

GO_Biological_Process_2021

Term

negative regulation of macrophage
differentiation (GO:0045650)

negative regulation of sodium ion
transmembrane transporter activity
(GO:2000650)

negative regulation of sodium ion
transmembrane transport (GO:1902306)

regulation of relaxation of cardiac muscle
(GO:1901897)

regulation of Rab protein signal
transduction (GO:0032483)

regulation of macrophage differentiation
(GO:0045649)

negative regulation of sodium ion transport
(GO:0010766)

regulation of granulocyte differentiation
(GO:0030852)

positive regulation of RIG-I signaling
pathway (GO:1900246)

negative regulation of ion transmembrane
transporter activity (GO:0032413)

0

2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021

Term

cytoskeleton of presynaptic active zone
(GO:0048788)

Smc5–Smc6 complex (GO:0030915)

G protein–coupled receptor dimeric complex
(GO:0038037)

serine C–palmitoyltransferase complex
(GO:0017059)

inhibitory synapse (GO:0060077)

SUMO ligase complex (GO:0106068)

podosome (GO:0002102)

clathrin–coated vesicle (GO:0030136)

keratin filament (GO:0045095)

coated vesicle (GO:0030135)

0

1

2

3

4

Enrichment log(combined score)

GO_Molecular_Function_2021

Term

cAMP binding (GO:0030552)

protein kinase C activity (GO:0004697)

serine C–palmitoyltransferase activity
(GO:0004758)

phosphatidylinositol trisphosphate
phosphatase activity (GO:0034594)

chondroitin sulfotransferase activity
(GO:0034481)

N–acetylgalactosamine 4–O–sulfotransferase
activity (GO:0001537)

C–palmitoyltransferase activity (GO:0016454)

neurotrophin TRKA receptor binding
(GO:0005168)

neurotrophin TRK receptor binding
(GO:0005167)

myosin light chain binding (GO:0032027)

G protein–coupled serotonin receptor binding
(GO:0031821)

CoA carboxylase activity (GO:0016421)

0

2

4

Enrichment log(combined score)