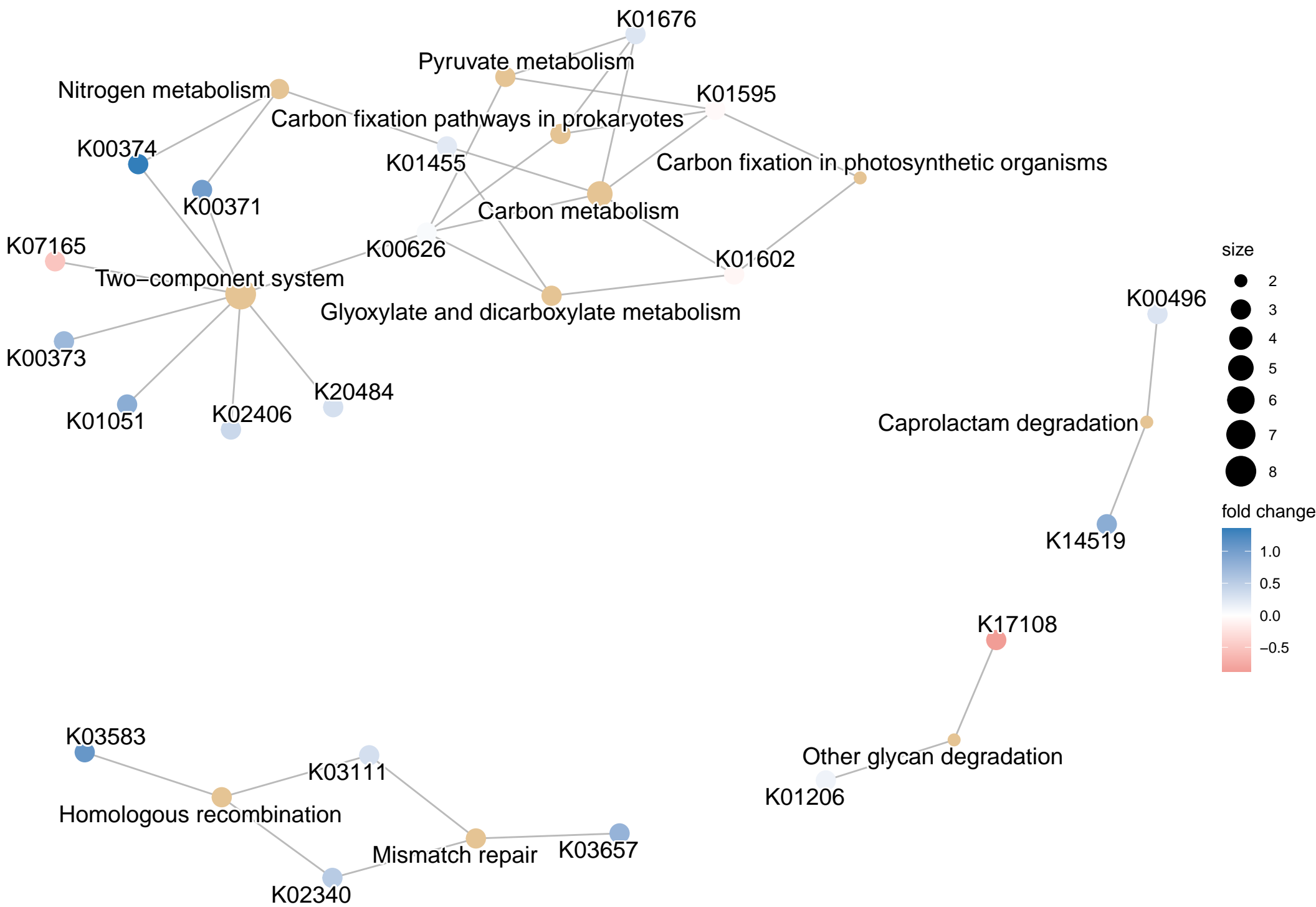
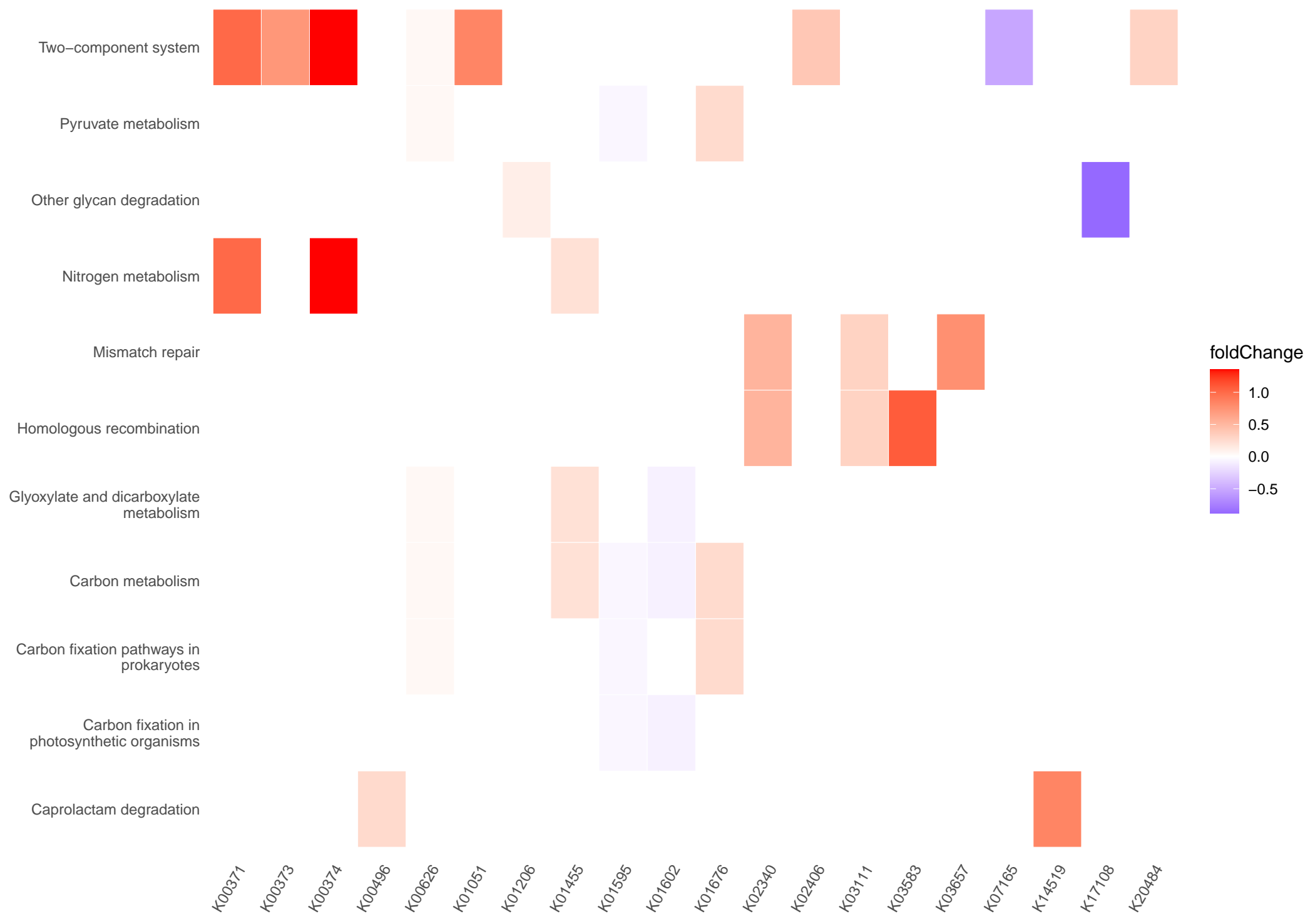
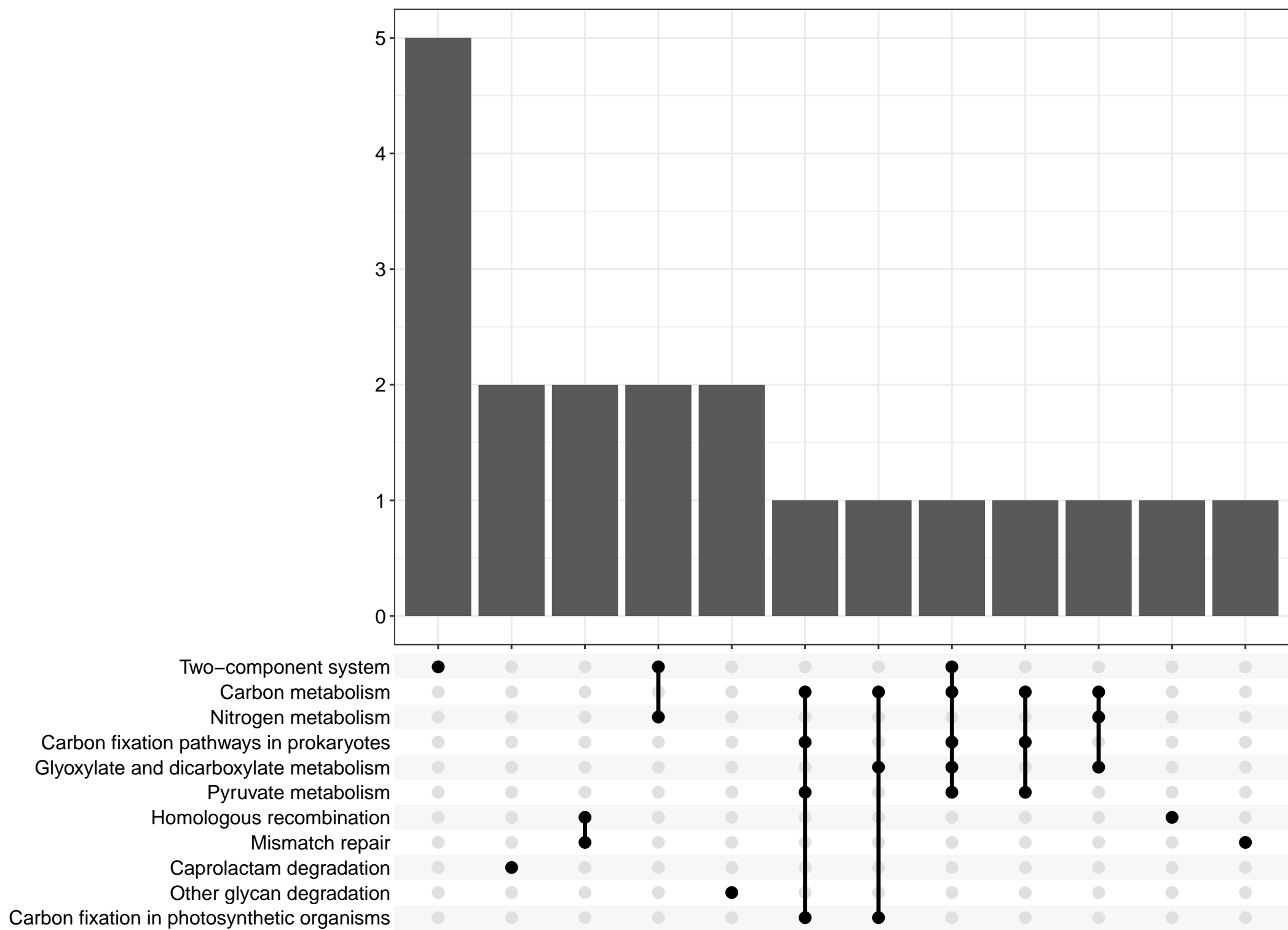


