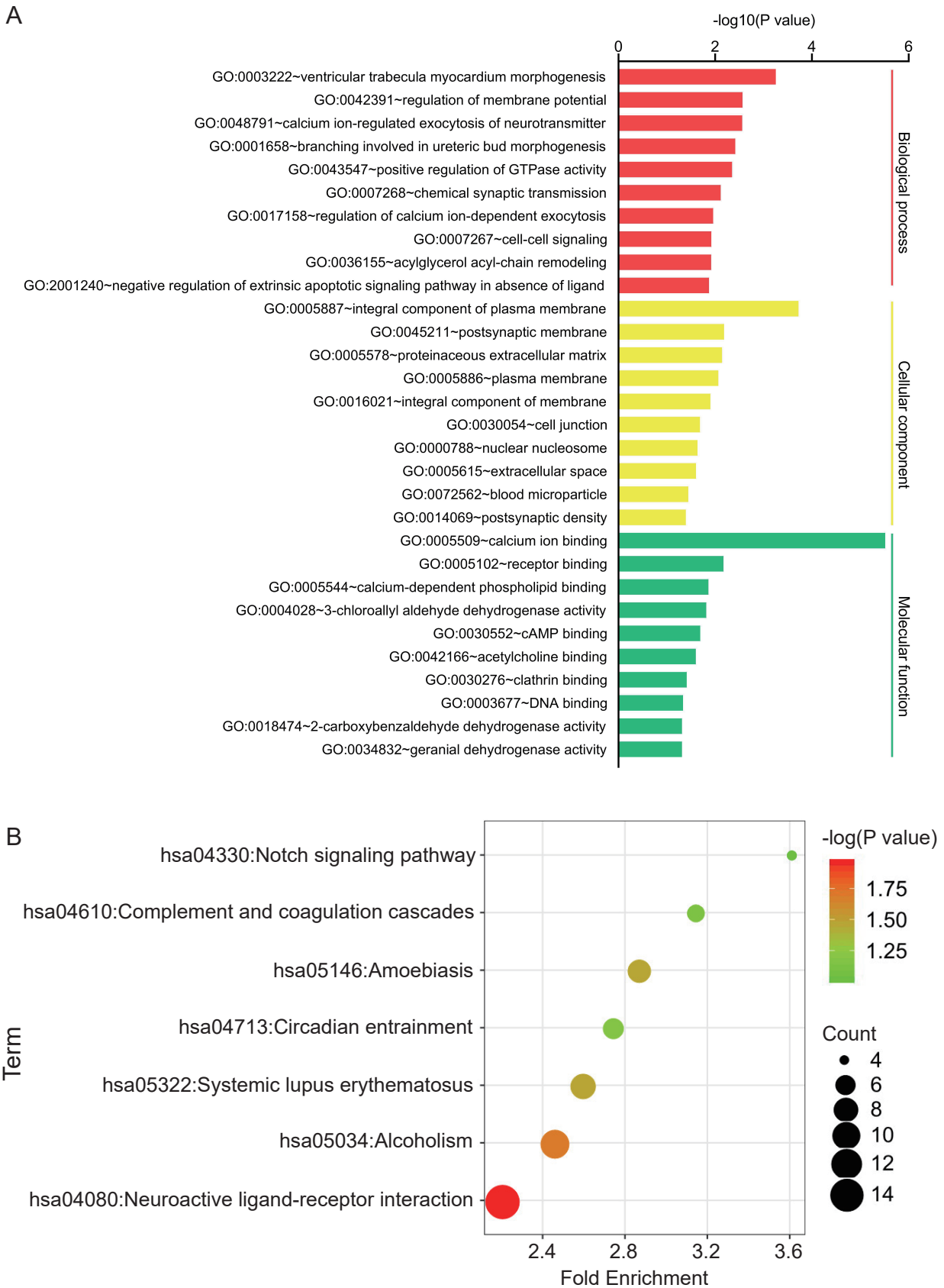
A
-log10(P value)
0
2
4
6
GO:0003222~ventricular trabecula myocardium morphogenesis
GO:0042391~regulation of membrane potential
GO:0048791~calcium ion-regulated exocytosis of neurotransmitter
GO:0001658~branching involved in ureteric bud morphogenesis
GO:0043547~positive regulation of GTPase activity
GO:0007268~chemical synaptic transmission
GO:0017158~regulation of calcium ion-dependent exocytosis
GO:0007267~cell-cell signaling
GO:0036155~acylglycerol acyl-chain remodeling
GO:2001240~negative regulation of extrinsic apoptotic signaling pathway in absence of ligand
GO:0005887~integral component of plasma membrane
GO:0045211~postsynaptic membrane
GO:0005578~proteinaceous extracellular matrix
GO:0005886~plasma membrane
GO:0016021~integral component of membrane
GO:0030054~cell junction
GO:0000788~nuclear nucleosome
GO:0005615~extracellular space
GO:0072562~blood microparticle
GO:0014069~postsynaptic density
GO:0005509~calcium ion binding
GO:0005102~receptor binding
GO:0005544~calcium-dependent phospholipid binding
GO:0004028~3-chloroallyl aldehyde dehydrogenase activity
GO:0030552~cAMP binding
GO:0042166~acetylcholine binding
GO:0030276~clathrin binding
GO:0003677~DNA binding
GO:0018474~2-carboxybenzaldehyde dehydrogenase activity
GO:0034832~geranial dehydrogenase activity
Biological process
Cellular component
Molecular function
B
Term
hsa04330:Notch signaling pathway
hsa04610:Complement and coagulation cascades
hsa05146:Amoebiasis
hsa04713:Circadian entrainment
hsa05322:Systemic lupus erythematosus
hsa05034:Alcoholism
hsa04080:Neuroactive ligand-receptor interaction
2.4
2.8
3.2
3.6
Fold Enrichment
-log(P value)
1.75
1.50
1.25
Count
4
6
8
10
12
14