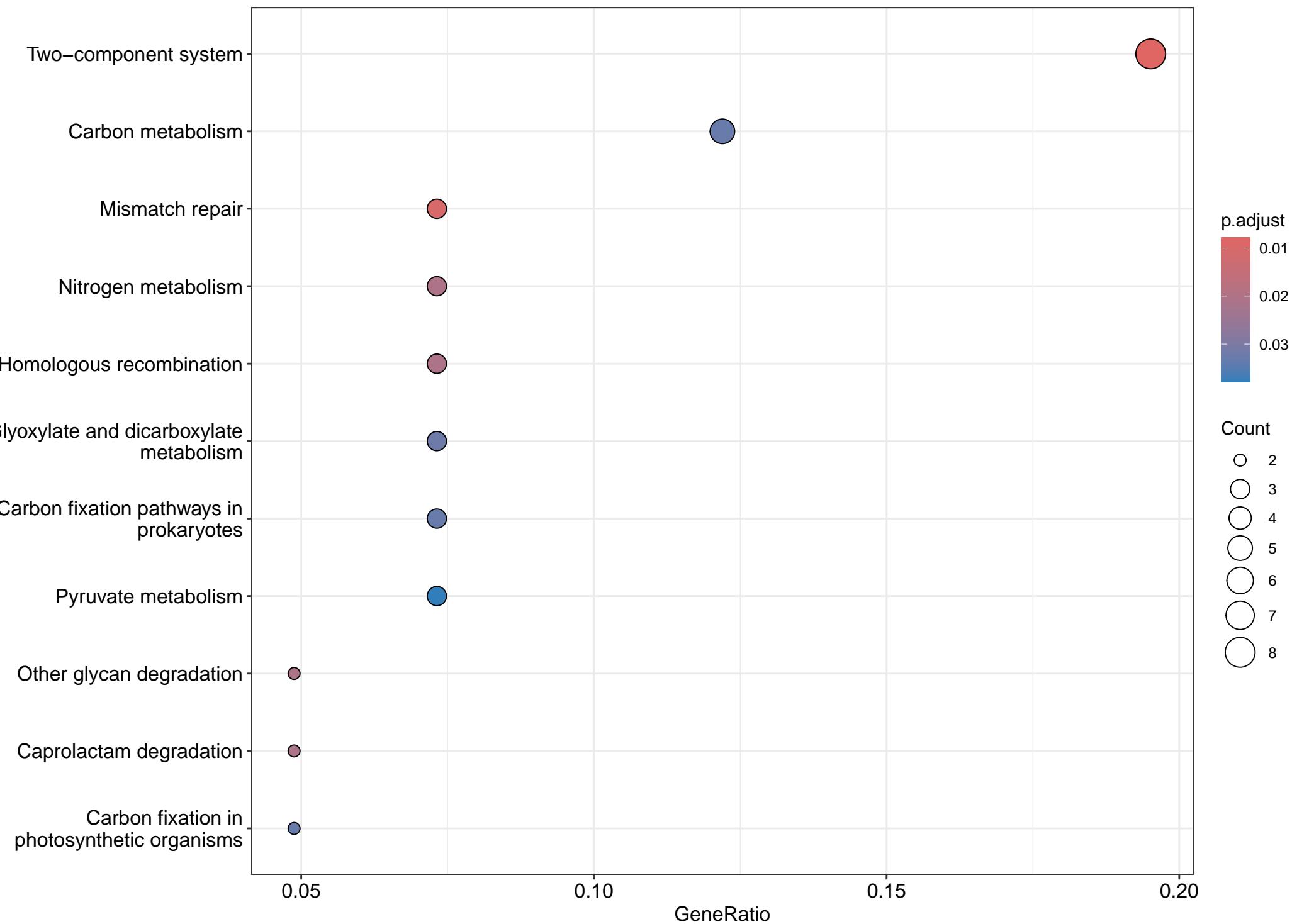
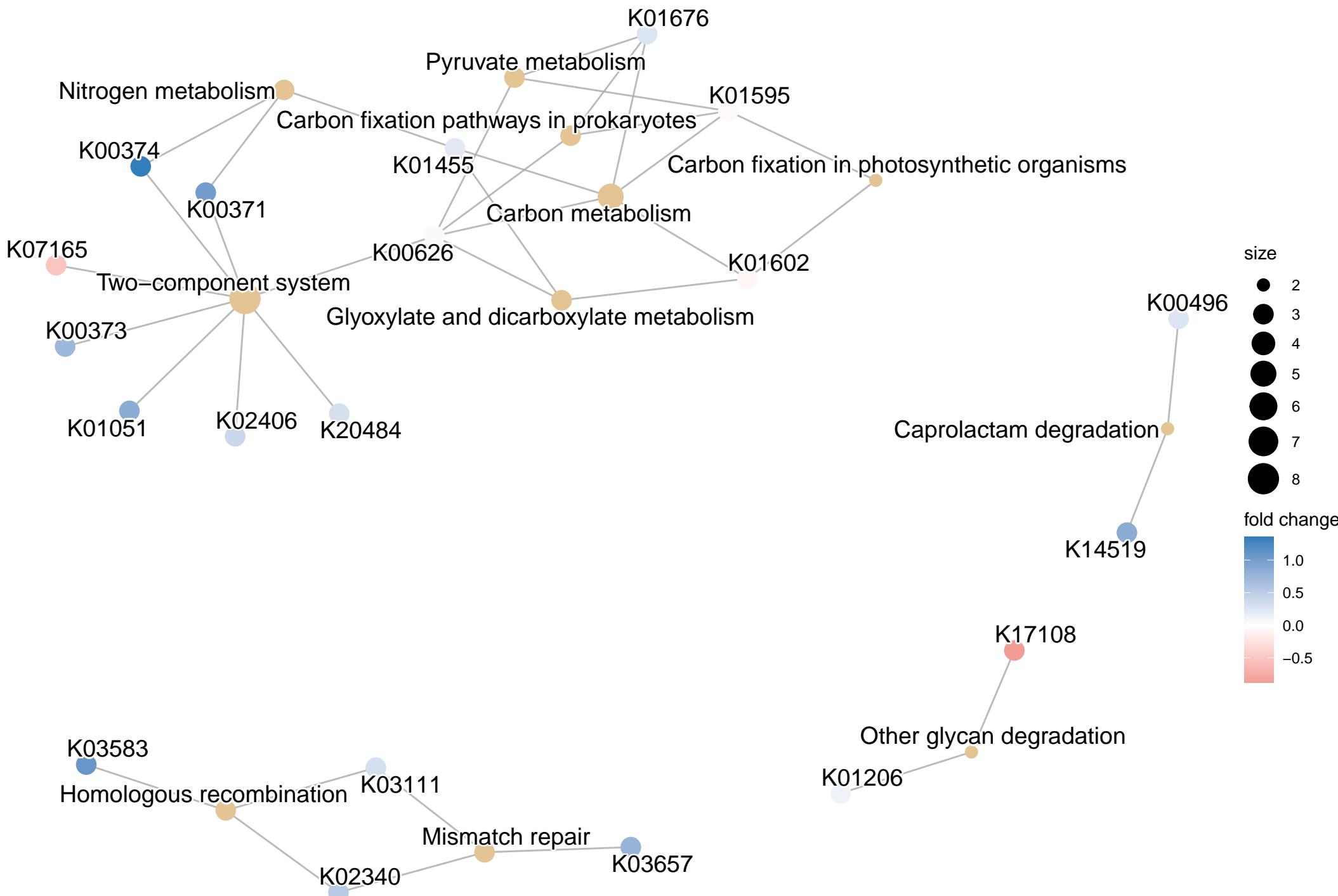