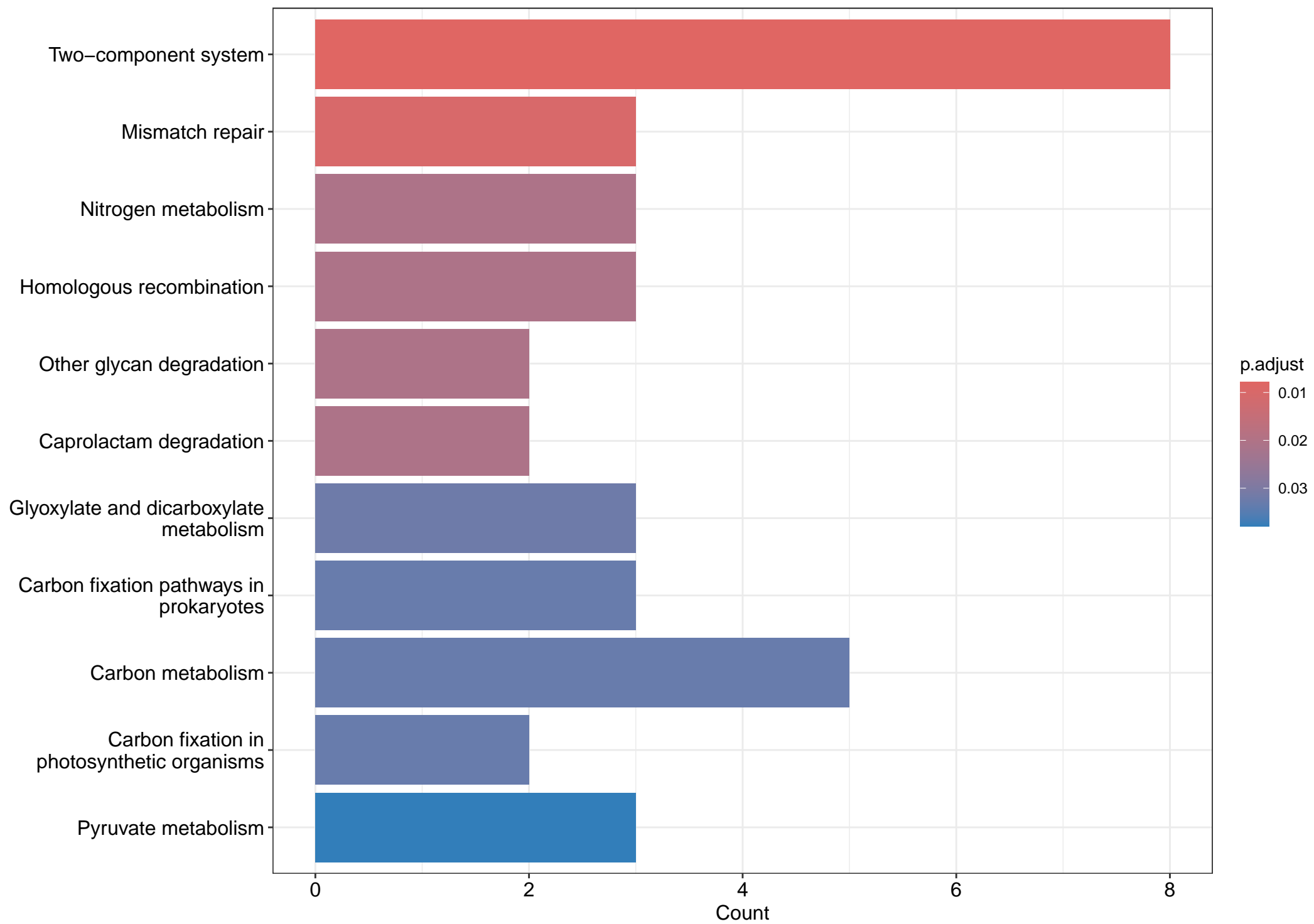
Site geom_path



Site geom_path







Site gsea_path

Flagellar assembly

p.adjust

0.001066

Count

21

0.60

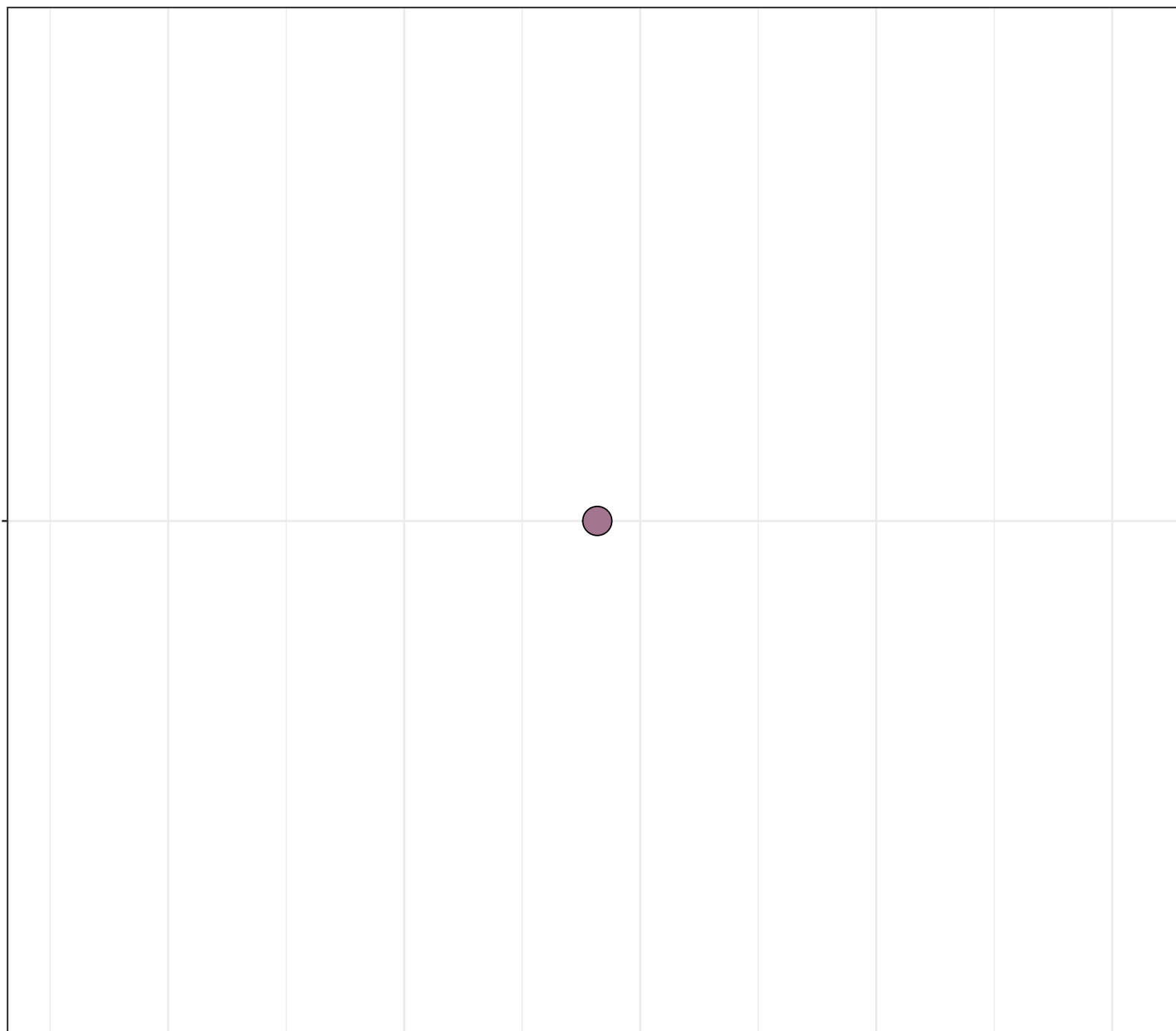
0.62

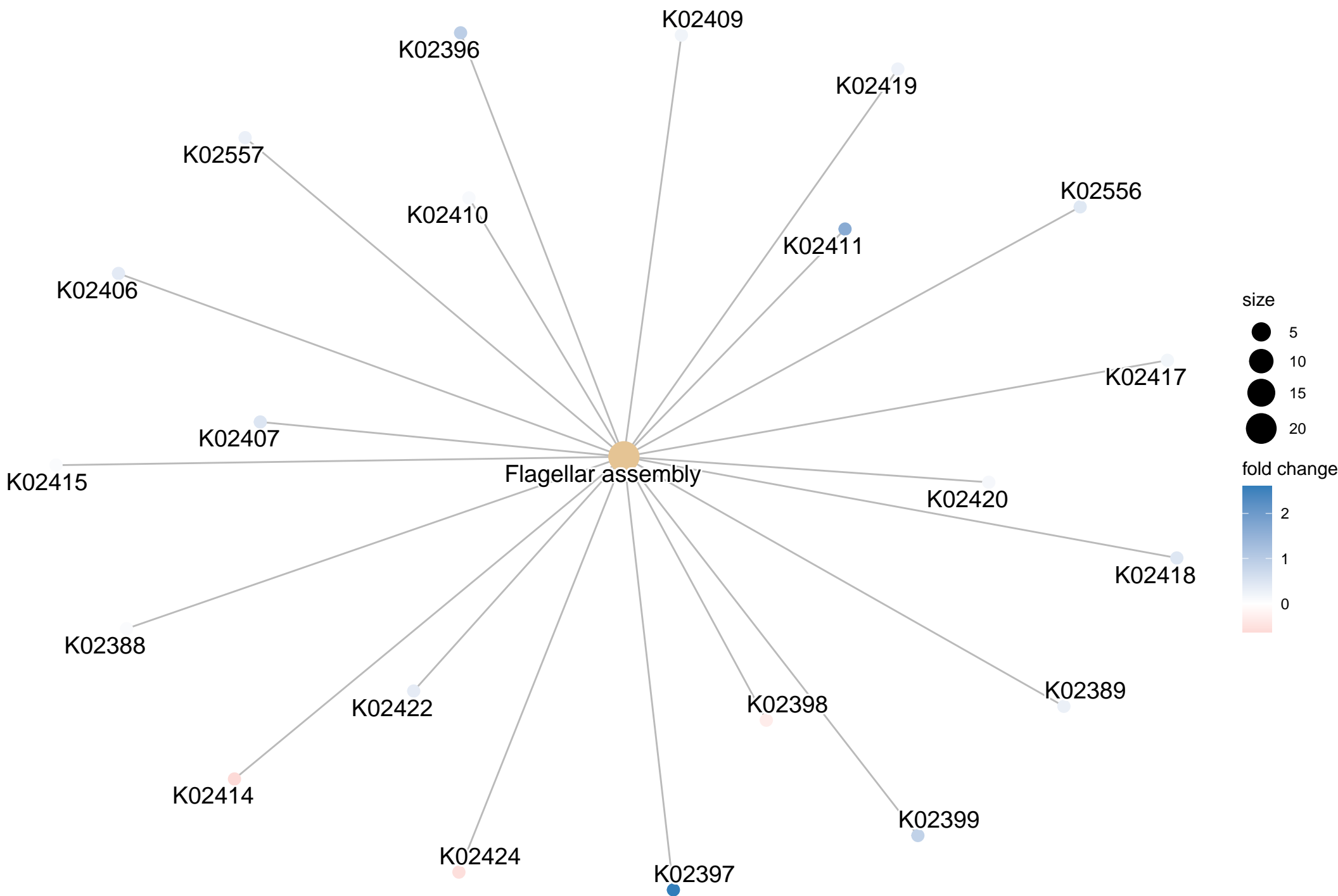
0.64

0.66

0.68

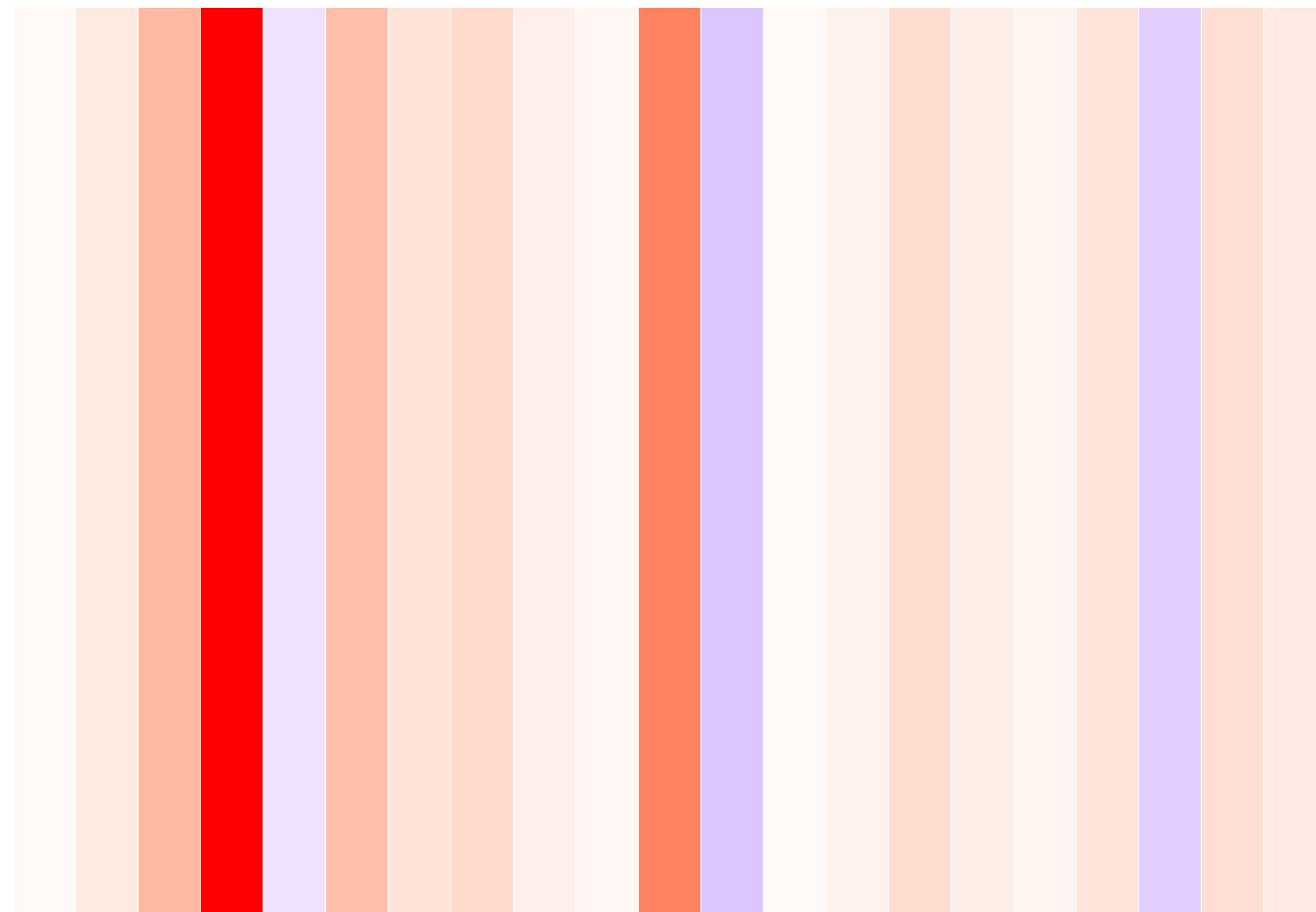
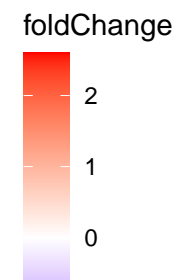
GeneRatio





