

# GO\_Biological\_Process\_2021

Term

regulation of dendritic spine maintenance  
(GO:1902950)

negative regulation of smooth muscle  
contraction (GO:0045986)

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

filopodium assembly (GO:0046847)

negative regulation of amyloid-beta  
formation (GO:1902430)

regulation of glutamate receptor signaling  
pathway (GO:1900449)

positive regulation of non-membrane spanning  
protein tyrosine kinase activity  
(GO:1903997)

negative regulation of cellular response to  
hypoxia (GO:1900038)

negative regulation of amyloid precursor  
protein catabolic process (GO:1902992)

cellular monovalent inorganic cation  
homeostasis (GO:0030004)

0

2

4

6

Enrichment log(combined score)

# GO\_Cellular\_Component\_2021

Term

glial cell projection (GO:0097386)

astrocyte projection (GO:0097449)

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

smooth endoplasmic reticulum membrane  
(GO:0030868)

phosphatidylinositol 3-kinase complex, class  
I (GO:0097651)

NURF complex (GO:0016589)

cation-transporting ATPase complex  
(GO:0090533)

catenin complex (GO:0016342)

platelet dense granule membrane (GO:0031088)

nuclear envelope lumen (GO:0005641)

0

2

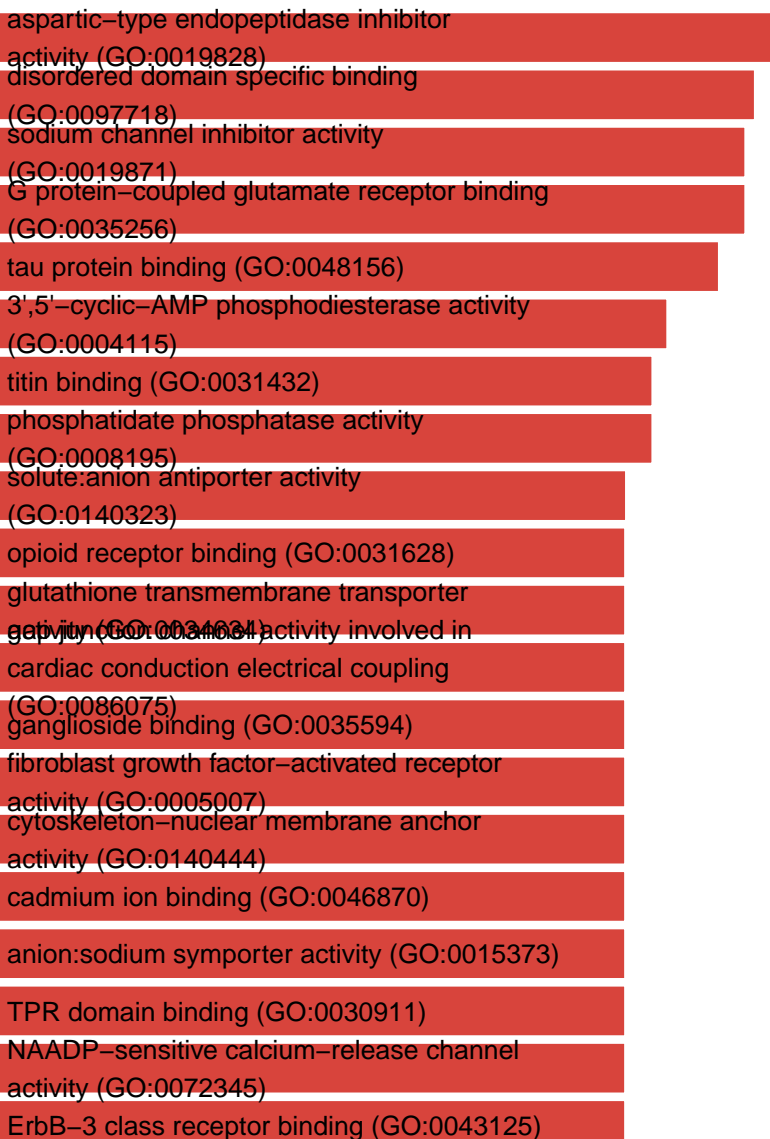
4

6

Enrichment log(combined score)

# GO\_Molecular\_Function\_2021

Term



0

2

4

6

Enrichment log(combined score)