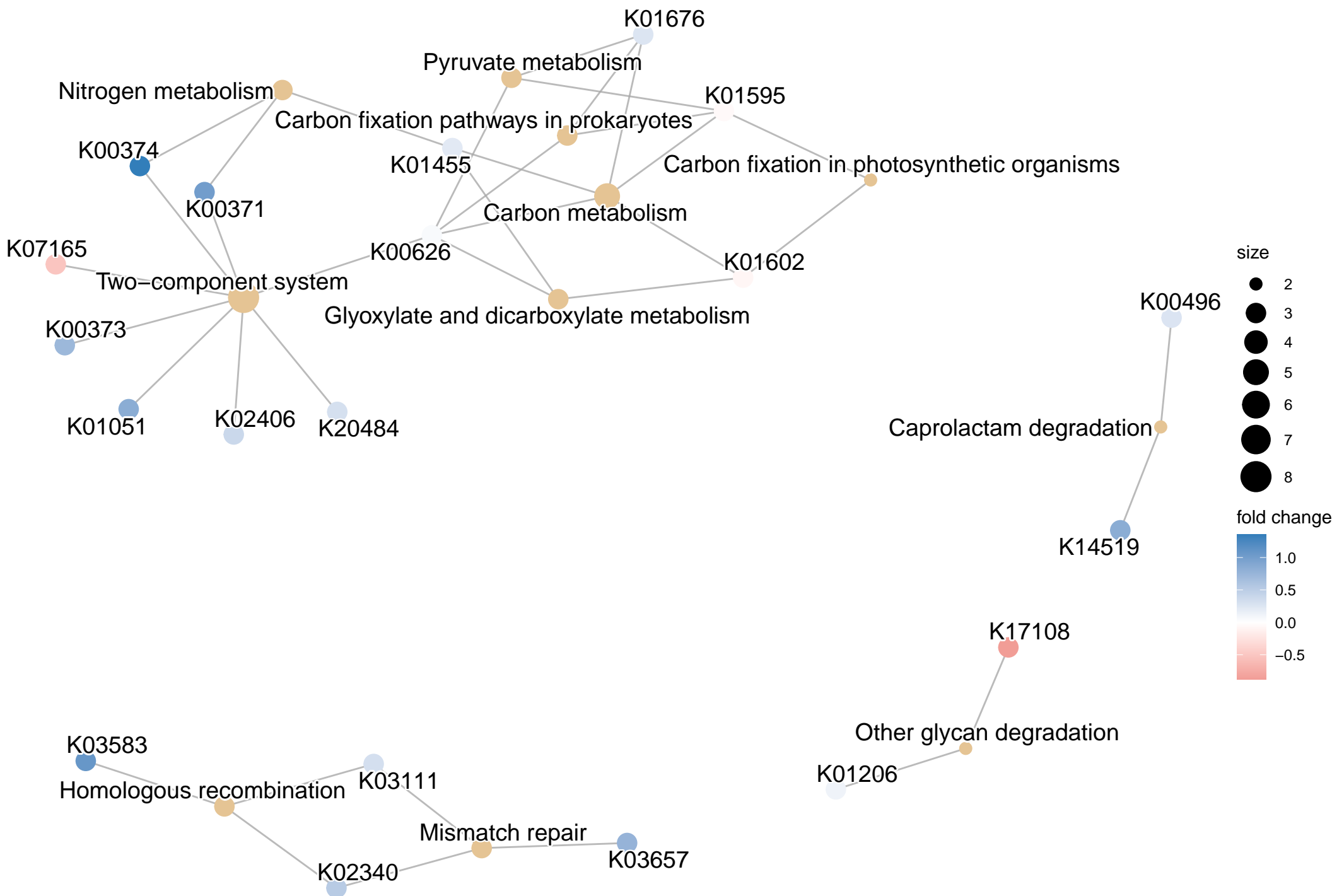
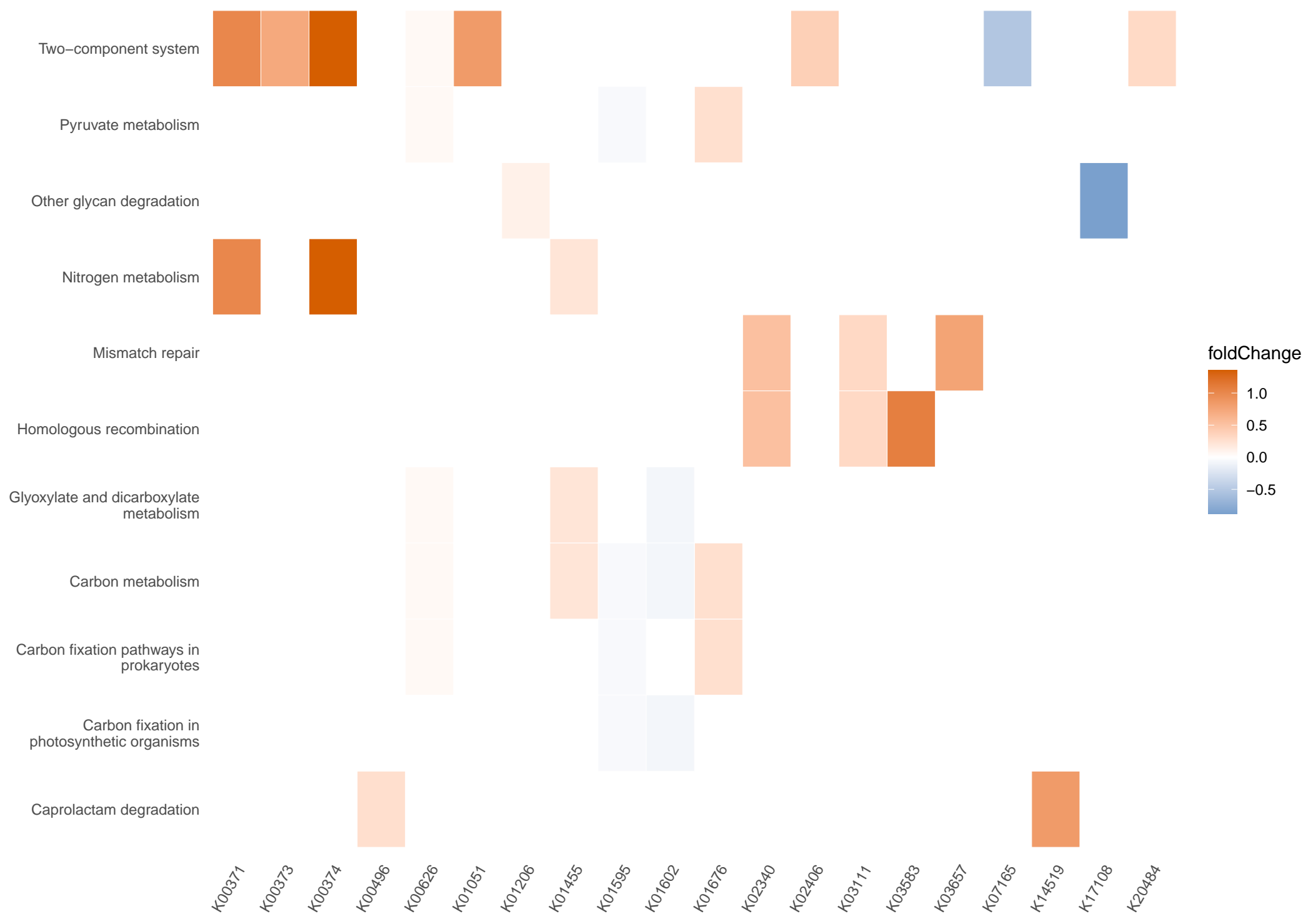
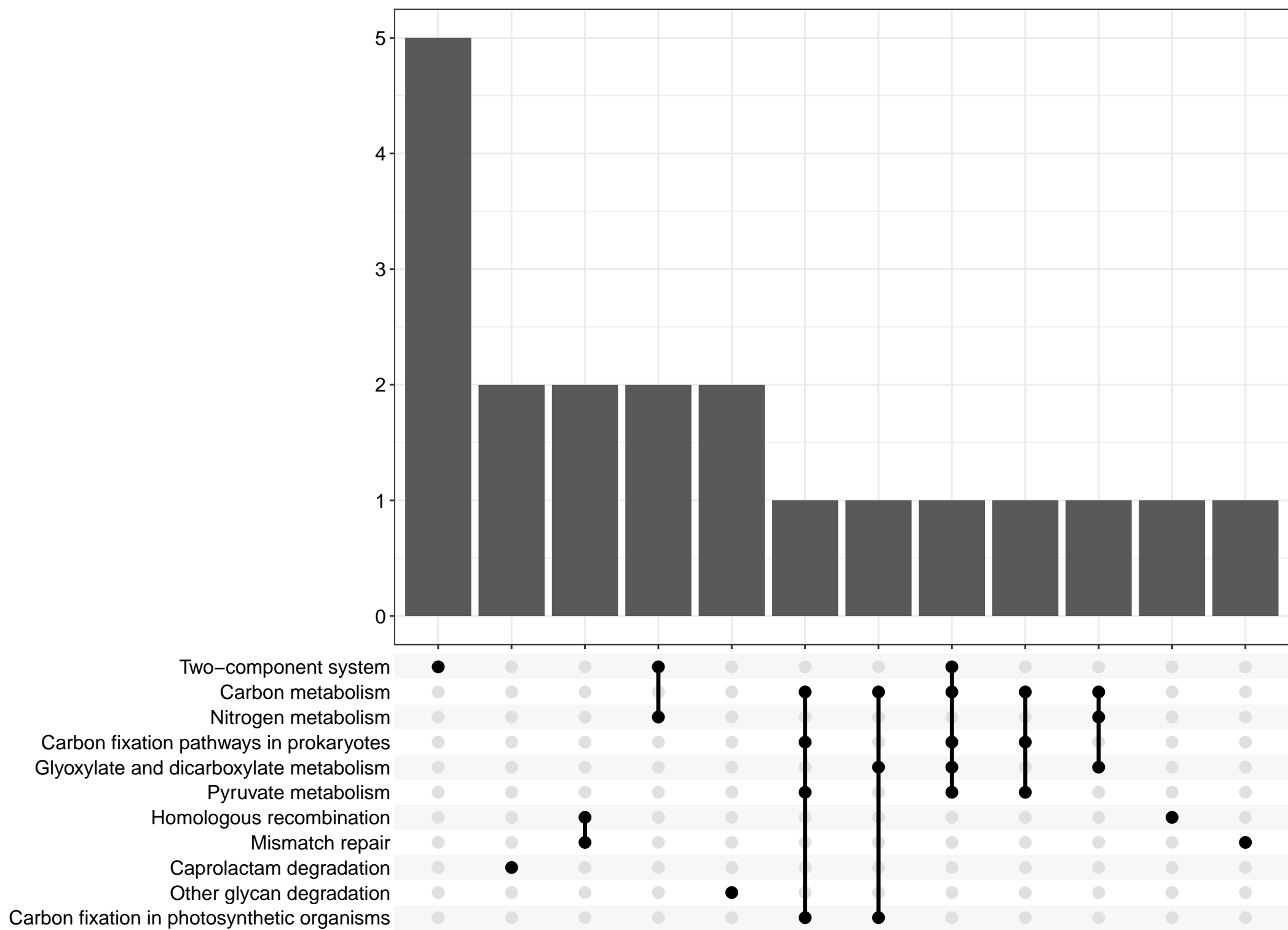