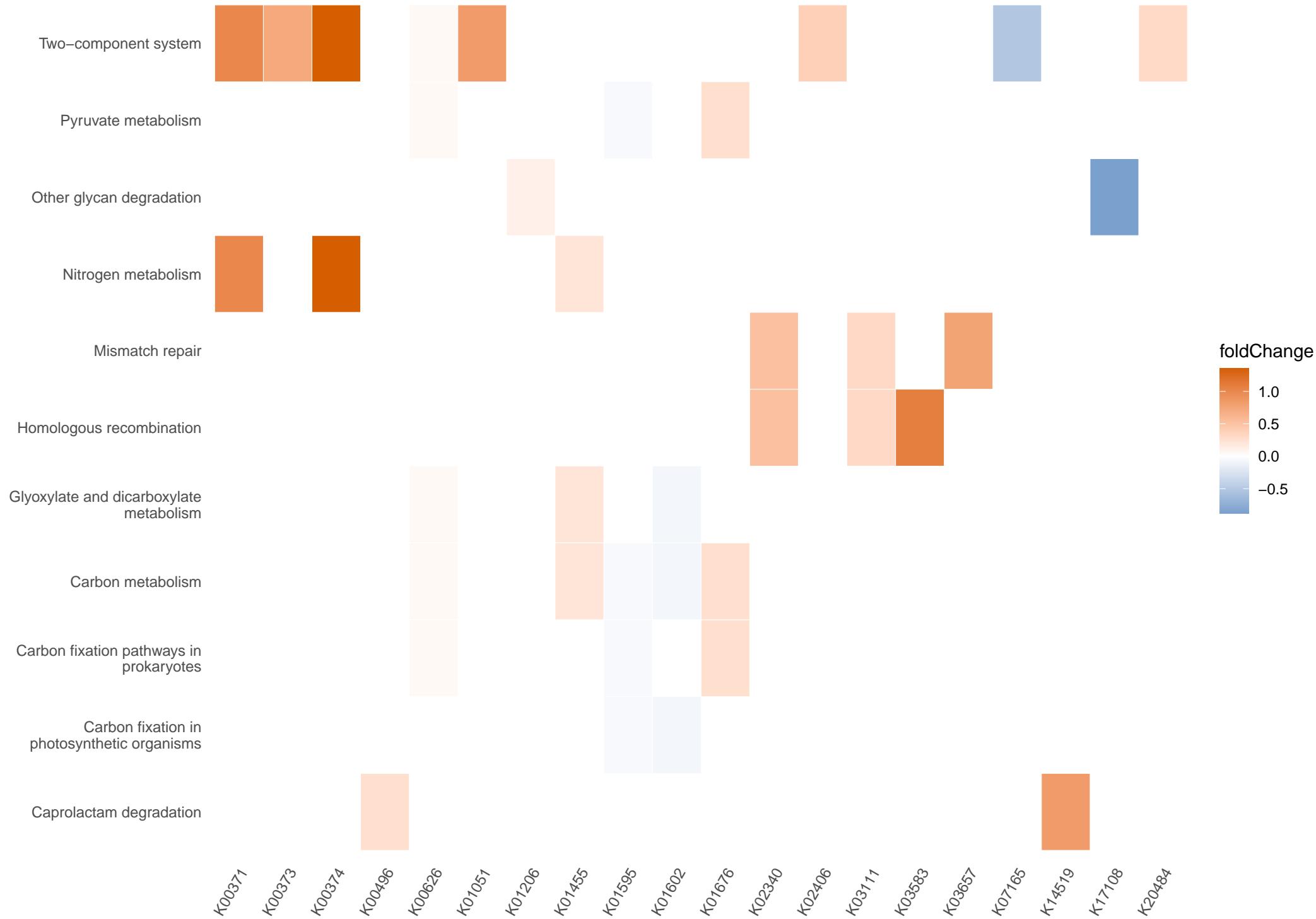
Site geom_path

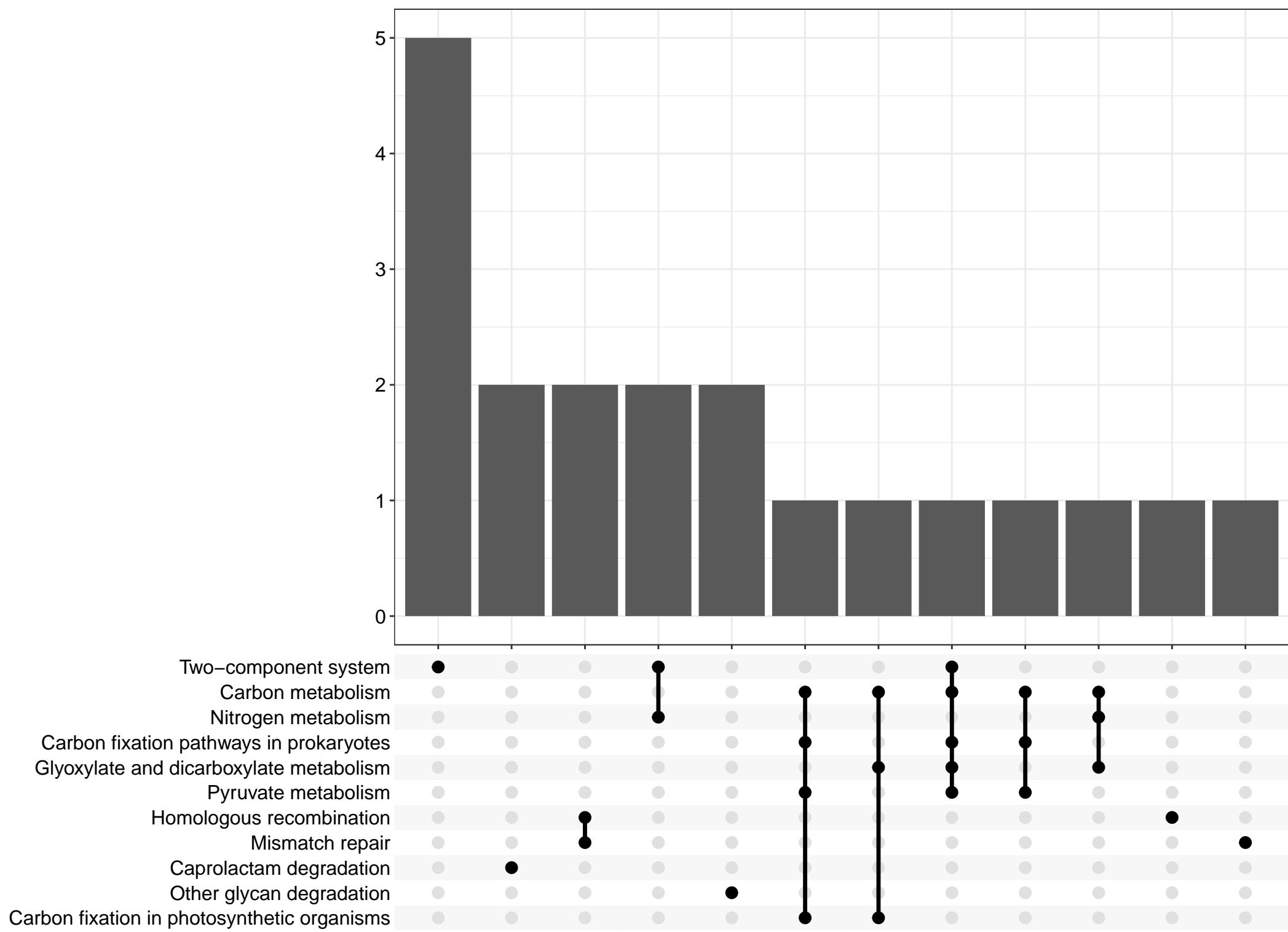


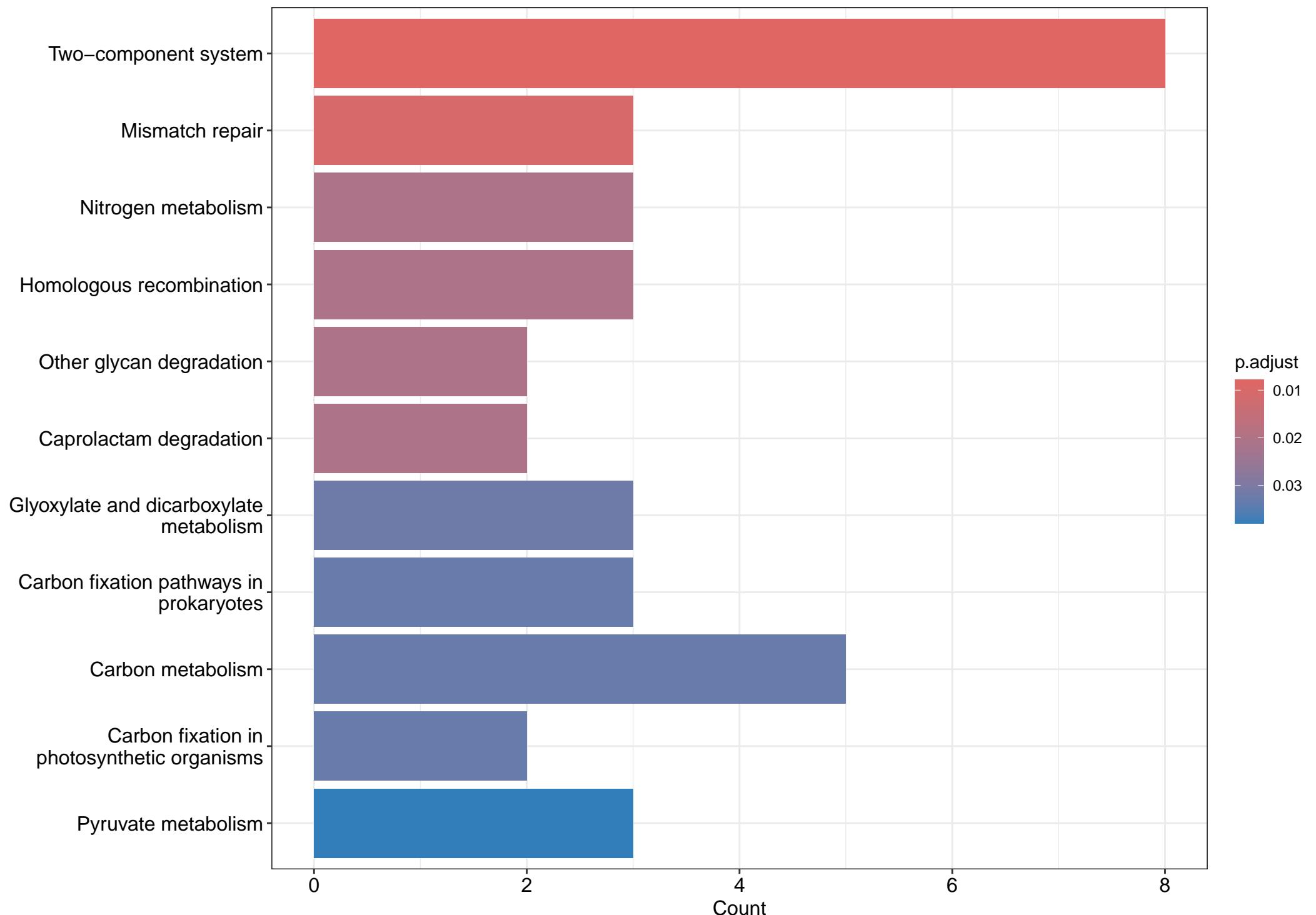
