

GO_Biological_Process_2021

Term

lipoprotein metabolic process (GO:0042157)

positive regulation of vascular endothelial
growth factor receptor signaling pathway
(GO:0030949)

positive regulation of monocyte
differentiation (GO:0045657)

positive regulation of macrophage
differentiation (GO:0045651)

lipoprotein catabolic process (GO:0042159)

regulation of pseudopodium assembly
(GO:0031272)

positive regulation of pseudopodium assembly
(GO:0031274)

positive regulation of leukocyte
proliferation (GO:0070665)

negative regulation of hemostasis
(GO:1900047)

positive regulation of macrophage chemotaxis
(GO:0010759)

positive regulation of cellular response to
insulin stimulus (GO:1900078)

negative regulation of coagulation
(GO:0050819)

0

2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021

Term

protein kinase complex (GO:1902911)

low-density lipoprotein particle
(GO:0034362)

ISWI-type complex (GO:0031010)

chylomicron (GO:0042627)

platelet dense granule lumen (GO:0031089)

pericentric heterochromatin (GO:0005721)

very-low-density lipoprotein particle
(GO:0034361)

triglyceride-rich plasma lipoprotein
particle (GO:0034385)

high-density lipoprotein particle
(GO:0034364)

H4/H2A histone acetyltransferase complex
(GO:0043189)

0

1

2

3

4

5

Enrichment log(combined score)

GO_Molecular_Function_2021

Term

oxidoreductase activity, acting on metal
ions, NAD or NADP as acceptor (GO:0016723)

cytoskeleton–nuclear membrane anchor
activity (GO:0140444)

RNA adenylyltransferase activity
(GO:1990817)

phosphatidylcholine–sterol O–acyltransferase
activator activity (GO:0060228)

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

sphingosine–1–phosphate phosphatase activity
(GO:0042392)

oxysterol binding (GO:0008142)

insulin–like growth factor II binding
(GO:0031995)

vascular endothelial growth factor receptor
2 binding (GO:0043184)

mitogen–activated protein kinase kinase
binding (GO:0031434)

L–glutamine transmembrane transporter
activity (GO:0015186)

0

2

4

Enrichment log(combined score)