

GO_Biological_Process_2021

Term

axon choice point recognition (GO:0016198)

presynaptic membrane assembly (GO:0097105)

presynaptic membrane organization (GO:0097090)

adherens junction organization (GO:0034332)

chondroitin sulfate catabolic process (GO:0030207)

aorta development (GO:0035904)

response to muscle stretch (GO:0035994)

ubiquitin-dependent glycoprotein ERAD pathway (GO:0097466)

response to morphine (GO:0043278)

regulation of homophilic cell adhesion (GO:1903385)

regulation of histone H4-K16 acetylation (GO:2000618)

protein deglycosylation involved in glycoprotein catabolic process (GO:0035977)

positive regulation of mitochondrial depolarization (GO:0051901)

positive regulation of CD4-positive, alpha-beta T cell proliferation (GO:2000563)

positive regulation by host of viral genome replication (GO:0044829)

negative regulation of synapse organization (GO:1905809)

negative regulation of metallopeptidase activity (GO:1905049)

negative regulation of T cell migration (GO:2000405)

negative regulation of RNA polymerase II regulatory region sequence-specific DNA binding (GO:1903026)

mannose trimming involved in glycoprotein ERAD pathway (GO:1904382)

cellular response to brain-derived neurotrophic factor stimulus (GO:1990416)

Roundabout signaling pathway (GO:0035385)

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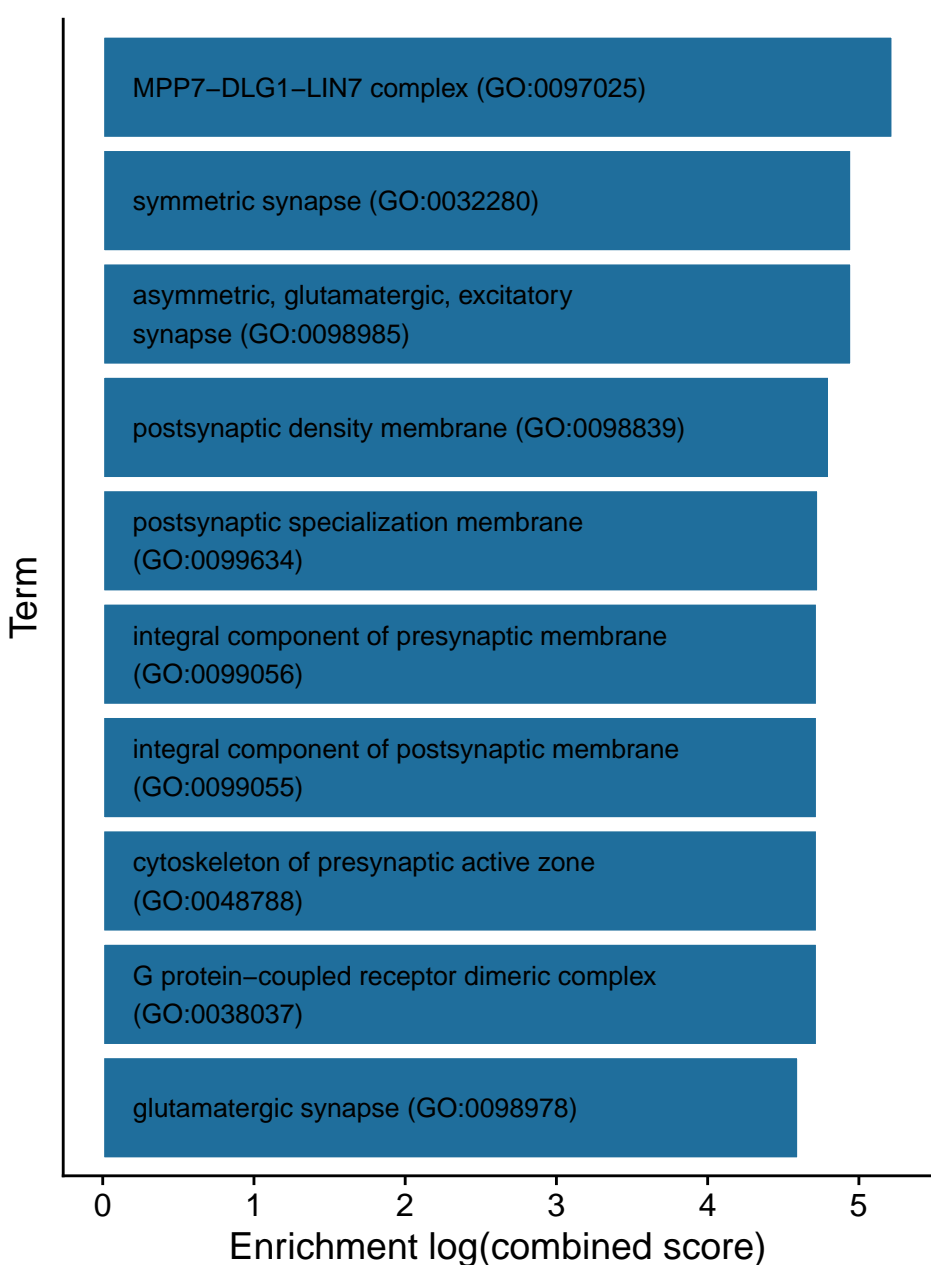
2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021



GO_Molecular_Function_2021

Term

nitric-oxide synthase binding (GO:0050998)

opioid receptor binding (GO:0031628)

neuroligin family protein binding
(GO:0097109)

transmembrane receptor protein tyrosine
phosphatase activity (GO:0005001)

transmembrane receptor protein phosphatase
activity (GO:0019198)

L-serine transmembrane transporter activity
(GO:0015194)

G protein-coupled serotonin receptor binding
(GO:0031821)

leak channel activity (GO:0022840)

calcium activated cation channel activity
(GO:0005227)

phosphatidylinositol phosphate 4-phosphatase
activity (GO:0034596)

mannosyl-oligosaccharide mannosidase
activity (GO:0015924)

mannosyl-oligosaccharide
1,2- α -mannosidase activity (GO:0004571)

UDP-xylosyltransferase activity (GO:0035252)

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4

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Enrichment log(combined score)