

GO_Biological_Process_2021

Term

presynaptic membrane assembly (GO:0097105)

presynaptic membrane organization
(GO:0097090)

presynapse assembly (GO:0099054)

postsynaptic membrane assembly (GO:0097104)

protein localization to axon (GO:0099612)

cerebellar granule cell differentiation
(GO:0021707)

regulation of actin filament-based movement
(GO:1903115)

regulation of release of sequestered calcium
ion into cytosol by sarcoplasmic reticulum
(GO:0010880)

postsynapse assembly (GO:0099068)

regulation of relaxation of cardiac muscle
(GO:1901897)

regulation of peptidyl-cysteine
S-nitrosylation (GO:2000169)

prepulse inhibition (GO:0060134)

neurotransmitter-gated ion channel
clustering (GO:0072578)

negative regulation of relaxation of muscle
(GO:1901078)

0

2

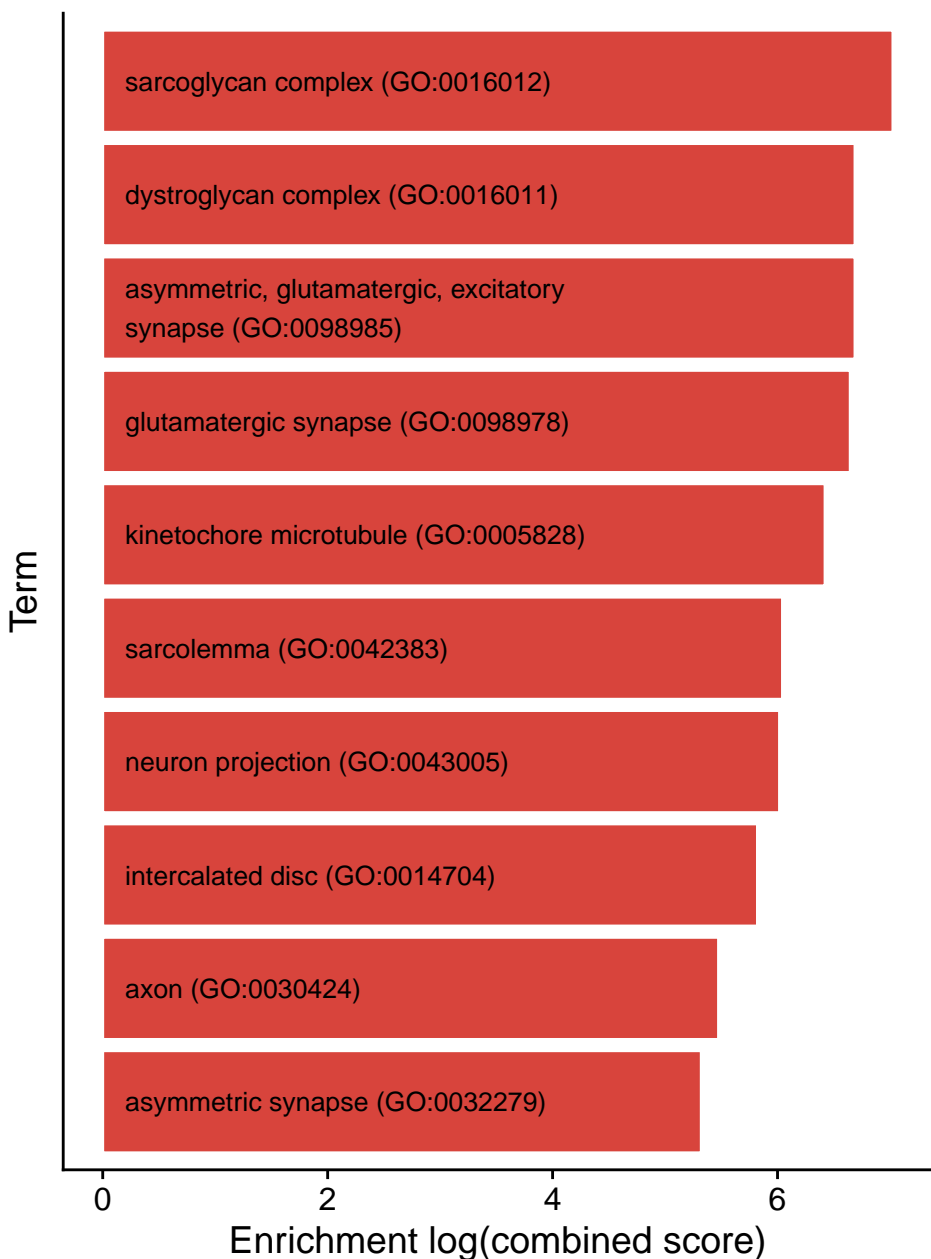
4

6

8

Enrichment log(combined score)

GO_Cellular_Component_2021



GO_Molecular_Function_2021

Term

transmembrane receptor protein tyrosine
phosphatase activity (GO:0005001)

transmembrane receptor protein phosphatase
activity (GO:0019198)

glutamate receptor activity (GO:0008066)

nitric-oxide synthase binding (GO:0050998)

adenylate cyclase inhibiting G
protein-coupled glutamate receptor activity
(GO:0001640)

G protein-coupled glutamate receptor
activity (GO:0098988)

neurexin family protein binding (GO:0042043)

cAMP binding (GO:0030552)

3',5'-cyclic-AMP phosphodiesterase activity
(GO:0004115)

PDZ domain binding (GO:0030165)

0

2

4

6

Enrichment log(combined score)