Site gsea_path

Flagellar assembly



K02388

K02389

K02396

K02397

K02398

K02399

K02406

K02407

K02409

K02410

K02411

K02414

K02415

K02417

K02418

K02419

K02420

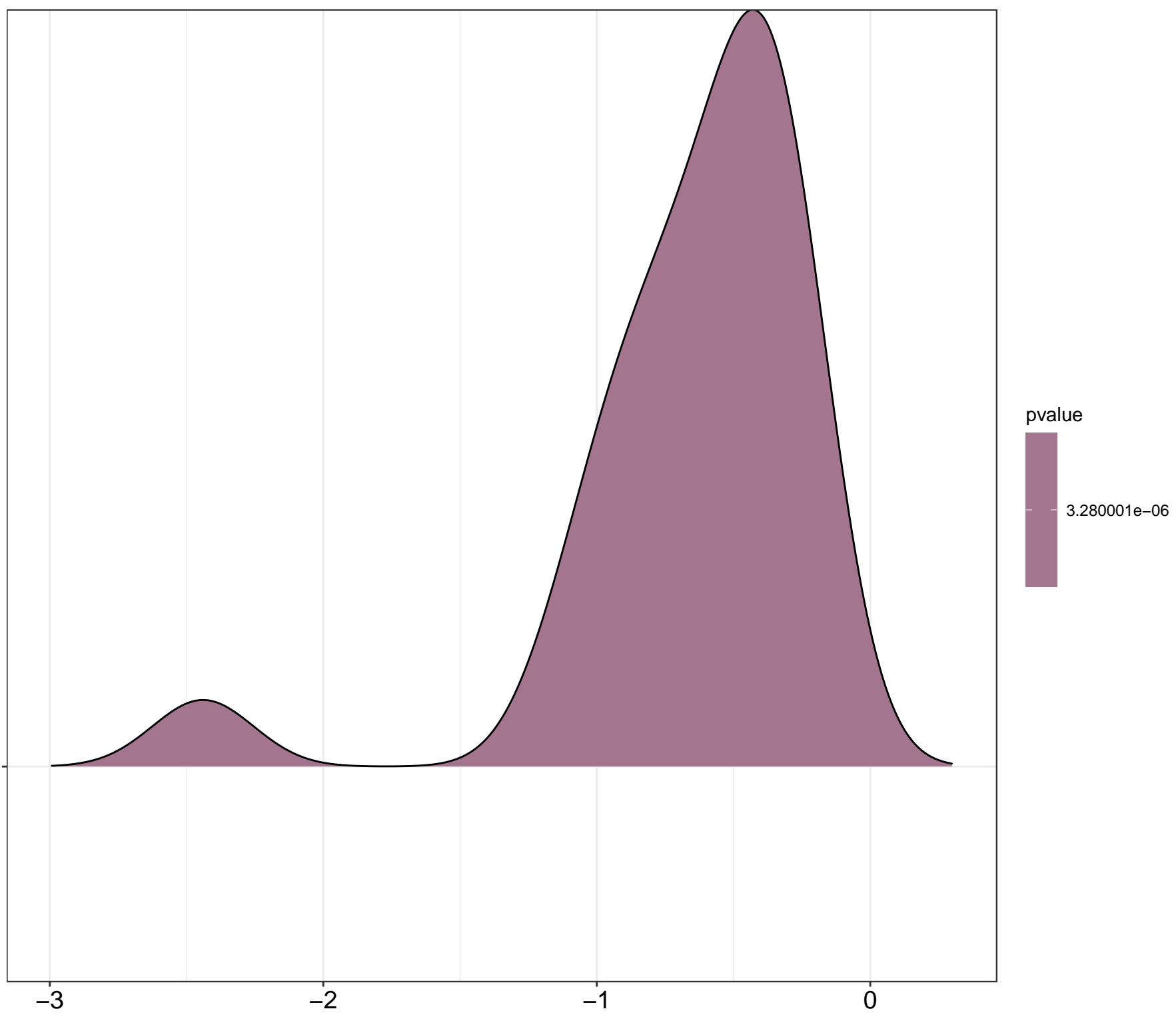
K02422

K02424

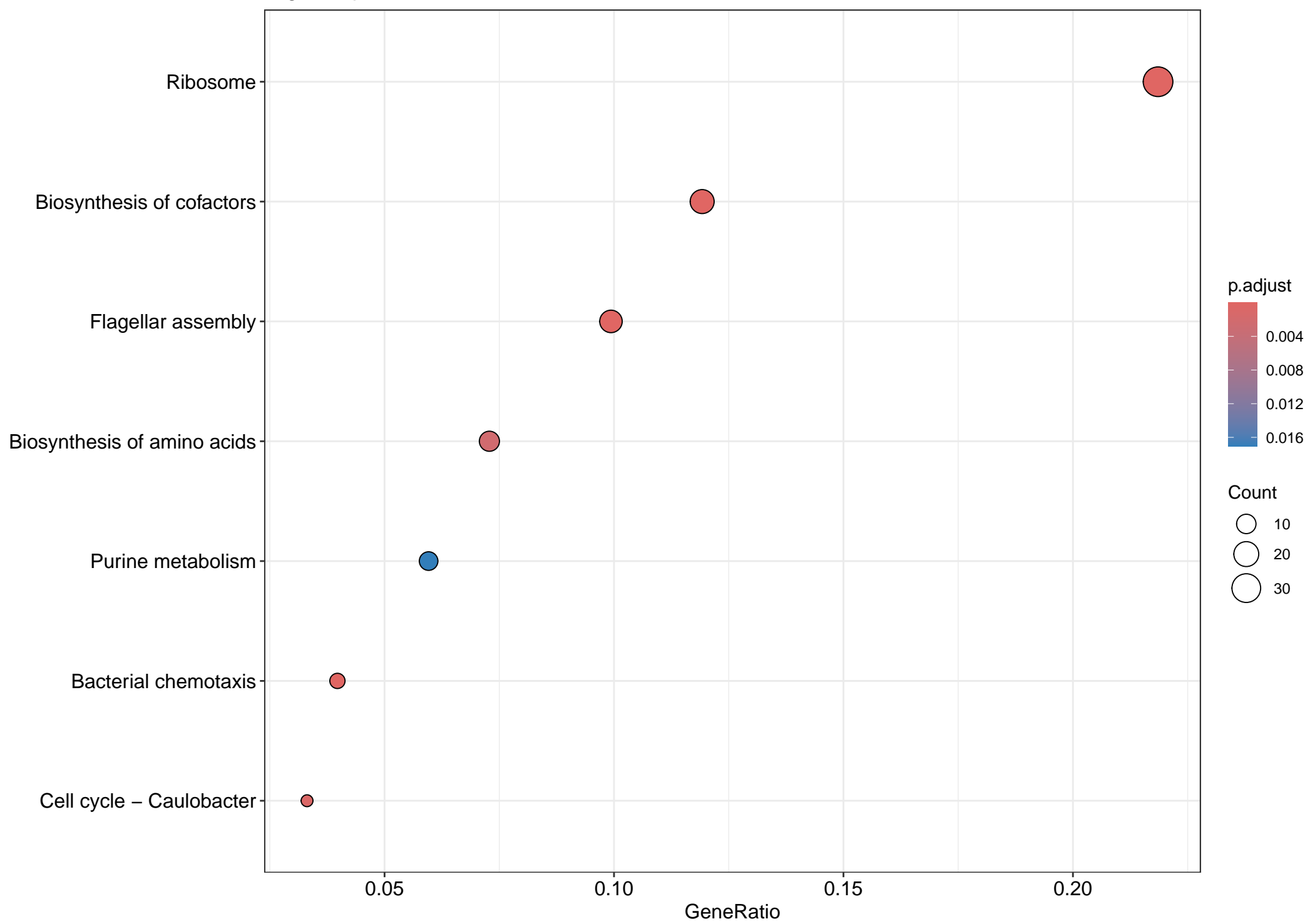
K02556

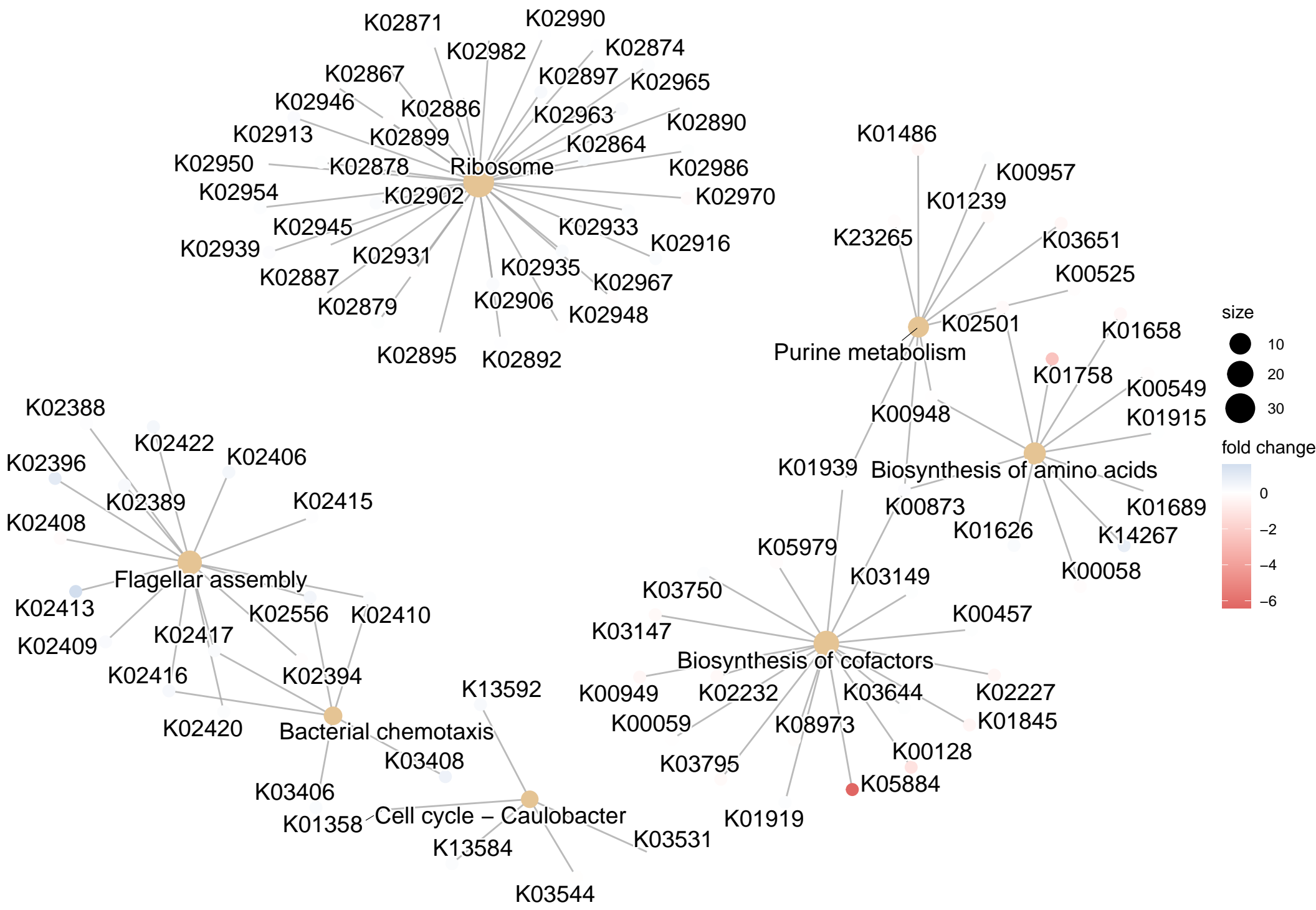
K02557

Flagellar assembly

