GeneRatio

0.02

regulation of DNA-binding transcription factor activity

reactive oxygen species metabolic process

response to fungus

axoneme assembly

cilium movement

cilium or flagellum-dependent cell motility

microtubule-based movement

microtubule-based process

positive regulation of NF-kappaB transcription factor activity

defense response to fungus

cilium-dependent cell motility

positive regulation of response to external stimulus

carboxylic acid transport

response to oxygen-containing compound

O 25 () 50

number of genes

p.adjust 0.0015

0.0010

0.0005

blood circulation

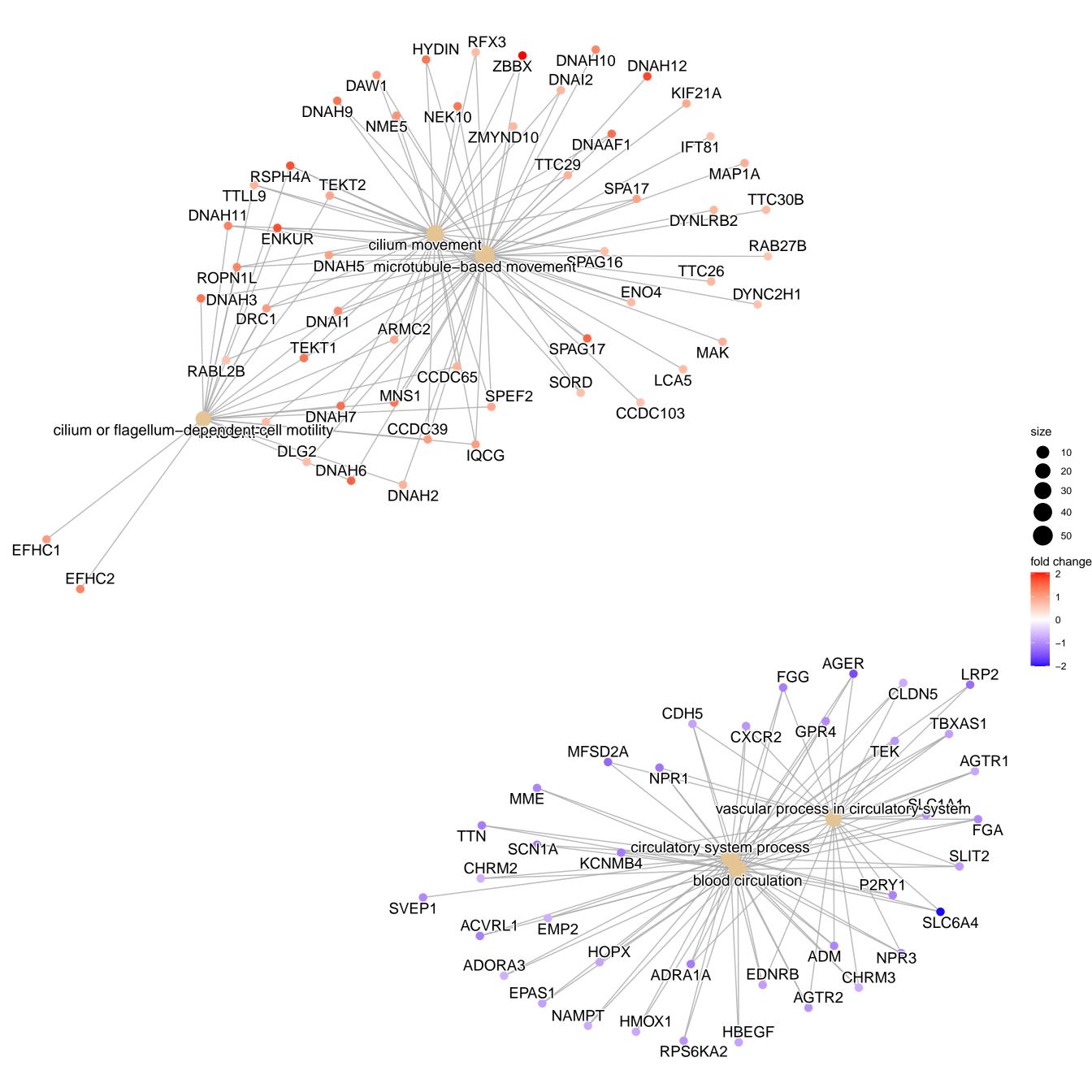
circulatory system process

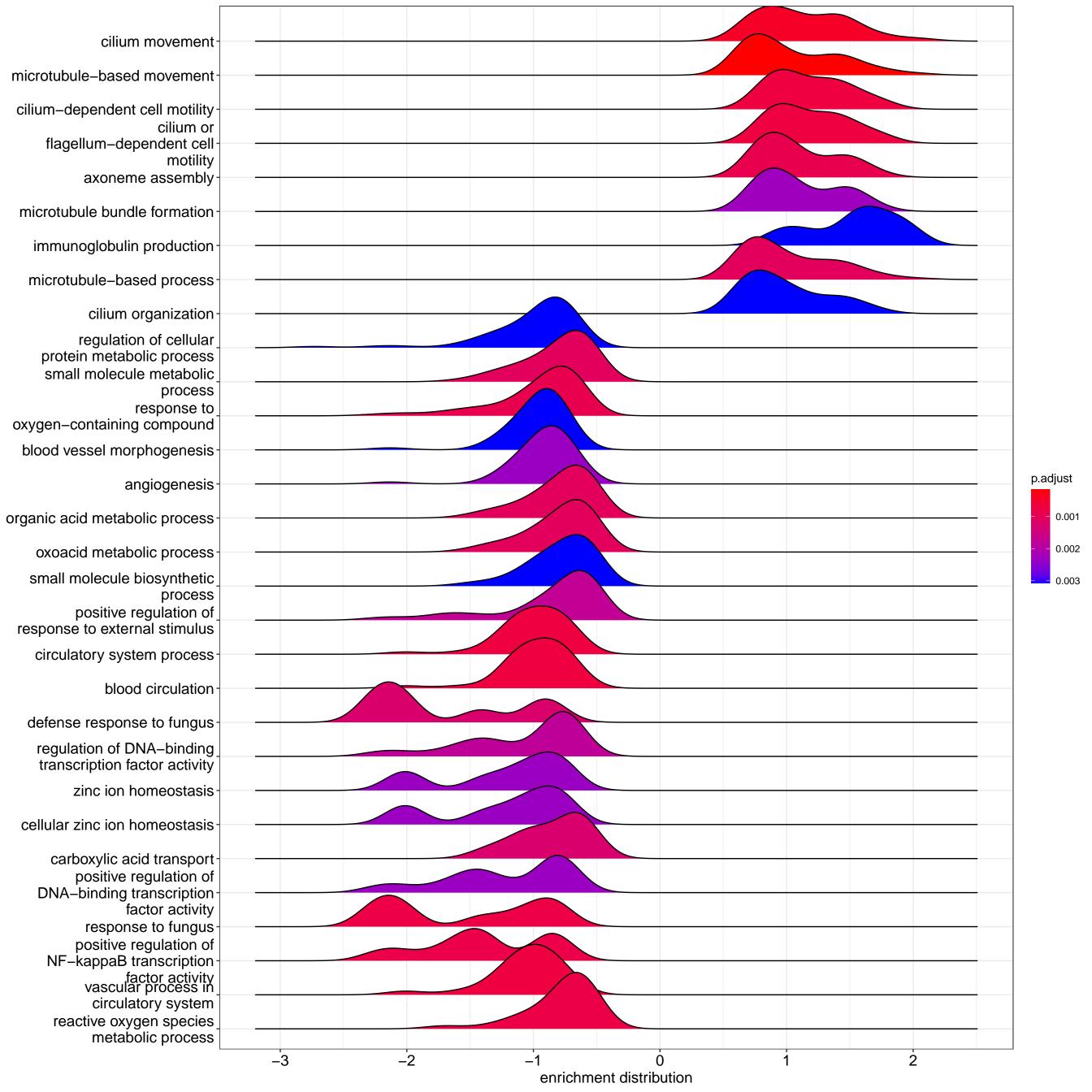
vascular process in circulatory system