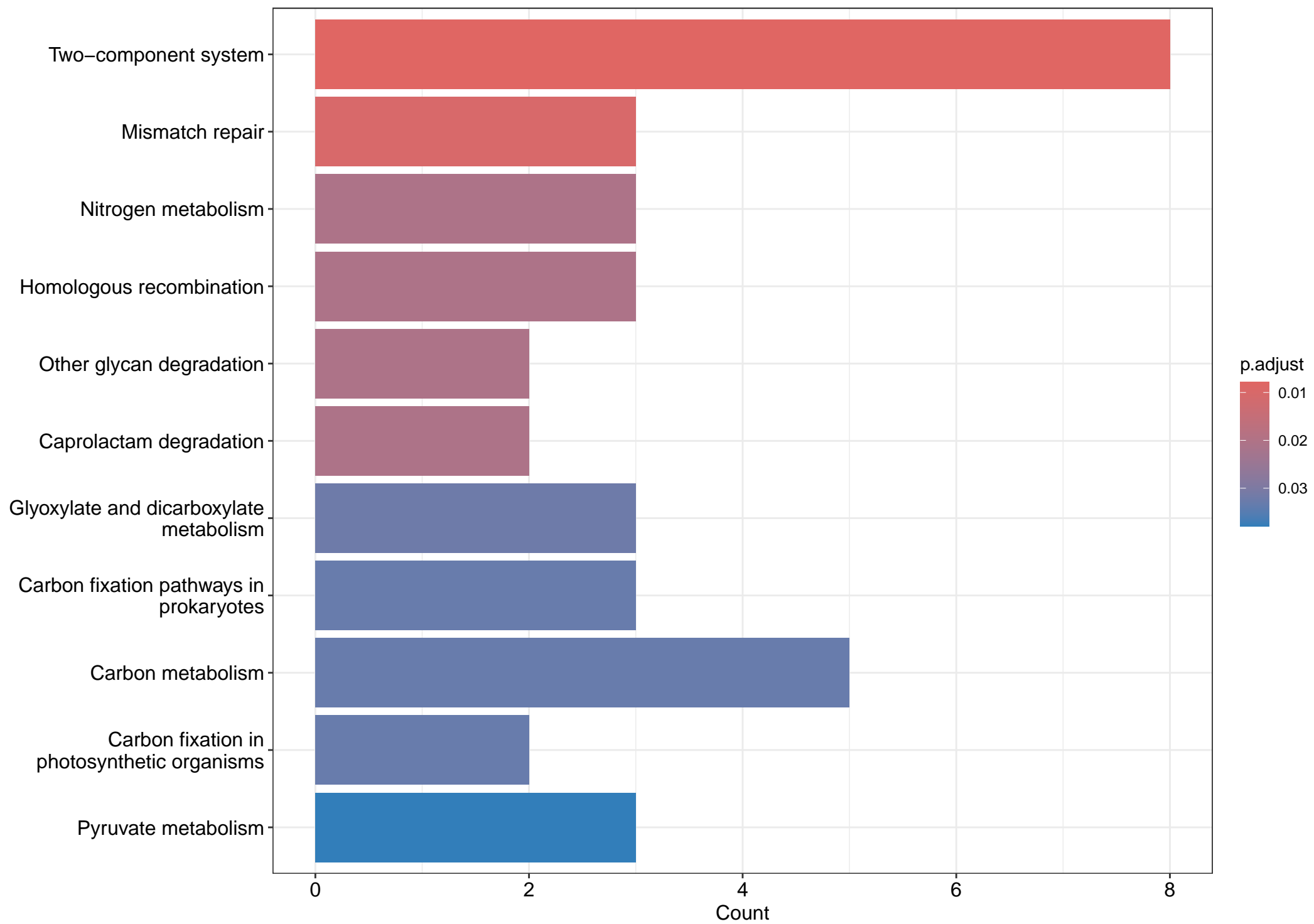
Site geom_path



Site geom_path







Site gsea_path

Flagellar assembly

p.adjust

0.005364302

Count

21

0.60

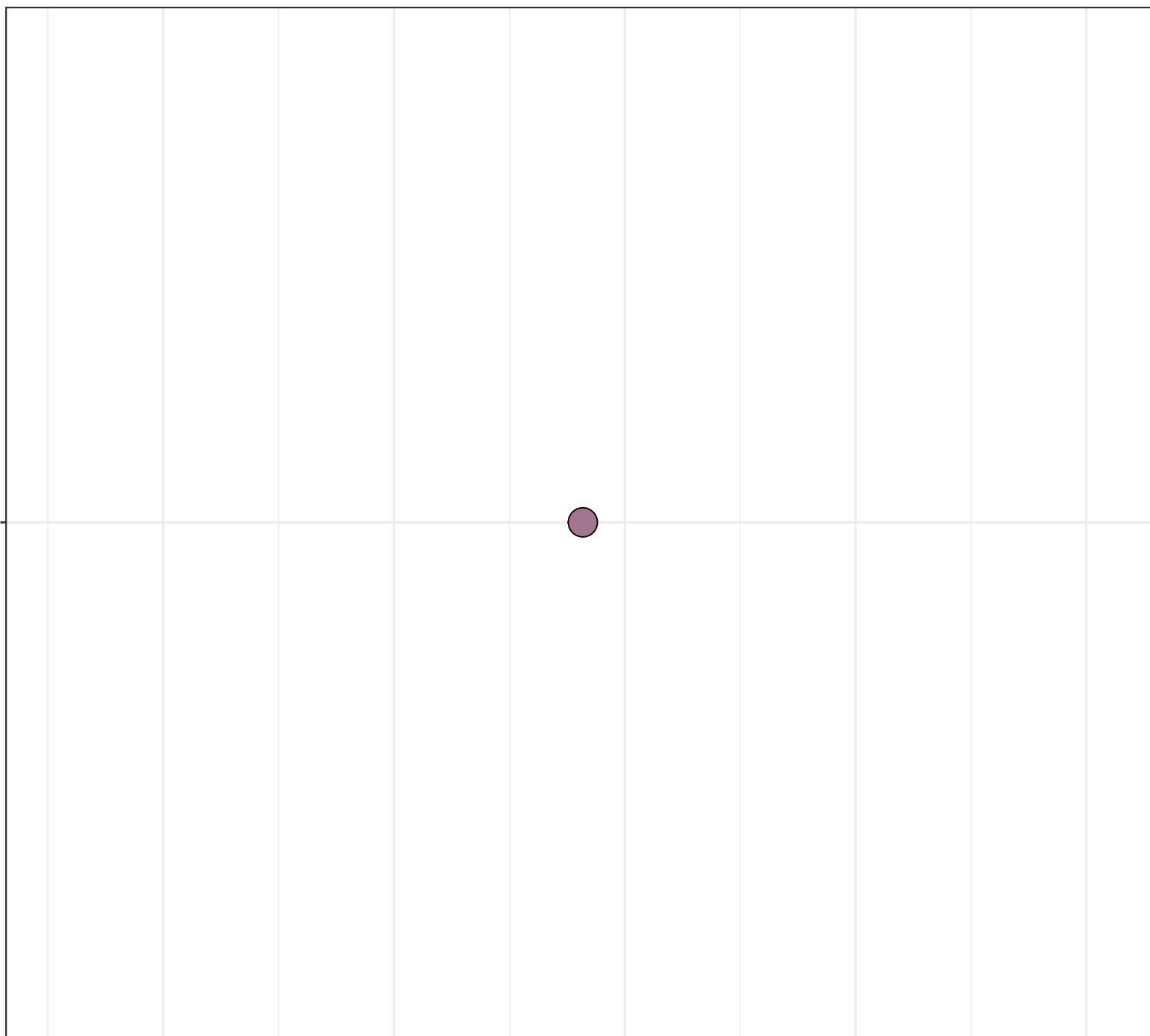
0.62

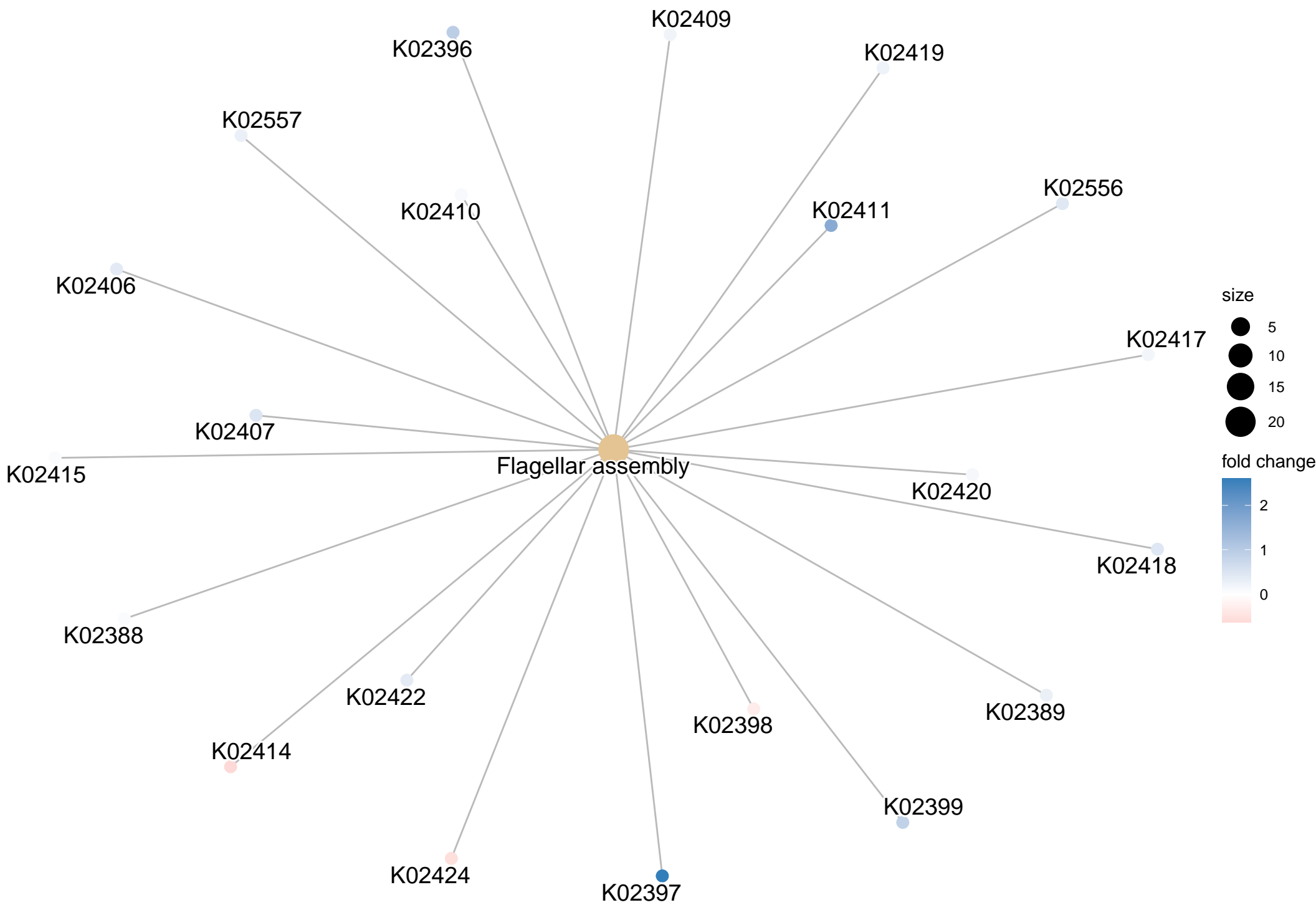
0.64

0.66

0.68

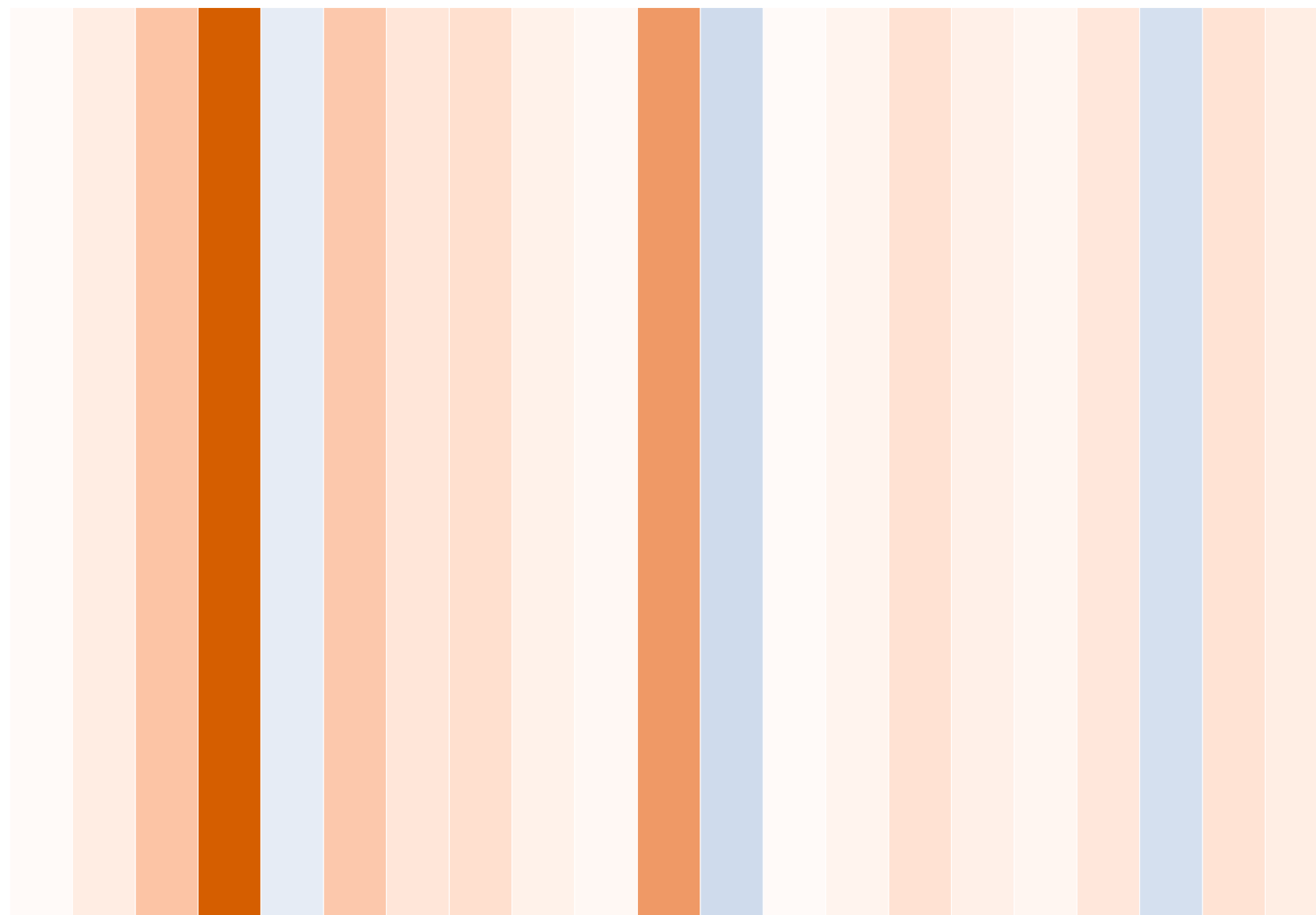
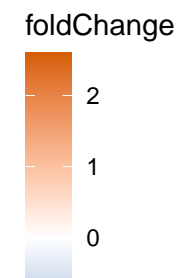
GeneRatio



