

GO_Biological_Process_2021

Term

protein deglutamylation (GO:0035608)
 organic heteropentacyclic compound metabolic
 process (GO:1901376)
 mycotoxin metabolic process (GO:0043385)
 aflatoxin metabolic process (GO:0046222)
 salivary gland morphogenesis (GO:0007435)
 salivary gland development (GO:0007431)
 response to indole-3-methanol (GO:0071680)
 regulation of skeletal muscle tissue
 development (GO:0048641)
 regulation of protein import (GO:1904589)
 receptor transactivation (GO:0035624)
 protein side chain deglutamylation
 (GO:0035610)
 negative regulation of posttranscriptional
 gene silencing (GO:0060148)
 negative regulation of metalloproteinase
 activity (GO:1905049)
 negative regulation of macrophage cytokine
 production (GO:0010936)
 negative regulation of gene silencing by RNA
 (GO:0060967)
 maintenance of animal organ identity
 (GO:0048496)
 lymph node development (GO:0048535)
 leukotriene B4 metabolic process
 (GO:0036102)
 glutathione transport (GO:0034635)
 extracellular exosome biogenesis
 (GO:0097734)
 exocyst localization (GO:0051601)
 cellular response to indole-3-methanol
 (GO:0071681)
 caveola assembly (GO:0070836)
 auditory receptor cell stereocilium
 organization (GO:0060088)
 auditory receptor cell morphogenesis
 (GO:0002093)
 alkaloid metabolic process (GO:0009820)
 T cell extravasation (GO:0072683)

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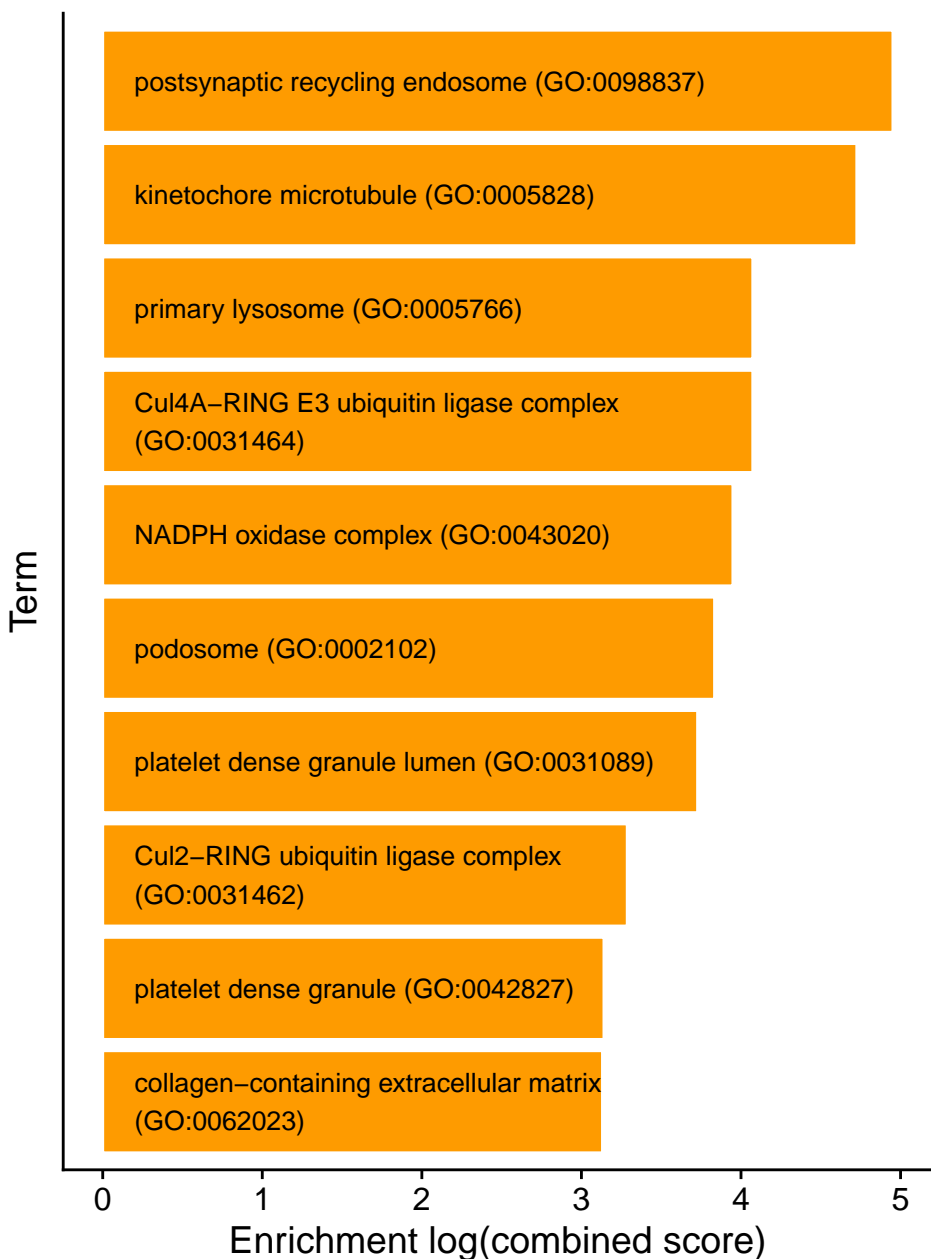
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4

6

Enrichment log(combined score)

GO_Cellular_Component_2021



GO_Molecular_Function_2021

Term

metalloendopeptidase inhibitor activity
(GO:0008191)

protein-arginine omega-N
monomethyltransferase activity (GO:0035241)

procollagen-proline 4-dioxygenase activity
(GO:0004656)

5'-deoxyribose-5-phosphate lyase activity
(GO:0051575)

1,4-alpha-oligoglucan phosphorylase activity
(GO:0004645)

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

arylesterase activity (GO:0004064)

amidine-lyase activity (GO:0016842)

DNA-directed DNA polymerase activity
(GO:0003887)

interleukin-6 receptor binding (GO:0005138)

fatty-acyl-CoA synthase activity
(GO:0004321)

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

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

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2

4

Enrichment log(combined score)