small molecule metabolic process

oxoacid metabolic process

organic acid metabolic process

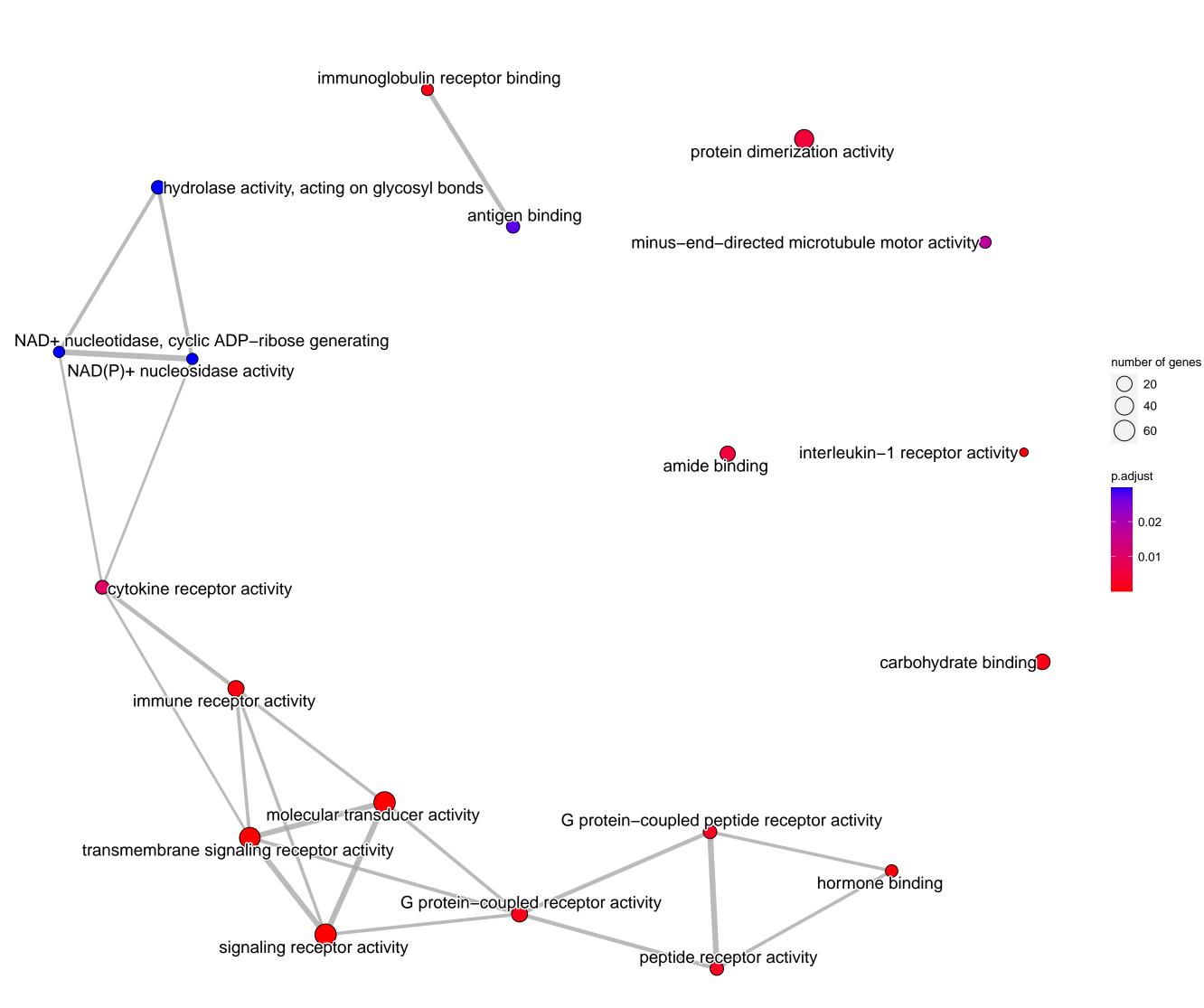


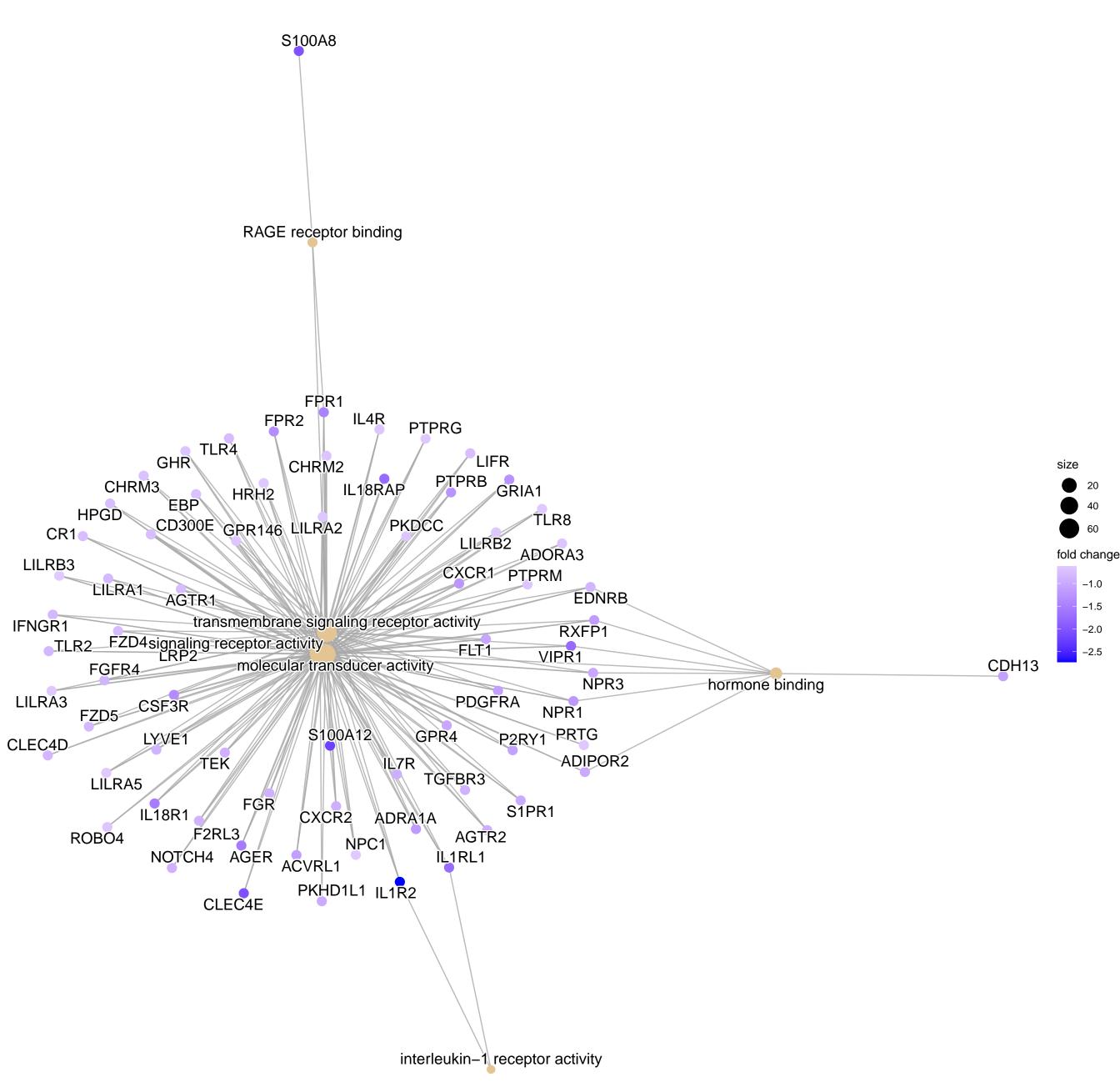