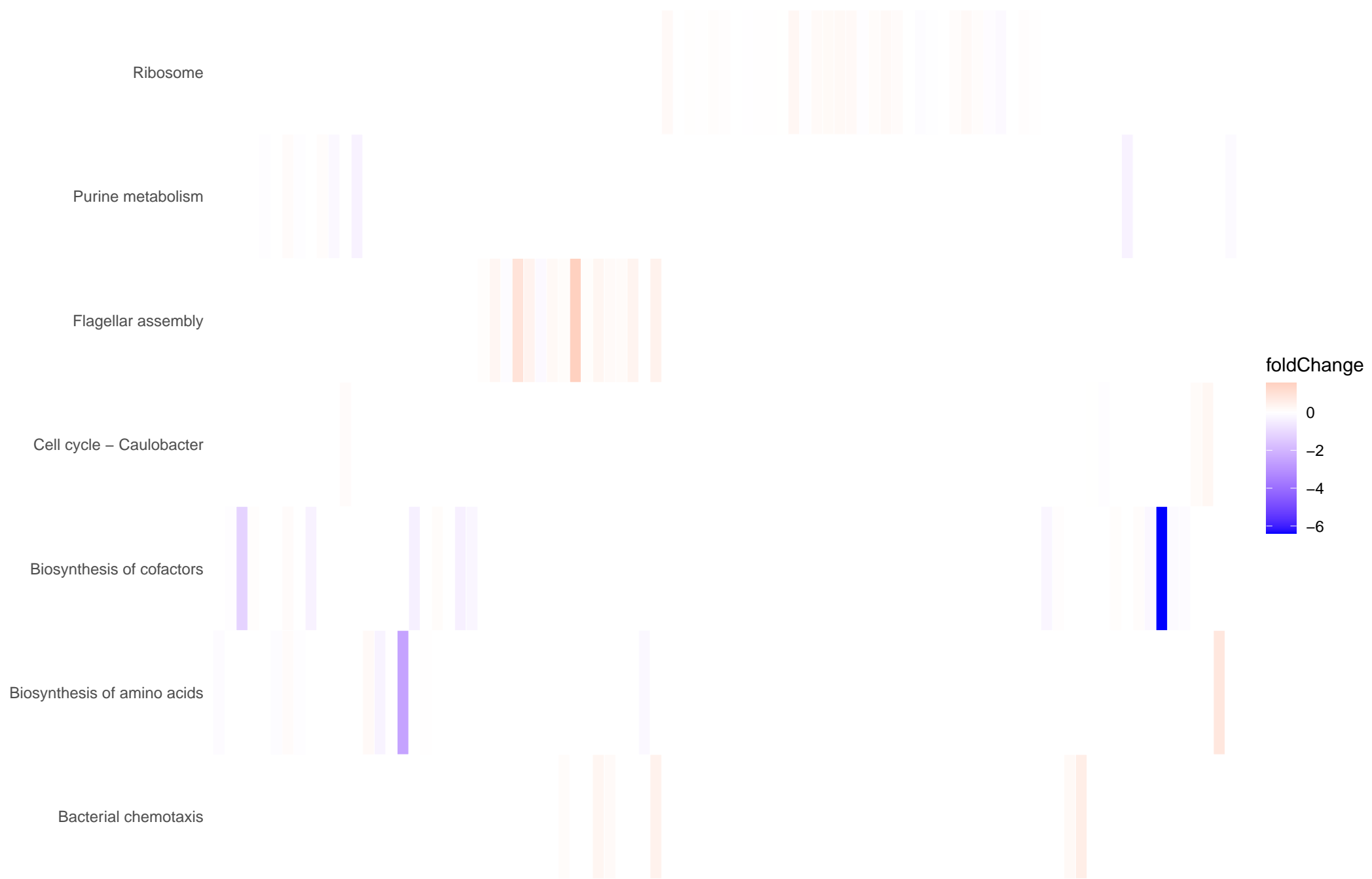


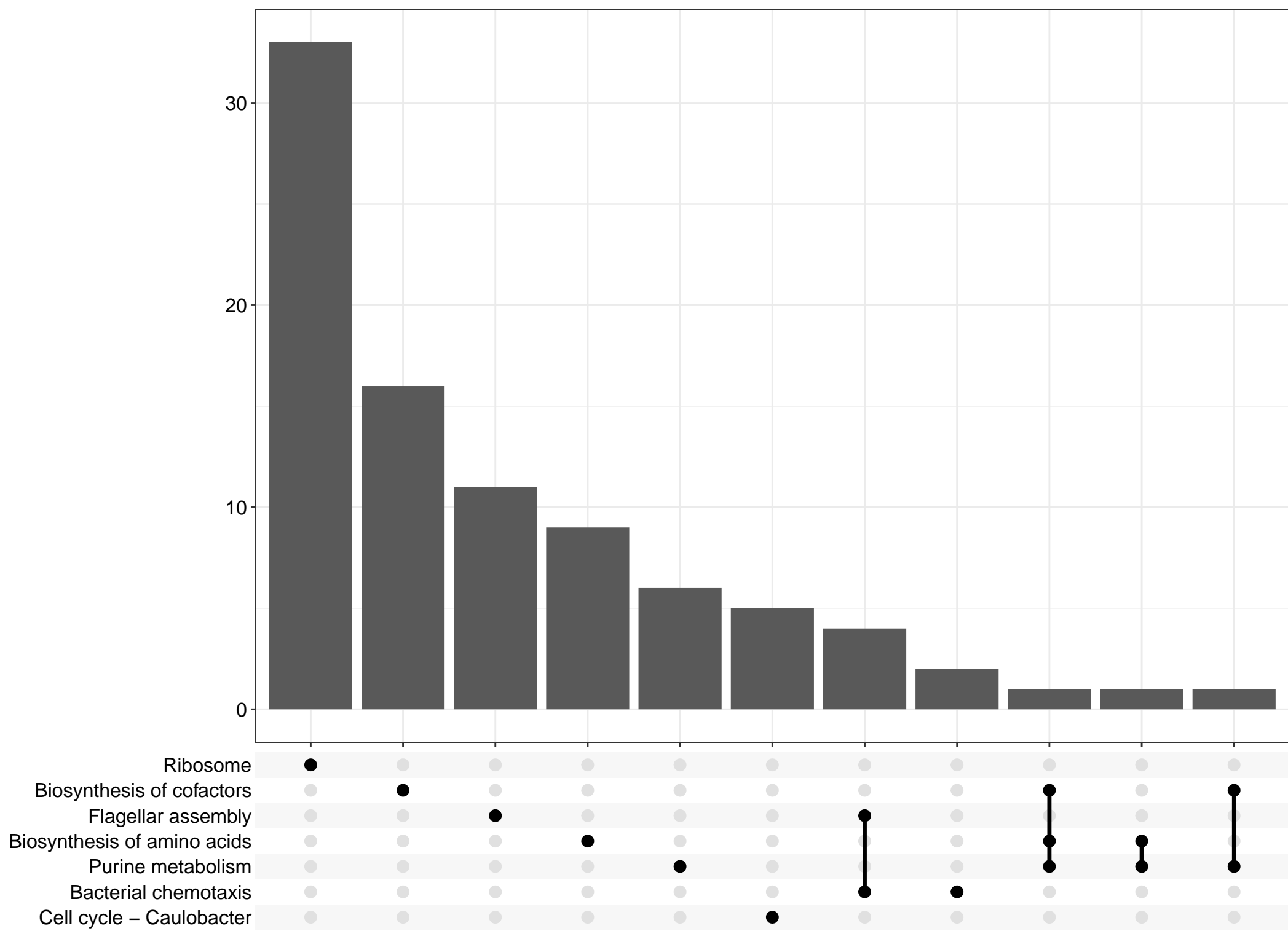
C geom_path

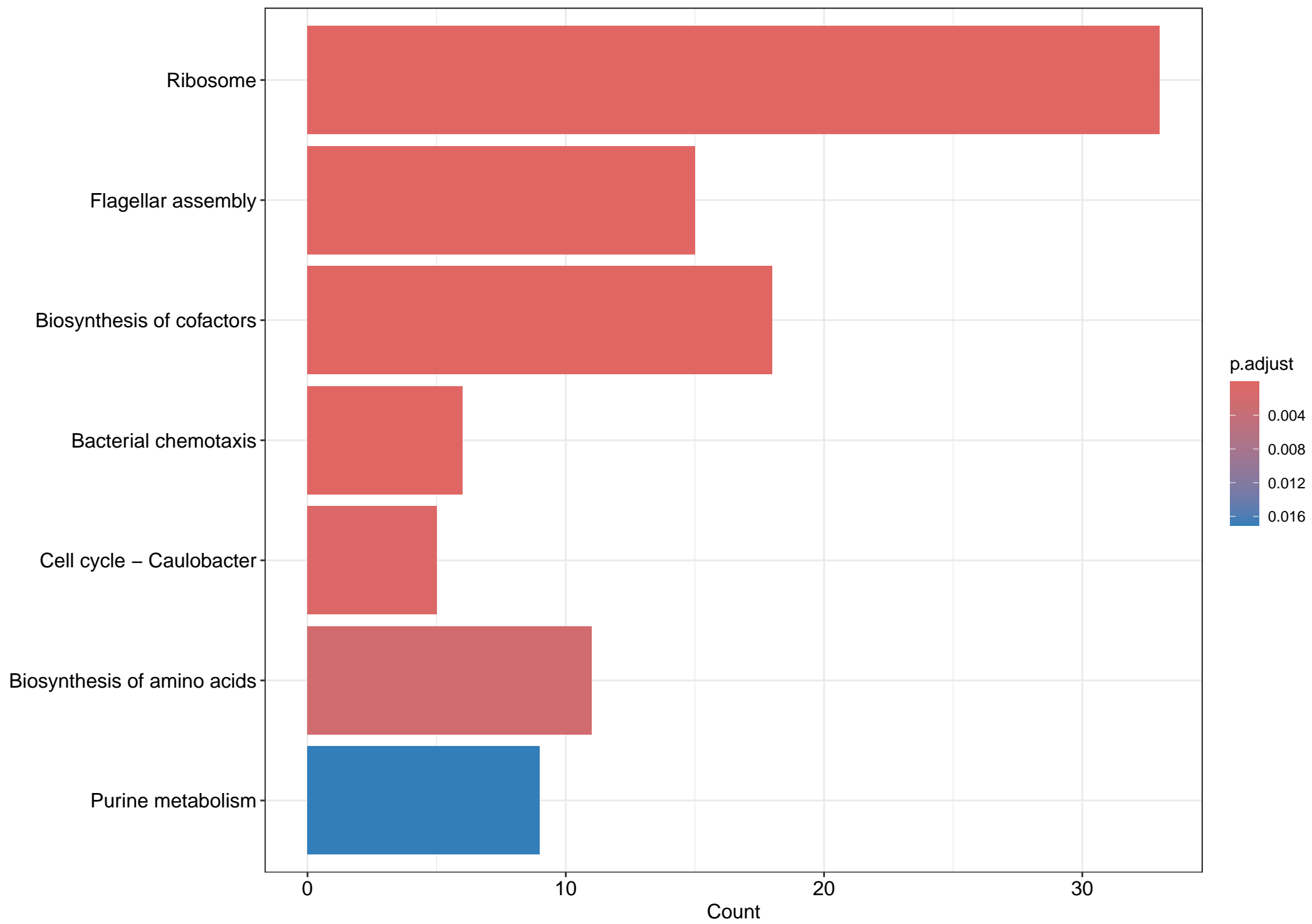




C geom_path







C gsea_path

Flagellar assembly

0.850

0.875

0.900

0.925

GeneRatio

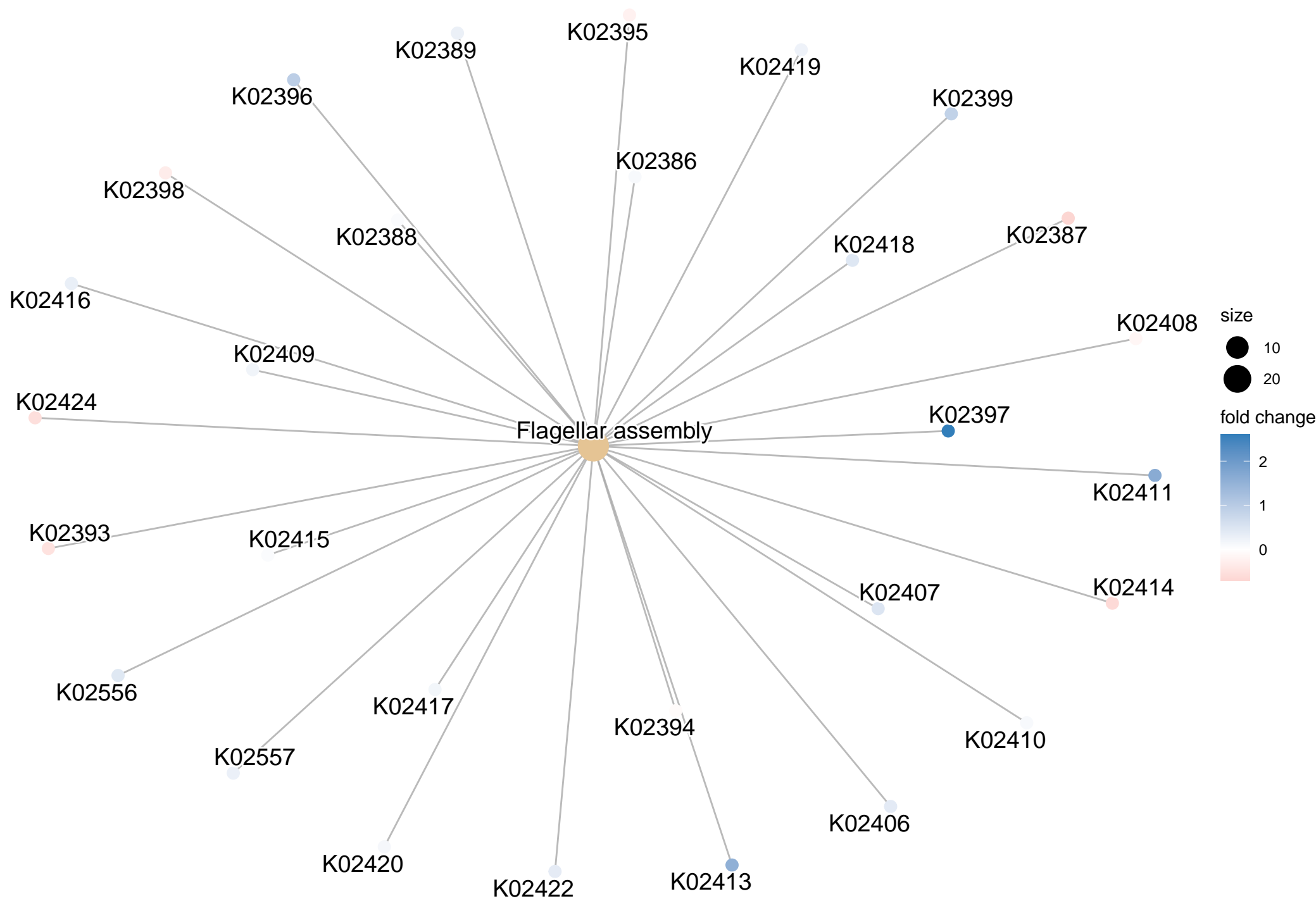
p.adjust

3.25e-08

Count

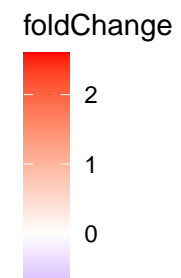
29





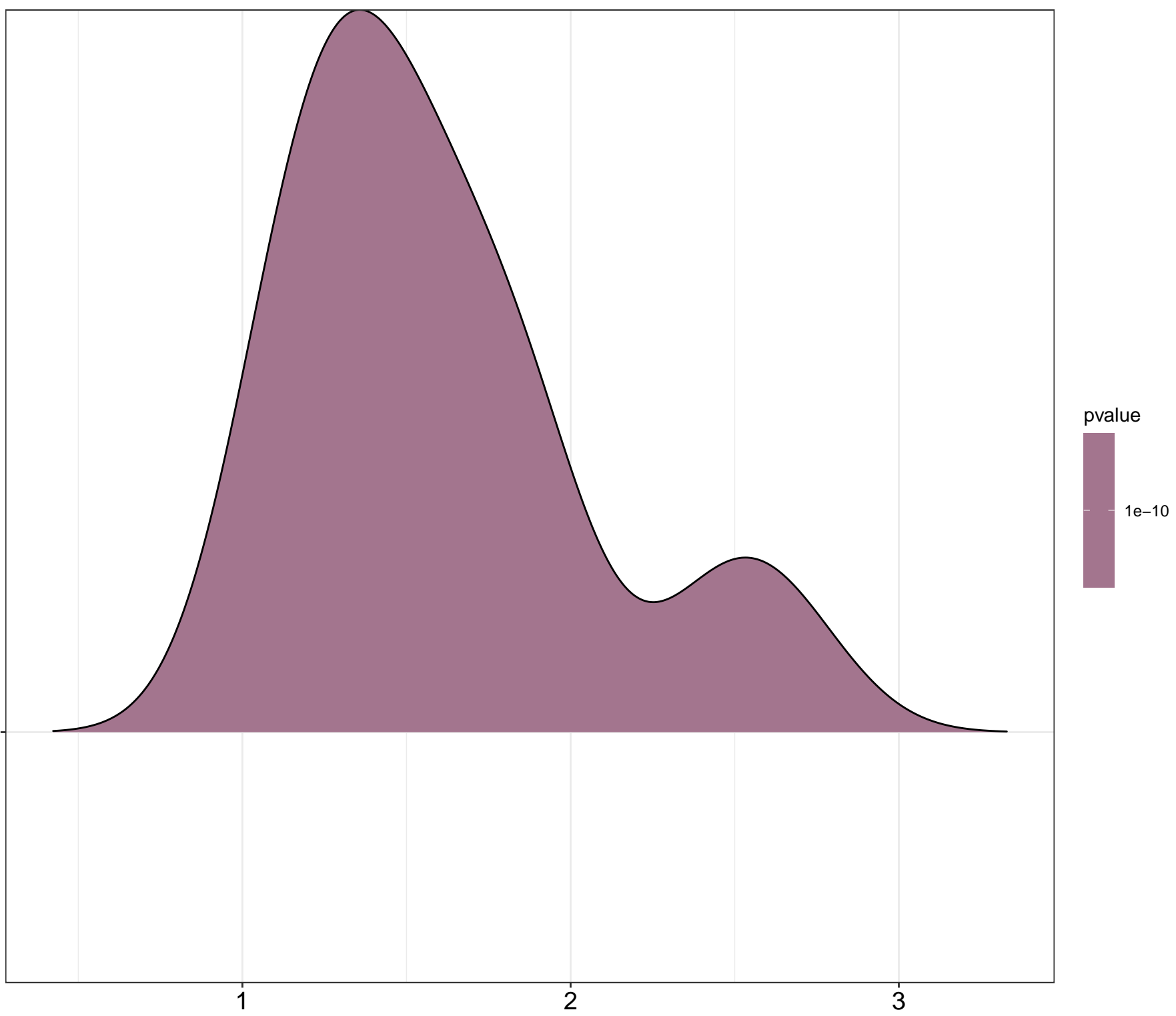
