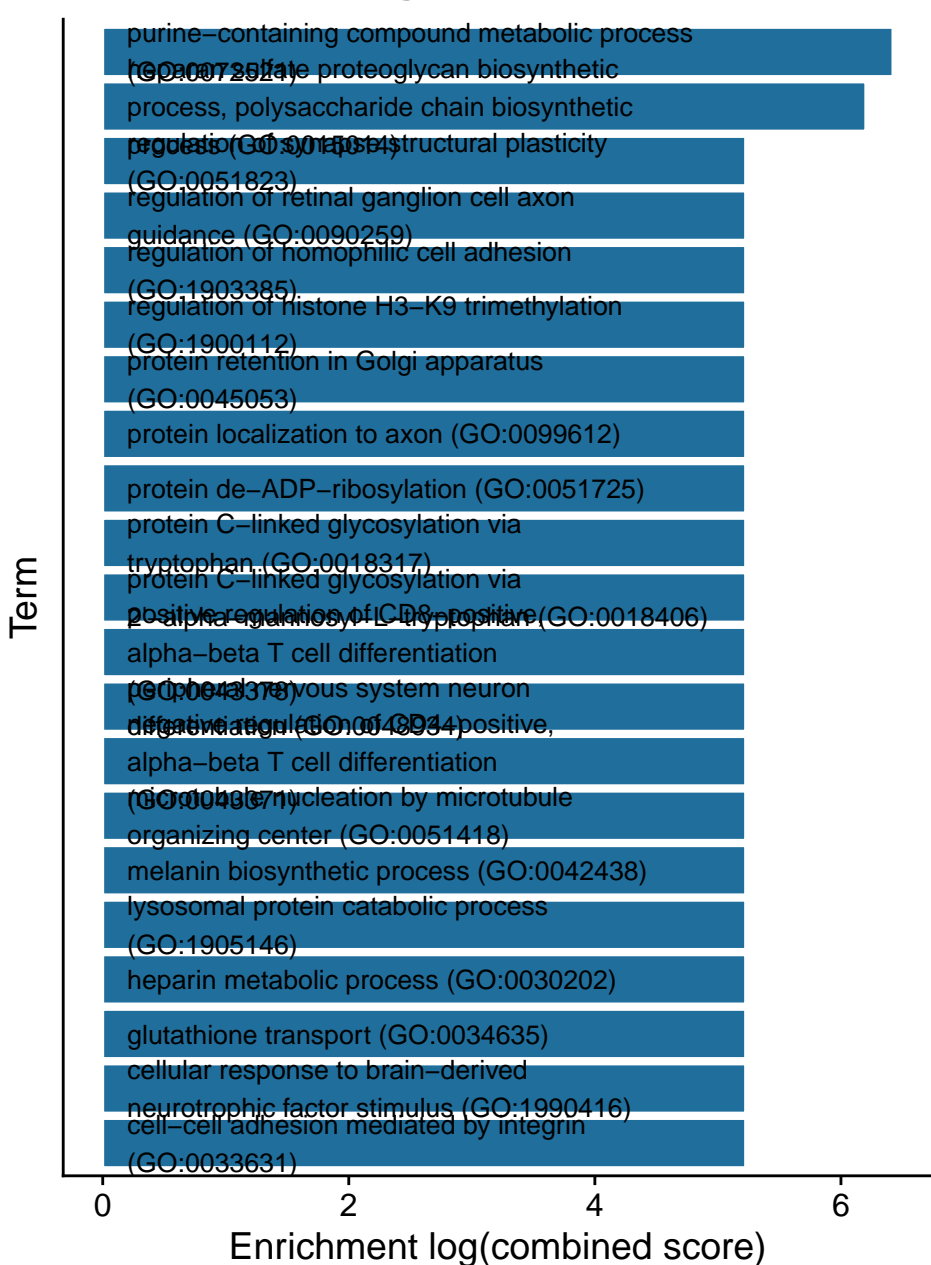


GO_Biological_Process_2021



GO_Cellular_Component_2021

Term

neuronal dense core vesicle (GO:0098992)

neurofibrillary tangle (GO:0097418)

cyclin/CDK positive transcription elongation
factor complex (GO:0008024)

cortical endoplasmic reticulum (GO:0032541)

PTW/PP1 phosphatase complex (GO:0072357)

NSL complex (GO:0044545)

protein serine/threonine phosphatase complex
(GO:0008287)

cyclin-dependent protein kinase holoenzyme
complex (GO:0000307)

nuclear inclusion body (GO:0042405)

STAGA complex (GO:0030914)

0

1

2

3

4

5

Enrichment log(combined score)

GO_Molecular_Function_2021

Term

guanylate kinase activity (GO:0004385)

oxo-acid-lyase activity (GO:0016833)

transmembrane receptor protein tyrosine

phosphatase activity (GO:0005001)

transmembrane receptor protein phosphatase

activity (GO:0019198)

myosin light chain binding (GO:0032027)

glucuronosyl-N-acetylgalactosaminyl-proteoglycan

4-beta-N-acetylgalactosaminyltransferase

activity (GO:0047238)

chromo shadow domain binding (GO:0070087)

arylesterase activity (GO:0004064)

nucleoside monophosphate kinase activity

(GO:0050145)

transmembrane receptor protein tyrosine

kinase activator activity (GO:0030297)

sphingosine-1-phosphate phosphatase activity

(GO:0042392)

oxysterol binding (GO:0008142)

oxidoreductase activity, acting on a sulfur

group of donors, oxygen as acceptor

hydrolase activity, acting on acid

(GO:0016570)

anhydrides, in phosphorus-containing

anhydrides (GO:0016818)

SUMO-specific protease activity (GO:0016929)

JUN kinase binding (GO:0008432)

3',5'-cyclic-GMP phosphodiesterase activity

(GO:0047555)

0

2

4

6

Enrichment log(combined score)