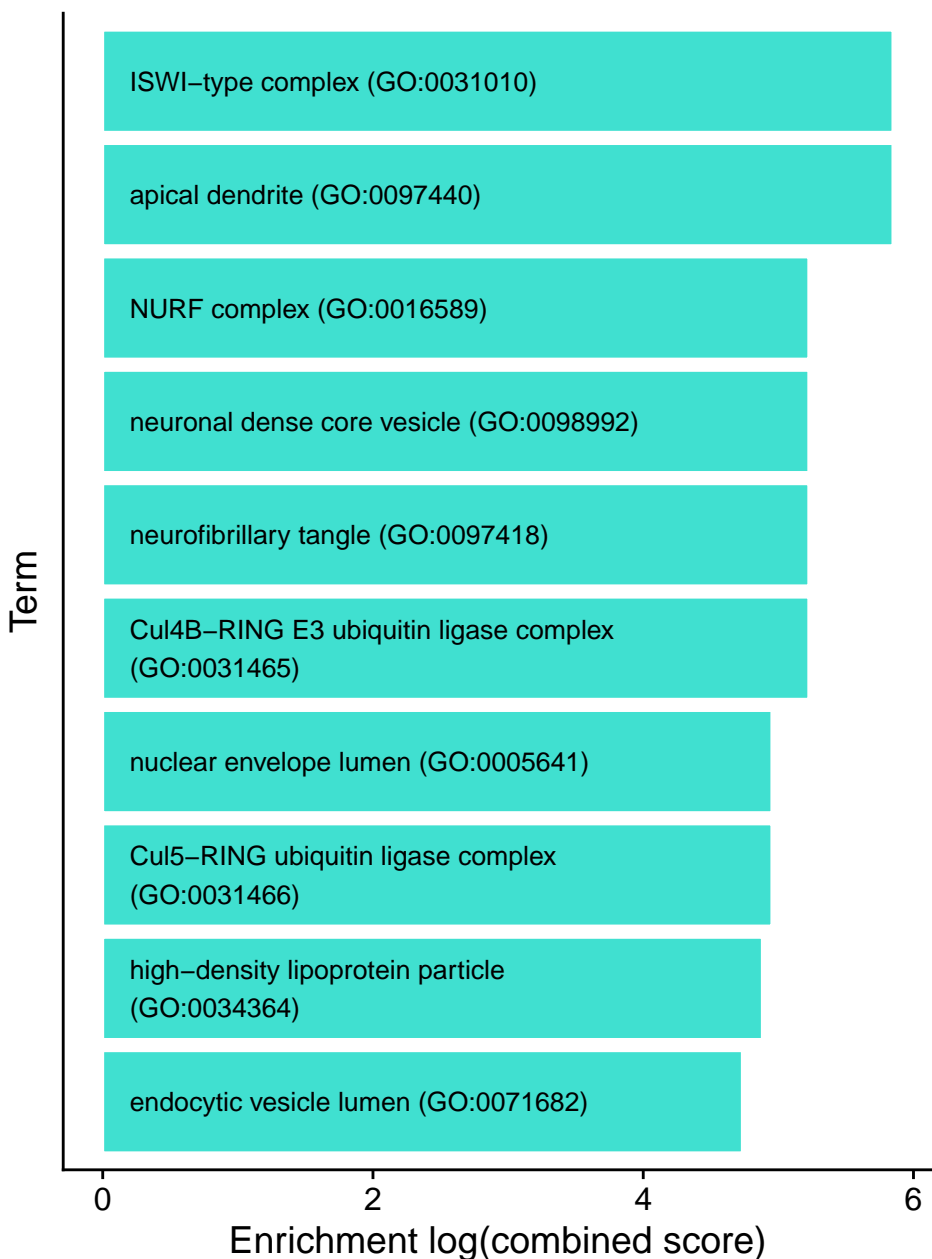
2

4

6

Enrichment log(combined score)

# GO\_Cellular\_Component\_2021



# GO\_Molecular\_Function\_2021

Term

TPR domain binding (GO:0030911)

RNA–DNA hybrid ribonuclease activity  
(GO:0004523)

phosphodiesterase I activity (GO:0004528)

NEDD8 transferase activity (GO:0019788)

glutathione transmembrane transporter  
activity (GO:0034634)

gap junction channel activity involved in  
cardiac conduction electrical coupling  
(GO:0086075)

MHC class II protein complex binding  
(GO:0023026)

tripeptide transmembrane transporter  
activity (GO:0042937)

phosphatidylcholine–sterol O–acyltransferase  
activator activity (GO:0060228)

gap junction hemi–channel activity  
(GO:0055077)

gap junction channel activity involved in  
cell communication by electrical coupling  
(GO:1903763)

dihydropyrimidinase activity (GO:0004157)

collagen receptor activity (GO:0038064)

C3HC4–type RING finger domain binding  
(GO:0055131)

arylesterase activity (GO:0004064)

apolipoprotein receptor binding (GO:0034190)

0

1

2

3

4

5

Enrichment log(combined score)