C gsea_path

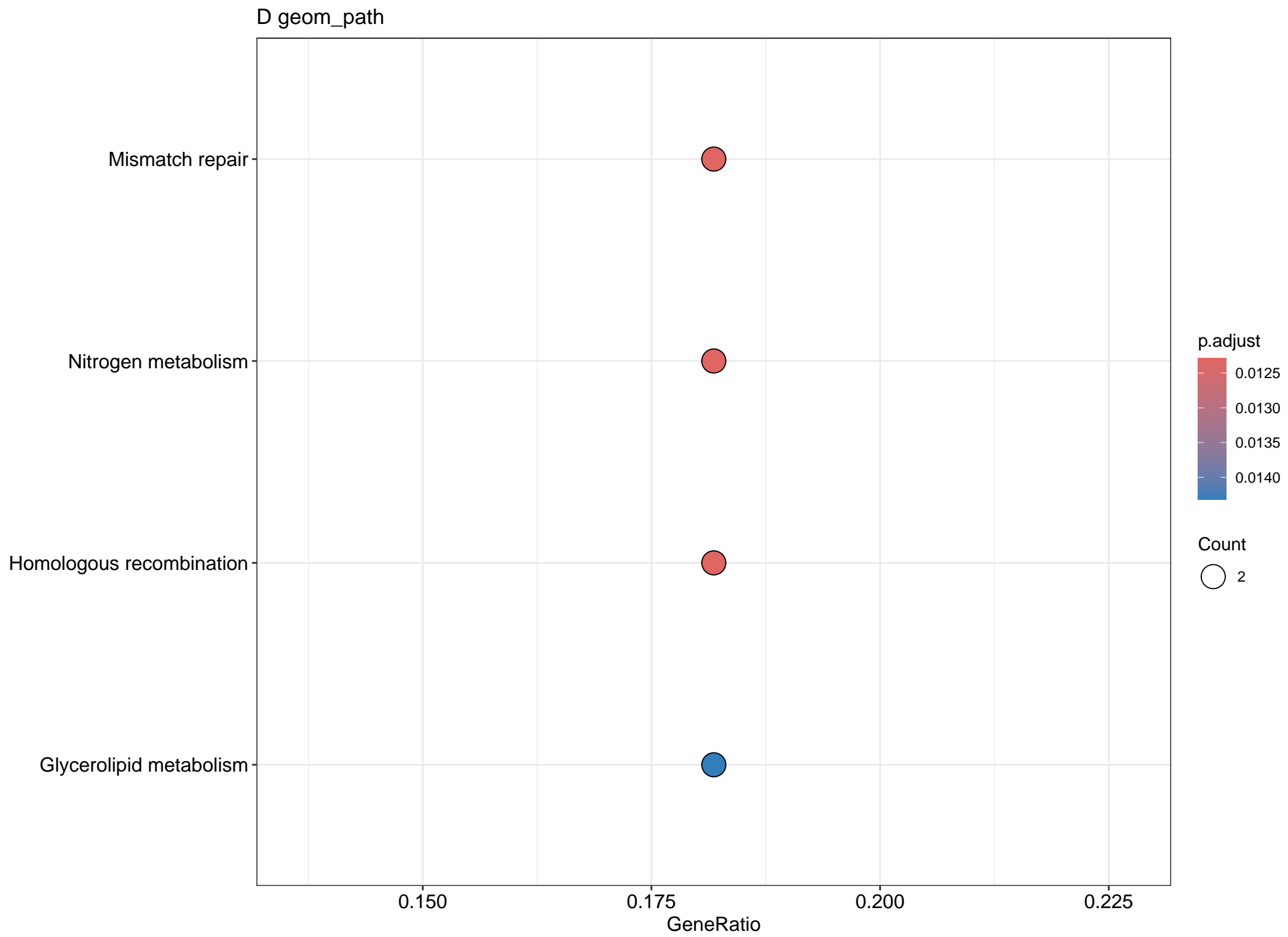
Flagellar assembly



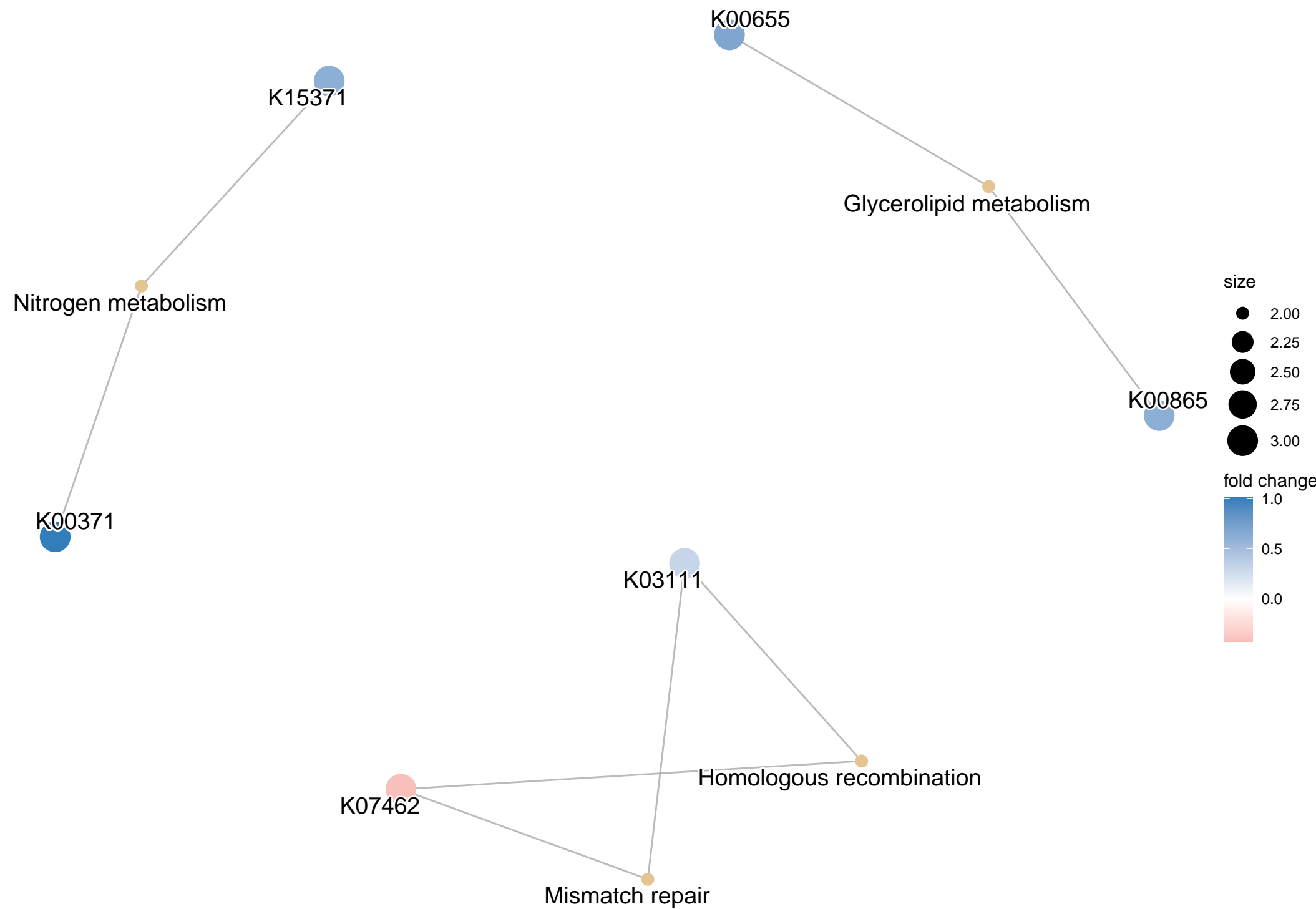
K02386
K02387
K02388
K02389
K02393
K02394
K02395
K02396
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K02398
K02399
K02406
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K02408
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K02416
K02417
K02418
K02419
K02420
K02422
K02424
K02556
K02557

Flagellar assembly





D geom_path



D geom_path

Nitrogen metabolism

Mismatch repair

Homologous recombination

Glycerolipid metabolism

foldChange

1.0

0.5

0.0

K00371

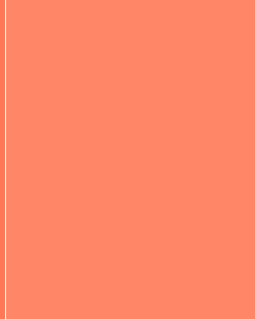
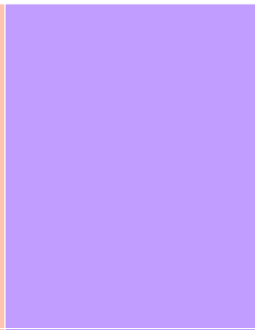
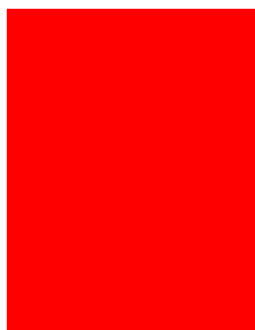
K00655

