

# GO\_Biological\_Process\_2021

Term

T cell extravasation (GO:0072683)

mitochondrion morphogenesis (GO:0070584)

cell part morphogenesis (GO:0032990)

vesicle fusion with Golgi apparatus  
(GO:0048280)

ribose phosphate metabolic process  
(GO:0019693)

regulation of lyase activity (GO:0051339)

regulation of circadian sleep/wake cycle  
(GO:0042749)

quinone catabolic process (GO:1901662)

positive regulation of neutrophil  
extravasation (GO:2000391)

positive regulation of metallopeptidase  
activity (GO:1905050)

negative regulation of vascular endothelial  
cell proliferation (GO:1905563)

negative regulation of lipid transport  
(GO:0032369)

negative regulation of gonadotropin  
secretion (GO:0032277)

glutathione transport (GO:0034635)

cellular response to erythropoietin  
(GO:0036018)

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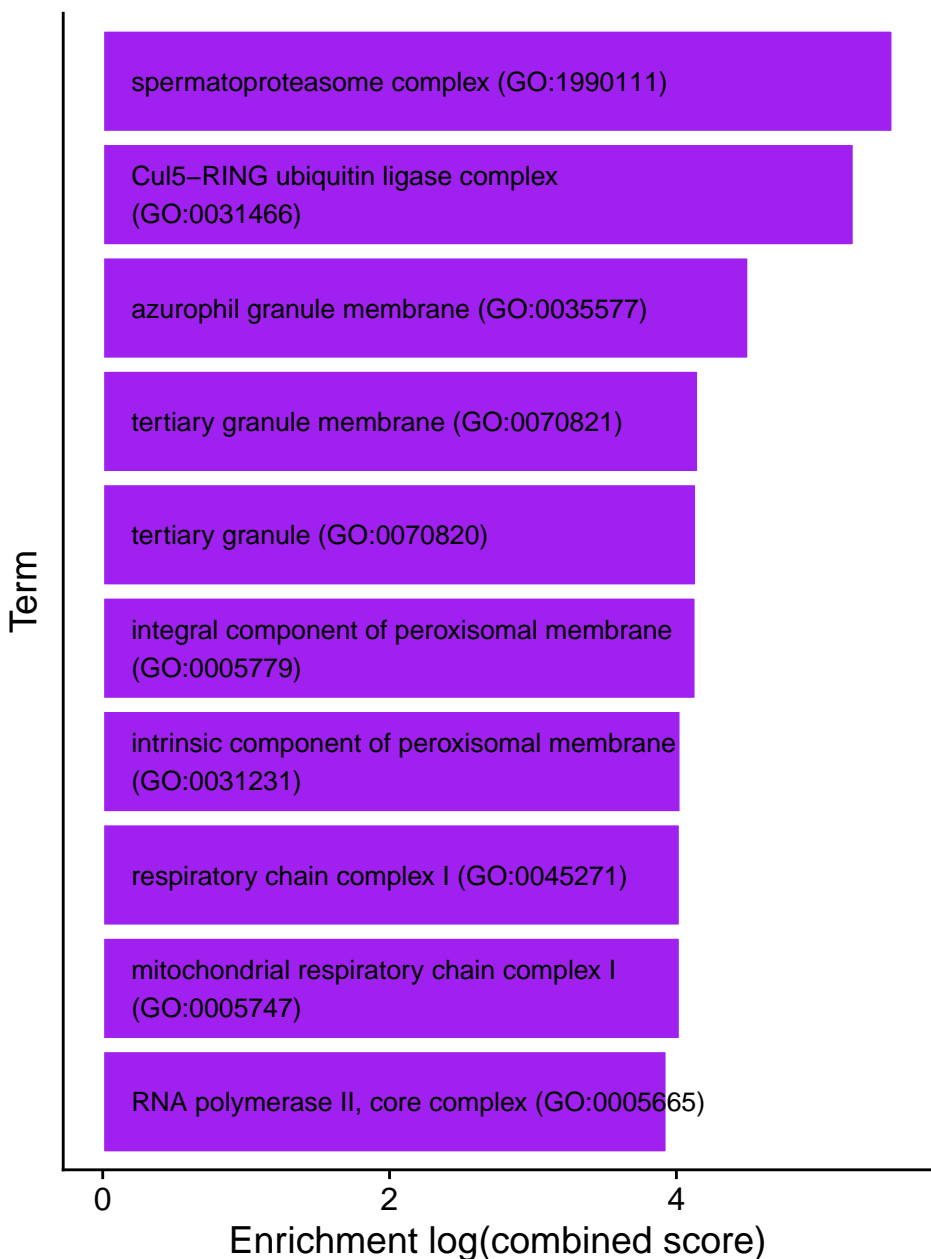
2

4

6

Enrichment log(combined score)

# GO\_Cellular\_Component\_2021



# GO\_Molecular\_Function\_2021

Term

IgG binding (GO:0019864)

glutathione transmembrane transporter  
activity (GO:0034634)

gap junction channel activity involved in  
cardiac conduction electrical coupling  
(GO:0086075)

oxidoreductase activity, acting on the CH-CH  
group of donors, NAD or NADP as acceptor  
(GO:0016628)

tripeptide transmembrane transporter  
activity (GO:0042937)

gap junction hemi-channel activity  
(GO:0055077)

gap junction channel activity involved in  
cell communication by electrical coupling  
(GO:1903763)

phosphatidylinositol-4,5-bisphosphate  
5-phosphatase activity (GO:0004439)

phosphatidylinositol phosphate 5-phosphatase  
activity (GO:0034595)

G protein-coupled glutamate receptor binding  
(GO:0035256)

cAMP-dependent protein kinase inhibitor  
activity (GO:0004862)

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2

4

Enrichment log(combined score)