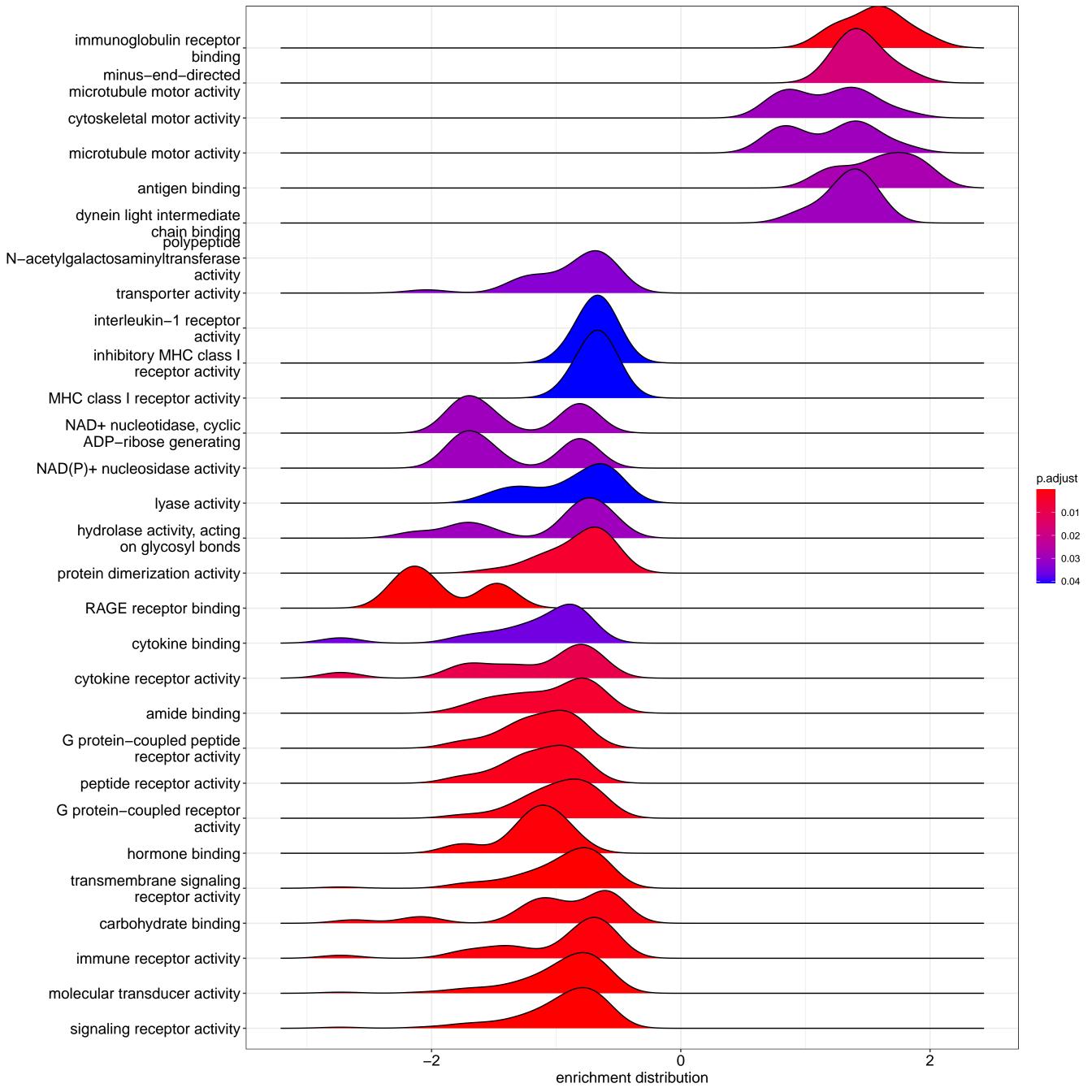
0.03

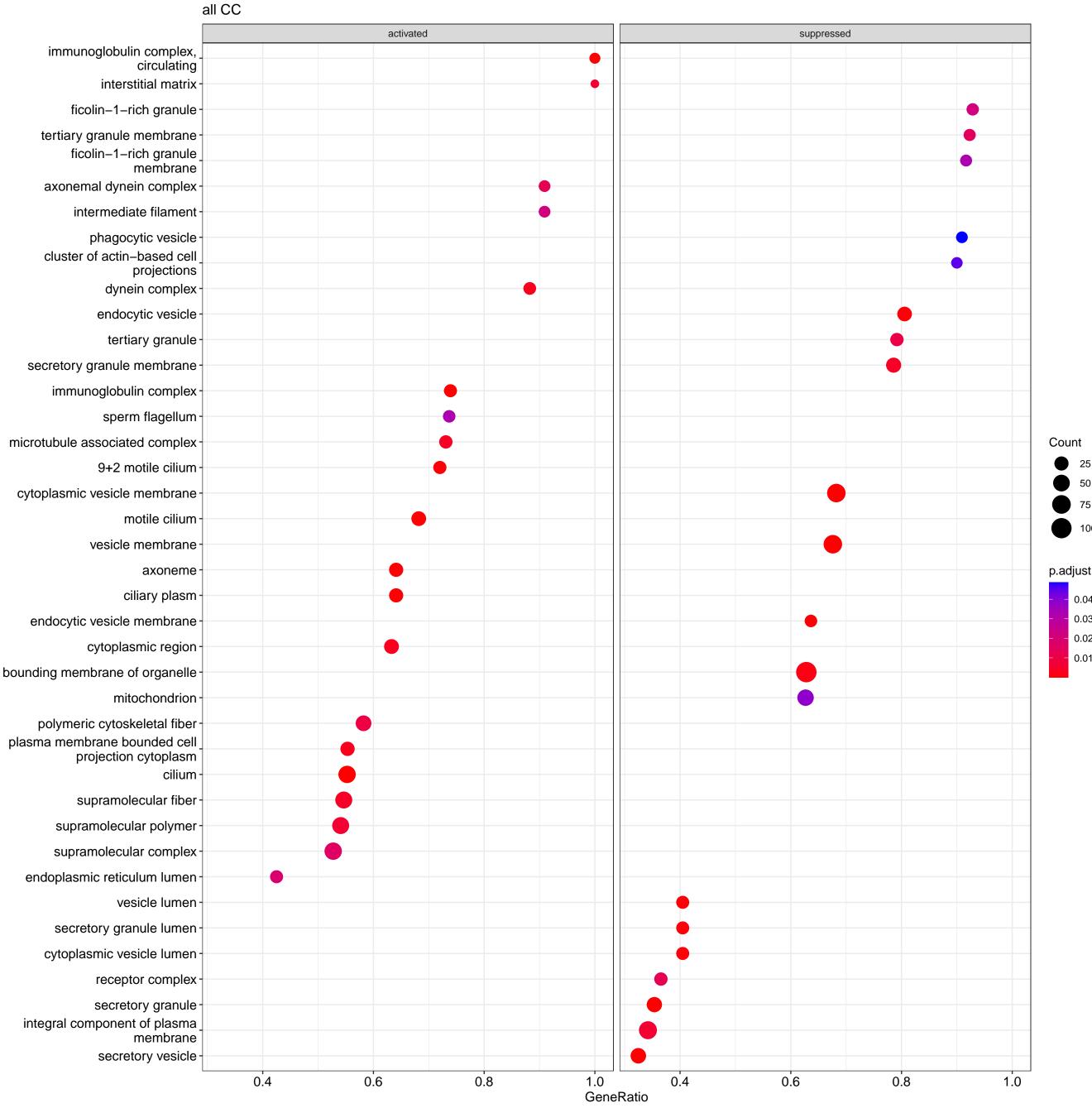
0.02

RAGE receptor binding



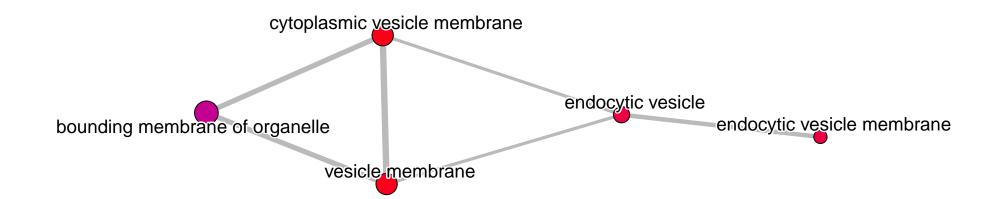


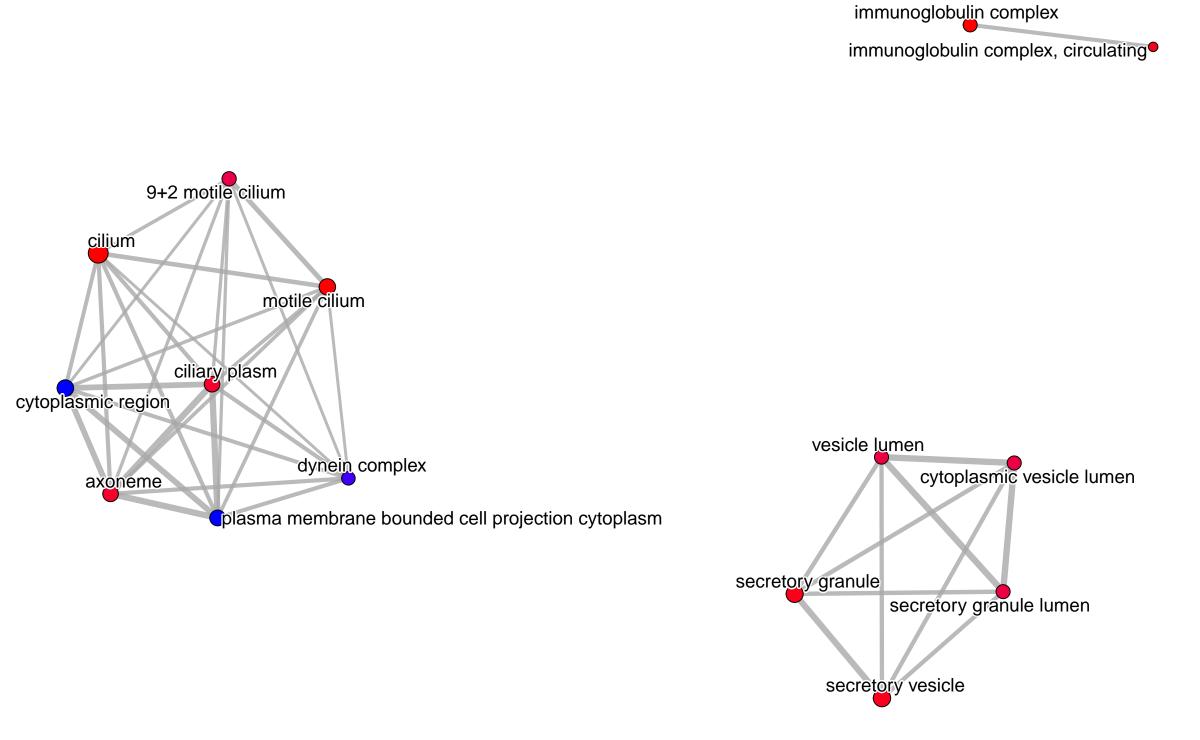