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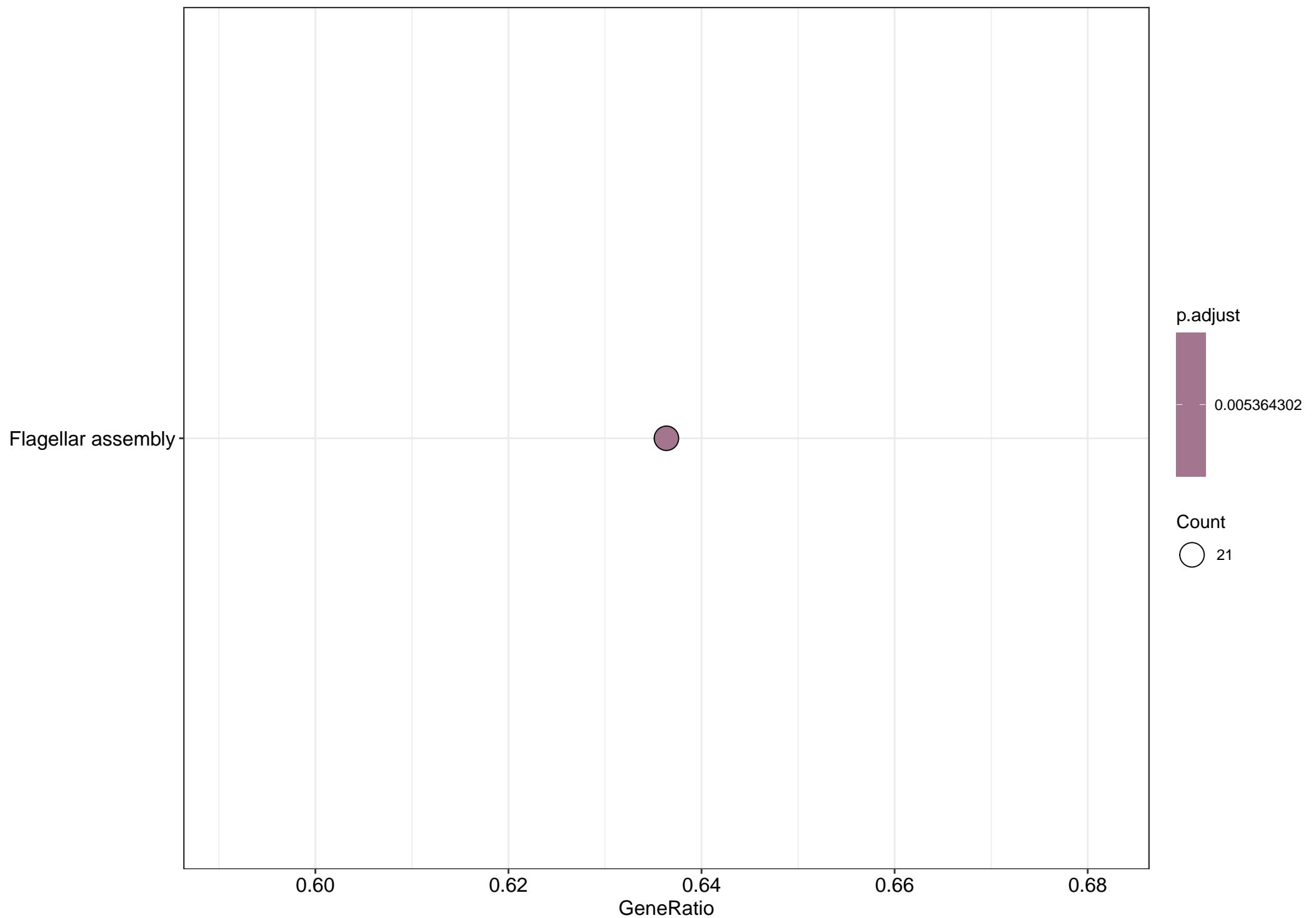
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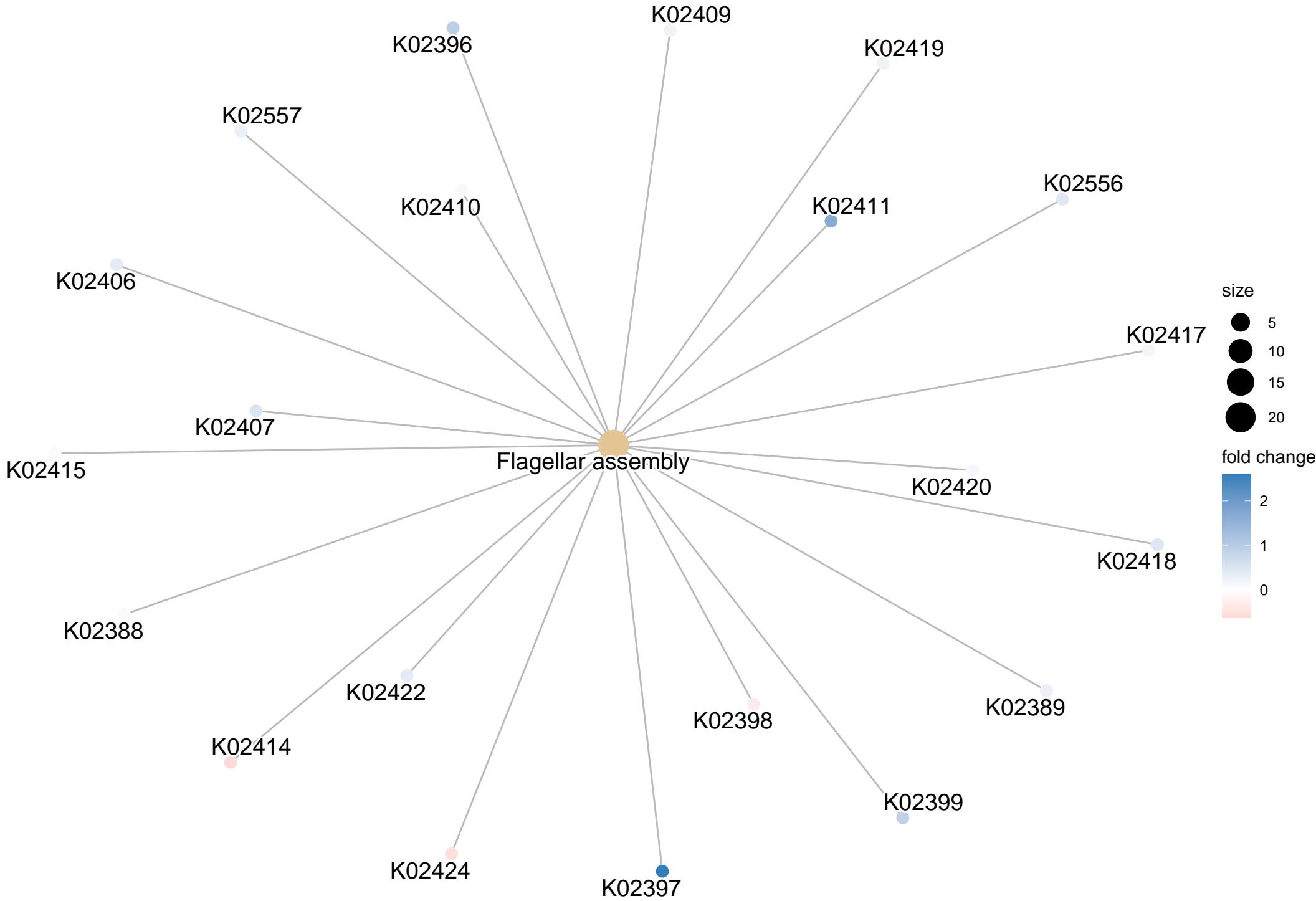




