

GO_Biological_Process_2021

Term

regulation of microvillus assembly
(GO:0032534)

regulation of cytochrome-c oxidase activity
(GO:1904959)

regulation of chloride transport
(GO:2001225)

regulation of Fc-gamma receptor signaling
pathway involved in phagocytosis
(GO:1905449)

positive regulation of histone H3-K27
methylation (GO:0061087)

negative regulation of sphingolipid
biosynthetic process (GO:0090155)

negative regulation of histone H3-K27
methylation (GO:0061086)

negative regulation of ceramide biosynthetic
process (GO:1900060)

monocyte extravasation (GO:0035696)

glutathione transport (GO:0034635)

SRP-dependent cotranslational protein
targeting to membrane, translocation
(GO:0006616)

0

1

2

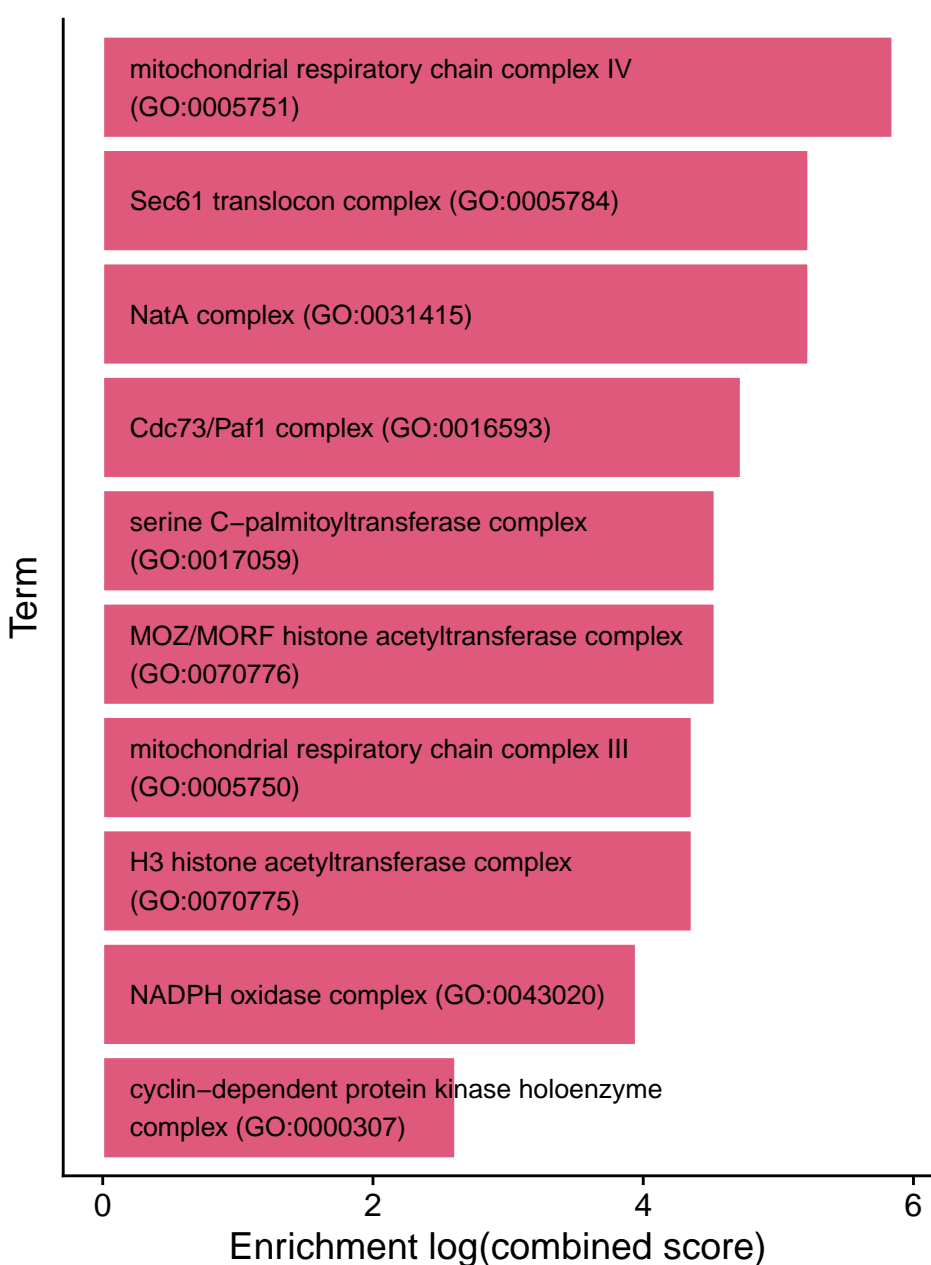
3

4

5

Enrichment log(combined score)

GO_Cellular_Component_2021



GO_Molecular_Function_2021

Term

pyruvate transmembrane transporter activity
(GO:0050833)

phosphatidylserine flippase activity
(GO:0140346)

phosphatidylcholine flippase activity
(GO:0140345)

histone acetyltransferase activity (H3-K23
specific) (GO:0043994)

aminophospholipid flippase activity
(GO:0015247)

NAD(P)H oxidase H₂O₂-forming activity
(GO:0016174)

3',5'-cyclic-GMP phosphodiesterase activity
(GO:0047555)

poly(G) binding (GO:0034046)

estradiol 17-beta-dehydrogenase activity
(GO:0004303)

aldehyde-lyase activity (GO:0016832)

5'-3' exodeoxyribonuclease activity
(GO:0035312)

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Enrichment log(combined score)