

GO_Biological_Process_2021

Term

proteasomal ubiquitin-independent protein
catabolic process (GO:0010499)

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

regulation of mRNA stability (GO:0043488)

nucleic acid metabolic process (GO:0090304)

rRNA-containing ribonucleoprotein complex
export from nucleus (GO:0071428)

ribosomal small subunit export from nucleus
(GO:0000056)

ribosomal large subunit export from nucleus
(GO:0000055)

protein localization to cytoplasmic stress
granule (GO:1903608)

regulation of chromosome organization
(GO:0033044)

regulation of mRNA catabolic process
(GO:0061013)

0

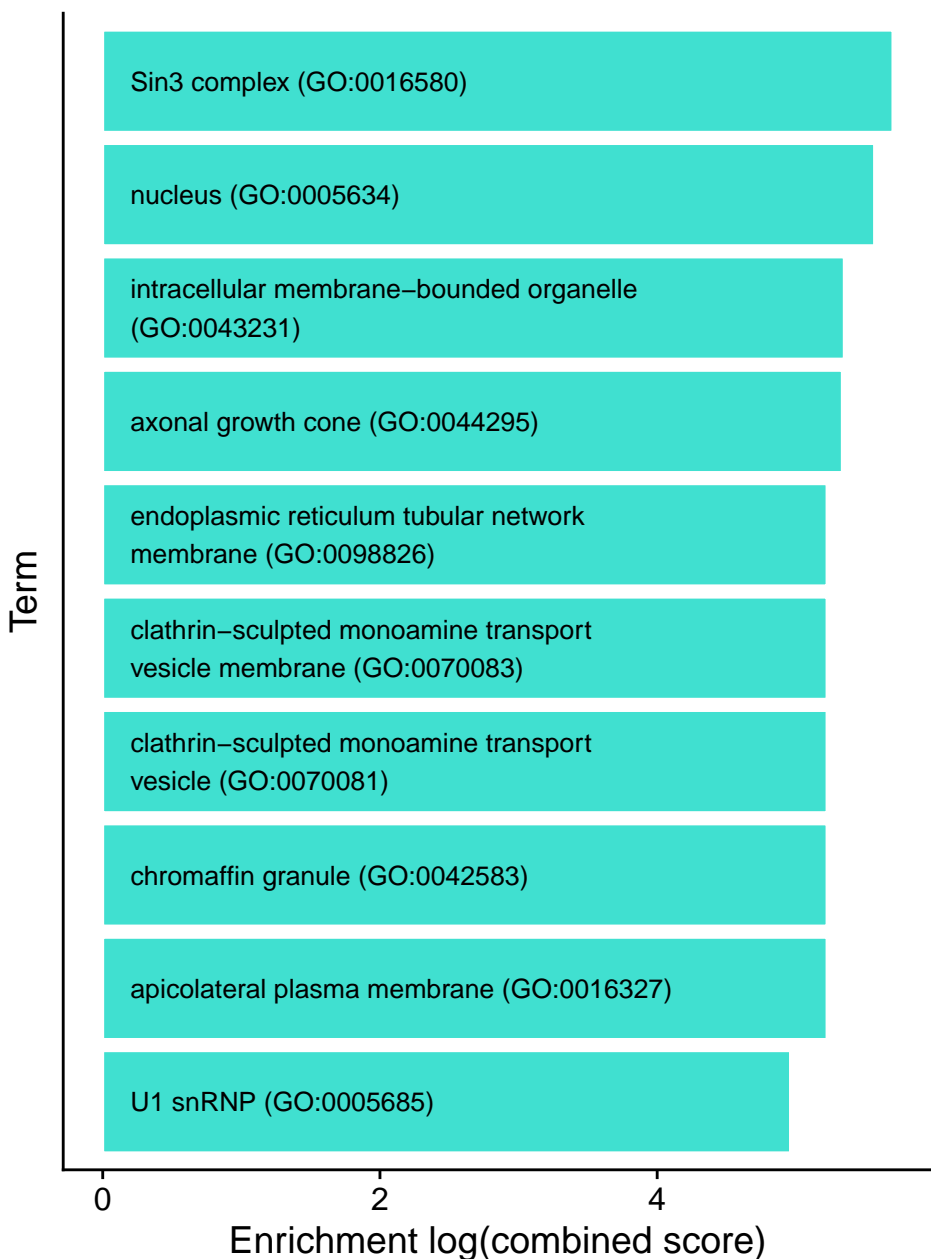
2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021



GO_Molecular_Function_2021

Term

TPR domain binding (GO:0030911)

G-rich strand telomeric DNA binding
(GO:0098505)

single-stranded telomeric DNA binding
(GO:0043047)

N6-methyladenosine-containing RNA binding
(GO:1990247)

telomeric DNA binding (GO:0042162)

MHC class II protein complex binding
(GO:0023026)

phosphoserine residue binding (GO:0050815)

miRNA binding (GO:0035198)

Tat protein binding (GO:0030957)

RNA binding (GO:0003723)

0

2

4

6

Enrichment log(combined score)