

# GO\_Biological\_Process\_2021

Term

positive regulation of astrocyte  
differentiation (GO:0048711)

bicarbonate transport (GO:0015701)

regulation of astrocyte differentiation  
(GO:0048710)

axonal fasciculation (GO:0007413)

Roundabout signaling pathway (GO:0035385)

regulation of inositol trisphosphate  
biosynthetic process (GO:0032960)

regulation of catecholamine uptake involved  
in synaptic transmission (GO:0051940)

protein C-linked glycosylation via  
tryptophan (GO:0018317)

protein C-linked glycosylation via  
2'-alpha-mannosyl-L-tryptophan (GO:0018406)

negative regulation of plasminogen  
activation (GO:0010757)

negative regulation of B cell  
differentiation (GO:0045578)

His-Purkinje system development (GO:0003164)

ciliary neurotrophic factor-mediated  
signaling pathway (GO:0070120)

cell fate commitment involved in formation  
of primary germ layer (GO:0060795)

carbon dioxide transport (GO:0015670)

0

2

4

6

Enrichment log(combined score)

# GO\_Cellular\_Component\_2021

Term

sarcoglycan complex (GO:0016012)

mitochondrial alpha-ketoglutarate  
dehydrogenase complex (GO:0005947)

dystroglycan complex (GO:0016011)

Cdc73/Paf1 complex (GO:0016593)

anchored component of plasma membrane  
(GO:0046658)

0

2

4

Enrichment log(combined score)

# GO\_Molecular\_Function\_2021

Term

carbonate dehydratase activity (GO:0004089)

arylesterase activity (GO:0004064)

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

GPI-linked ephrin receptor activity  
(GO:0005004)

glutamate-gated calcium ion channel activity  
(GO:0022849)

dopamine neurotransmitter receptor activity  
(GO:0004952)

0

2

4

6

Enrichment log(combined score)