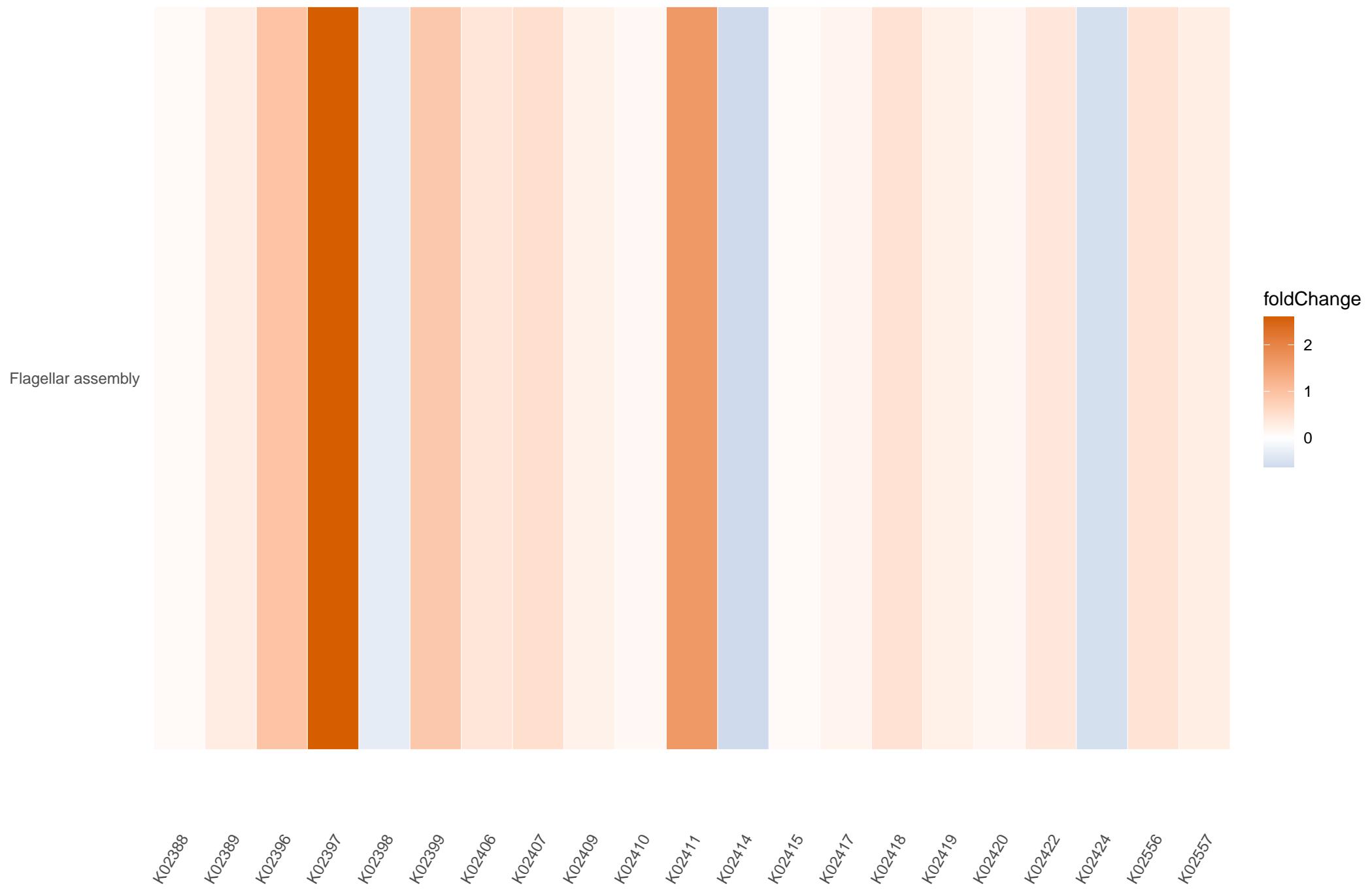


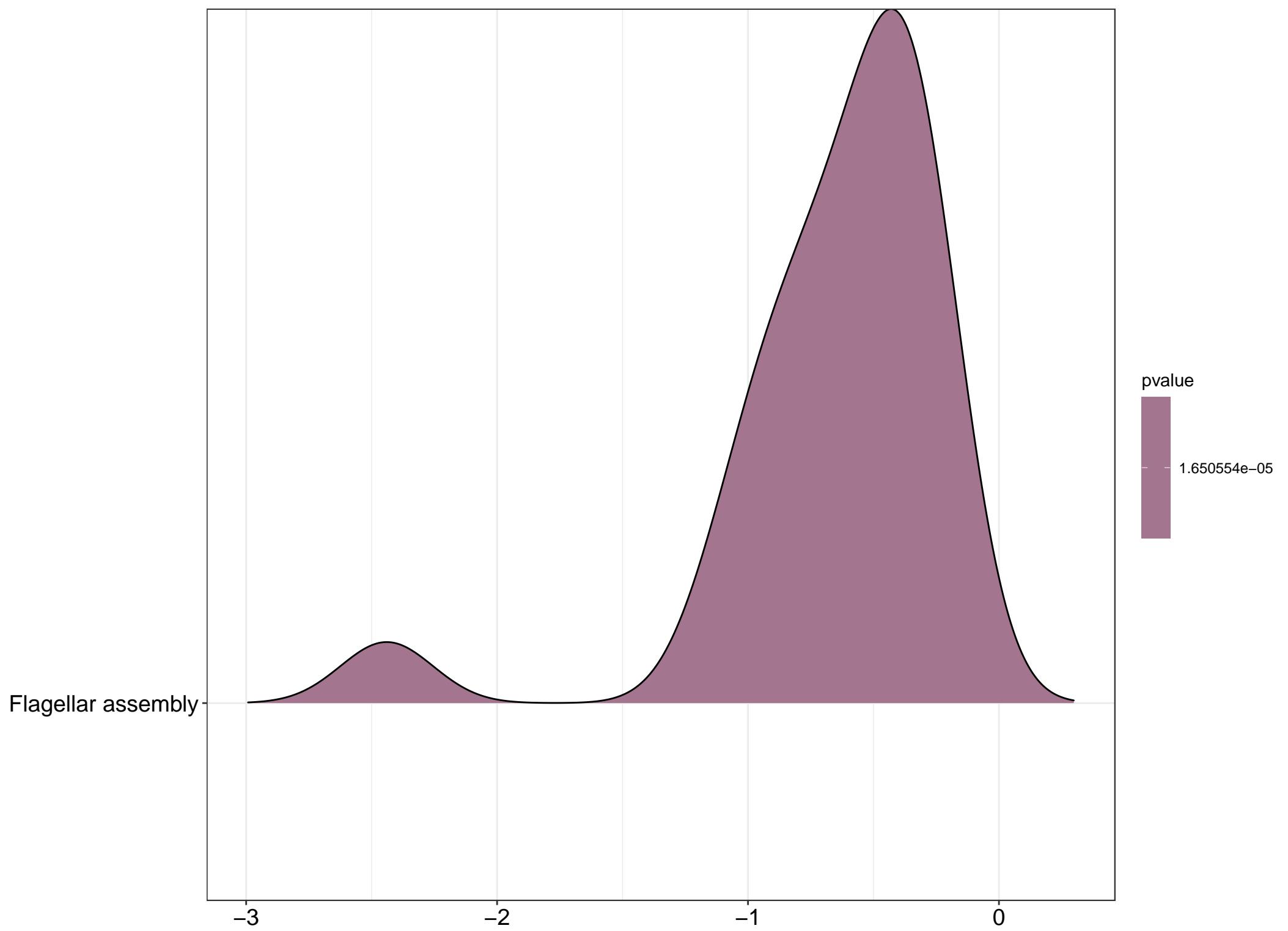
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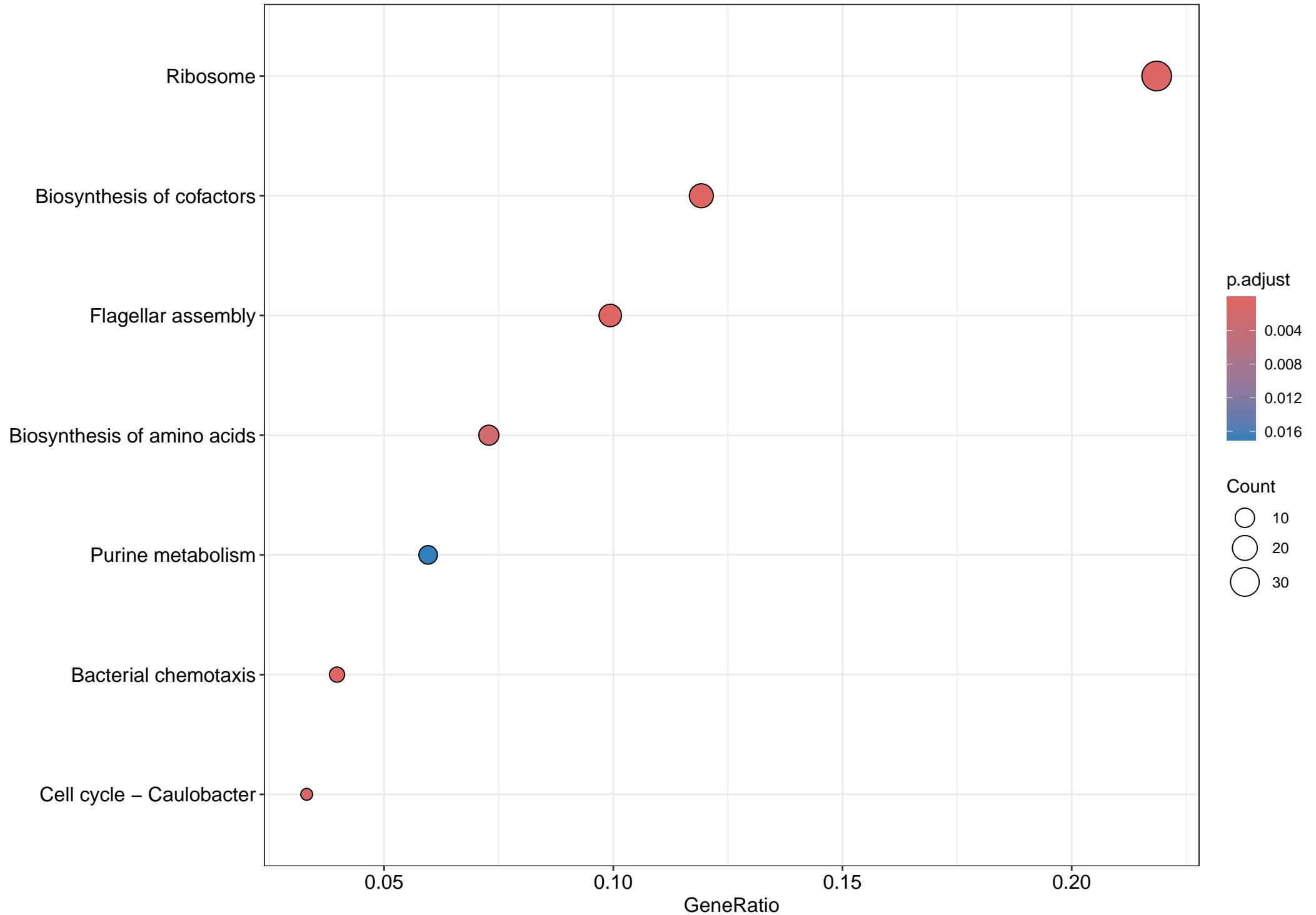


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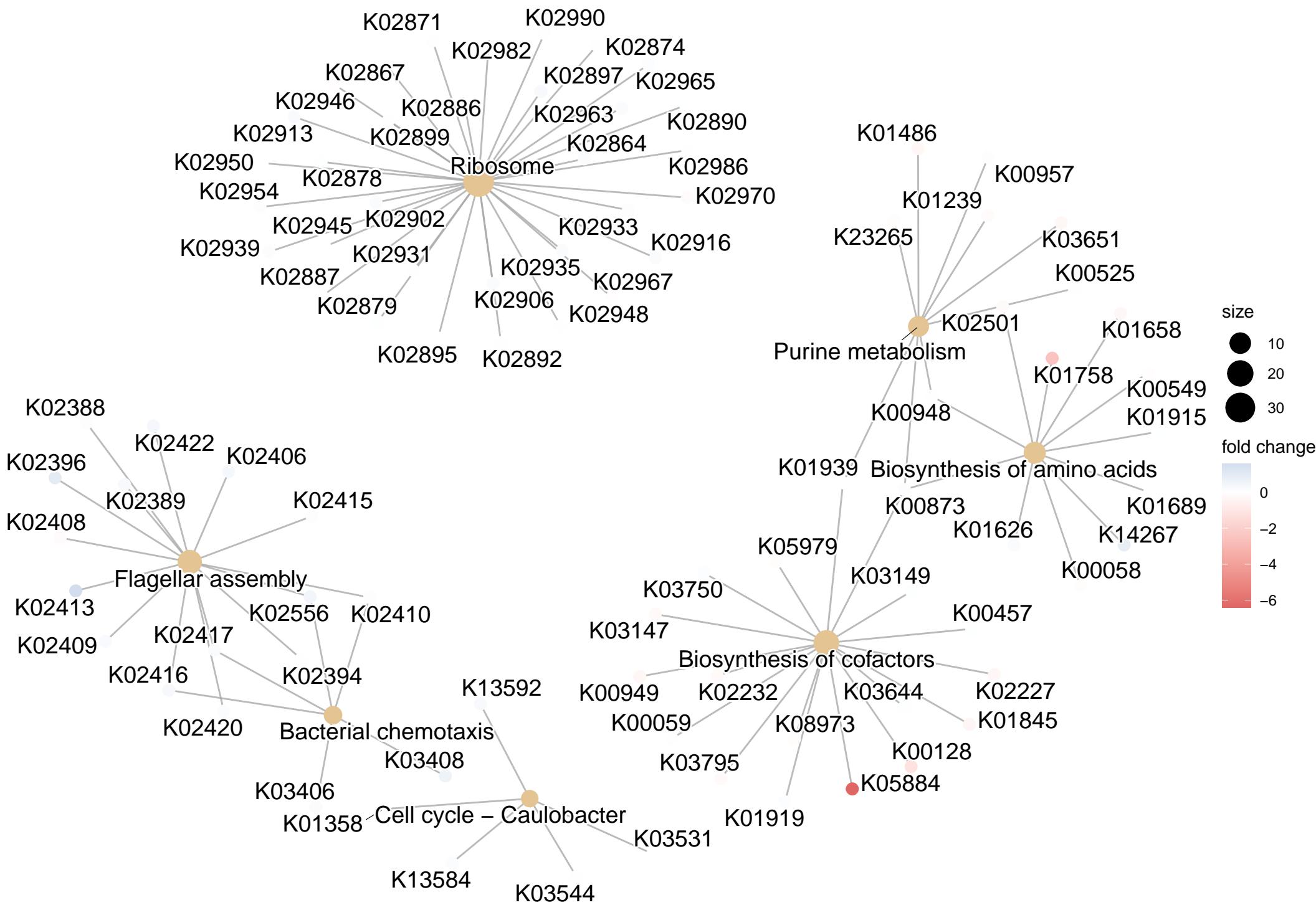




