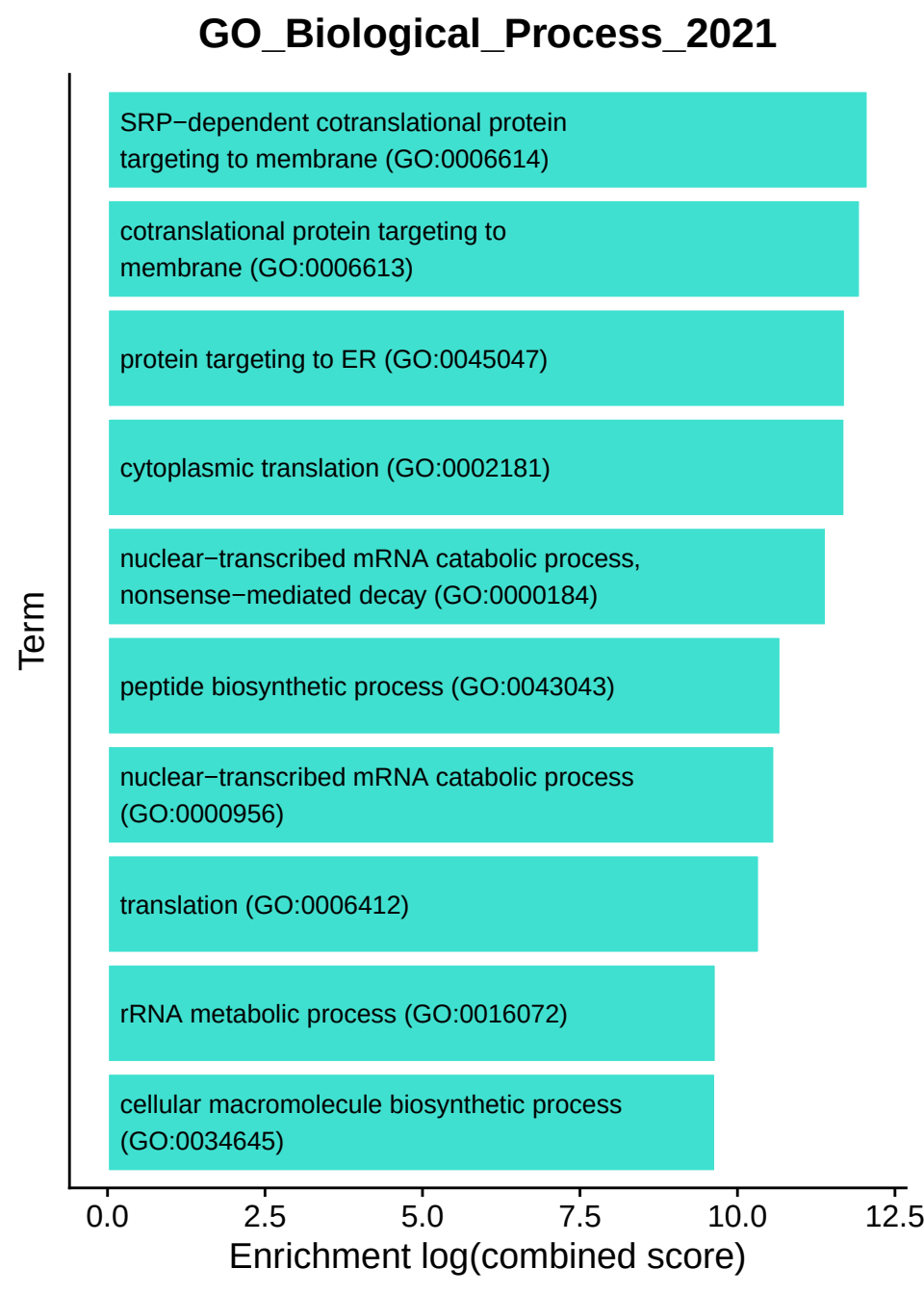
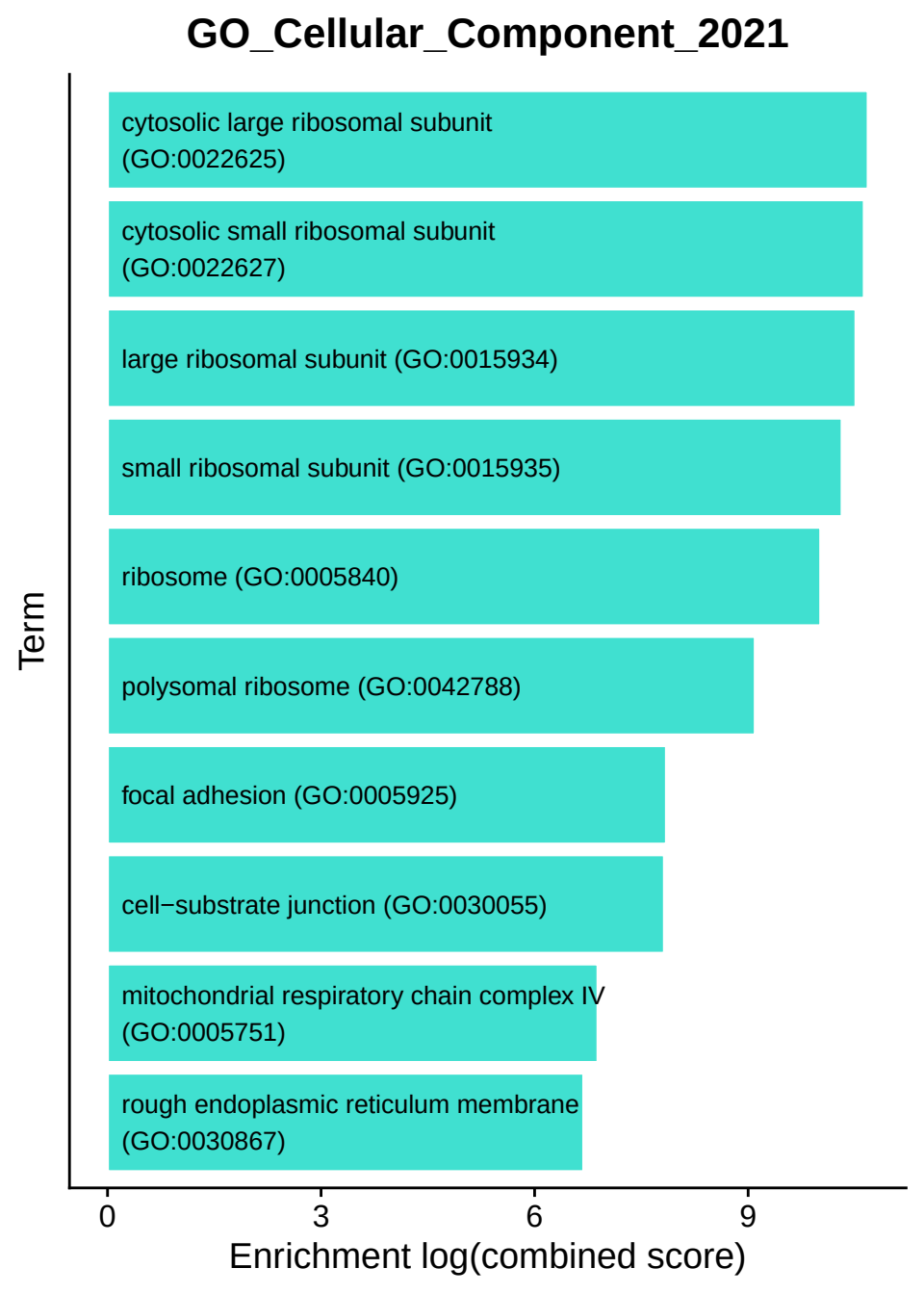