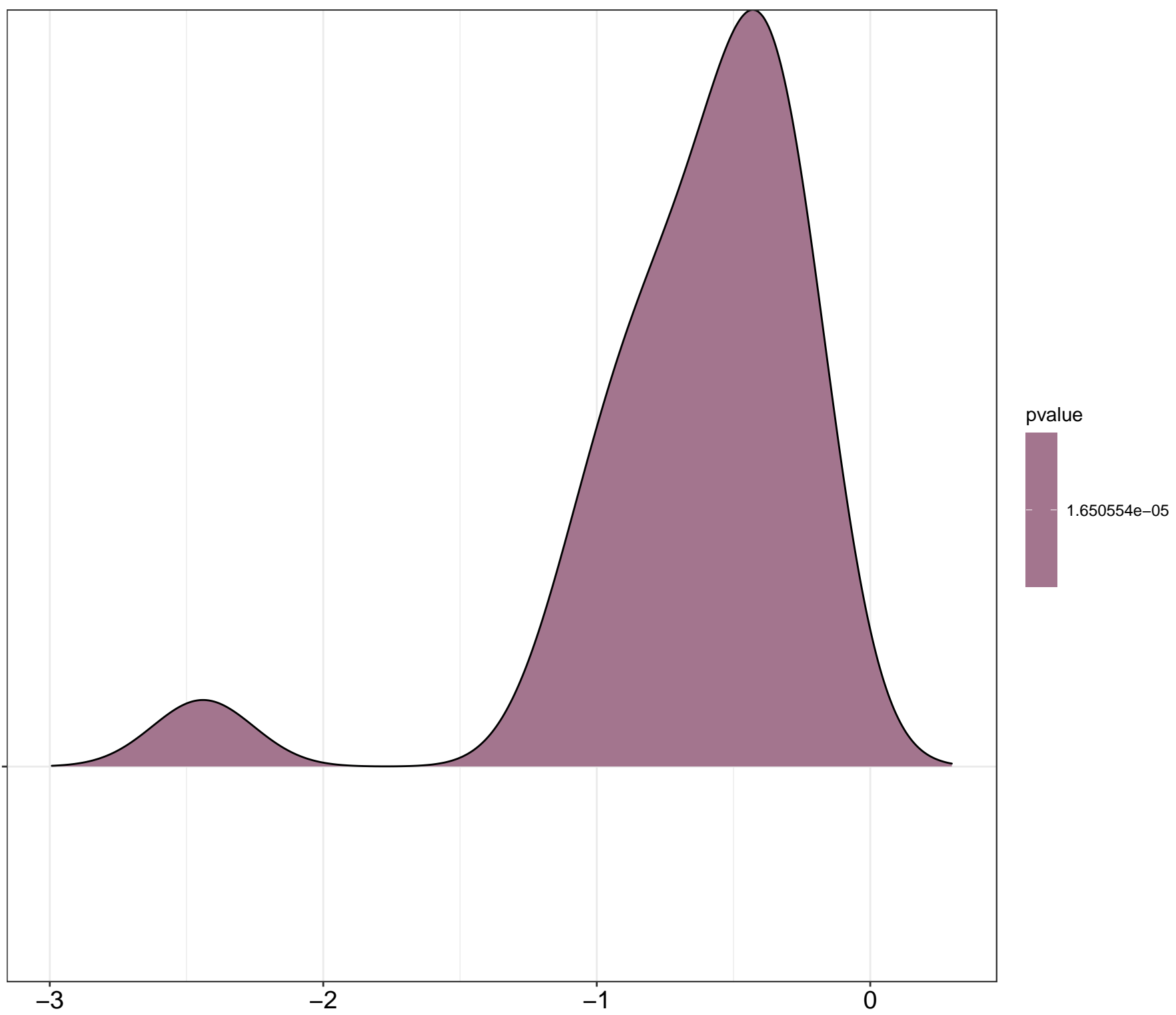


Site gsea_path

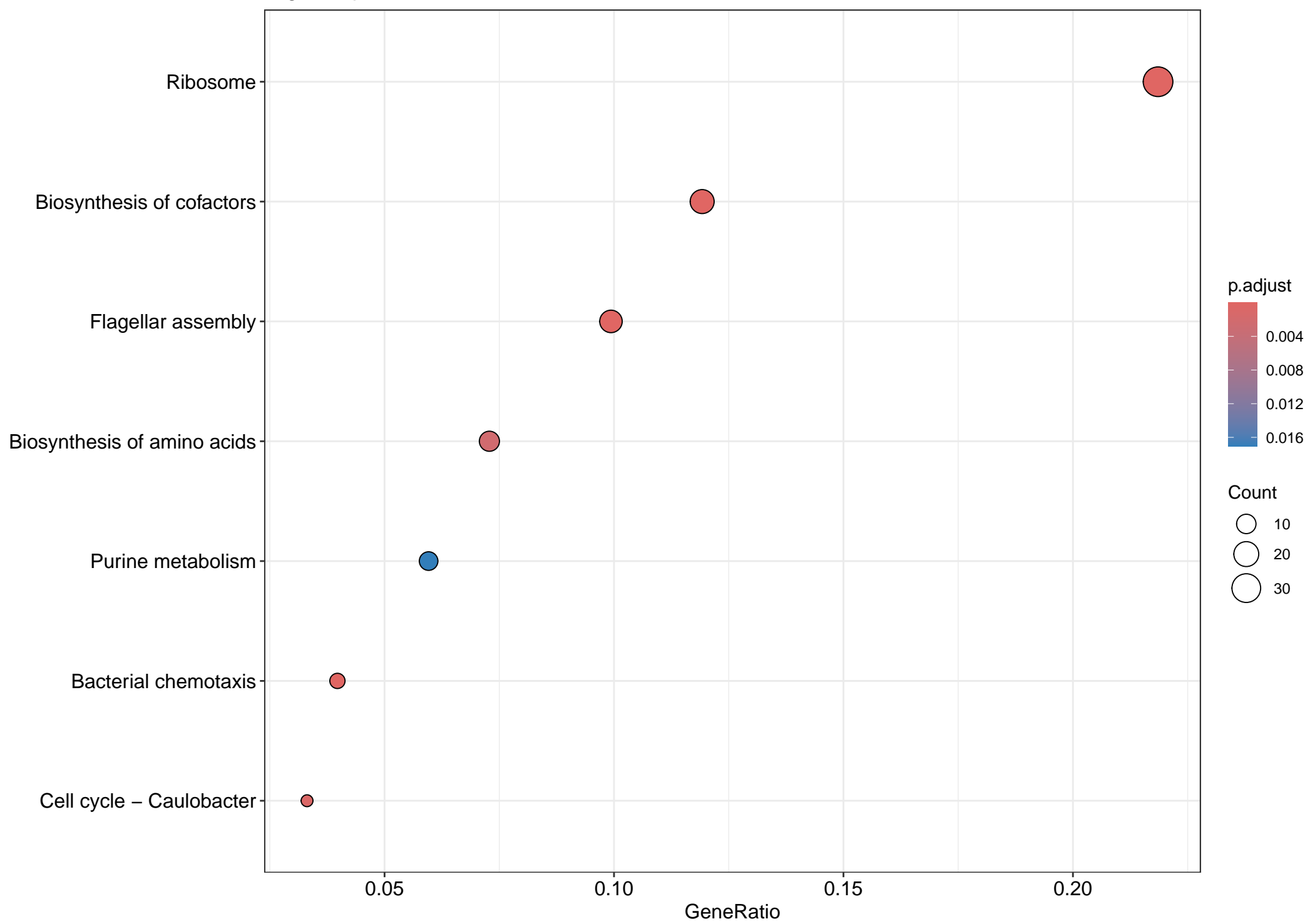
Flagellar assembly



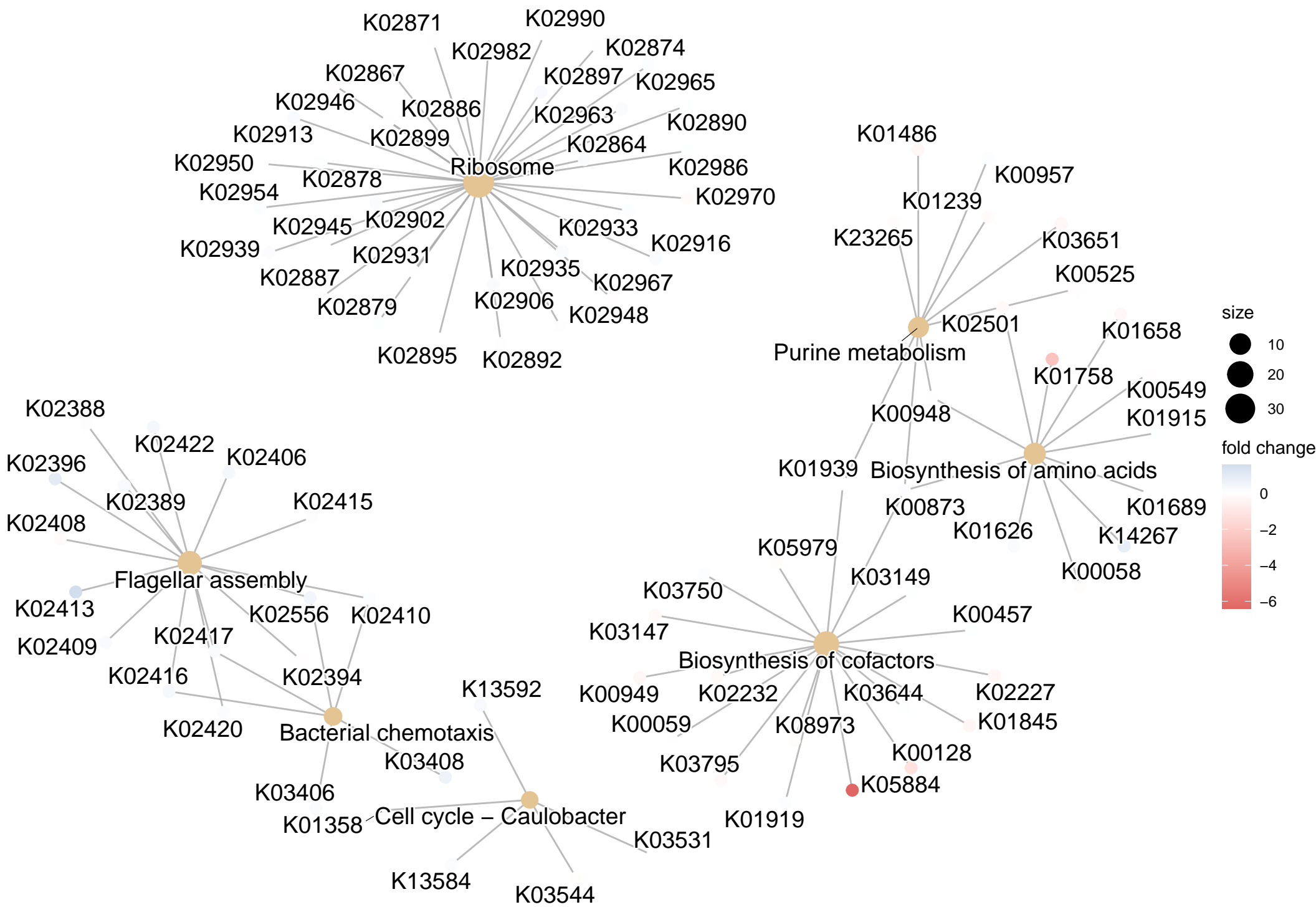
Flagellar assembly



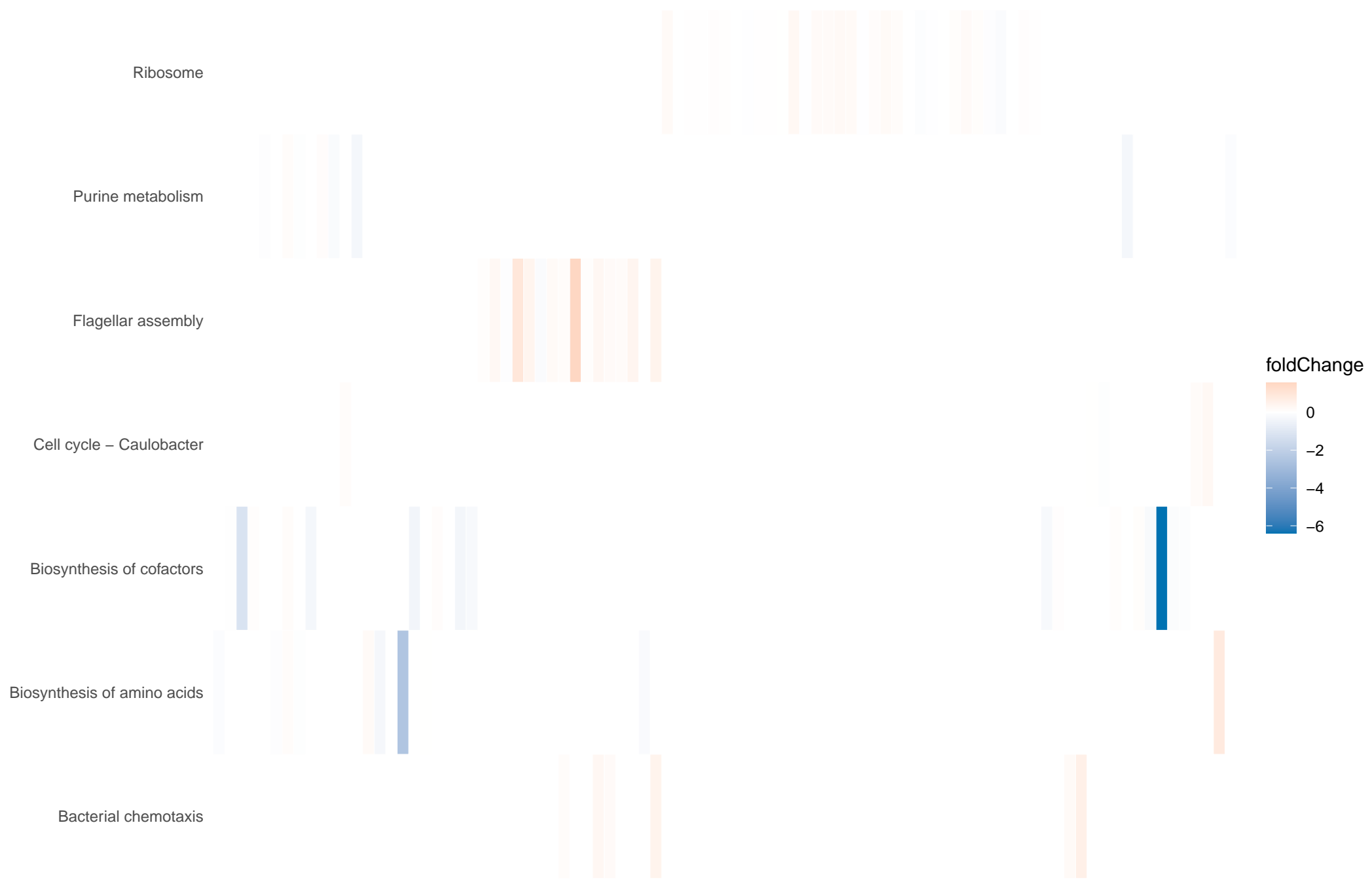