C geom_path



C geom path



C geom_path

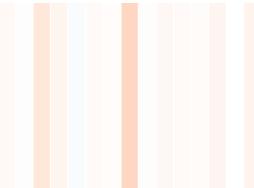
Ribosome



Purine metabolism



Flagellar assembly



Cell cycle – Caulobacter



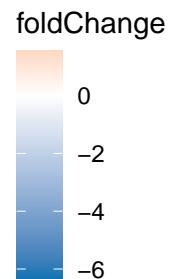
Biosynthesis of cofactors

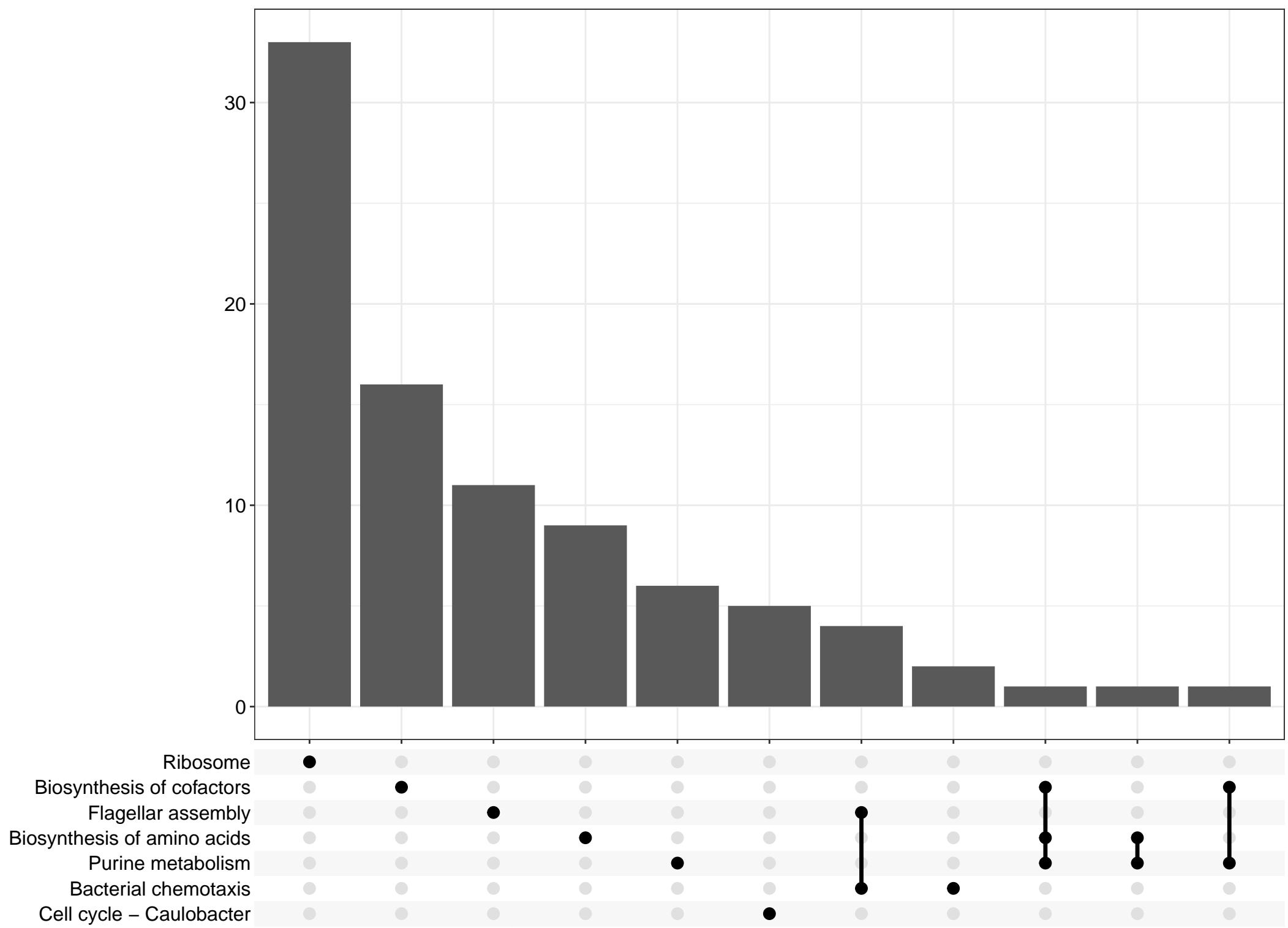


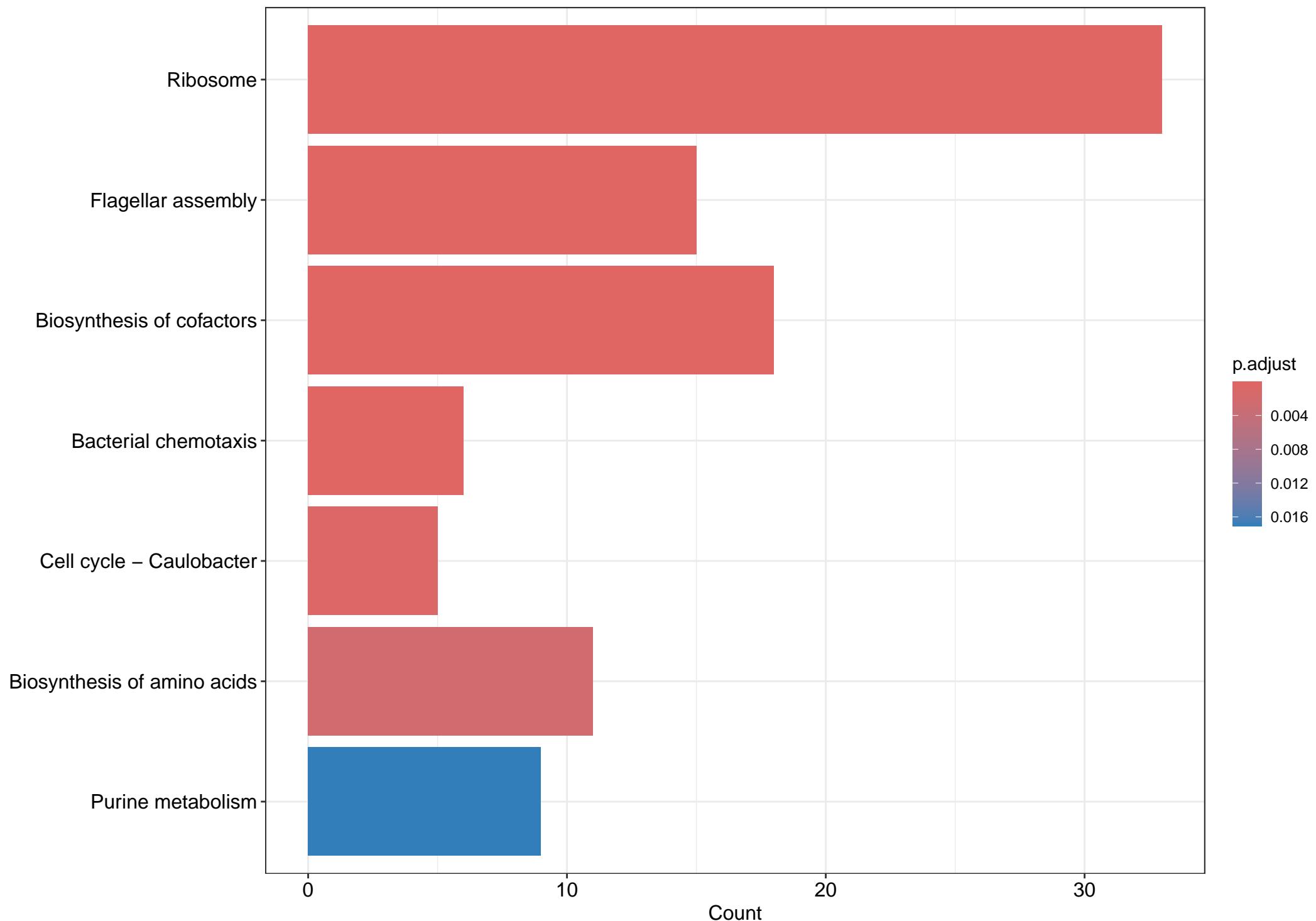
Biosynthesis of amino acids



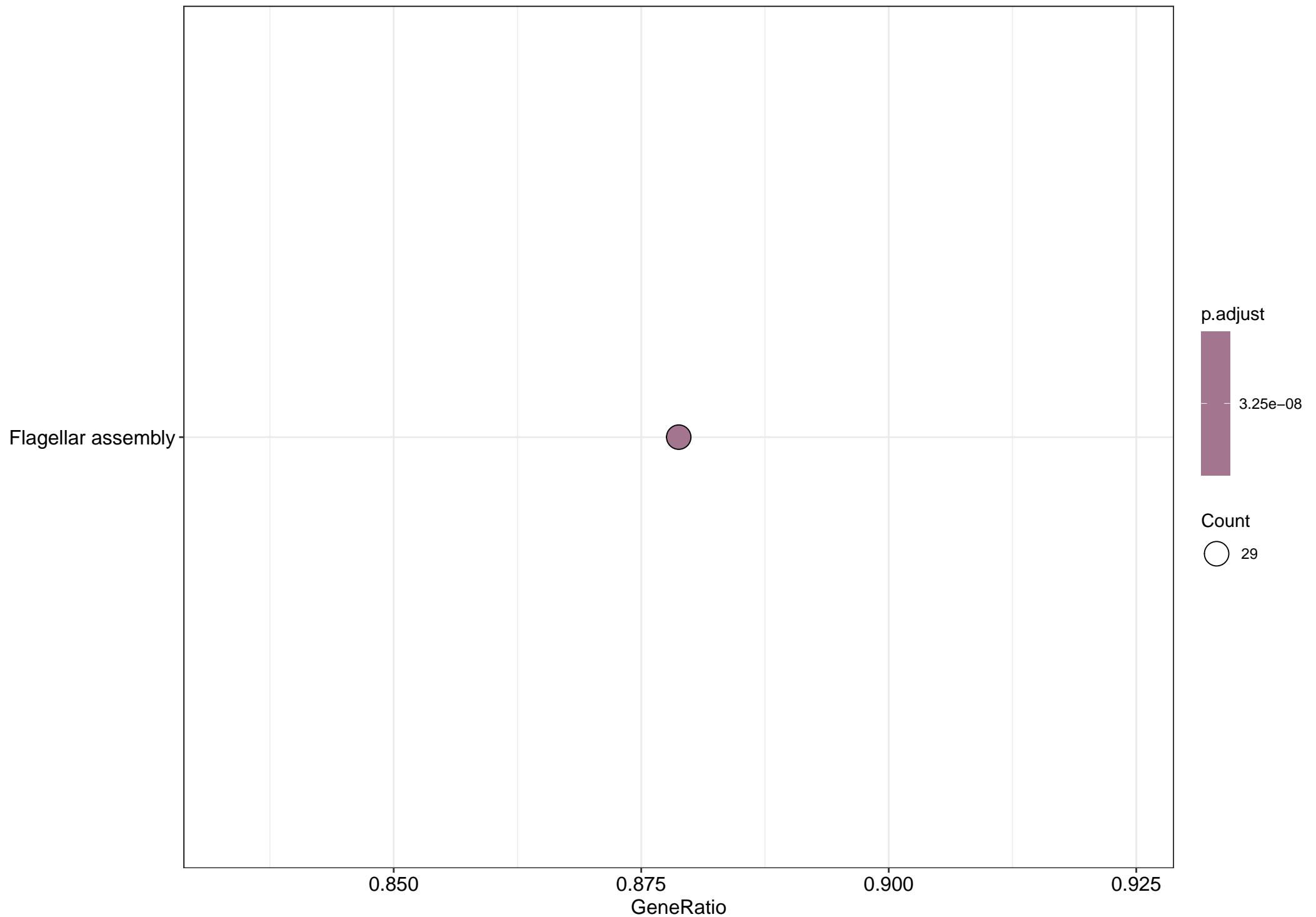
Bacterial chemotaxis



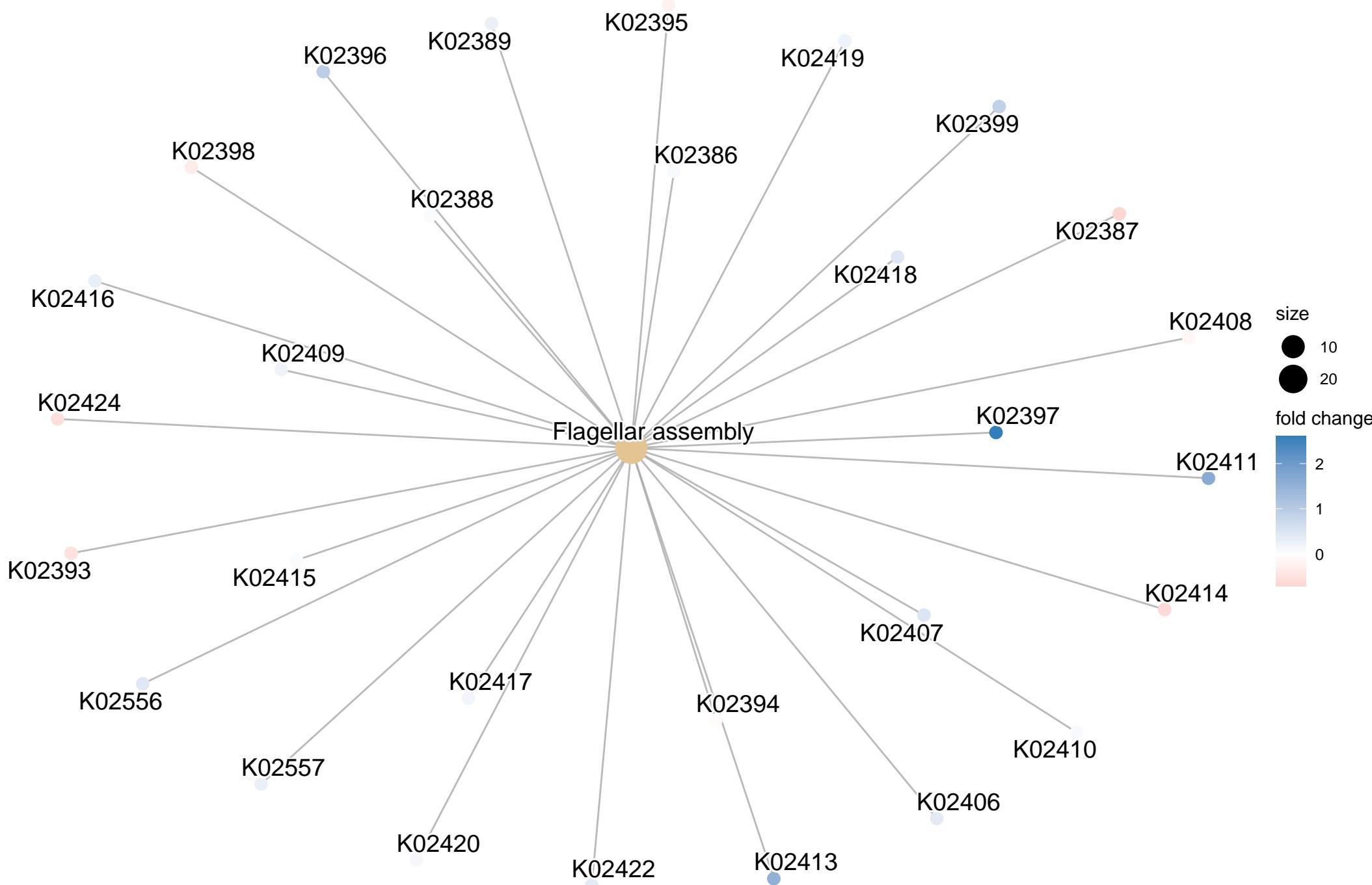




