

# GO\_Biological\_Process\_2021

Term

GPI anchor metabolic process (GO:0006505)

inner dynein arm assembly (GO:0036159)

GPI anchor biosynthetic process (GO:0006506)

preassembly of GPI anchor in ER membrane  
(GO:0016254)

glycolipid biosynthetic process (GO:0009247)

fatty acid beta-oxidation using acyl-CoA  
dehydrogenase (GO:0033539)

glycerophospholipid metabolic process  
(GO:0006650)

central nervous system projection neuron  
axonogenesis (GO:0021952)

cellular component maintenance (GO:0043954)

protein lipidation (GO:0006497)

0

2

4

6

Enrichment log(combined score)

# GO\_Cellular\_Component\_2021

Term

transcription factor TFIIH holo complex  
(GO:0005675)

tRNA methyltransferase complex (GO:0043527)

nuclear outer membrane (GO:0005640)

transcription factor TFIIIC complex  
(GO:0000127)

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

gamma-secretase complex (GO:0070765)

astral microtubule (GO:0000235)

extrinsic component of mitochondrial inner  
membrane (GO:0031314)

MOZ/MORF histone acetyltransferase complex  
(GO:0070776)

MLL1/2 complex (GO:0044665)

MLL1 complex (GO:0071339)

0

2

4

Enrichment log(combined score)

# GO\_Molecular\_Function\_2021

Term

H3 histone acetyltransferase activity  
(GO:0010484)

aspartic endopeptidase activity,  
intramembrane cleaving (GO:0042500)

acyl-CoA dehydrogenase activity (GO:0003995)

mRNA methyltransferase activity (GO:0008174)

microtubule minus-end binding (GO:0051011)

aspartic-type endopeptidase activity  
(GO:0004190)

nucleotide phosphatase activity, acting on  
free nucleotides (GO:0098519)

histone acetyltransferase activity (H3-K23  
specific) (GO:0043994)

cAMP-dependent protein kinase activity  
(GO:0004691)

cyclic nucleotide-dependent protein kinase  
activity (GO:0004690)

acetyl-CoA C-acyltransferase activity  
(GO:0003988)

1-acylglycerophosphocholine  
O-acyltransferase activity (GO:0047184)

0

2

4

6

Enrichment log(combined score)