C geom_path



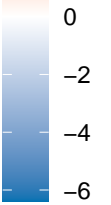
C geom_path

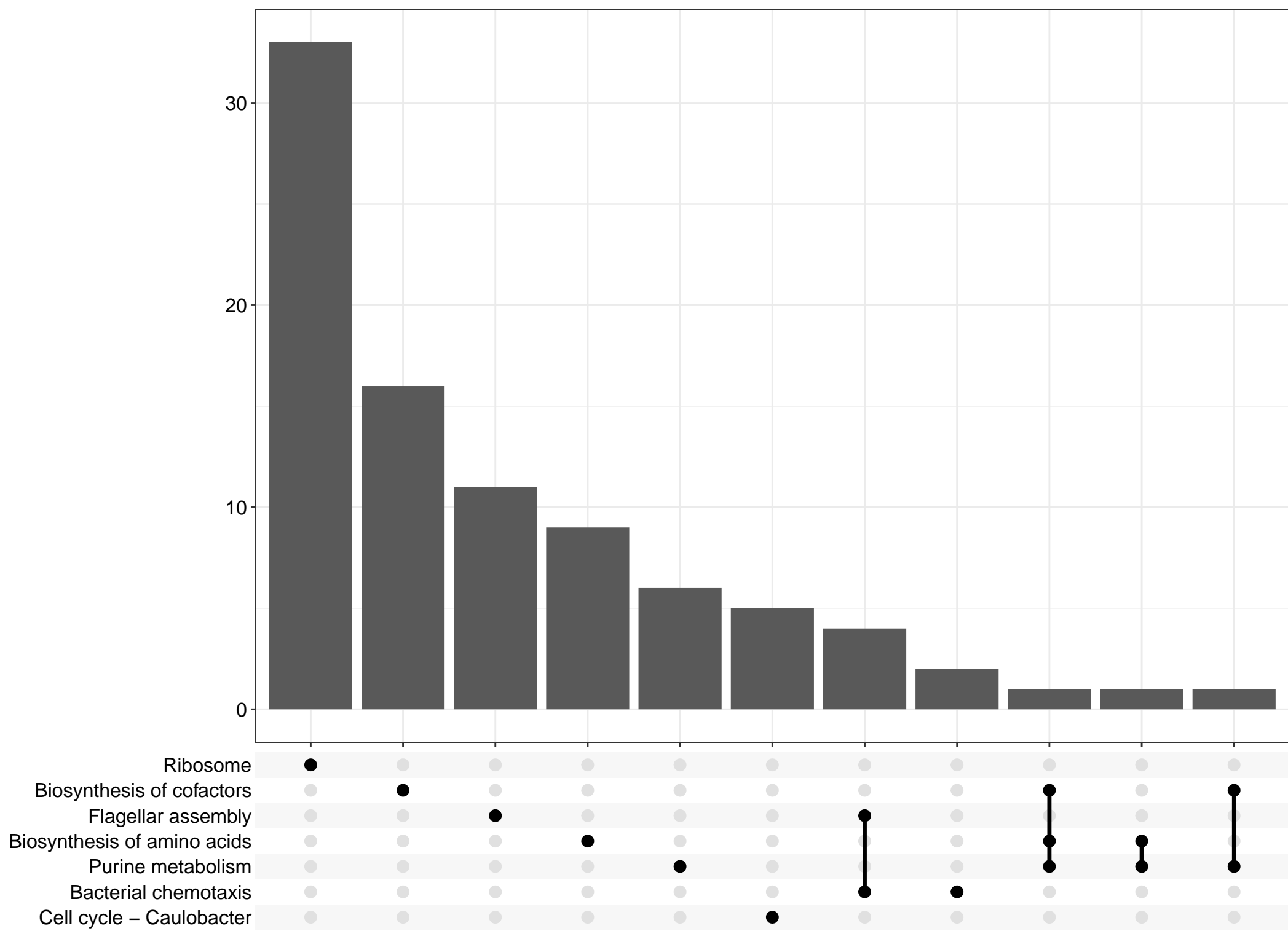


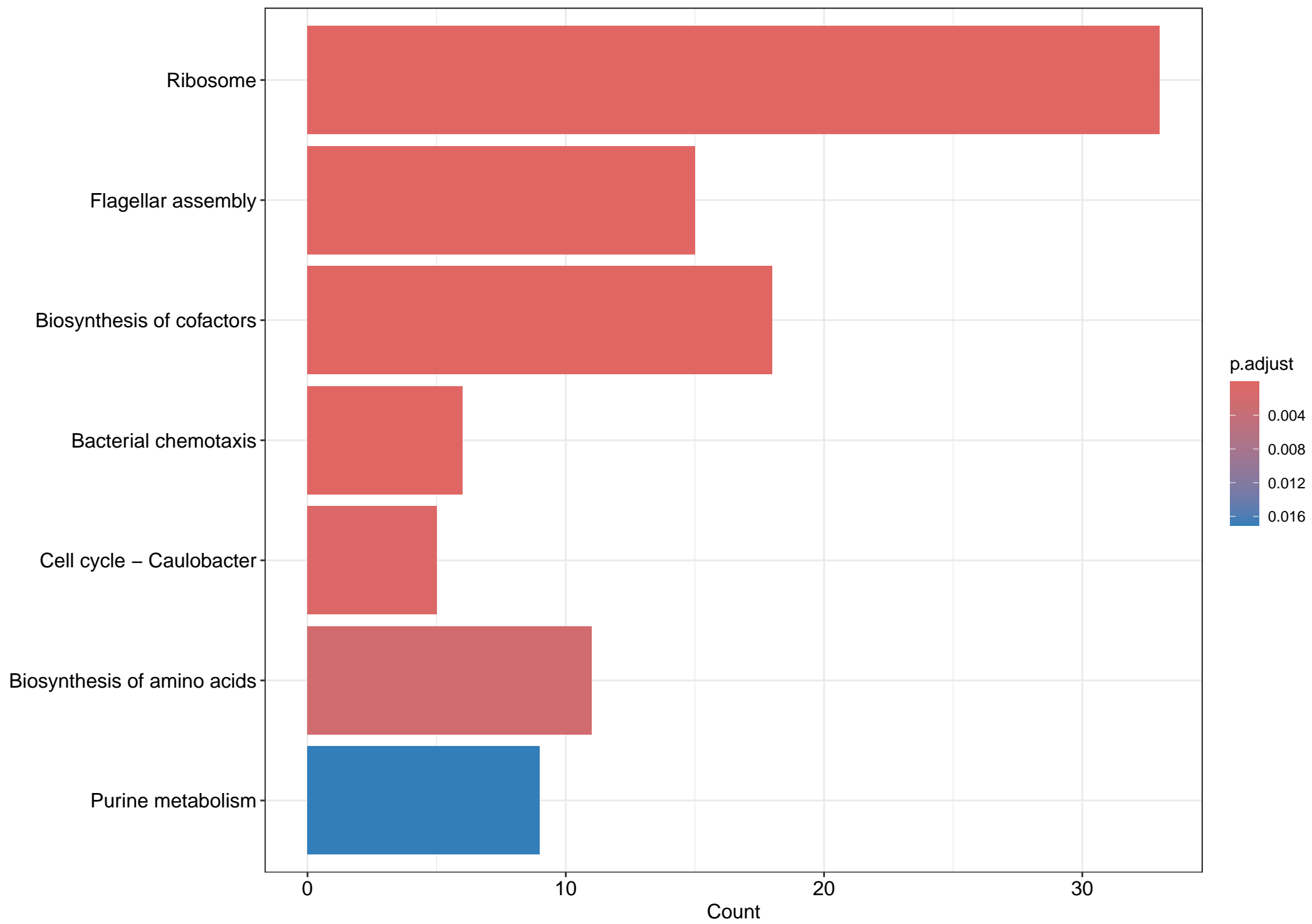
C geom_path



foldChange







C gsea_path

