

GO_Biological_Process_2021

Term

axon regeneration (GO:0031103)

neuron projection regeneration (GO:0031102)

synaptic transmission, GABAergic
(GO:0051932)

neurotransmitter uptake (GO:0001504)

regulation of cell communication by
electrical coupling (GO:0010649)

regulation of synaptic vesicle priming
(GO:0010807)

regulation of morphogenesis of a branching
structure (GO:0060688)

regulation of chaperone-mediated protein
folding (GO:1903644)

regulation of cell maturation (GO:1903429)

positive regulation of the force of heart
contraction (GO:0098735)

positive regulation of protein localization
positive regulation of muscle fiber

oligomerization domain containing signaling
pathway (GO:0070428)

positive regulation of neurotransmitter
transport (GO:0051590)

positive regulation of dopamine receptor
signaling pathway (GO:0060161)

negative regulation of anion transmembrane
transport (GO:1903960)

mRNA 3'-end processing by stem-loop binding
and cleavage (GO:0006398)

kidney morphogenesis (GO:0060993)

intracellular distribution of mitochondria
(GO:0048312)

glutamate catabolic process (GO:0006538)

axon choice point recognition (GO:0016198)

RNA import into mitochondrion (GO:0035927)

0

2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021

Term

autolysosome (GO:0044754)

dendrite membrane (GO:0032590)

secondary lysosome (GO:0005767)

platelet dense granule membrane (GO:0031088)

phosphatidylinositol 3-kinase complex, class III (GO:0035032)

asymmetric, glutamatergic, excitatory synapse (GO:0098985)

GABA-A receptor complex (GO:1902711)

U7 snRNP (GO:0005683)

vacuolar proton-transporting V-type ATPase, V1 domain (GO:0000221)

proton-transporting V-type ATPase, V1 domain (GO:0033180)

dendritic spine membrane (GO:0032591)

0

2

4

6

Enrichment log(combined score)

GO_Molecular_Function_2021

Term

volume-sensitive anion channel activity
(GO:0005225)

benzodiazepine receptor activity
(GO:0008503)

neurotransmitter receptor activity involved
in regulation of postsynaptic membrane
potential (GO:0099529)

extracellular ligand-gated ion channel
activity (GO:0005230)

syntaxin-1 binding (GO:0017075)

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

inhibitory extracellular ligand-gated ion
channel activity (GO:0005237)

syntaxin binding (GO:0019905)

phosphatidylserine flippase activity
(GO:0140346)

kainate selective glutamate receptor
activity (GO:0015277)

collagen binding involved in cell-matrix
adhesion (GO:0098639)

GPI-linked ephrin receptor activity
(GO:0005004)

0

2

4

6

Enrichment log(combined score)