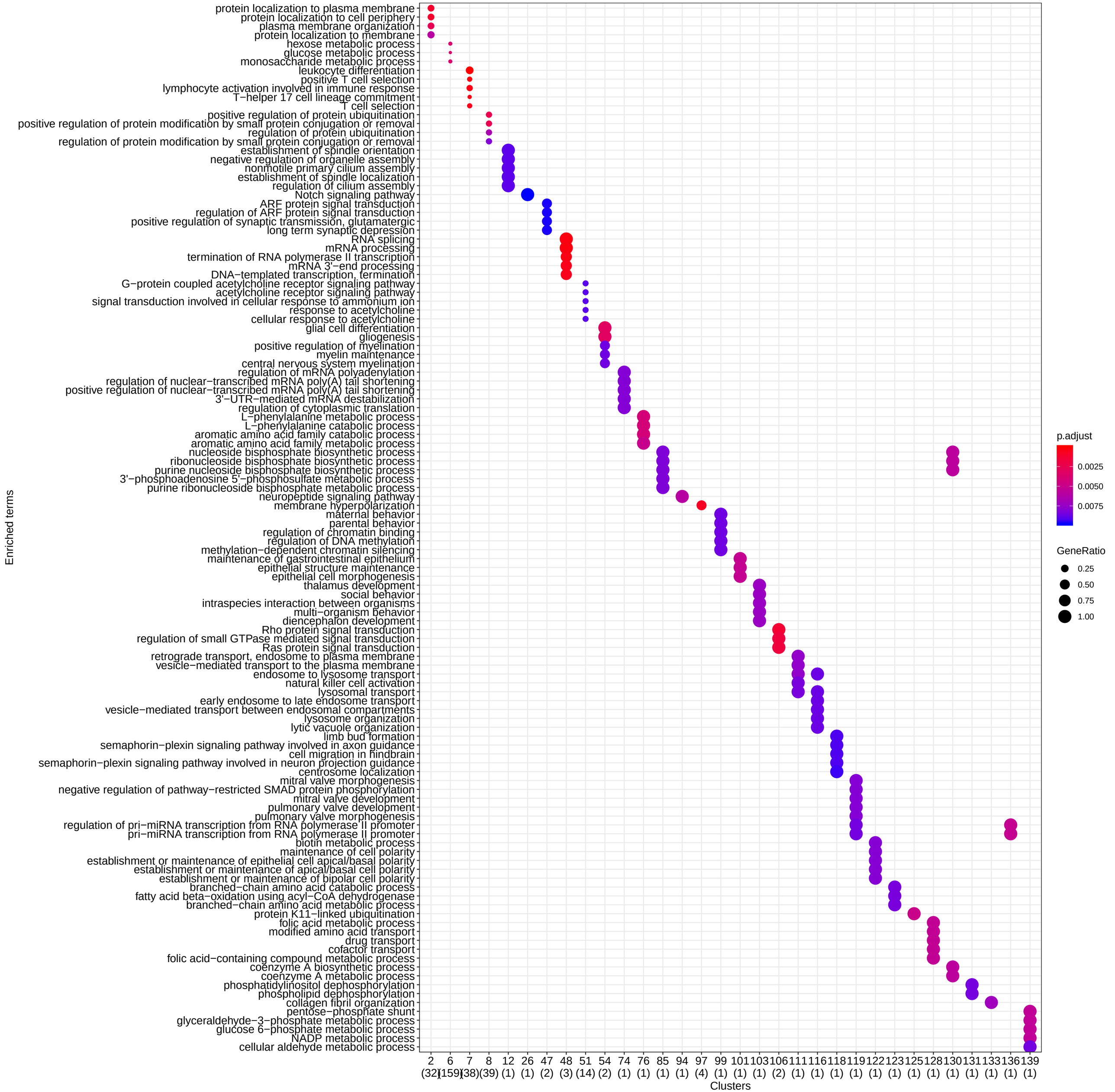
Enriched terms
protein localization to plasma membrane
protein localization to cell periphery
plasma membrane organization
protein localization to membrane
hexose metabolic process
glucose metabolic process
monosaccharide metabolic process
leukocyte differentiation
positive T cell selection
lymphocyte activation involved in immune response
T−helper 17 cell lineage commitment
T cell selection
positive regulation of protein ubiquitination
positive regulation of protein modification by small protein conjugation or removal
regulation of protein ubiquitination
regulation of protein modification by small protein conjugation or removal
establishment of spindle orientation
negative regulation of organelle assembly
nonmotile primary cilium assembly
establishment of spindle localization
regulation of cilium assembly
Notch signaling pathway
ARF protein signal transduction
regulation of ARF protein signal transduction
positive regulation of synaptic transmission, glutamatergic
long term synaptic depression
RNA splicing
mRNA processing
termination of RNA polymerase II transcription
mRNA 3'−end processing
DNA−templated transcription, termination
G−protein coupled acetylcholine receptor signaling pathway
acetylcholine receptor signaling pathway
signal transduction involved in cellular response to ammonium ion
response to acetylcholine
cellular response to acetylcholine
glial cell differentiation
gliogenesis
positive regulation of myelination
myelin maintenance
central nervous system myelination
regulation of mRNA polyadenylation
regulation of nuclear−transcribed mRNA poly(A) tail shortening
positive regulation of nuclear−transcribed mRNA poly(A) tail shortening
3'−UTR−mediated mRNA destabilization
regulation of cytoplasmic translation
L−phenylalanine metabolic process
L−phenylalanine catabolic process
aromatic amino acid family catabolic process
aromatic amino acid family metabolic process
nucleoside bisphosphate biosynthetic process
ribonucleoside bisphosphate biosynthetic process
purine nucleoside bisphosphate biosynthetic process
3'−phosphoadenosine 5'−phosphosulfate metabolic process
purine ribonucleoside bisphosphate metabolic process
neuropeptide signaling pathway
membrane hyperpolarization
maternal behavior
parental behavior
regulation of chromatin binding
regulation of DNA methylation
methylation−dependent chromatin silencing
