

GO_Biological_Process_2021

Term

retinal ganglion cell axon guidance
(GO:0031290)

forebrain neuron differentiation
(GO:0021879)

membrane assembly (GO:0071709)

negative regulation of focal adhesion
assembly (GO:0051895)

negative regulation of cell–substrate
junction organization (GO:0150118)

postsynaptic membrane assembly (GO:0097104)

presynaptic membrane assembly (GO:0097105)

nerve growth factor signaling pathway
(GO:0038180)

presynaptic membrane organization
(GO:0097090)

postsynapse assembly (GO:0099068)

camera–type eye morphogenesis (GO:0048593)

0

2

4

6

8

Enrichment log(combined score)

GO_Cellular_Component_2021

Term

slit diaphragm (GO:0036057)

neurofibrillary tangle (GO:0097418)

mitochondrial alpha-ketoglutarate
dehydrogenase complex (GO:0005947)

intrinsic component of presynaptic membrane
(GO:0098889)

filtration diaphragm (GO:0036056)

MPP7-DLG1-LIN7 complex (GO:0097025)

glutamatergic synapse (GO:0098978)

uniplex complex (GO:1990246)

symmetric synapse (GO:0032280)

cyclin/CDK positive transcription elongation
factor complex (GO:0008024)

chromocenter (GO:0010369)

asymmetric, glutamatergic, excitatory
synapse (GO:0098985)

0

1

2

3

4

5

Enrichment log(combined score)

GO_Molecular_Function_2021

Term

JUN kinase binding (GO:0008432)

transmembrane receptor protein tyrosine
phosphatase activity (GO:0005001)

transmembrane receptor protein phosphatase
activity (GO:0019198)

adrenergic receptor binding (GO:0031690)

cAMP binding (GO:0030552)

GABA-gated chloride ion channel activity
(GO:0022851)

epidermal growth factor receptor binding
(GO:0005154)

neuroligin family protein binding
(GO:0097109)

choline transmembrane transporter activity
(GO:0015220)

beta-2 adrenergic receptor binding
(GO:0031698)

0

2

4

6

Enrichment log(combined score)