

# GO\_Biological\_Process\_2021

Term

transferrin transport (GO:0033572)

positive regulation of tau-protein kinase  
activity (GO:1902949)

phagosome acidification (GO:0090383)

relaxation of cardiac muscle (GO:0055119)

response to host defenses (GO:0052200)

modulation by symbiont of host defense  
response (GO:0052031)

intermediate filament bundle assembly  
(GO:0045110)

intracellular pH reduction (GO:0051452)

iron ion transport (GO:0006826)

regulation of cyclic-nucleotide  
phosphodiesterase activity (GO:0051342)

positive regulation of ryanodine-sensitive  
calcium-release channel activity  
(GO:0060316)

0

2

4

6

Enrichment log(combined score)

# GO\_Cellular\_Component\_2021

Term

neurofibrillary tangle (GO:0097418)

vacuolar proton-transporting V-type ATPase,  
V1 domain (GO:0000221)

proton-transporting V-type ATPase, V1 domain  
(GO:0033180)

mitochondrial respiratory chain complex III  
(GO:0005750)

vacuolar proton-transporting V-type ATPase  
complex (GO:0016471)

sodium:potassium-exchanging ATPase complex  
(GO:0005890)

apical dendrite (GO:0097440)

proton-transporting V-type ATPase complex  
(GO:0033176)

neuronal dense core vesicle (GO:0098992)

endoplasmic reticulum tubular network  
membrane (GO:0098826)

chromaffin granule (GO:0042583)

0

2

4

6

8

Enrichment log(combined score)

# GO\_Molecular\_Function\_2021