maintenance of gastrointestinal epithelium
epithelial structure maintenance
epithelial cell morphogenesis
thalamus development
social behavior
intraspecies interaction between organisms
multi−organism behavior
diencephalon development
Rho protein signal transduction
regulation of small GTPase mediated signal transduction
Ras protein signal transduction
retrograde transport, endosome to plasma membrane
vesicle−mediated transport to the plasma membrane
endosome to lysosome transport
natural killer cell activation
lysosomal transport
early endosome to late endosome transport
vesicle−mediated transport between endosomal compartments
lysosome organization
lytic vacuole organization
limb bud formation
semaphorin−plexin signaling pathway involved in axon guidance
cell migration in hindbrain
semaphorin−plexin signaling pathway involved in neuron projection guidance
centrosome localization
mitral valve morphogenesis
negative regulation of pathway−restricted SMAD protein phosphorylation
mitral valve development
pulmonary valve development
pulmonary valve morphogenesis
regulation of pri−miRNA transcription from RNA polymerase II promoter
pri−miRNA transcription from RNA polymerase II promoter
biotin metabolic process
maintenance of cell polarity
establishment or maintenance of epithelial cell apical/basal polarity
establishment or maintenance of apical/basal cell polarity
establishment or maintenance of bipolar cell polarity
branched−chain amino acid catabolic process
fatty acid beta−oxidation using acyl−CoA dehydrogenase
branched−chain amino acid metabolic process
protein K11−linked ubiquitination
folic acid metabolic process
modified amino acid transport
drug transport
cofactor transport
folic acid−containing compound metabolic process
coenzyme A biosynthetic process
coenzyme A metabolic process
phosphatidylinositol dephosphorylation
phospholipid dephosphorylation
collagen fibril organization
pentose−phosphate shunt
glyceraldehyde−3−phosphate metabolic process
glucose 6−phosphate metabolic process
NADP metabolic process
cellular aldehyde metabolic process
2 (32) 6 (159) 7 (38) 8 (39) 12 (1) 26 (1) 47 (2) 48 (3) 51 (14) 54 (2) 74 (1) 76 (1) 85 (1) 94 (1) 97 (4) 99 (1) 101 (1) 103 (1) 106 (2) 111 (1) 116 (1) 118 (1) 119 (1) 122 (1) 123 (1) 125 (1) 128 (1) 130 (1) 131 (1) 133 (1) 136 (1) 139 (1)
Clusters
p.adjust
0.0025
0.0050
0.0075
GeneRatio
0.25
0.50
0.75
1.00