

GO_Biological_Process_2021

Term

SRP-dependent cotranslational protein
targeting to membrane (GO:0006614)

cotranslational protein targeting to
membrane (GO:0006613)

protein targeting to ER (GO:0045047)

cytoplasmic translation (GO:0002181)

nuclear-transcribed mRNA catabolic process,
nonsense-mediated decay (GO:0000184)

peptide biosynthetic process (GO:0043043)

positive regulation of protein catabolic
process in the vacuole (GO:1904352)

positive regulation of nucleotide-binding
oligomerization domain containing signaling
pathway (GO:0070426)

neutral lipid catabolic process (GO:0046461)

nuclear-transcribed mRNA catabolic process
(GO:0000956)

0

2

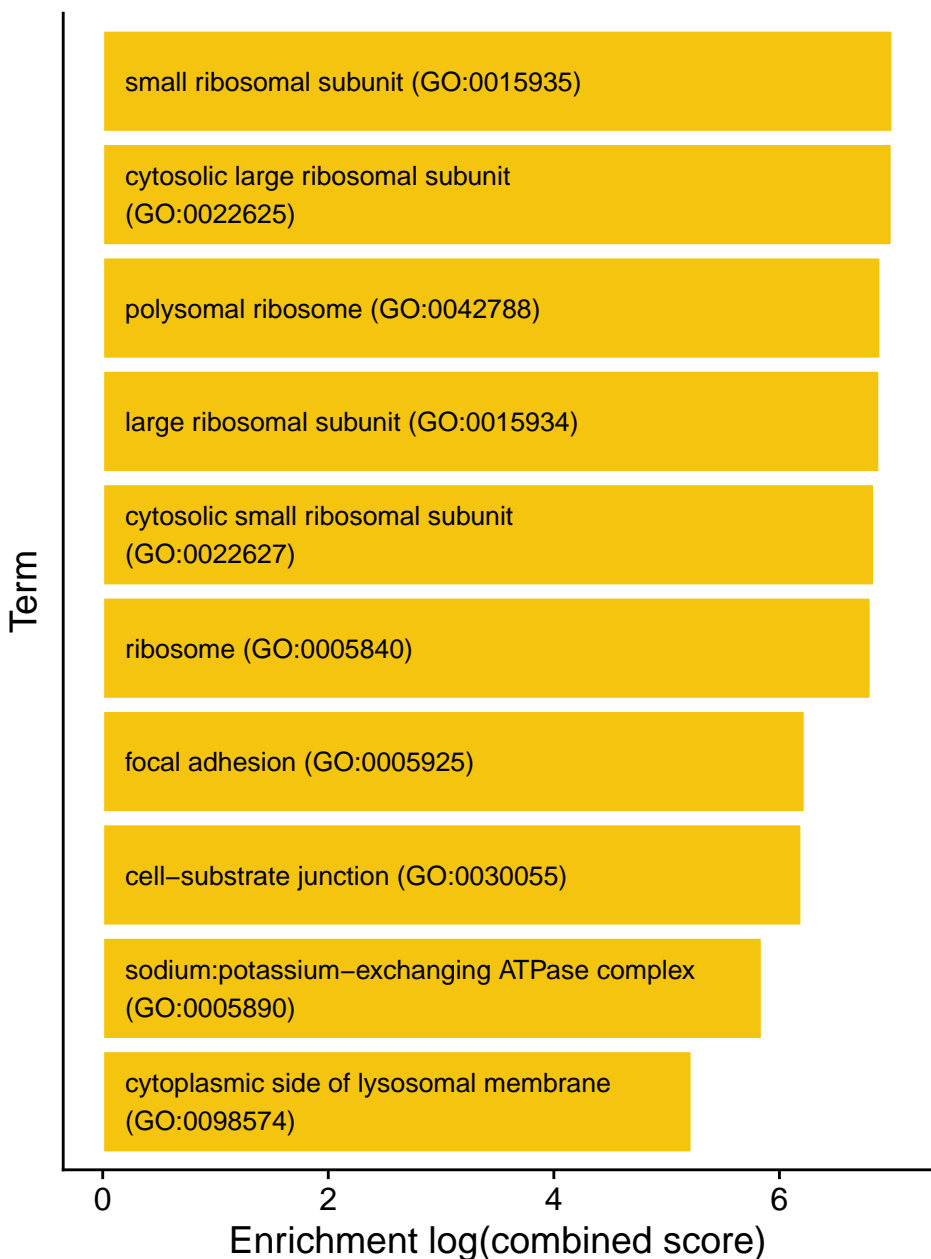
4

6

8

Enrichment log(combined score)

GO_Cellular_Component_2021



GO_Molecular_Function_2021

Term

ubiquitin ligase inhibitor activity

(GO:1990948)

ubiquitin-protein transferase inhibitor

activity (GO:0055105)

C3HC4-type RING finger domain binding

(GO:0055131)

acylglycerol lipase activity (GO:0047372)

voltage-gated calcium channel activity

involved in cardiac muscle cell action

potential (GO:0086007)

nitrite reductase activity (GO:0098809)

fructose-2,6-bisphosphate 2-phosphatase

activity (GO:0004331)

fibroblast growth factor-activated receptor

activity (GO:0005007)

endopolyphosphatase activity (GO:0000298)

diphosphoinositol-polyphosphate

diphosphatase activity (GO:0008486)

calcitonin family receptor activity

(GO:0097642)

bis(5'-adenosyl)-pentaphosphatase activity

(GO:0034432)

bis(5'-adenosyl)-hexaphosphatase activity

(GO:0034431)

benzodiazepine receptor binding (GO:0030156)

TPR domain binding (GO:0030911)

L-ornithine transmembrane transporter

activity (GO:0000064)

L-lysine transmembrane transporter activity

(GO:0015189)

0

2

4

6

Enrichment log(combined score)