

# GO\_Biological\_Process\_2021

Term

positive regulation of type I  
interferon-mediated signaling pathway  
(GO:0060340)

positive regulation of monocyte  
differentiation (GO:0045657)

regulation of trans-synaptic signaling  
(GO:0099177)

purine ribonucleoside triphosphate metabolic  
process (GO:0009205)

central nervous system projection neuron  
axonogenesis (GO:0021952)

regulation of cAMP-mediated signaling  
(GO:0043949)

positive regulation of ion transmembrane  
transporter activity (GO:0032414)

negative regulation of pri-miRNA  
transcription by RNA polymerase II  
(GO:1902894)

regulation of aerobic respiration  
(GO:1903715)

GTP metabolic process (GO:0046039)

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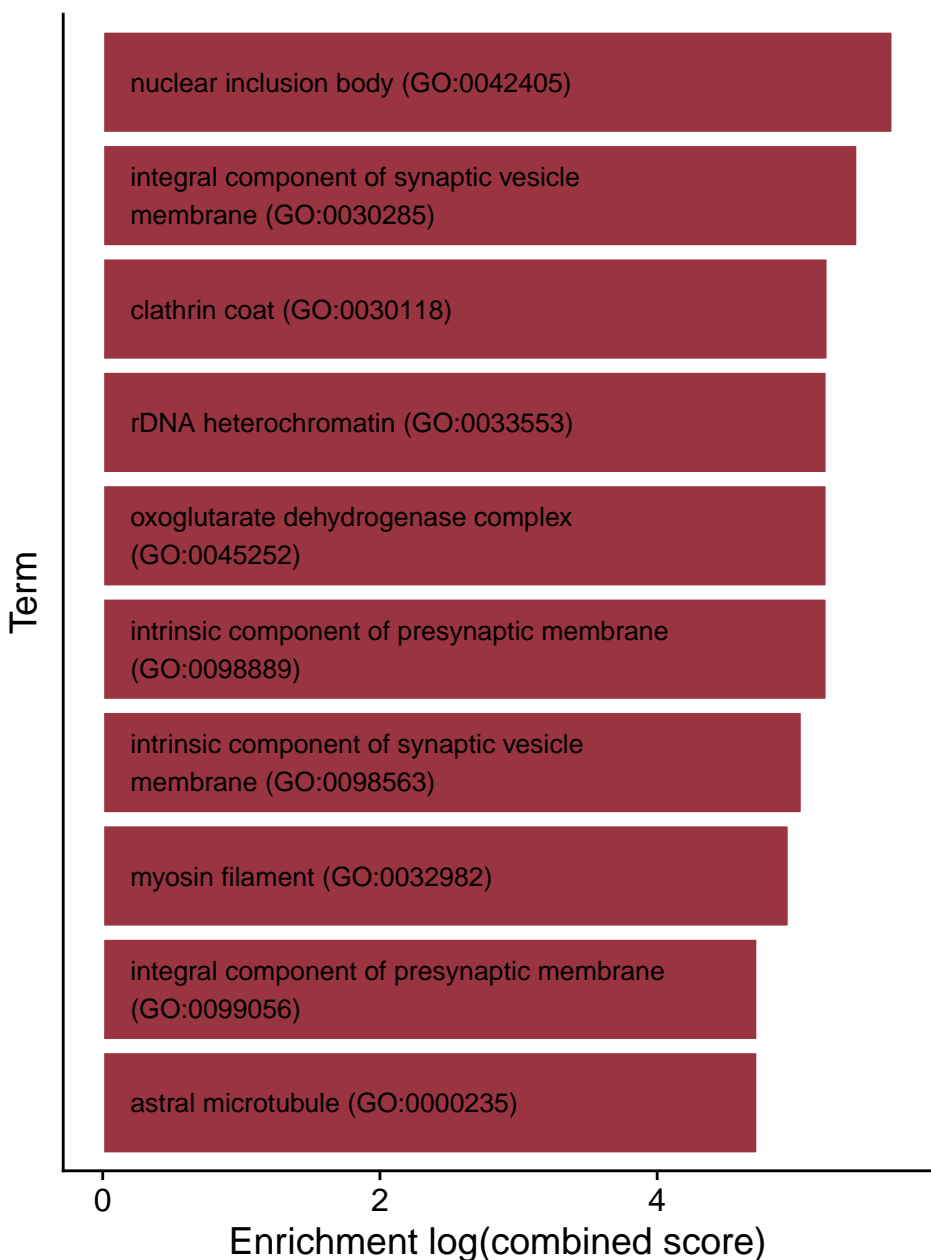
2

4

6

Enrichment log(combined score)

# GO\_Cellular\_Component\_2021



# GO\_Molecular\_Function\_2021

Term

kainate selective glutamate receptor  
activity (GO:0015277)

unmethylated CpG binding (GO:0045322)

RNA polymerase I transcription regulatory  
region sequence-specific DNA binding  
(GO:0001163)

RNA polymerase I core promoter  
sequence-specific DNA binding (GO:0001164)

lysine-acetylated histone binding  
(GO:0070577)

acetylation-dependent protein binding  
(GO:0140033)

RNA 7-methylguanosine cap binding  
(GO:0000340)

opioid receptor binding (GO:0031628)

inorganic phosphate transmembrane  
transporter activity (GO:0005315)

beta-2 adrenergic receptor binding  
(GO:0031698)

acyl carnitine transmembrane transporter  
activity (GO:0015227)

L-lysine transmembrane transporter activity  
(GO:0015189)

0

2

4

6

Enrichment log(combined score)