

GO_Biological_Process_2021

Term

determination of liver left/right asymmetry
(GO:0071910)

translational readthrough (GO:0006451)

selenocysteine incorporation (GO:0001514)

positive regulation of homotypic cell–cell
adhesion (GO:0034112)

cardiac atrium development (GO:0003230)

DNA geometric change (GO:0032392)

presynaptic membrane assembly (GO:0097105)

presynaptic membrane organization
(GO:0097090)

mitochondrion distribution (GO:0048311)

regulation of homotypic cell–cell adhesion
(GO:0034110)

negative regulation of histone methylation
(GO:0031061)

negative regulation of DNA–dependent DNA
replication (GO:2000104)

cytoplasmic sequestering of protein
(GO:0051220)

0

2

4

6

Enrichment log(combined score)

GO_Cellular_Component_2021

Term

apicolateral plasma membrane (GO:0016327)

Sec61 translocon complex (GO:0005784)

Lewy body (GO:0097413)

symmetric synapse (GO:0032280)

integral component of cytoplasmic side of
endoplasmic reticulum membrane (GO:0071458)

asymmetric, glutamatergic, excitatory
synapse (GO:0098985)

Cul5-RING ubiquitin ligase complex
(GO:0031466)

spanning component of membrane (GO:0089717)

spectrin-associated cytoskeleton
(GO:0014731)

inhibitory synapse (GO:0060077)

astrocyte projection (GO:0097449)

0

1

2

3

4

5

Enrichment log(combined score)

GO_Molecular_Function_2021