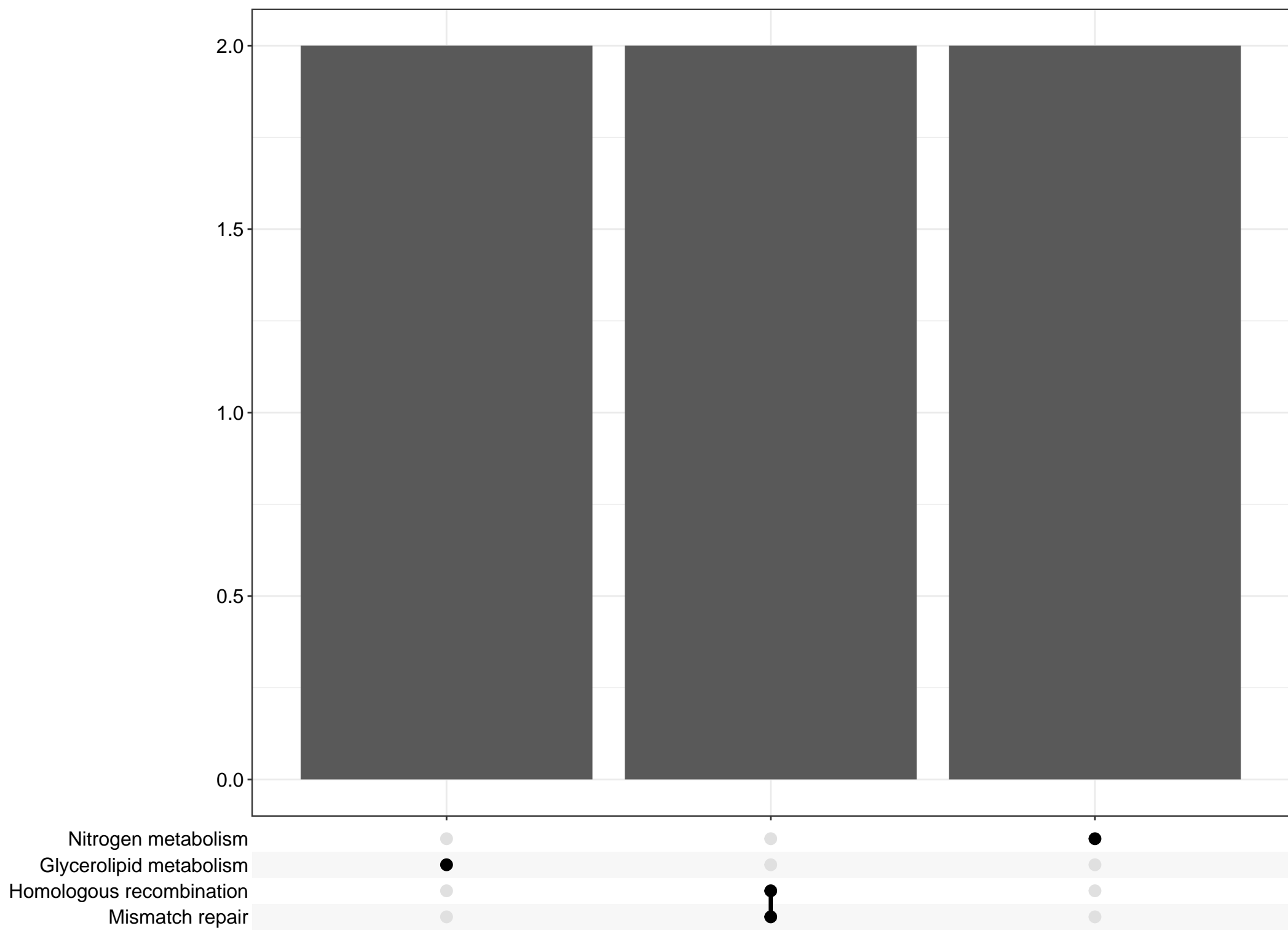
K00865

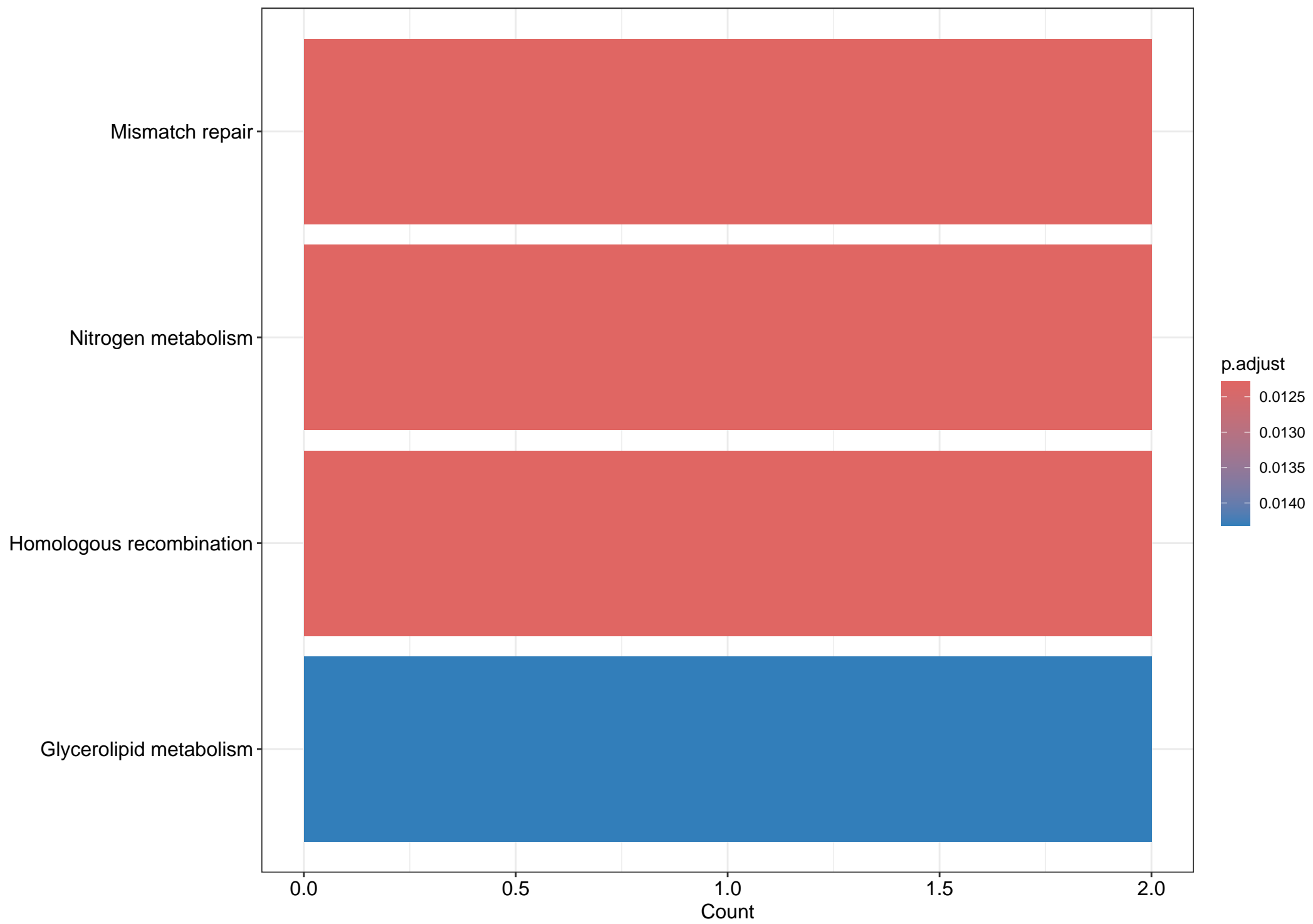
K03111

K07462

K15371







D gsea_path

Coronavirus disease – COVID-19

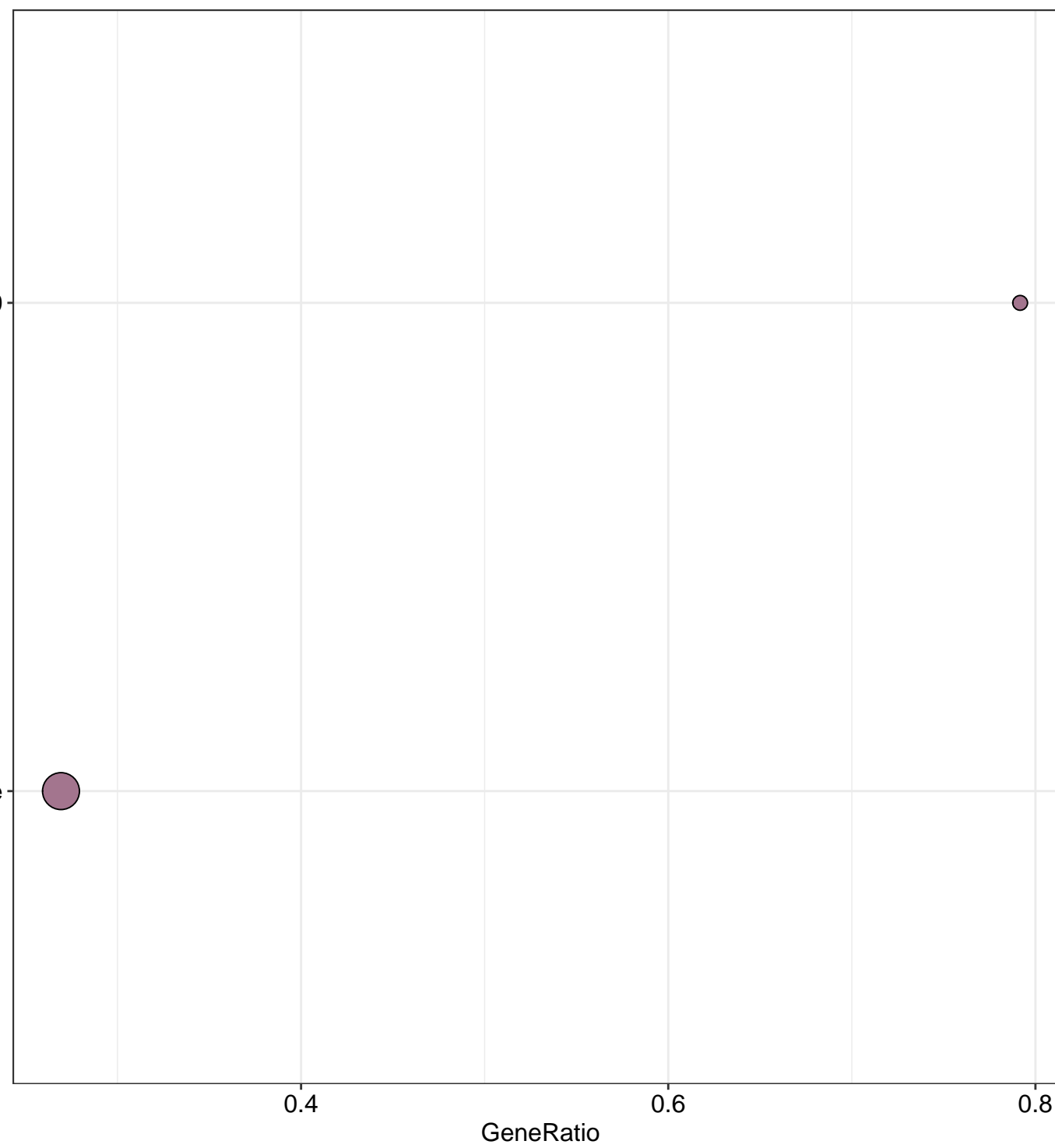
Ribosome

