

# GO\_Biological\_Process\_2021

Term

negative regulation of sodium ion  
transmembrane transporter activity  
(GO:2000650)

negative regulation of sodium ion  
transmembrane transport (GO:1902306)

regulation of relaxation of cardiac muscle  
(GO:1901897)

positive regulation of ion transmembrane  
transport (GO:0034767)

regulation of protein localization to early  
endosome (GO:1902965)

positive regulation of protein localization  
to endosome (GO:1905668)

positive regulation of protein localization  
to early endosome (GO:1902966)

filopodium assembly (GO:0046847)

negative regulation of actin filament bundle  
assembly (GO:0032232)

actin-mediated cell contraction (GO:0070252)

0

2

4

6

Enrichment log(combined score)

# GO\_Cellular\_Component\_2021

Term

sodium channel complex (GO:0034706)

sarcoglycan complex (GO:0016012)

intrinsic component of presynaptic membrane  
(GO:0098889)

chloride channel complex (GO:0034707)

Lewy body (GO:0097413)

dystroglycan complex (GO:0016011)

integral component of presynaptic membrane  
(GO:0099056)

activin receptor complex (GO:0048179)

integral component of synaptic membrane  
(GO:0099699)

astrocyte projection (GO:0097449)

0

2

4

Enrichment log(combined score)

# GO\_Molecular\_Function\_2021

Term

protein kinase B binding (GO:0043422)

ligand-gated calcium channel activity  
(GO:0099604)

intracellular ligand-gated ion channel  
activity (GO:0005217)

cAMP binding (GO:0030552)

3',5'-cyclic-AMP phosphodiesterase activity  
(GO:0004115)

titin binding (GO:0031432)

GTPase activator activity (GO:0005096)

kainate selective glutamate receptor  
activity (GO:0015277)

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

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

ADP-ribose diphosphatase activity  
(GO:0047631)

3-chloroallyl aldehyde dehydrogenase  
activity (GO:0004028)

0

2

4

6

Enrichment log(combined score)