GO_Biological_Process_2021
SRP−dependent cotranslational protein targeting to membrane (GO:0006614)
cotranslational protein targeting to membrane (GO:0006613)
protein targeting to ER (GO:0045047)
cytoplasmic translation (GO:0002181)
nuclear−transcribed mRNA catabolic process, nonsense−mediated decay (GO:0000184)
peptide biosynthetic process (GO:0043043)
nuclear−transcribed mRNA catabolic process (GO:0000956)
translation (GO:0006412)
rRNA metabolic process (GO:0016072)
cellular macromolecule biosynthetic process (GO:0034645)
Term
0.0
2.5
5.0
7.5
10.0
12.5
Enrichment log(combined score)


GO_Cellular_Component_2021
cytosolic large ribosomal subunit (GO:0022625)
cytosolic small ribosomal subunit (GO:0022627)
large ribosomal subunit (GO:0015934)
small ribosomal subunit (GO:0015935)
ribosome (GO:0005840)
polysomal ribosome (GO:0042788)
focal adhesion (GO:0005925)
cell−substrate junction (GO:0030055)
mitochondrial respiratory chain complex IV (GO:0005751)
rough endoplasmic reticulum membrane (GO:0030867)
Term
0
3
6
9
Enrichment log(combined score)

GO_Molecular_Function_2021
ubiquitin ligase inhibitor activity (GO:1990948)
ubiquitin−protein transferase inhibitor activity (GO:0055105)
RNA binding (GO:0003723)
cytochrome−c oxidase activity (GO:0004129)
proton transmembrane transporter activity (GO:0015078)
rRNA binding (GO:0019843)
small ribosomal subunit rRNA binding (GO:0070181)
large ribosomal subunit rRNA binding (GO:0070180)
active ion transmembrane transporter activity (GO:0022853)
ubiquinol−cytochrome−c reductase activity (GO:0008121)
Term
0.0
2.5
5.0
7.5
Enrichment log(combined score)