

GO_Biological_Process_2021

Term

positive regulation of prostaglandin
biosynthetic process (GO:0031394)

positive regulation of CD4-positive,
CD25-positive, alpha-beta regulatory T cell
differentiation (GO:0032831)

carbon dioxide transport (GO:0015670)

positive regulation of viral entry into host
cell (GO:0046598)

positive regulation by symbiont of entry
into host (GO:0075294)

regulation of T-helper cell differentiation
(GO:0045622)

peptide antigen assembly with MHC protein
complex (GO:0002501)

membrane raft assembly (GO:0001765)

cellular water homeostasis (GO:0009992)

protein insertion into mitochondrial inner
membrane (GO:0045039)

0

2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021

Term

MHC class II protein complex (GO:0042613)

luminal side of endoplasmic reticulum
membrane (GO:0098553)

integral component of luminal side of
endoplasmic reticulum membrane (GO:0071556)

MHC protein complex (GO:0042611)

ER to Golgi transport vesicle membrane
(GO:0012507)

coated vesicle membrane (GO:0030662)

transport vesicle membrane (GO:0030658)

TIM23 mitochondrial import inner membrane
translocase complex (GO:0005744)

phagocytic vesicle lumen (GO:0097013)

Lewy body (GO:0097413)

early endosome lumen (GO:0031905)

0

2

4

6

Enrichment log(combined score)

GO_Molecular_Function_2021

Term

leukotriene-C4 synthase activity
(GO:0004464)

MHC class II receptor activity (GO:0032395)

MHC class II protein complex binding
(GO:0023026)

MHC class II protein binding (GO:0042289)

CD4 receptor binding (GO:0042609)

RAGE receptor binding (GO:0050786)

cadherin binding involved in cell-cell
adhesion (GO:0098641)

phospholipase inhibitor activity
(GO:0004859)

water channel activity (GO:0015250)

water transmembrane transporter activity
(GO:0005372)

0

2

4

6

Enrichment log(combined score)