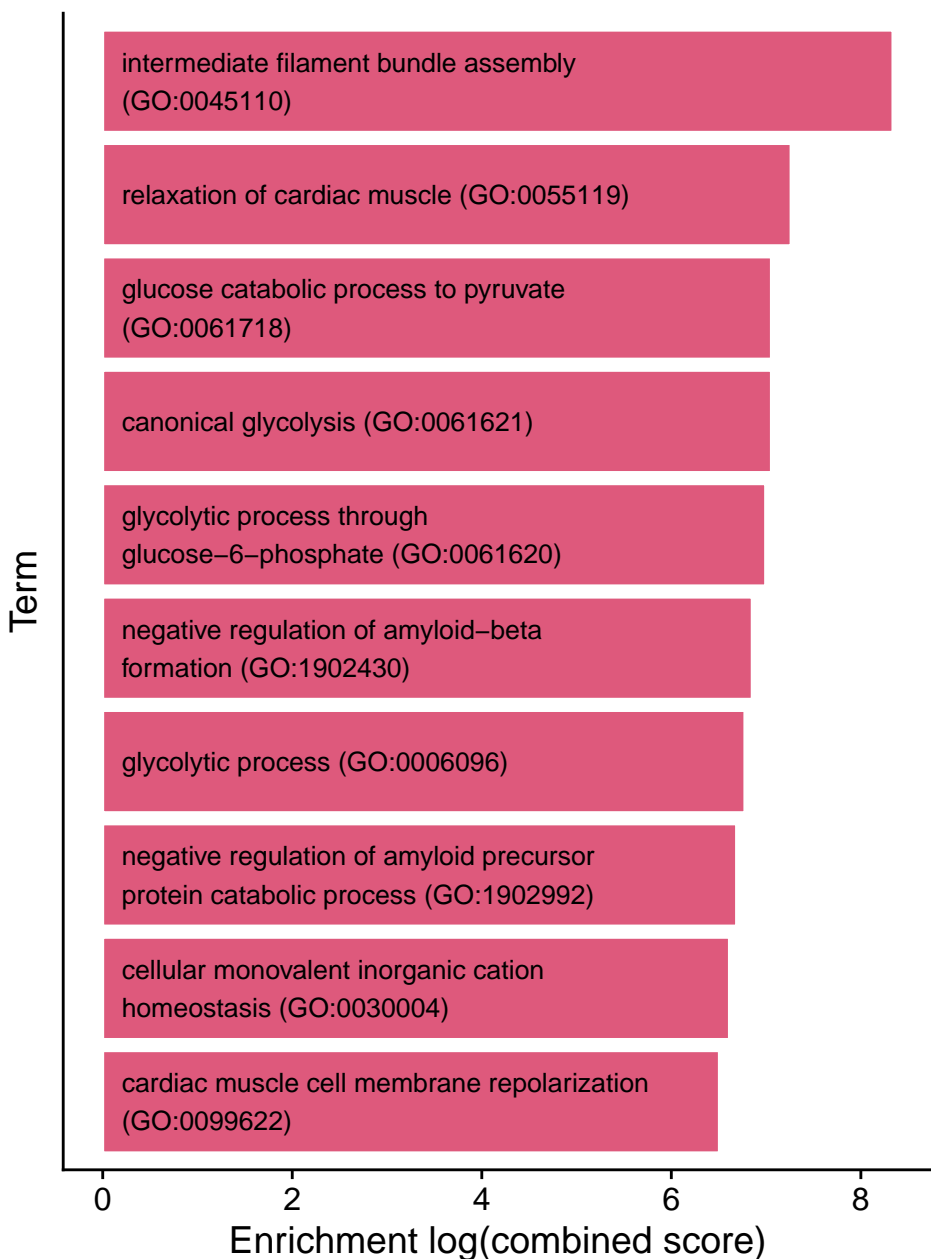


GO_Biological_Process_2021



GO_Cellular_Component_2021



GO_Molecular_Function_2021

Term

ubiquinol–cytochrome–c reductase activity

(GO:0008121)

pyrophosphate hydrolysis–driven proton
transmembrane transporter activity

(GO:0009678)

phosphotransferase activity, nitrogenous
group as acceptor (GO:0016775)

ganglioside binding (GO:0035594)

creatine kinase activity (GO:0004111)

alpha–2A adrenergic receptor binding
(GO:0031694)

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

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

sodium ion binding (GO:0031402)

fructose binding (GO:0070061)

dihydropyrimidinase activity (GO:0004157)

P–type sodium:potassium–exchanging
transporter activity (GO:0005391)

P–type sodium transporter activity
(GO:0008554)

0

2

4

6

Enrichment log(combined score)