0.03

0.02



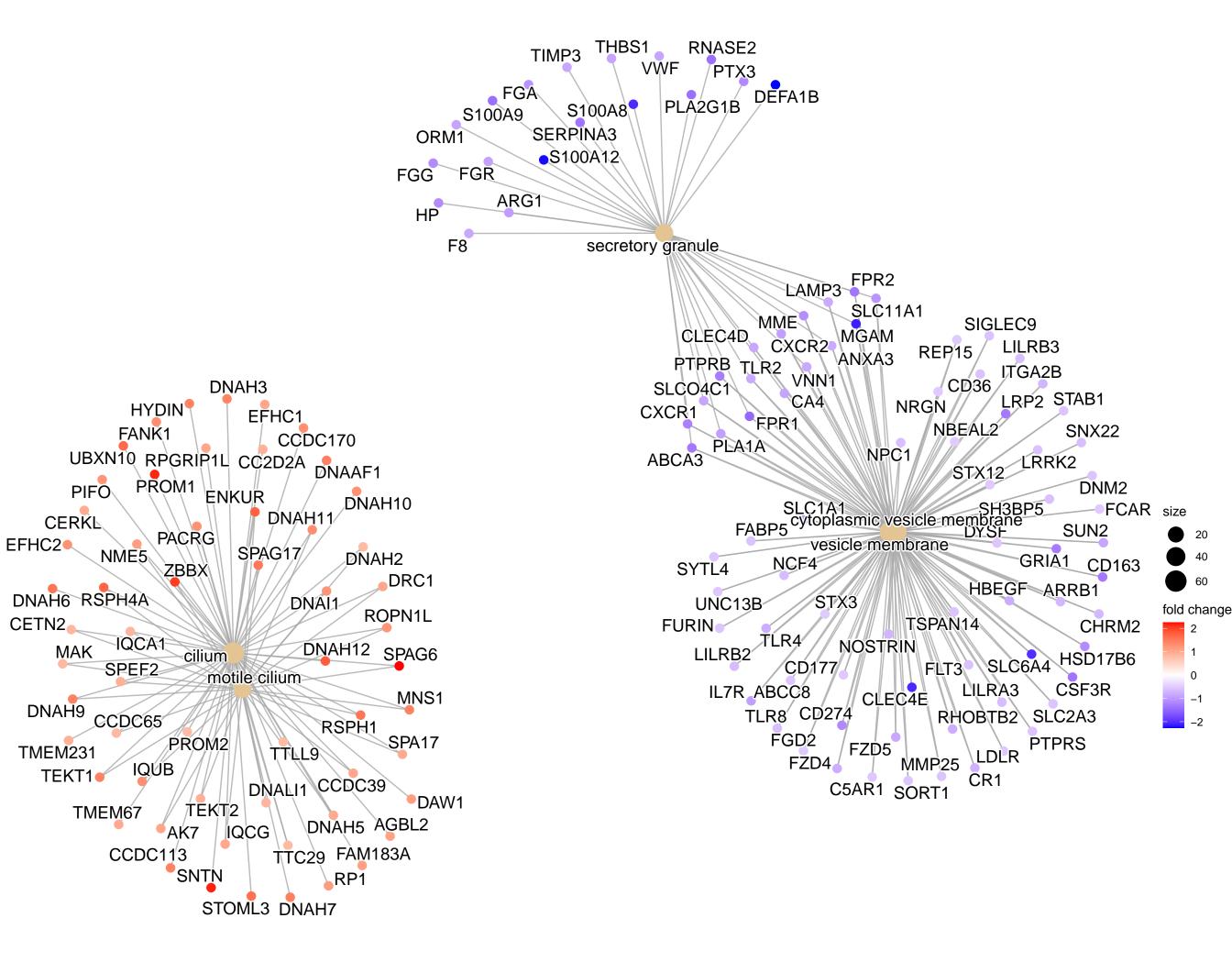


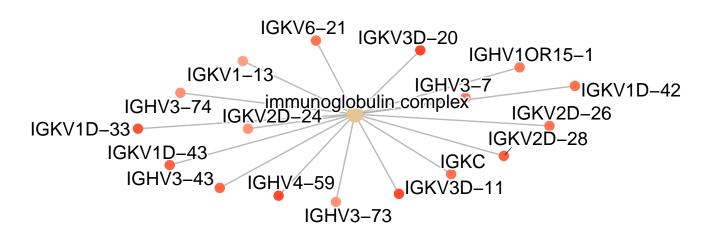
number of genes

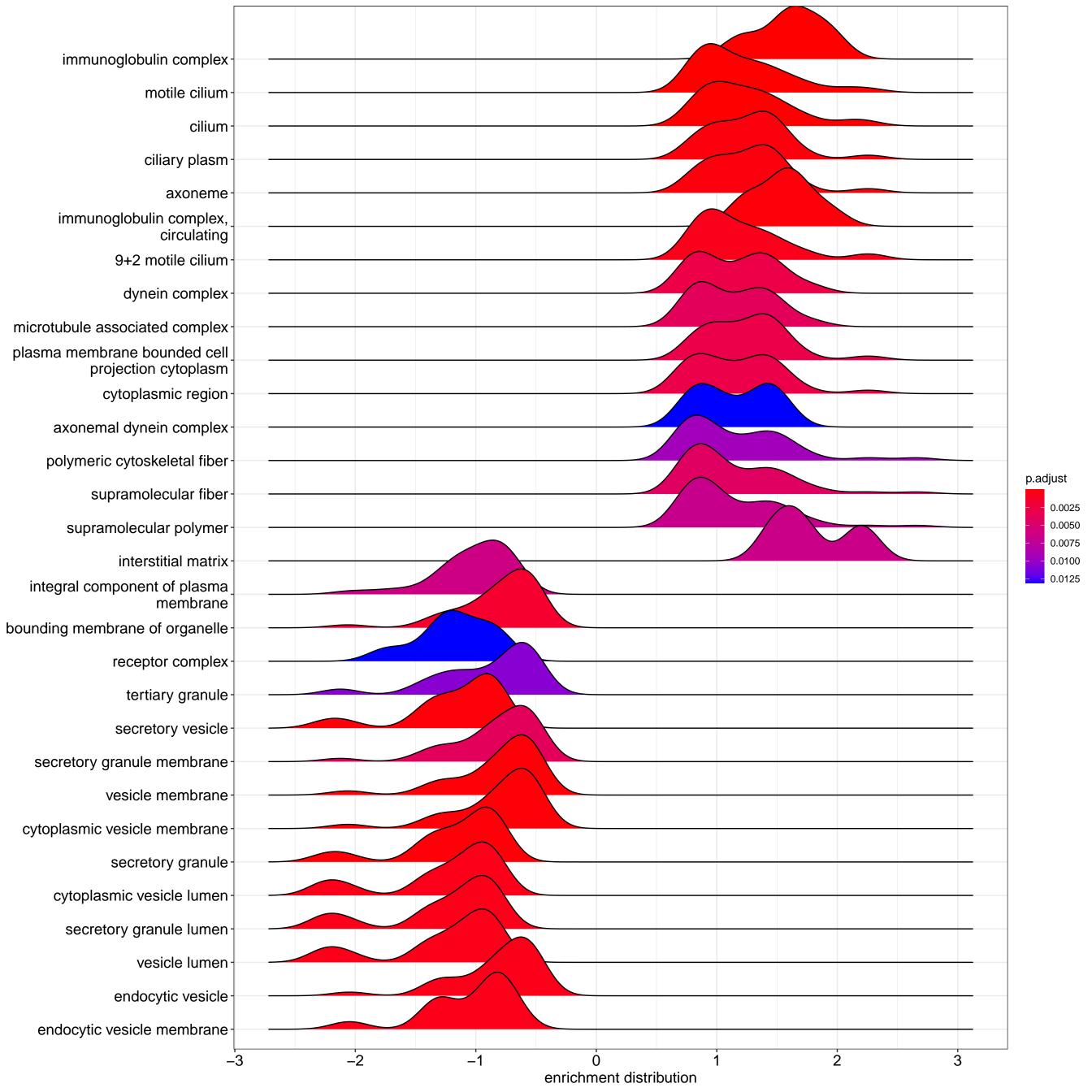
255075

