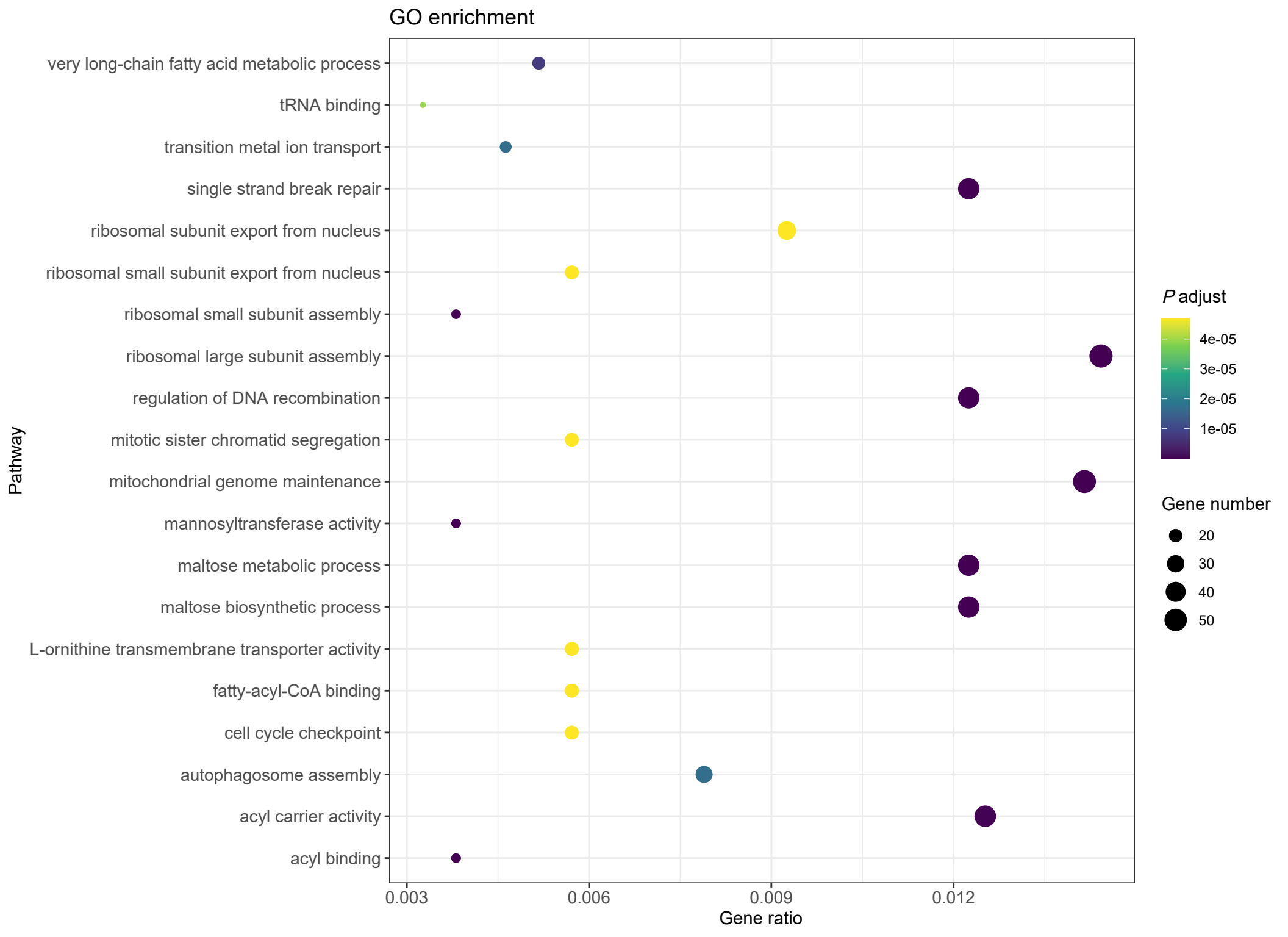
GO enrichment
very long-chain fatty acid metabolic process
tRNA binding
transition metal ion transport
single strand break repair
ribosomal subunit export from nucleus
ribosomal small subunit export from nucleus
ribosomal small subunit assembly
ribosomal large subunit assembly
regulation of DNA recombination
mitotic sister chromatid segregation
mitochondrial genome maintenance
mannosyltransferase activity
maltose metabolic process
maltose biosynthetic process
L-ornithine transmembrane transporter activity
fatty-acyl-CoA binding
cell cycle checkpoint
autophagosome assembly
acyl carrier activity
acyl binding
Pathway
0.003
0.006
0.009
0.012
Gene ratio
P adjust
4e-05
3e-05
2e-05
1e-05
Gene number
20
30
40
50