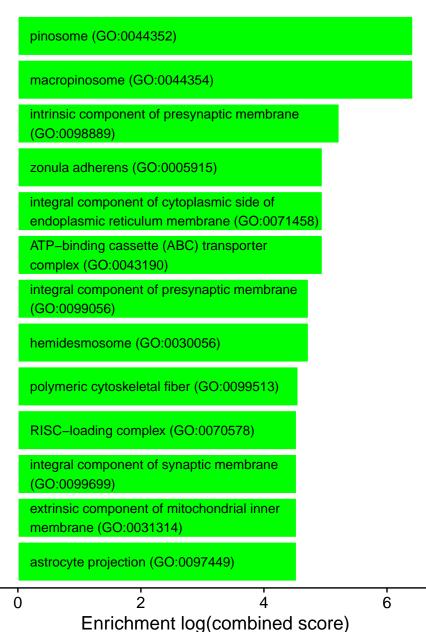
GO_Biological_Process_2021

```
regulation of sterol biosynthetic process
(GO:0106118)
myelin assembly (GO:0032288)
regulation of cholesterol metabolic process
(GO:0090181)
Golgi to lysosome transport (GO:0090160)
regulation of alcohol biosynthetic process
(GO:1902930)
glycogen catabolic process (GO:0005980)
glucan catabolic process (GO:0009251)
SA node cell to atrial cardiac muscle cell
signaling (GO:0086018)
response to acetylcholine (GO:1905144) regulation of low-density lipoprotein
particle receptor catabolic process
(@@la0ga&ogntracellular cholesterol
transport (GO:0032383) regulation of germinal ce
response to vascular
endothelial growth factor stimulus regulation of basement membrane assembly
(GO:1902547)
myolyed in embryonic body morphogenesis
positive registration of histone H3-K27
methylation (GO:0061087)
positive regulation of cAMP-dependent
postewekings lettorivotybasenzeno onembrane
assembly involved in embryonic body
negrative regulation of soldingolipid
biosynthetic process (GO:0090155)
membrane depolarization during SA node cell
action potential (GO:0086046)
L-aspartate transmembrane h
                                e transport
(GO:0070778)
beta–catenin destruction complex assembly
(GO:1904885)
                                                                          6
```

Enrichment log(combined score)



GO_Molecular_Function_2021

