

GO_Biological_Process_2021

Term

vesicle fusion with endoplasmic
reticulum–Golgi intermediate compartment
(ERGIC) membrane (GO:1990668)

response to host defenses (GO:0052200)

modulation by symbiont of host defense
response (GO:0052031)

oxidative DNA demethylation (GO:0035511)

negative regulation of IRE1–mediated
unfolded protein response (GO:1903895)

regulation of endoplasmic reticulum unfolded
protein response (GO:1900101)

modulation by symbiont of host process
(GO:0044003)

pyrimidine deoxyribonucleotide catabolic
process (GO:0009223)

negative regulation of endoplasmic reticulum
unfolded protein response (GO:1900102)

response to increased oxygen levels
(GO:0036296)

0

2

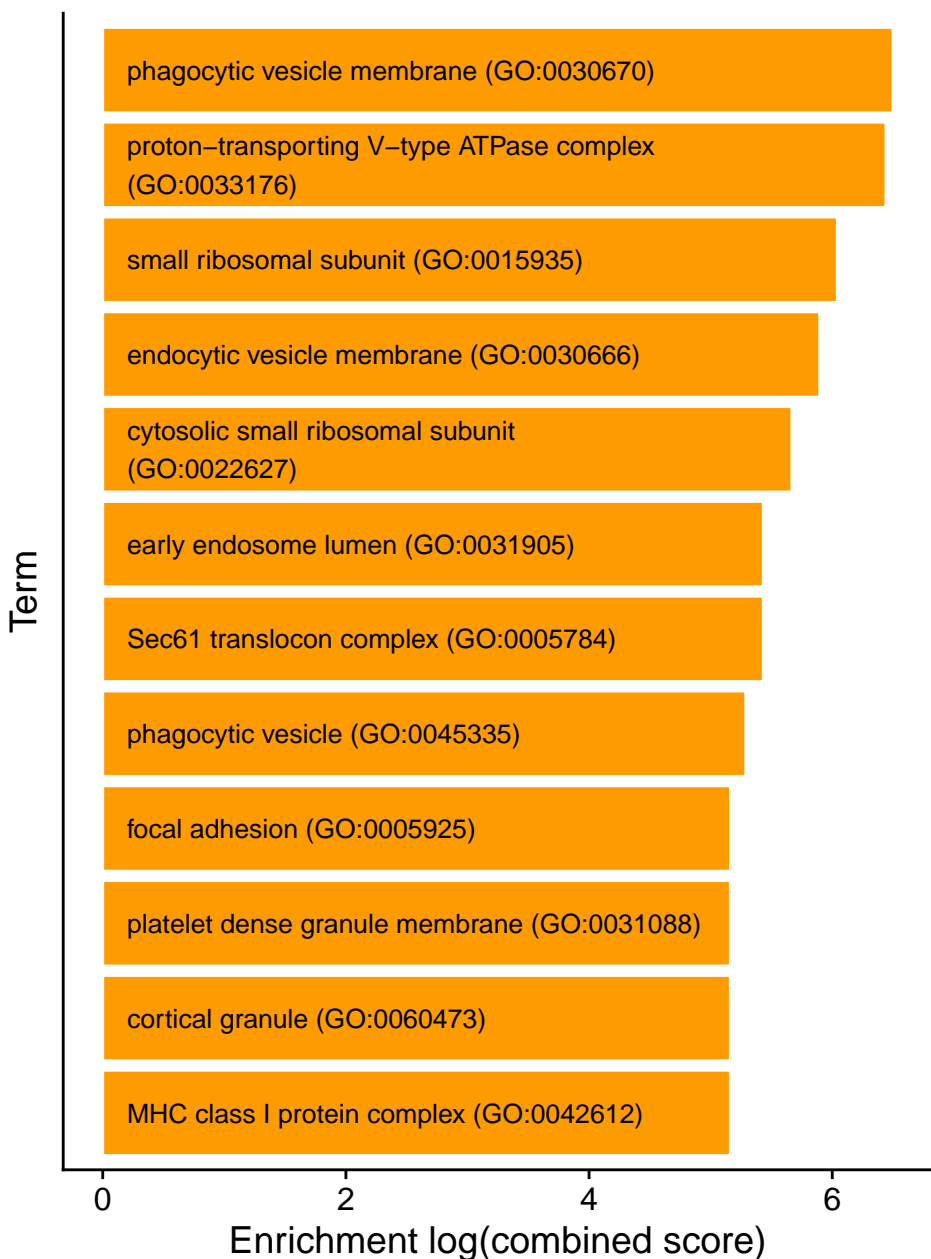
4

6

8

Enrichment log(combined score)

GO_Cellular_Component_2021



GO_Molecular_Function_2021

Term

TPR domain binding (GO:0030911)

TAP1 binding (GO:0046978)

BH3 domain binding (GO:0051434)

BH domain binding (GO:0051400)

deoxyribonuclease activity (GO:0004536)

DNA N-glycosylase activity (GO:0019104)

proton-transporting ATPase activity,
rotational mechanism (GO:0046961)

ATPase activity, coupled to transmembrane
movement of ions, rotational mechanism
(GO:0044769)

small ribosomal subunit rRNA binding
(GO:0070181)

endodeoxyribonuclease activity (GO:0004520)

0

2

4

6

Enrichment log(combined score)