Flagellar assembly

0.850

0.875

0.900

0.925

GeneRatio

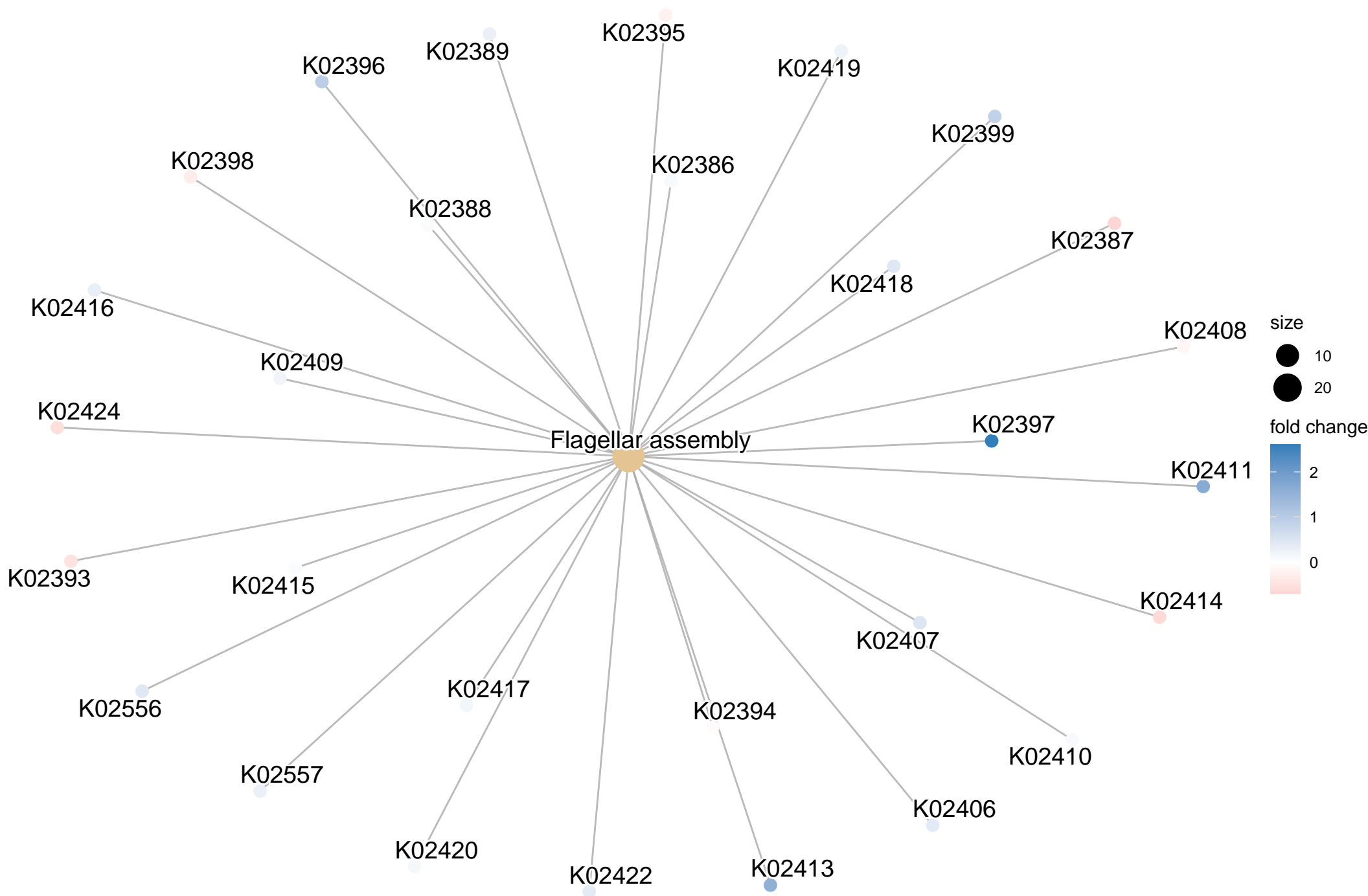
p.adjust

3.25e-08

Count

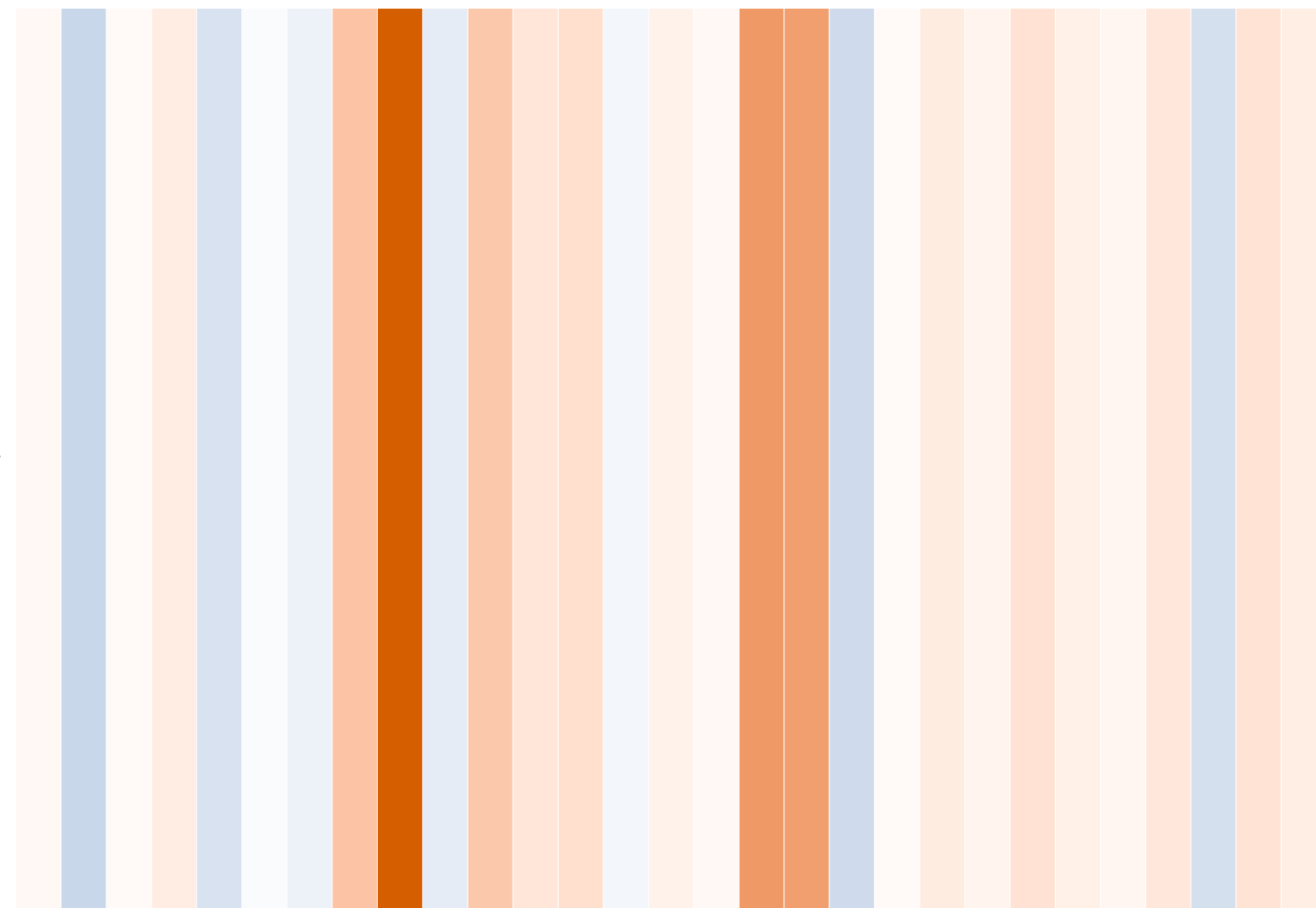
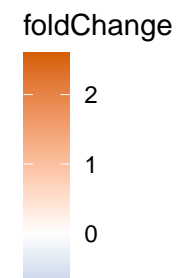
29



