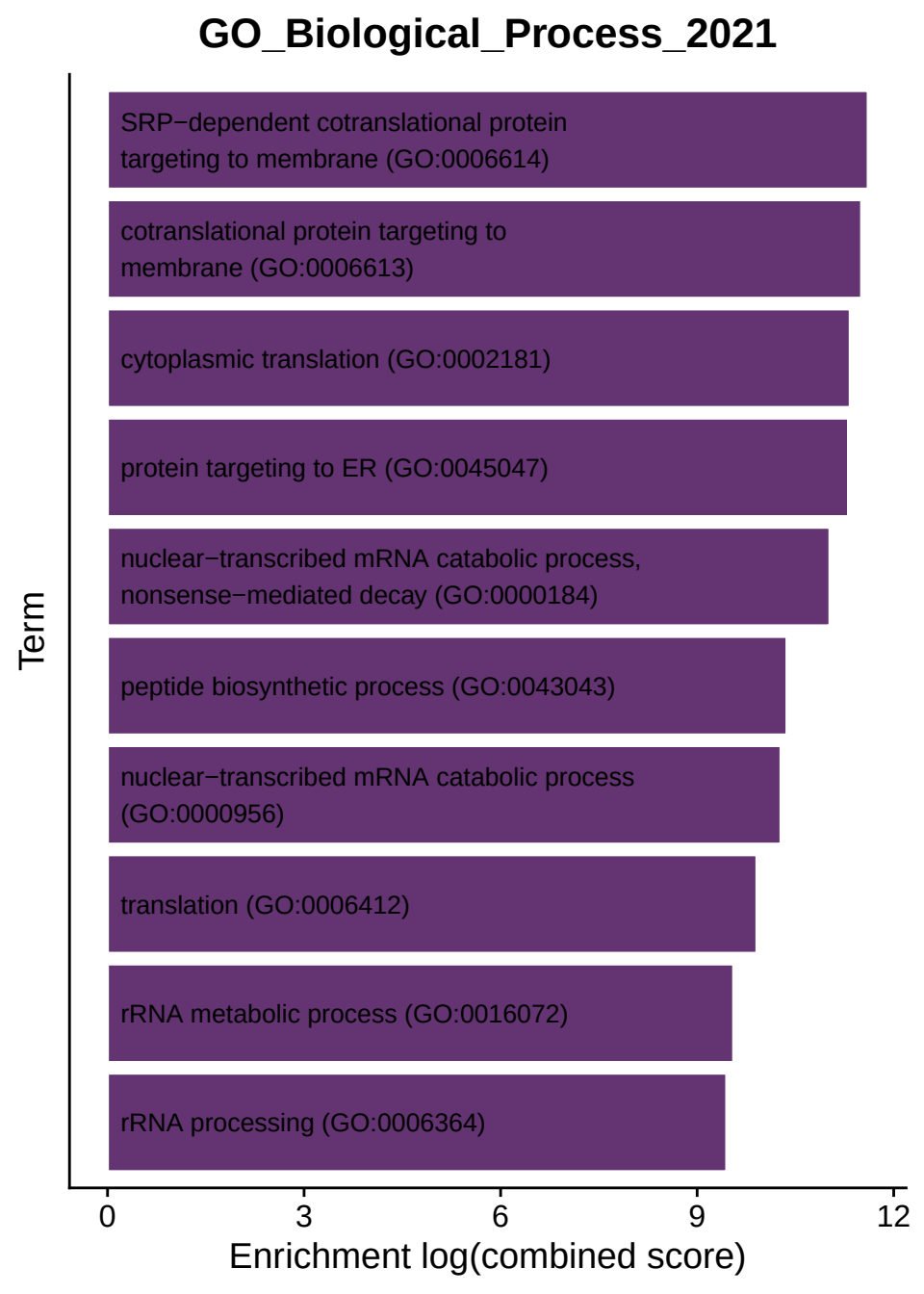
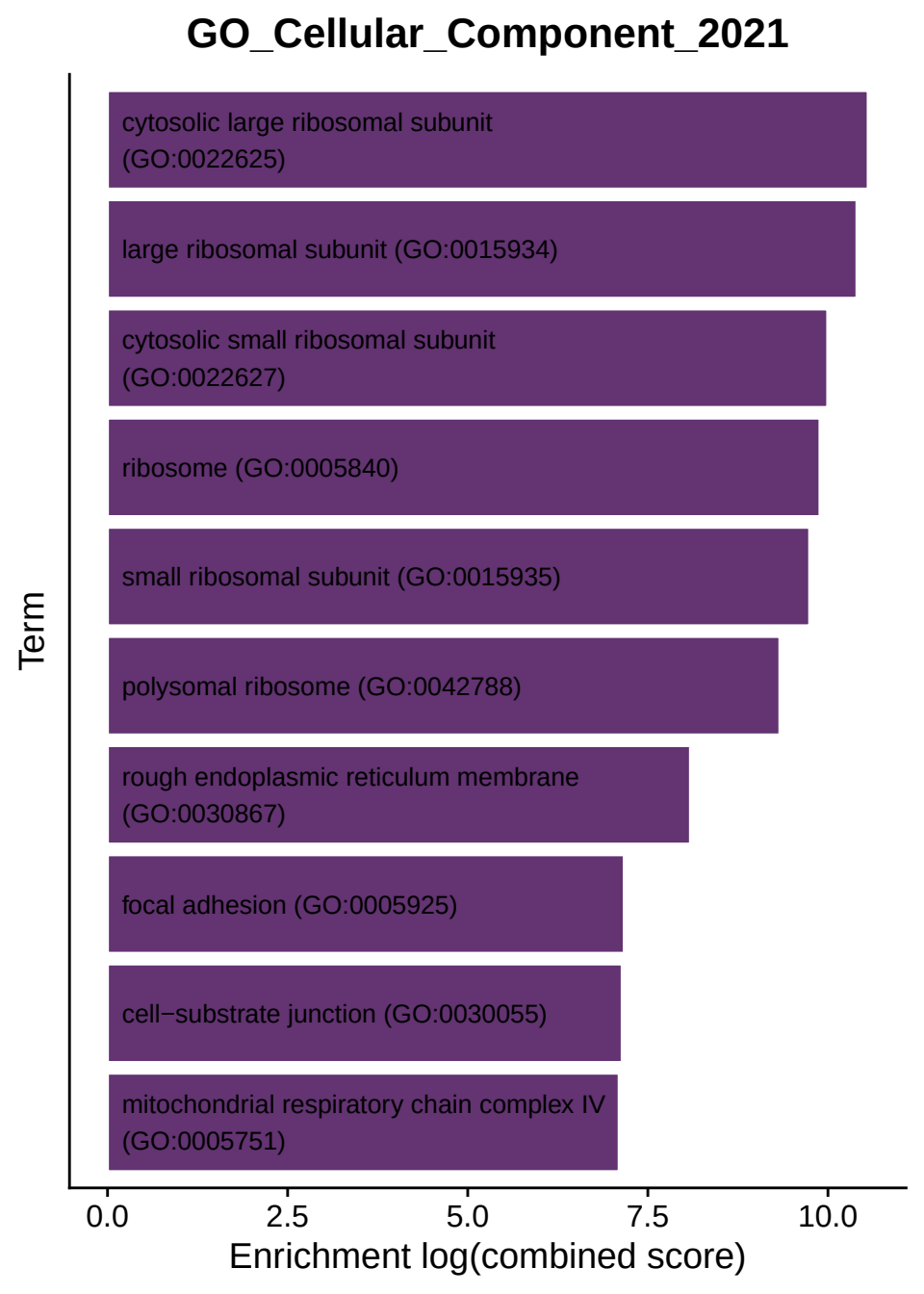
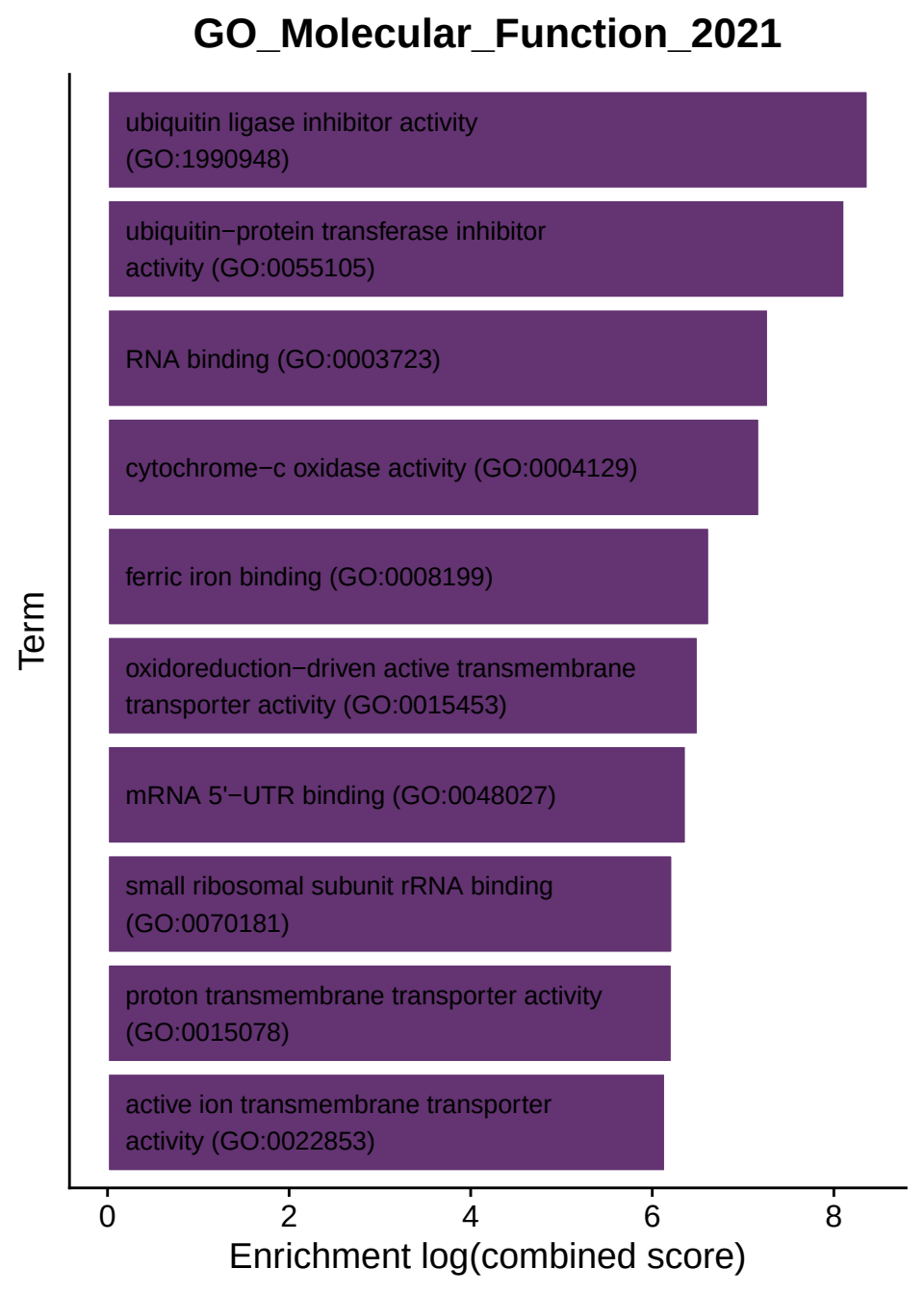

GO_Biological_Process_2021
SRP−dependent cotranslational protein targeting to membrane (GO:0006614)
cotranslational protein targeting to membrane (GO:0006613)
cytoplasmic translation (GO:0002181)
protein targeting to ER (GO:0045047)
nuclear−transcribed mRNA catabolic process, nonsense−mediated decay (GO:0000184)
peptide biosynthetic process (GO:0043043)
nuclear−transcribed mRNA catabolic process (GO:0000956)
translation (GO:0006412)
rRNA metabolic process (GO:0016072)
rRNA processing (GO:0006364)
Term
0
3
6
9
12
Enrichment log(combined score)


GO_Cellular_Component_2021
cytosolic large ribosomal subunit (GO:0022625)
large ribosomal subunit (GO:0015934)
cytosolic small ribosomal subunit (GO:0022627)
ribosome (GO:0005840)
small ribosomal subunit (GO:0015935)
polysomal ribosome (GO:0042788)
rough endoplasmic reticulum membrane (GO:0030867)
focal adhesion (GO:0005925)
cell−substrate junction (GO:0030055)
mitochondrial respiratory chain complex IV (GO:0005751)
Term
0.0
2.5
5.0
7.5
10.0
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)
ferric iron binding (GO:0008199)
oxidoreduction−driven active transmembrane transporter activity (GO:0015453)
mRNA 5'−UTR binding (GO:0048027)
small ribosomal subunit rRNA binding (GO:0070181)
proton transmembrane transporter activity (GO:0015078)
active ion transmembrane transporter activity (GO:0022853)
Term
0
2
4
6
8
Enrichment log(combined score)