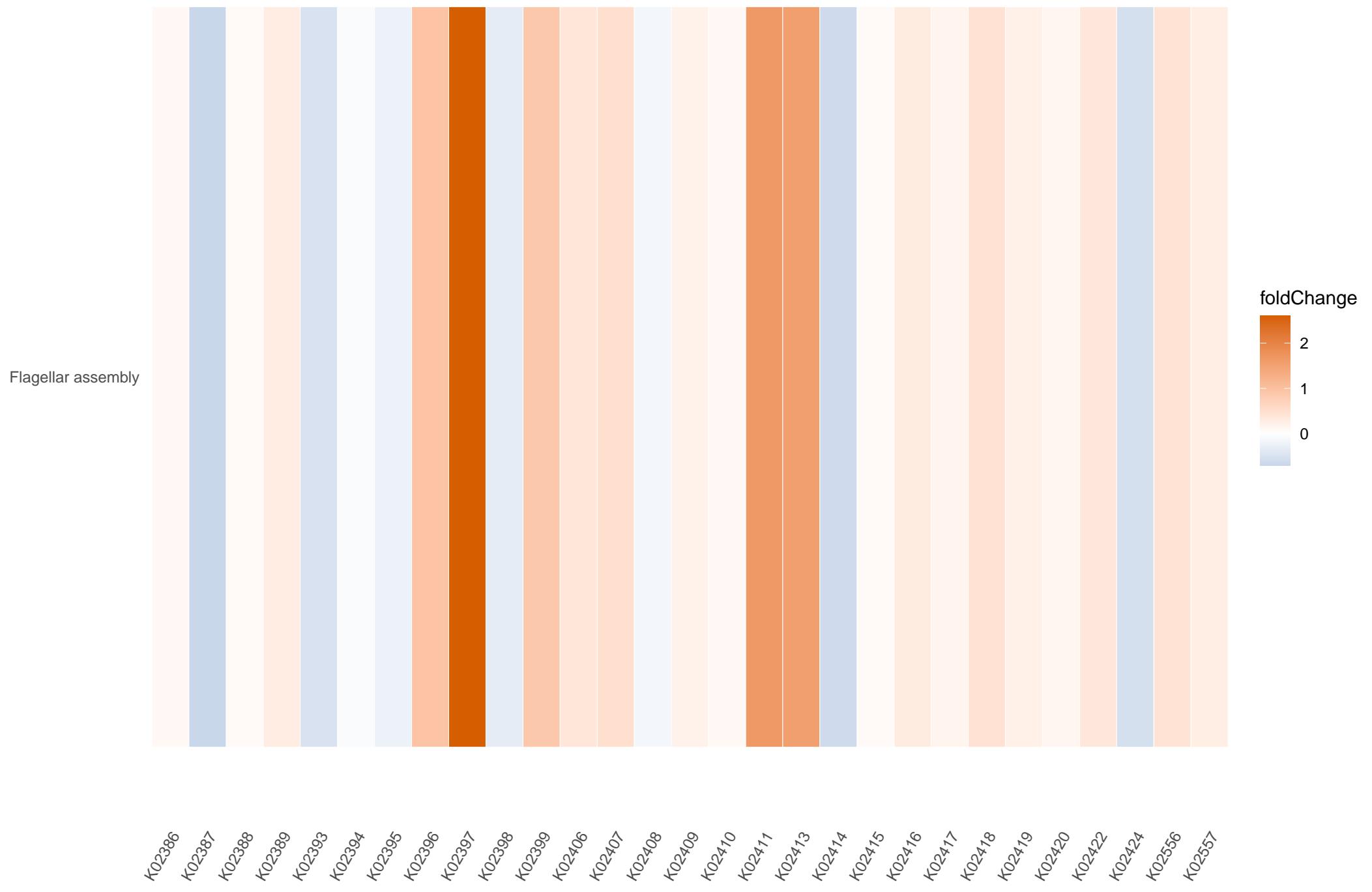
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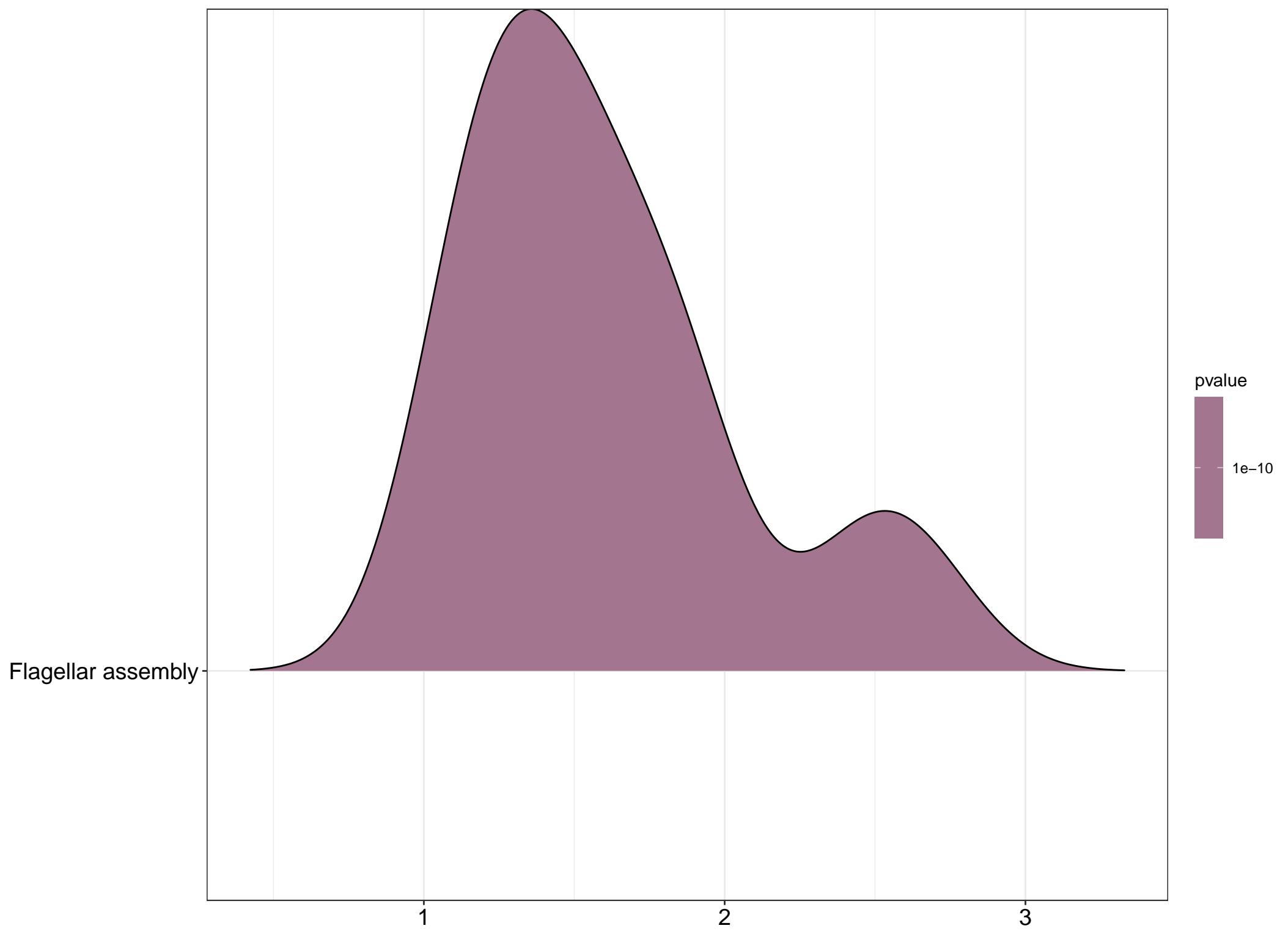


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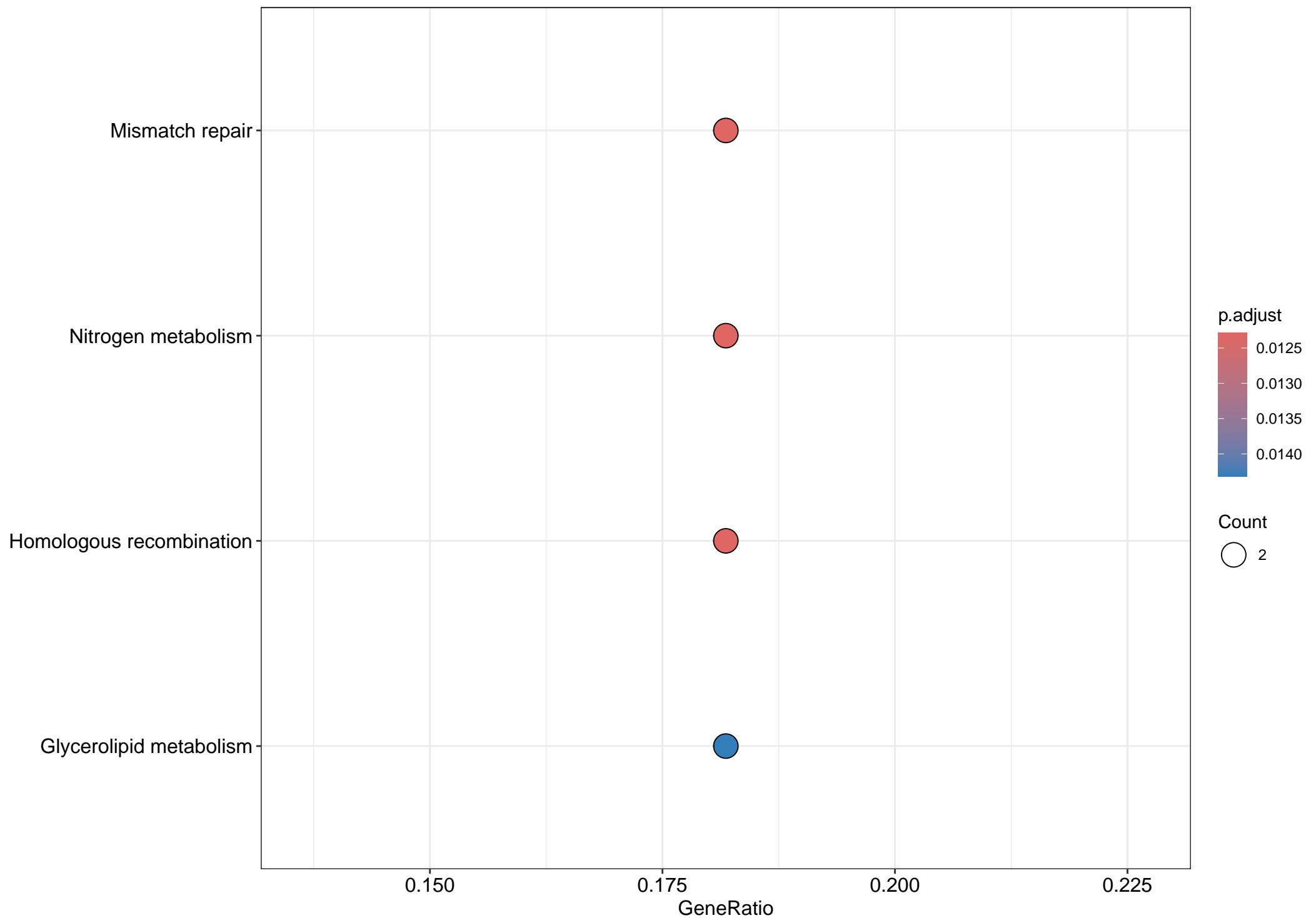


