

# GO\_Biological\_Process\_2021

Term

serine family amino acid catabolic process  
(GO:0009071)

negative regulation of establishment of  
protein localization (GO:1904950)

regulation of protein transport (GO:0051223)

vascular endothelial growth factor  
receptor-2 signaling pathway (GO:0036324)

regulation of sterol transport (GO:0032371)

protein lipoylation (GO:0009249)

mitochondrial tRNA methylation (GO:0070901)

mitochondrial RNA 3'-end processing  
(GO:0000965)

late endosome to Golgi transport  
(GO:0034499)

His-Purkinje system development (GO:0003164)

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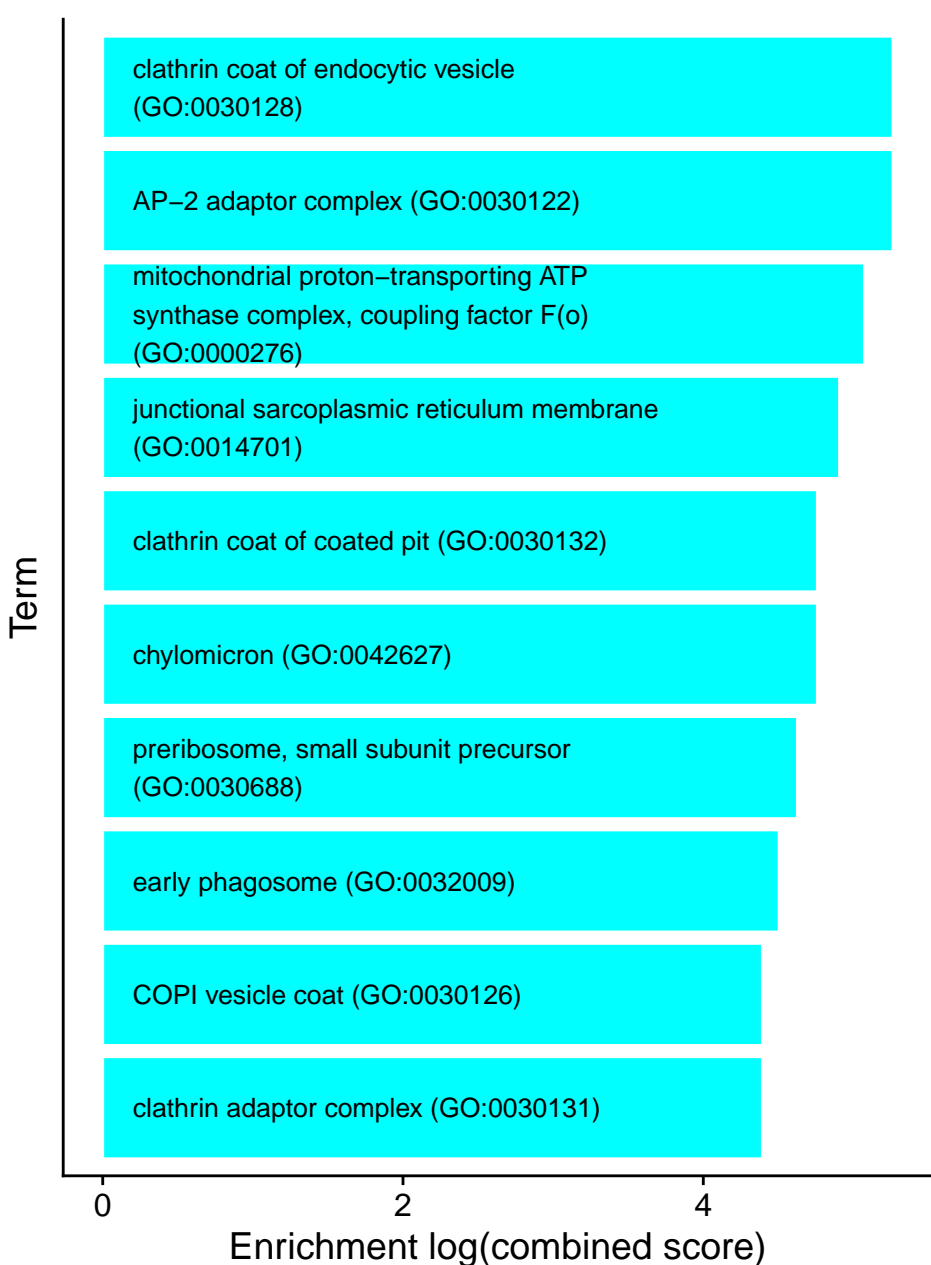
2

4

6

Enrichment log(combined score)

# GO\_Cellular\_Component\_2021



# GO\_Molecular\_Function\_2021

Term

testosterone dehydrogenase [NAD(P)] activity  
(GO:0030283)

GPI-linked ephrin receptor activity  
(GO:0005004)

benzodiazepine receptor binding (GO:0030156)

phosphatidylcholine-sterol O-acyltransferase  
activator activity (GO:0060228)

peroxisome targeting sequence binding  
(GO:0000268)

long-chain fatty acyl-CoA binding  
(GO:0036042)

3-hydroxyacyl-CoA dehydrogenase activity  
(GO:0003857)

procollagen-proline dioxygenase activity  
(GO:0019798)

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

phospholipase inhibitor activity  
(GO:0004859)

lipase inhibitor activity (GO:0055102)

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2

4

6

Enrichment log(combined score)