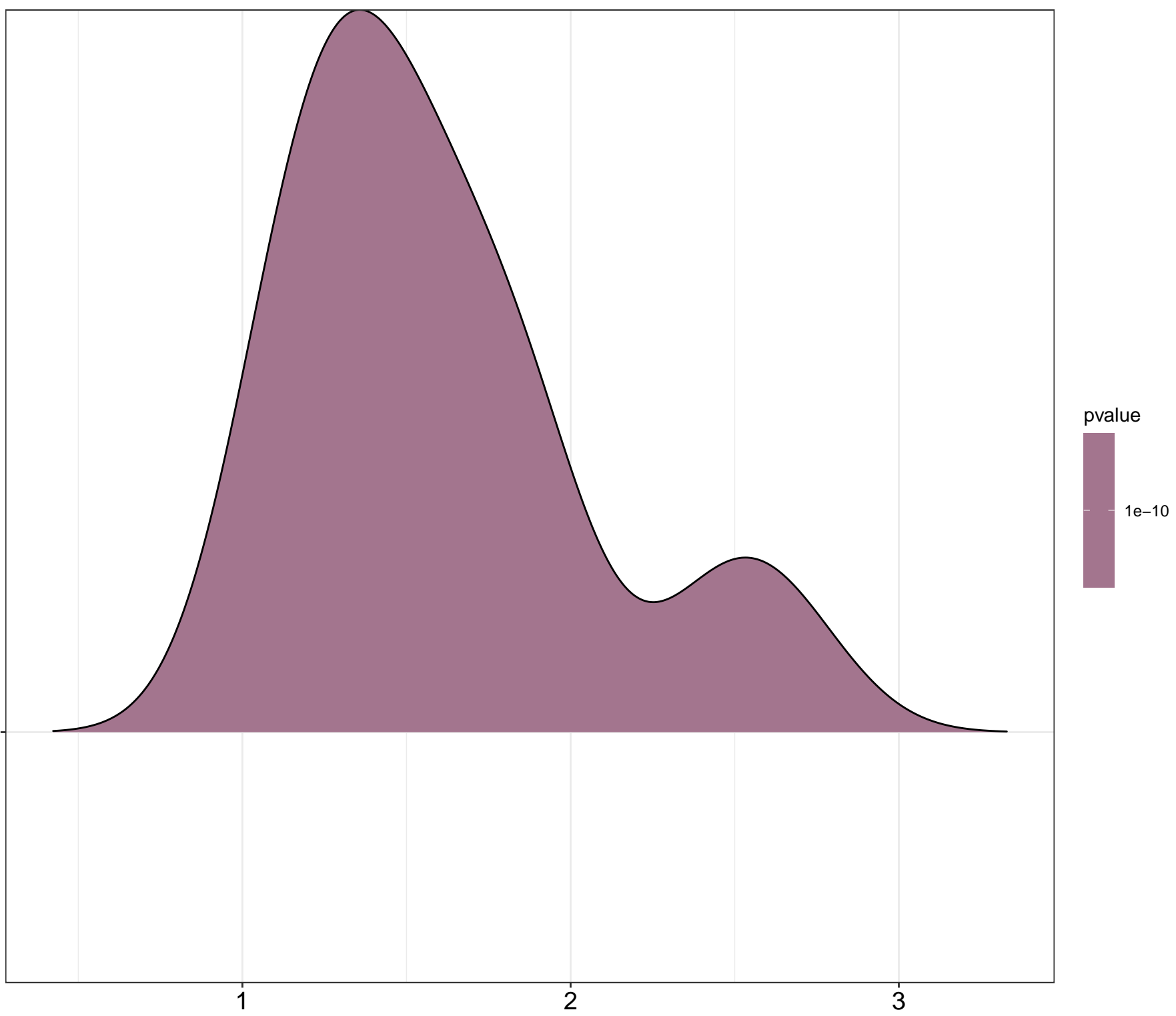


C gsea_path

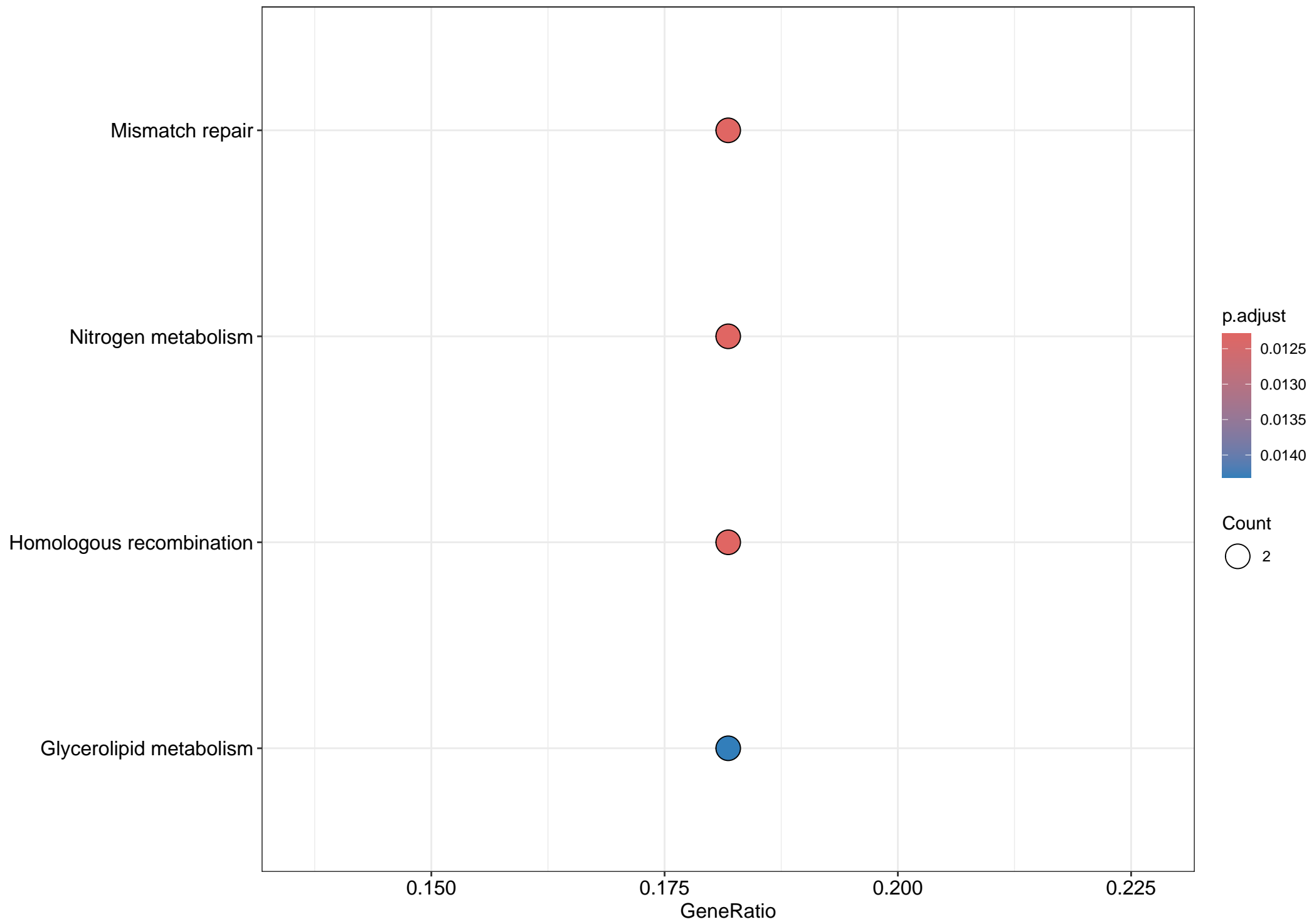
Flagellar assembly



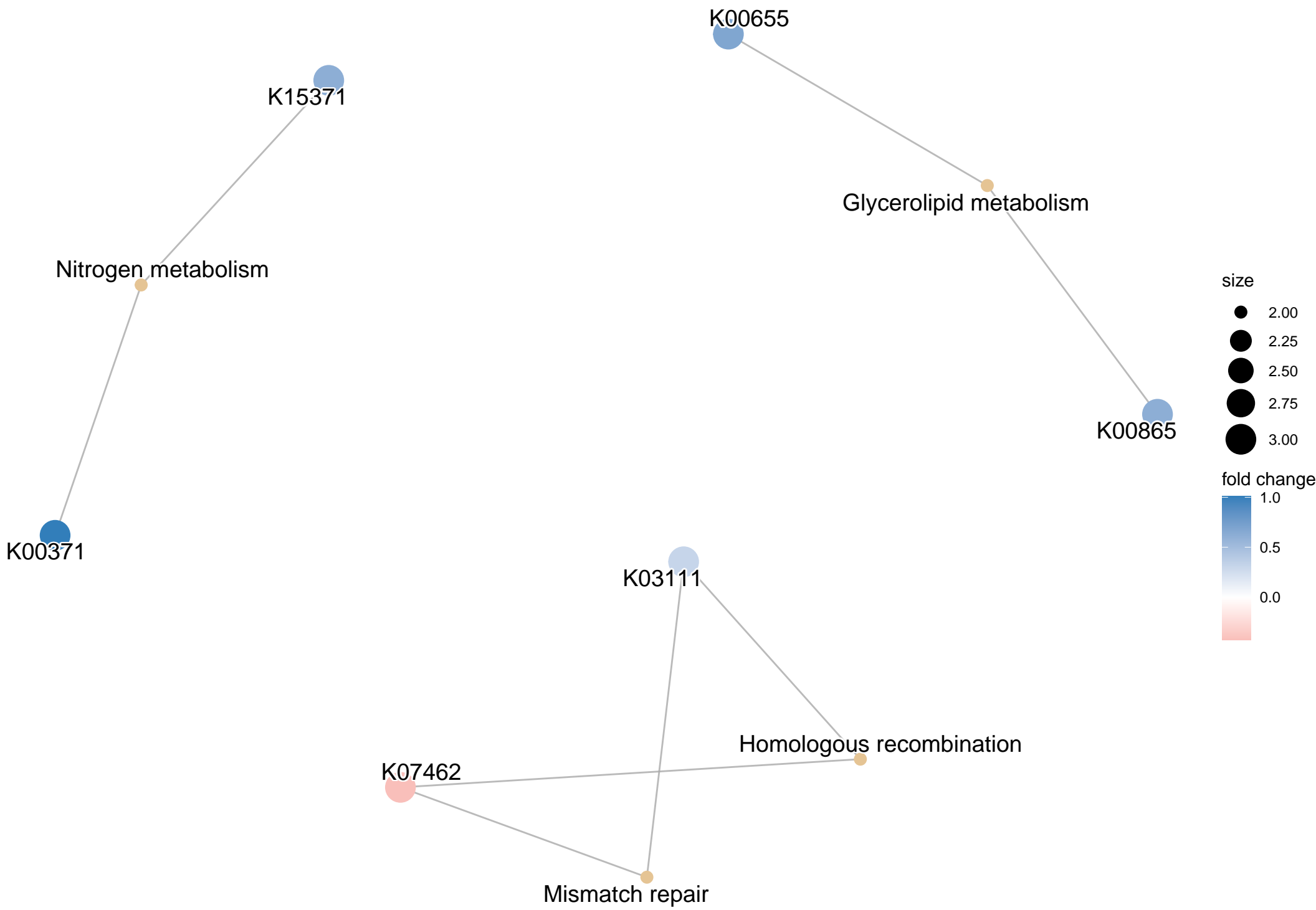
Flagellar assembly



D geom_path



D geom_path



D geom_path

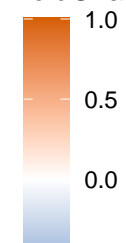
