

# GO\_Biological\_Process\_2021

Term

regulation of focal adhesion disassembly  
(GO:0120182)

positive regulation of focal adhesion  
disassembly (GO:0120183)

positive regulation of cardiac muscle cell  
proliferation (GO:0060045)

response to epinephrine (GO:0071871)

plasma membrane phospholipid scrambling  
(GO:0017121)

cellular response to epinephrine stimulus  
(GO:0071872)

positive regulation of cardiac muscle tissue  
growth (GO:0055023)

regulation of transmembrane transporter  
activity (GO:0022898)

positive regulation of keratinocyte  
migration (GO:0051549)

negative regulation of purine nucleotide  
metabolic process (GO:1900543)

negative regulation of glycolytic process  
(GO:0045820)

0

2

4

6

Enrichment log(combined score)

# GO\_Cellular\_Component\_2021

Term

protein kinase complex (GO:1902911)

HFE–transferrin receptor complex  
(GO:1990712)

transferase complex, transferring  
phosphorus–containing groups (GO:0061695)

intermediate filament cytoskeleton  
(GO:0045111)

chloride channel complex (GO:0034707)

zonula adherens (GO:0005915)

pinosome (GO:0044352)

macropinosome (GO:0044354)

hemidesmosome (GO:0030056)

excitatory synapse (GO:0060076)

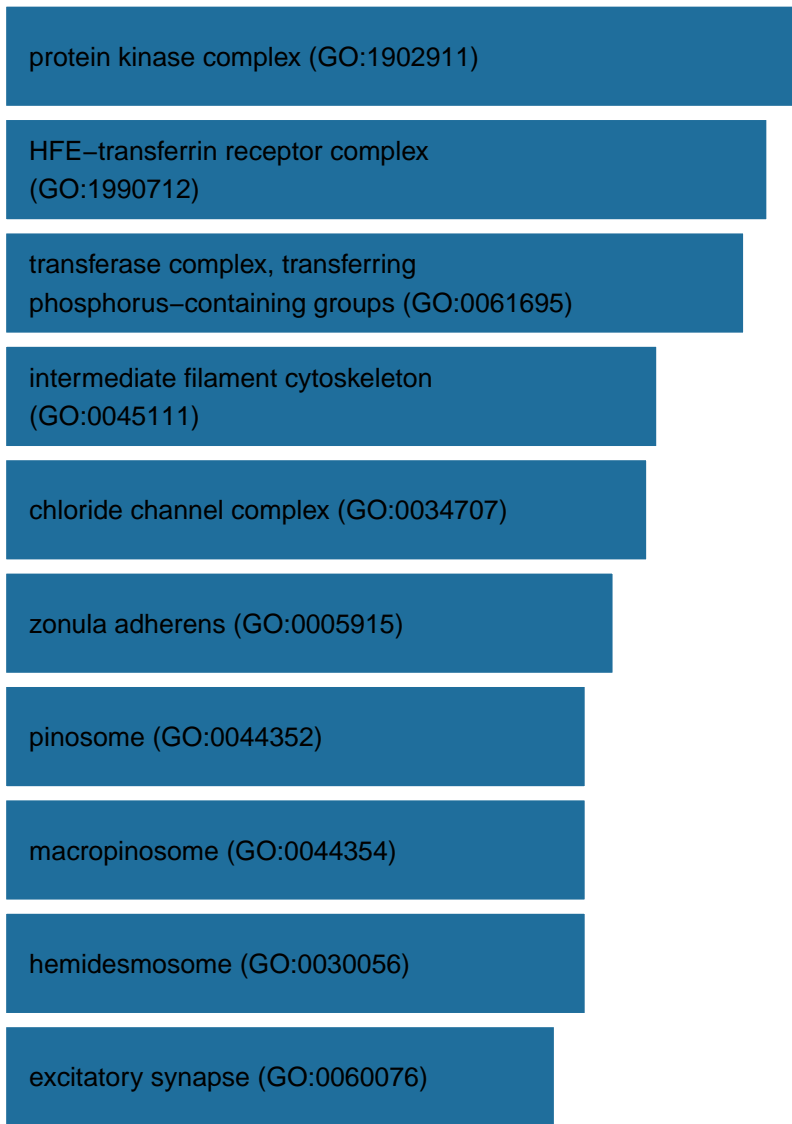
0

2

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6

Enrichment log(combined score)



# GO\_Molecular\_Function\_2021

Term

transforming growth factor beta receptor  
activity, type I (GO:0005025)

microtubule plus-end binding (GO:0051010)

phospholipid scramblase activity  
(GO:0017128)

transforming growth factor beta-activated  
receptor activity (GO:0005024)

protein kinase C activity (GO:0004697)

phosphotransferase activity, nitrogenous  
group as acceptor (GO:0016775)

fibroblast growth factor-activated receptor  
activity (GO:0005007)

creatine kinase activity (GO:0004111)

calmodulin-dependent protein phosphatase  
activity (GO:0033192)

C-X-C chemokine binding (GO:0019958)

0

2

4

6

Enrichment log(combined score)