

GO_Biological_Process_2021

Term

regulation of respiratory gaseous exchange
by nervous system process (GO:0002087)

regulation of platelet-derived growth factor
receptor-beta signaling pathway (GO:2000586)

positive regulation of protein catabolic
process in the vacuole (GO:1904352)

regulation of keratinocyte apoptotic process
(GO:1902172)

negative regulation of cellular response to
hypoxia (GO:1900038)

negative regulation of amyloid precursor
protein biosynthetic process (GO:0042985)

dermatan sulfate biosynthetic process
(GO:0030208)

sodium ion export across plasma membrane
(GO:0036376)

relaxation of cardiac muscle (GO:0055119)

dermatan sulfate metabolic process
(GO:0030205)

chondroitin sulfate catabolic process
(GO:0030207)

cellular potassium ion homeostasis
(GO:0030007)

0

2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021

Term

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

inhibitory synapse (GO:0060077)

cation-transporting ATPase complex
(GO:0090533)

dendrite cytoplasm (GO:0032839)

neuron projection cytoplasm (GO:0120111)

smooth endoplasmic reticulum membrane
(GO:0030868)

neurofibrillary tangle (GO:0097418)

GABA-ergic synapse (GO:0098982)

transcription factor TFIIIC complex
(GO:0000127)

symmetric synapse (GO:0032280)

rough endoplasmic reticulum membrane
(GO:0030867)

asymmetric, glutamatergic, excitatory
synapse (GO:0098985)

0

2

4

6

Enrichment log(combined score)

GO_Molecular_Function_2021

Term

sodium ion binding (GO:0031402)

P-type sodium:potassium-exchanging
transporter activity (GO:0005391)

P-type sodium transporter activity
(GO:0008554)

P-type potassium transmembrane transporter
activity (GO:0008556)

potassium ion binding (GO:0030955)

neurexin family protein binding (GO:0042043)

alkali metal ion binding (GO:0031420)

GTPase inhibitor activity (GO:0005095)

epidermal growth factor receptor binding
(GO:0005154)

amyloid-beta binding (GO:0001540)

0

2

4

6

Enrichment log(combined score)