Nitrogen metabolism

Mismatch repair

Homologous recombination

Glycerolipid metabolism

foldChange



K00371

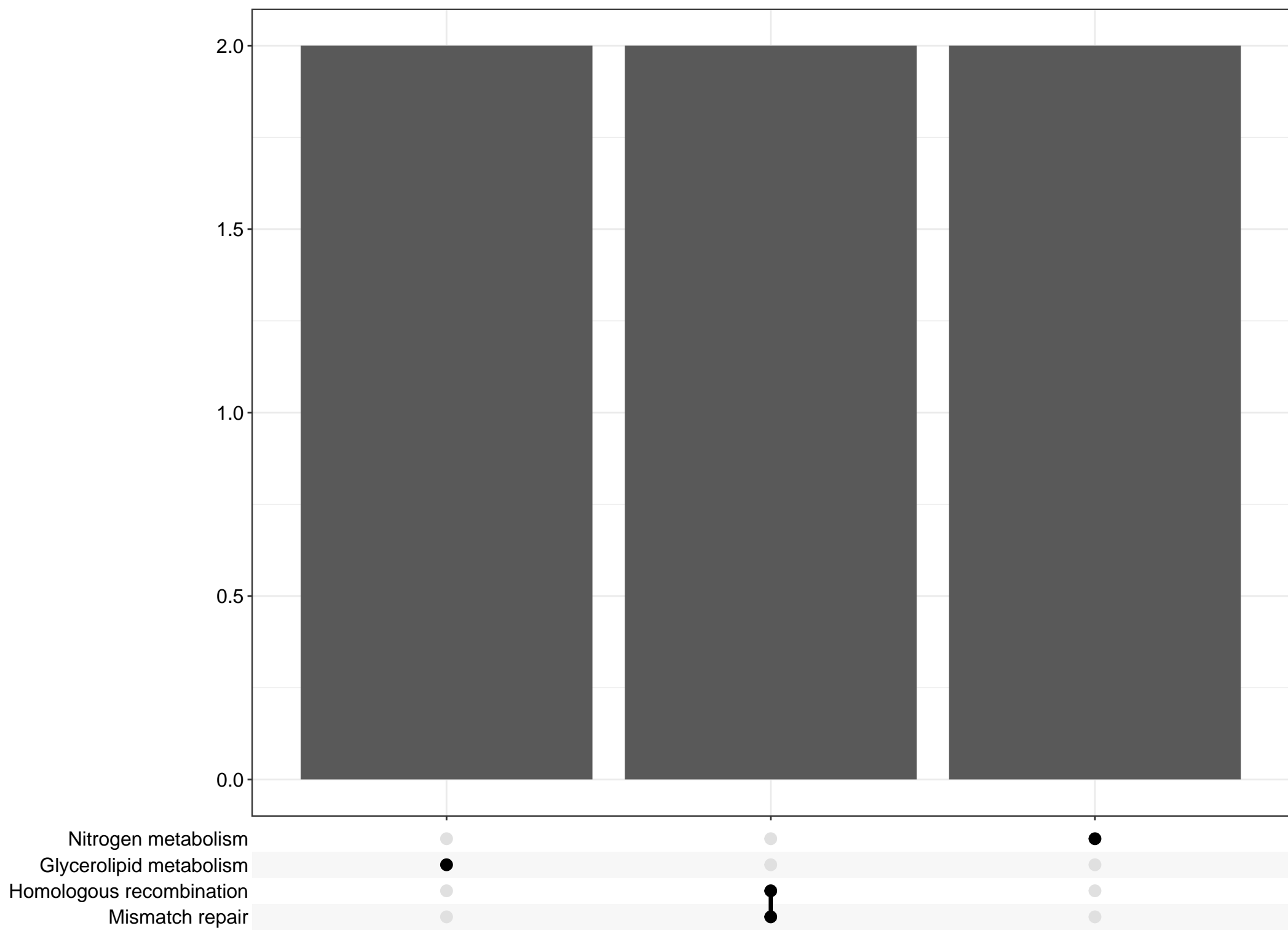
K00655

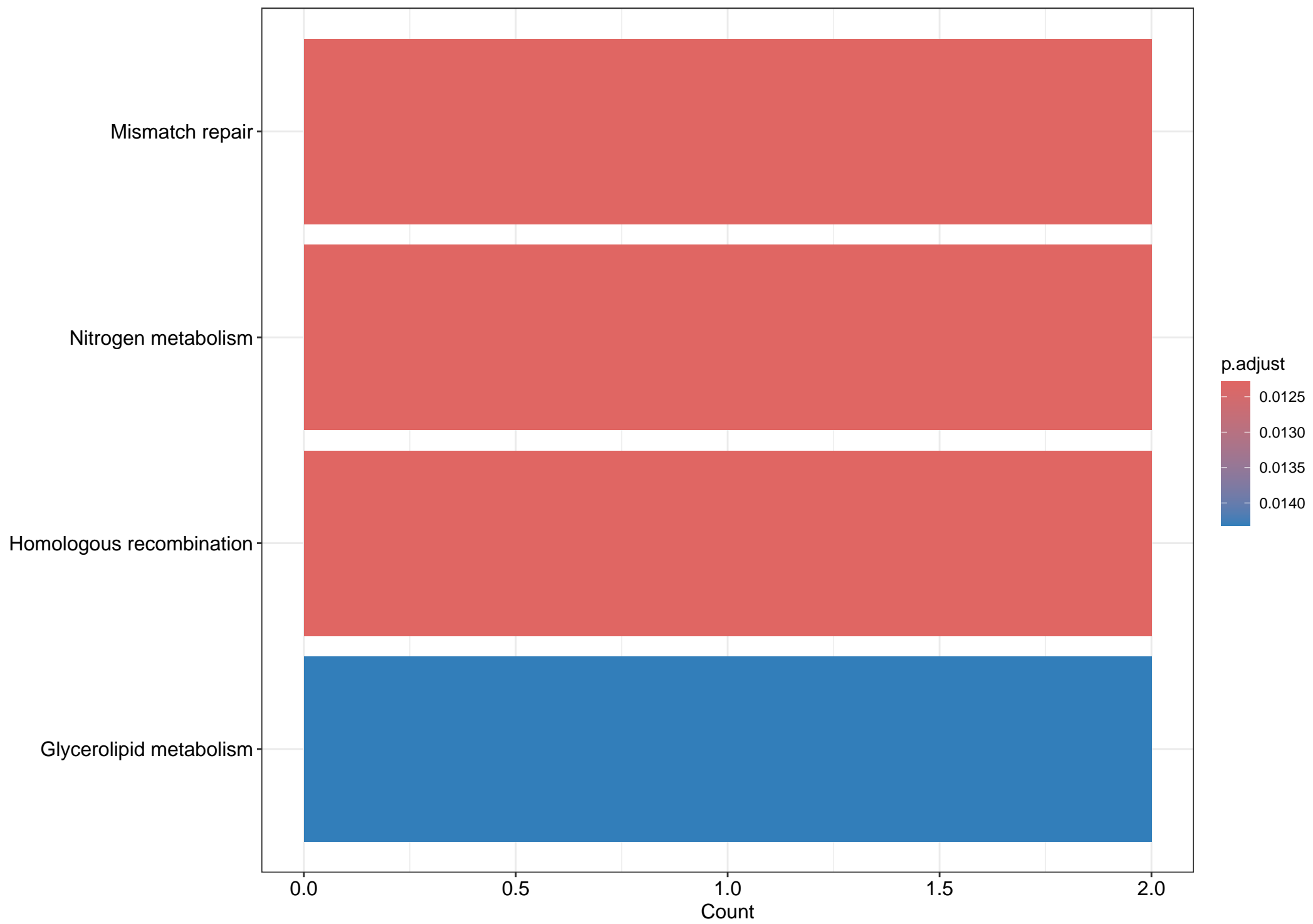
K00865

K03111

K07462

K15371





D gsea_path

Coronavirus disease – COVID-19