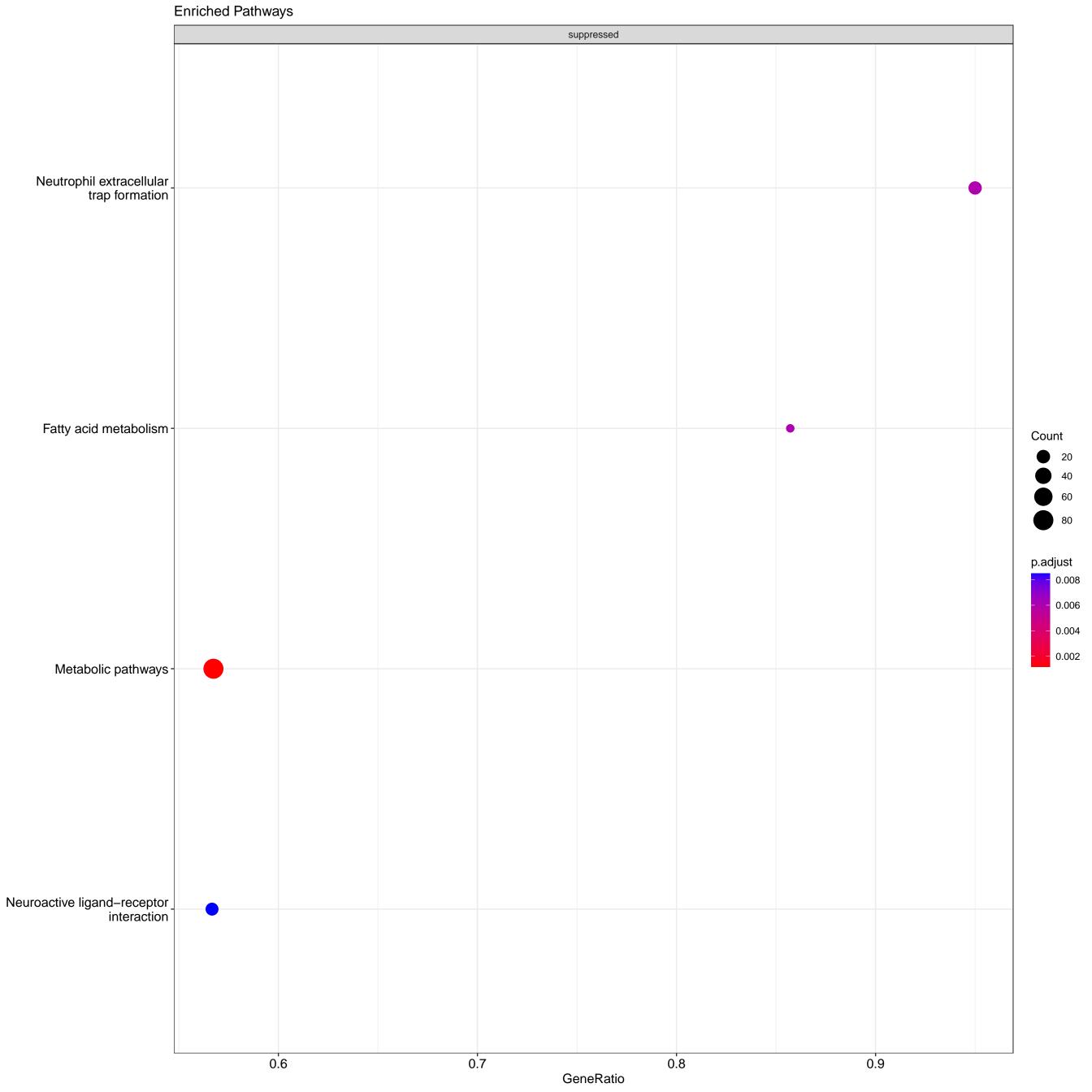
p.adjust

0.002









Neutrophil extracellular trap formation

number of genes
20
40
60
80

p.adjust
0.008
0.006
0.004
0.002

Metabolic pathways

