GO_Biological_Process_2021

postsynaptic membrane assembly (GO:0097104)

postsynaptic density protein 95 clustering (GO:0097119)

intracellular sequestering of iron ion (GO:0006880)

postsynapse assembly (GO:0099068)

regulation of extracellular matrix constituent secretion (GO:0003330)

postsynaptic density organization (GO:0097106)

excitatory synapse assembly (GO:1904861)

neuron cell−cell adhesion (GO:0007158)

mitochondrial fragmentation involved in apoptotic process (GO:0043653)
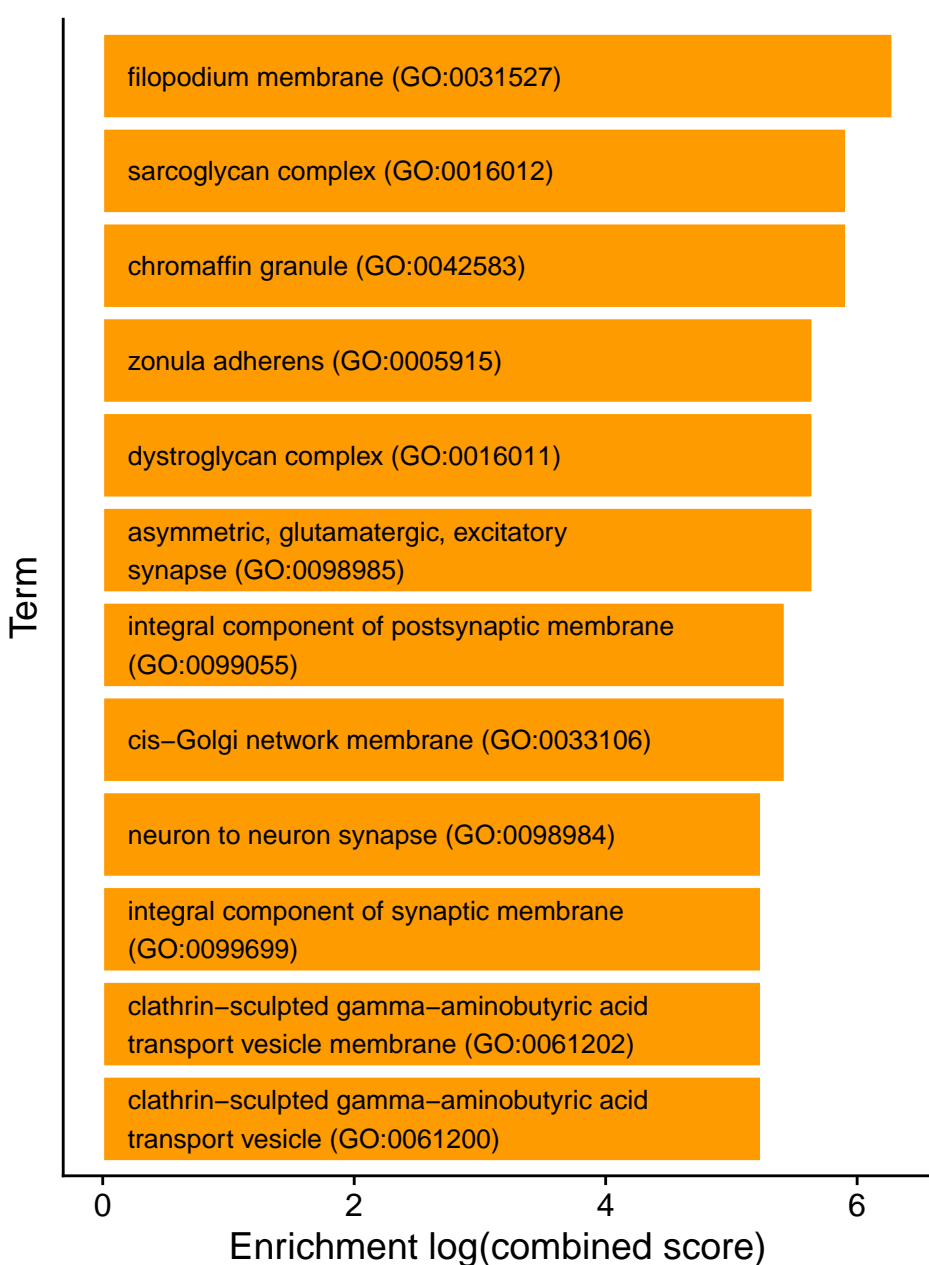
postsynaptic membrane organization (GO:0001941)

Term

Enrichment log(combined score)

GO_Cellular_Component_2021

Bar chart titled "GO_Cellular_Component_2021" with y-axis labeled "Term" and x-axis labeled "Enrichment log(combined score)". Bars from top to bottom:

- filopodium membrane (GO:0031527)
- sarcoglycan complex (GO:0016012)
- chromaffin granule (GO:0042583)
- zonula adherens (GO:0005915)
- dystroglycan complex (GO:0016011)
- asymmetric, glutamatergic, excitatory synapse (GO:0098985)
- integral component of postsynaptic membrane (GO:0099055)
- cis−Golgi network membrane (GO:0033106)
- neuron to neuron synapse (GO:0098984)
- integral component of synaptic membrane (GO:0099699)
- clathrin−sculpted gamma−aminobutyric acid transport vesicle membrane (GO:0061202)
- clathrin−sculpted gamma−aminobutyric acid transport vesicle (GO:0061200)

GO_Molecular_Function_2021

syndecan binding (GO:0045545)

neuroligin family protein binding
(GO:0097109)

neuregulin binding (GO:0038132)

C−X3−C chemokine binding (GO:0019960)

C3HC4−type RING finger domain binding
(GO:0055131)

glutamate receptor activity (GO:0008066)

nitric−oxide synthase binding (GO:0050998)

manganese ion transmembrane transporter
activity (GO:0005384)

ferric iron binding (GO:0008199)

N−acylsphingosine amidohydrolase activity
(GO:0017040)

Term

Enrichment log(combined score)

0   2   4   6