

GO_Biological_Process_2021

Term

regulation of sequestering of calcium ion
(GO:0051282)

SA node cell to atrial cardiac muscle cell
signaling (GO:0086018)

regulation of synapse structural plasticity
(GO:0051823)

regulation of inositol trisphosphate
biosynthetic process (GO:0032960)

protein side chain deglutamylation
(GO:0035610)

ornithine transport (GO:0015822)

membrane depolarization during SA node cell
action potential (GO:0086046)

maintenance of animal organ identity
(GO:0048496)

endothelin receptor signaling pathway
(GO:0086100)

C-terminal protein deglutamylation
(GO:0035609)

anion homeostasis (GO:0055081)

0

2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021

Term

slit diaphragm (GO:0036057)

NURF complex (GO:0016589)

filtration diaphragm (GO:0036056)

CD95 death-inducing signaling complex
(GO:0031265)

zonula adherens (GO:0005915)

0

2

4

6

Enrichment log(combined score)

GO_Molecular_Function_2021

Term

RNA polymerase II general transcription
initiation factor binding (GO:0001091)

L-lysine transmembrane transporter activity
(GO:0015189)

acyl carnitine transmembrane transporter
activity (GO:0015227)

1-acylglycerophosphocholine
O-acyltransferase activity (GO:0047184)

transmembrane receptor protein tyrosine
kinase activator activity (GO:0030297)

RNA polymerase II CTD heptapeptide repeat
phosphatase activity (GO:0008420)

phosphatidic acid transfer activity
(GO:1990050)

calcium-dependent protein serine/threonine
phosphatase activity (GO:0004723)

0

2

4

6

Enrichment log(combined score)