

GO_Biological_Process_2021

Term

peroxisome proliferator activated receptor
signaling pathway (GO:0035357)

positive regulation of respiratory burst
(GO:0060267)

regulation of respiratory burst (GO:0060263)

retinoic acid receptor signaling pathway
(GO:0048384)

digestive system development (GO:0055123)

negative regulation of substrate
adhesion-dependent cell spreading
(GO:1900025)

negative regulation of locomotion
(GO:0040013)

negative regulation of cell morphogenesis
involved in differentiation (GO:0010771)

neurotrophin TRK receptor signaling pathway
(GO:0048011)

negative regulation of cellular component
movement (GO:0051271)

ganglioside biosynthetic process
(GO:0001574)

0

2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021

Term

protein kinase complex (GO:1902911)

spectrin-associated cytoskeleton
(GO:0014731)

intrinsic component of the cytoplasmic side
of the plasma membrane (GO:0031235)

pseudopodium (GO:0031143)

platelet dense tubular network membrane
(GO:0031095)

insulin-responsive compartment (GO:0032593)

apical dendrite (GO:0097440)

platelet dense tubular network (GO:0031094)

dendrite cytoplasm (GO:0032839)

Cul3-RING ubiquitin ligase complex
(GO:0031463)

0

1

2

3

4

5

Enrichment log(combined score)

GO_Molecular_Function_2021

Term

nerve growth factor binding (GO:0048406)

choline transmembrane transporter activity
(GO:0015220)

calcitonin family receptor activity
(GO:0097642)

C-X-C chemokine binding (GO:0019958)

retinoic acid receptor binding (GO:0042974)

DNA polymerase binding (GO:0070182)

telomerase activity (GO:0003720)

supercoiled DNA binding (GO:0097100)

alpha-N-acetylgalactosaminide
alpha-2,6-sialyltransferase activity
(GO:0001665)

RNA-directed DNA polymerase activity
(GO:0003964)

LBD domain binding (GO:0050693)

0

2

4

6

Enrichment log(combined score)