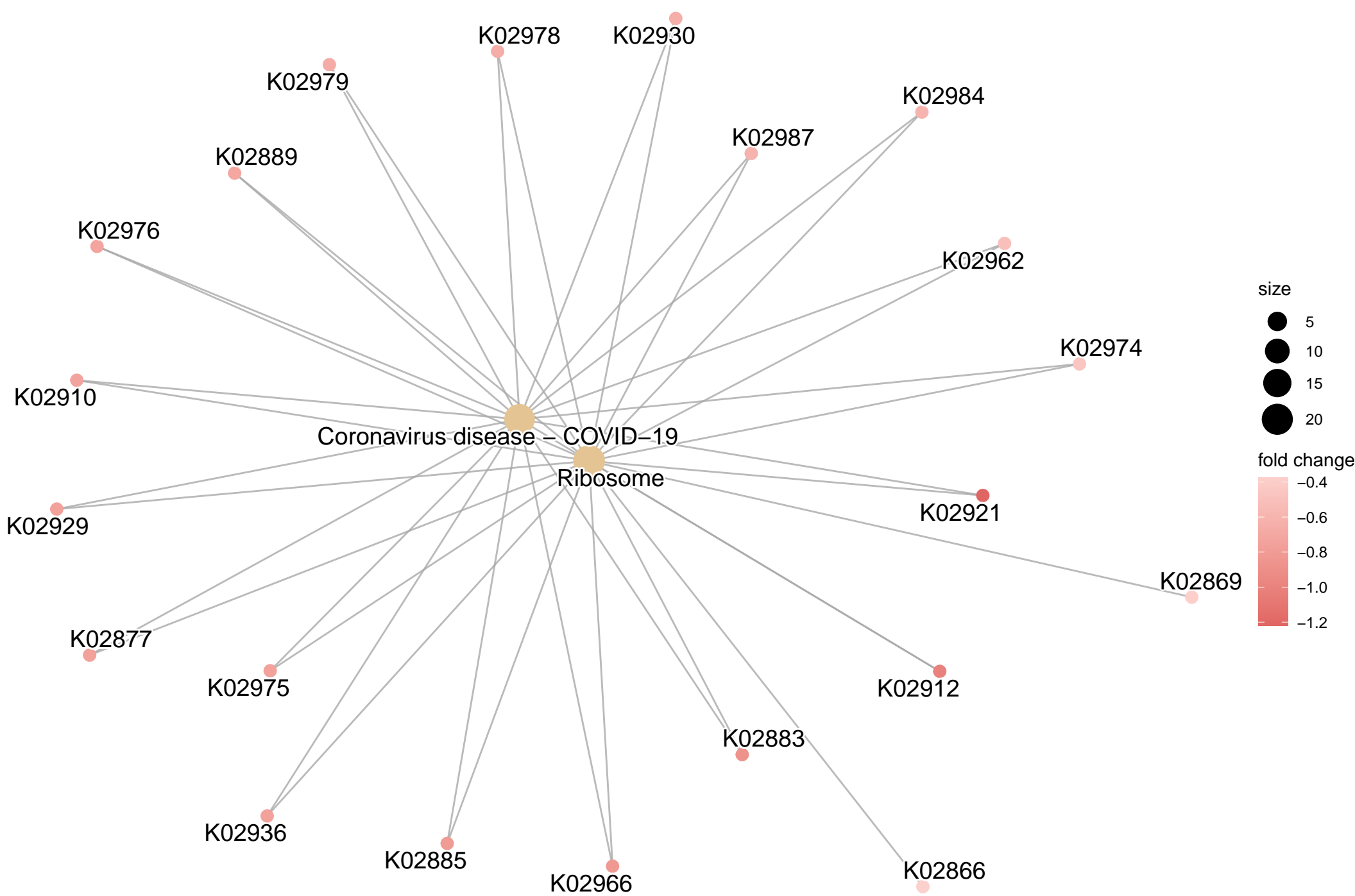
p.adjust

0.0434248

Count

- 19.0
- 19.5
- 20.0
- 20.5
- 21.0

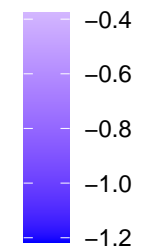




D gsea_path

Ribosome

foldChange



Coronavirus disease – COVID-19

