

GO_Biological_Process_2021

Term

regulation of protein depolymerization
(GO:1901879)

very long-chain fatty acid biosynthetic
process (GO:0042761)

fatty acid elongation (GO:0030497)

mitochondrial electron transport, NADH to
ubiquinone (GO:0006120)

aerobic electron transport chain
(GO:0019646)

regulation of microtubule polymerization or
depolymerization (GO:0031110)

mitochondrial ATP synthesis coupled electron
transport (GO:0042775)

positive regulation of viral release from
host cell (GO:1902188)

actin filament depolymerization (GO:0030042)

regulation of supramolecular fiber
organization (GO:1902903)

0

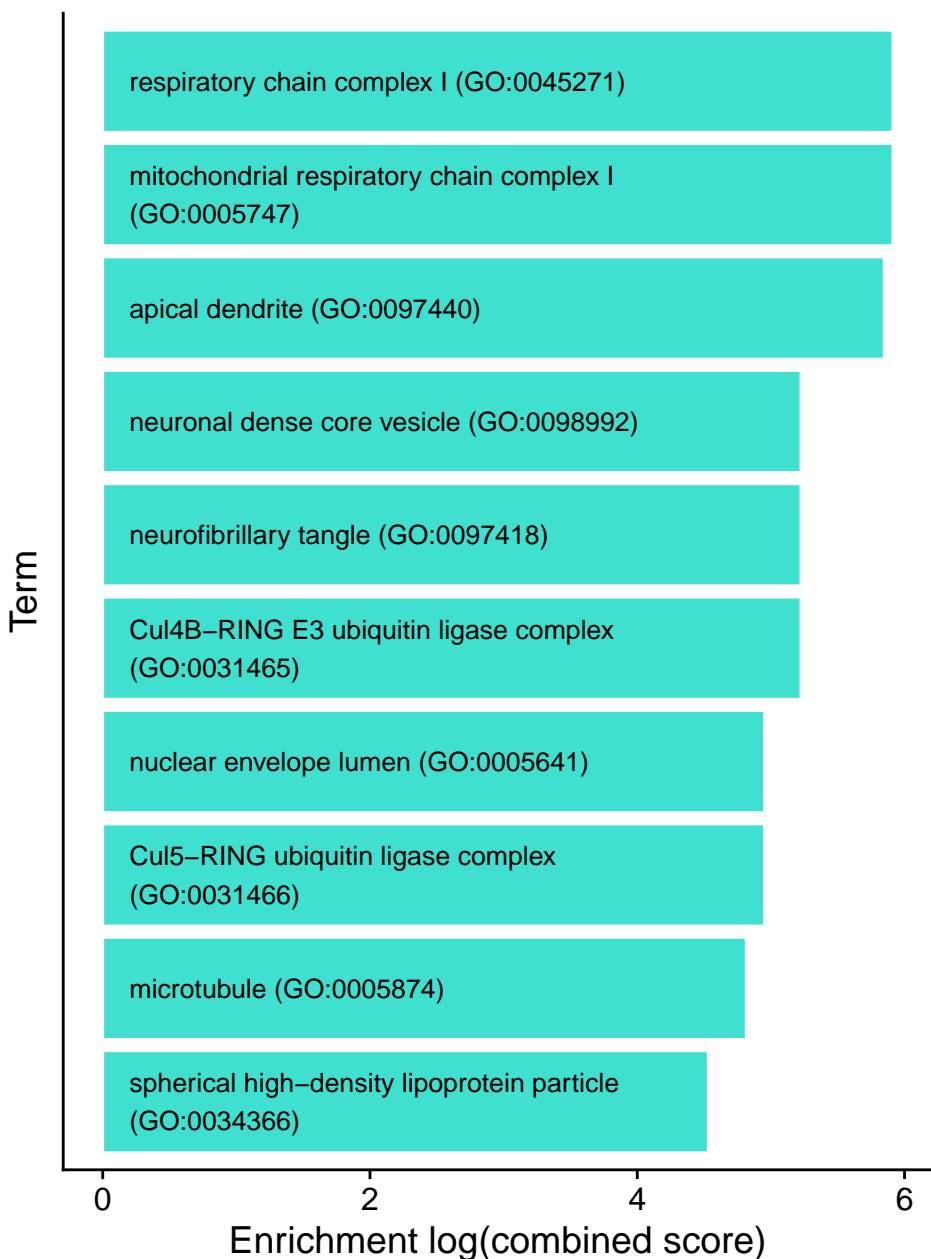
2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021



GO_Molecular_Function_2021

Term

RAGE receptor binding (GO:0050786)

oxidoreduction-driven active transmembrane
transporter activity (GO:0015453)

NADH dehydrogenase (ubiquinone) activity
(GO:0008137)

NADH dehydrogenase (quinone) activity
(GO:0050136)

tau protein binding (GO:0048156)

cytochrome-c oxidase activity (GO:0004129)

thiolester hydrolase activity (GO:0016790)

protein geranylgeranyltransferase activity
(GO:0004661)

NEDD8 transferase activity (GO:0019788)

leukotriene-C4 synthase activity
(GO:0004464)

isocitrate dehydrogenase activity
(GO:0004448)

0

2

4

6

Enrichment log(combined score)