

GO_Biological_Process_2021

Term

regulation of long-term neuronal synaptic
plasticity (GO:0048169)

negative regulation of microtubule
polymerization (GO:0031115)

regulation of neuronal synaptic plasticity
(GO:0048168)

neuron projection maintenance (GO:1990535)

glucose catabolic process to pyruvate
(GO:0061718)

canonical glycolysis (GO:0061621)

positive regulation by host of viral process
(GO:0044794)

glycolytic process through
glucose-6-phosphate (GO:0061620)

regulation of protein depolymerization
(GO:1901879)

positive regulation of tau-protein kinase
activity (GO:1902949)

negative regulation of cytosolic calcium ion
concentration (GO:0051481)

0

2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021

Term

apical dendrite (GO:0097440)

integral component of synaptic vesicle
membrane (GO:0030285)

intrinsic component of synaptic vesicle
membrane (GO:0098563)

neuron projection (GO:0043005)

sodium:potassium-exchanging ATPase complex
(GO:0005890)

axon (GO:0030424)

axonal growth cone (GO:0044295)

vesicle (GO:0031982)

neurofibrillary tangle (GO:0097418)

intrinsic component of presynaptic membrane
(GO:0098889)

flotillin complex (GO:0016600)

clathrin-sculpted monoamine transport
vesicle membrane (GO:0070083)

clathrin-sculpted monoamine transport
vesicle (GO:0070081)

0

2

4

6

8

Enrichment log(combined score)

GO_Molecular_Function_2021

Term

cuprous ion binding (GO:1903136)

P-type calcium transporter activity
(GO:0005388)

opioid receptor binding (GO:0031628)

alpha-2A adrenergic receptor binding
(GO:0031694)

adenylate cyclase binding (GO:0008179)

P-type ion transporter activity (GO:0015662)

tau protein binding (GO:0048156)

aspartic-type endopeptidase inhibitor
activity (GO:0019828)

disordered domain specific binding
(GO:0097718)

adrenergic receptor binding (GO:0031690)

0

2

4

6

Enrichment log(combined score)