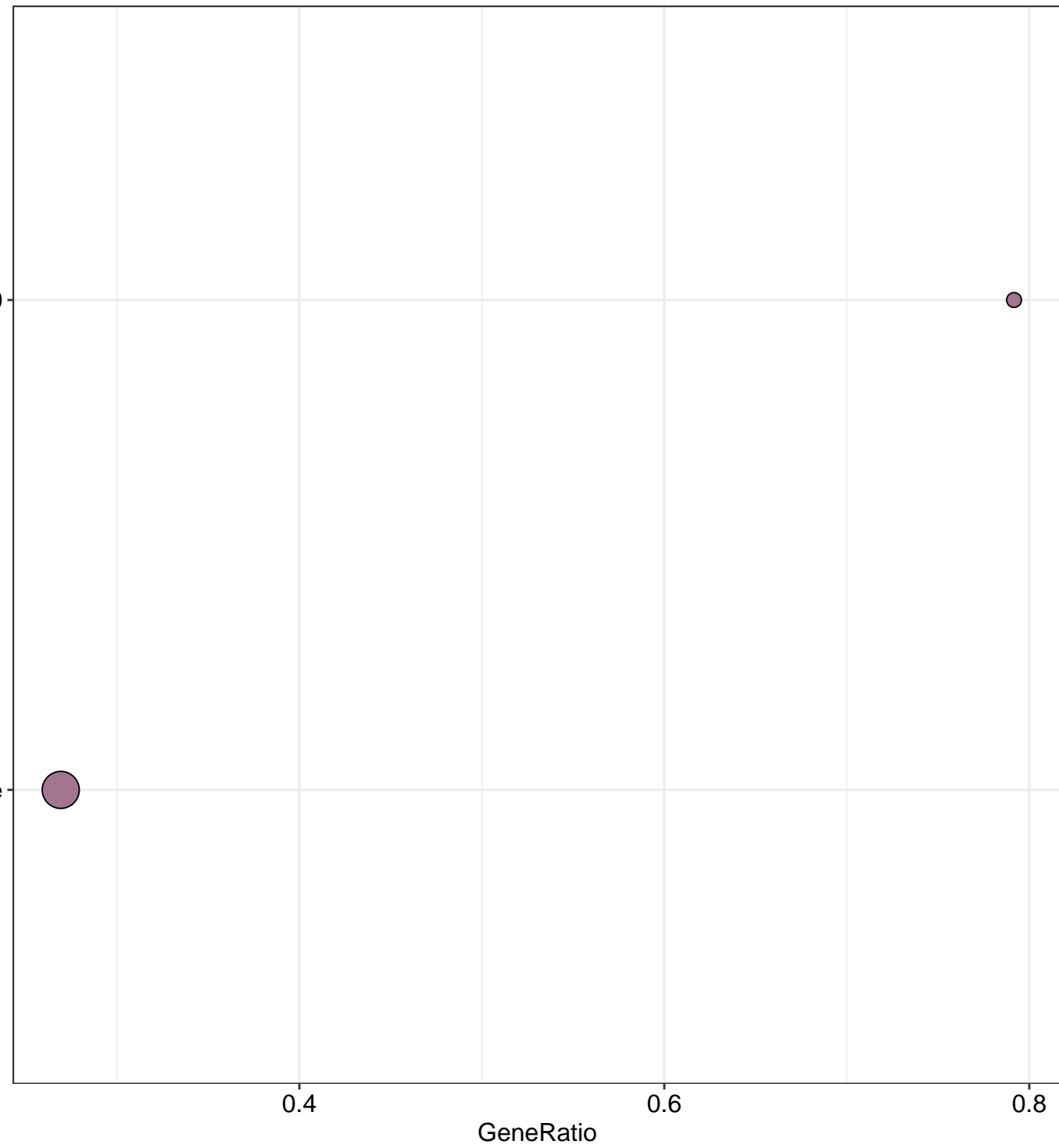
Ribosome

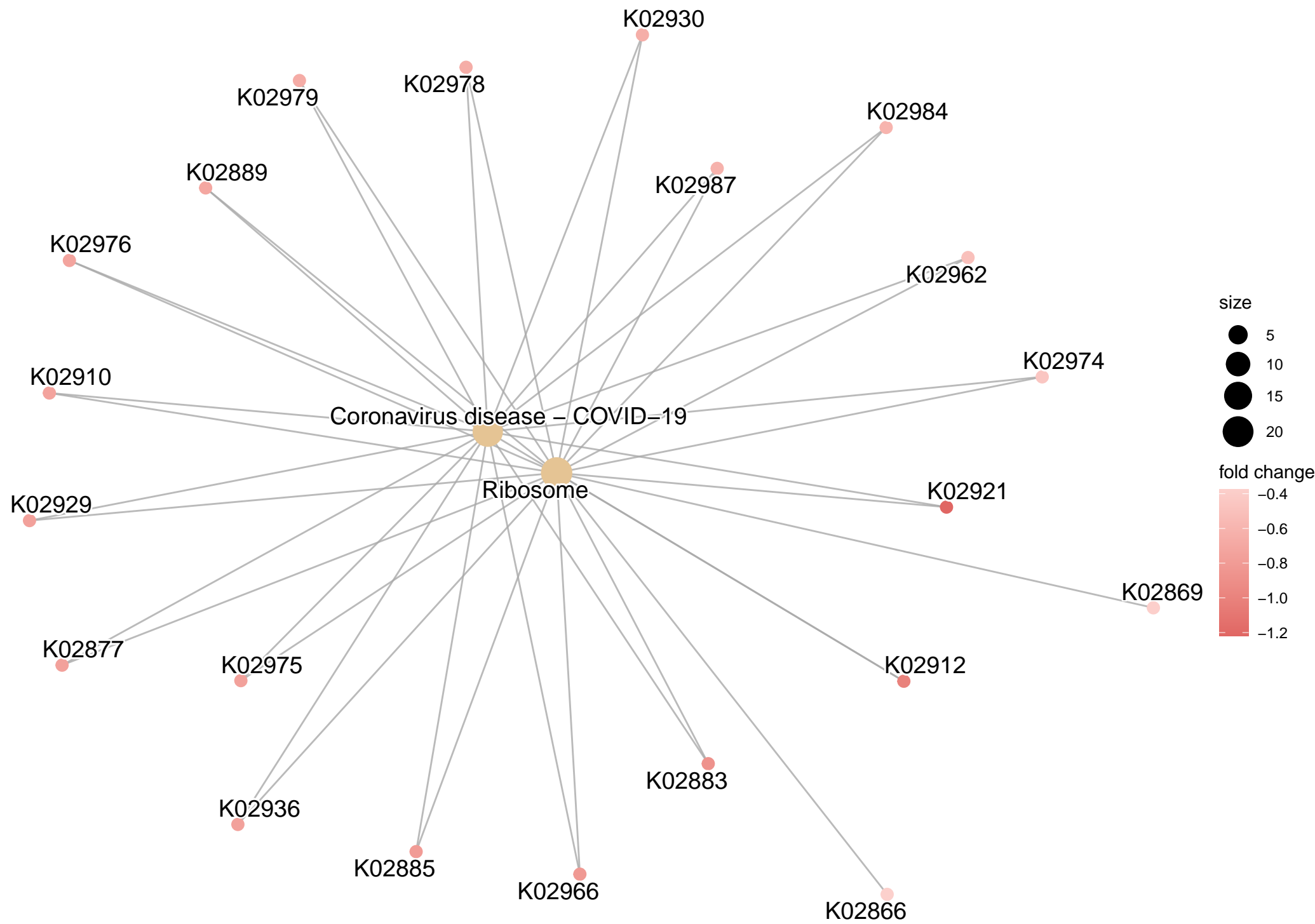
p.adjust

0.02470505

Count

- 19.0
- 19.5
- 20.0
- 20.5
- 21.0

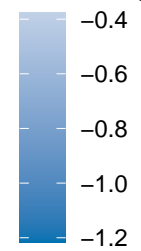




D gsea_path

Ribosome

foldChange



Coronavirus disease – COVID-19