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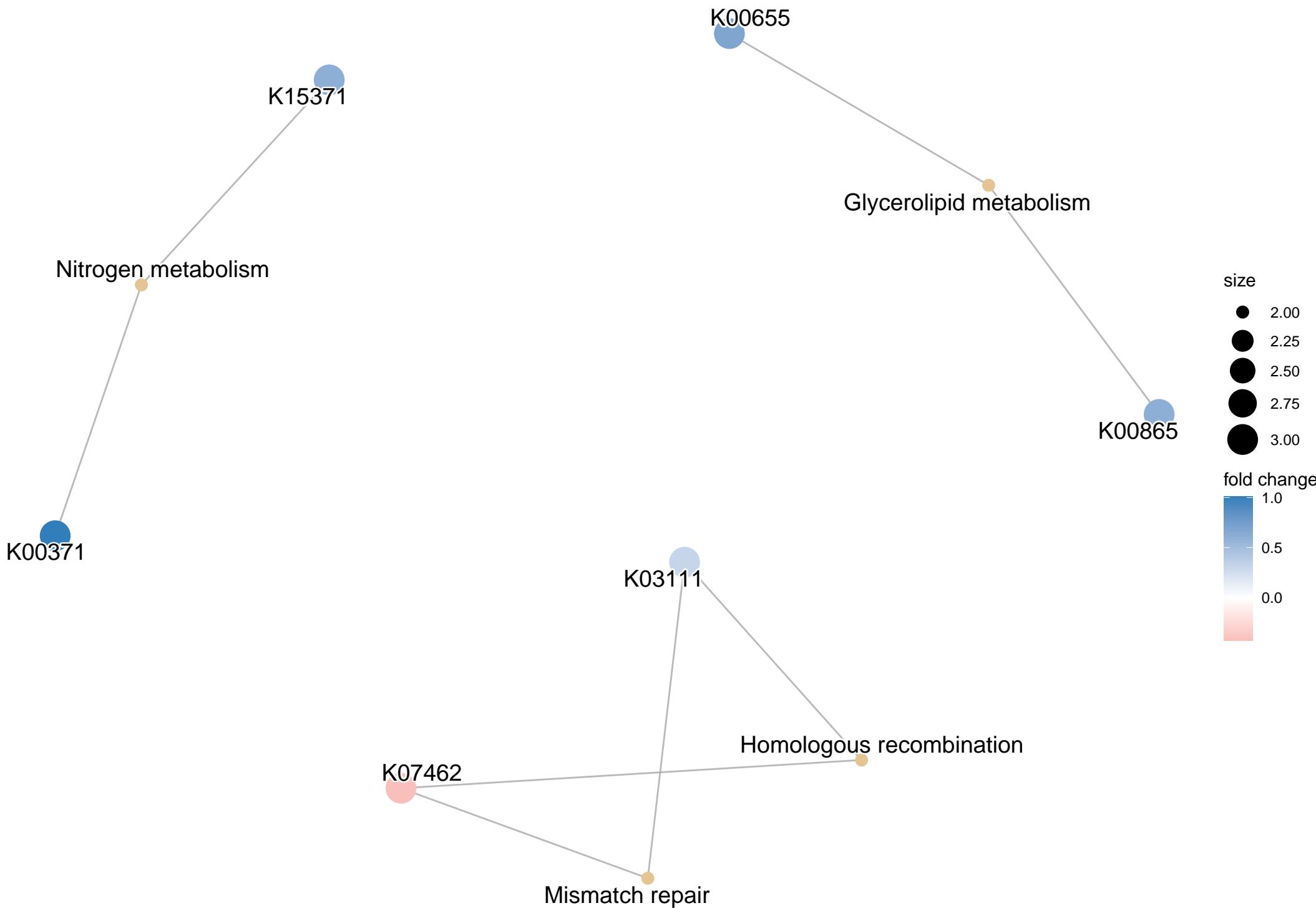




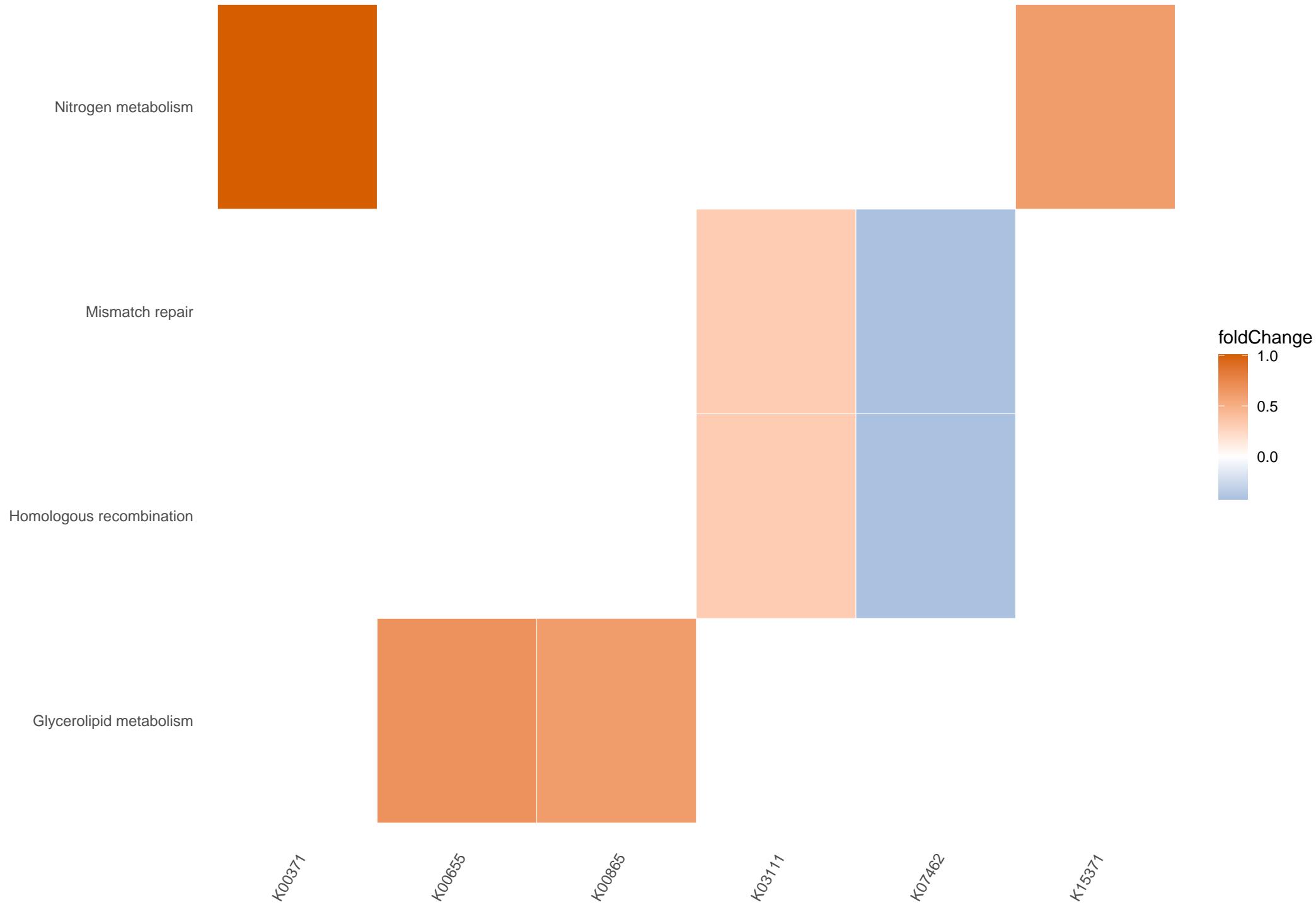
D geom_path

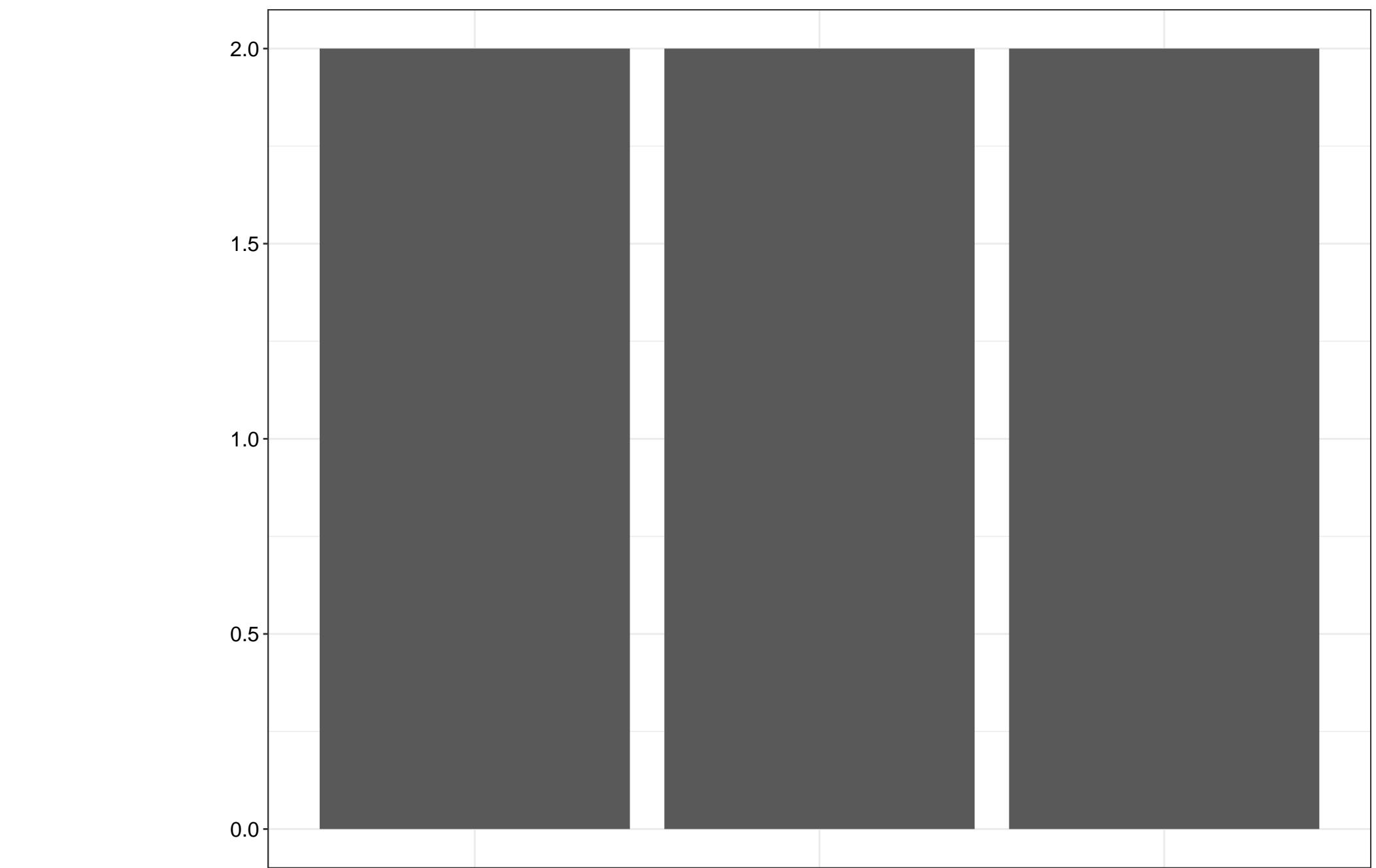


D geom_path

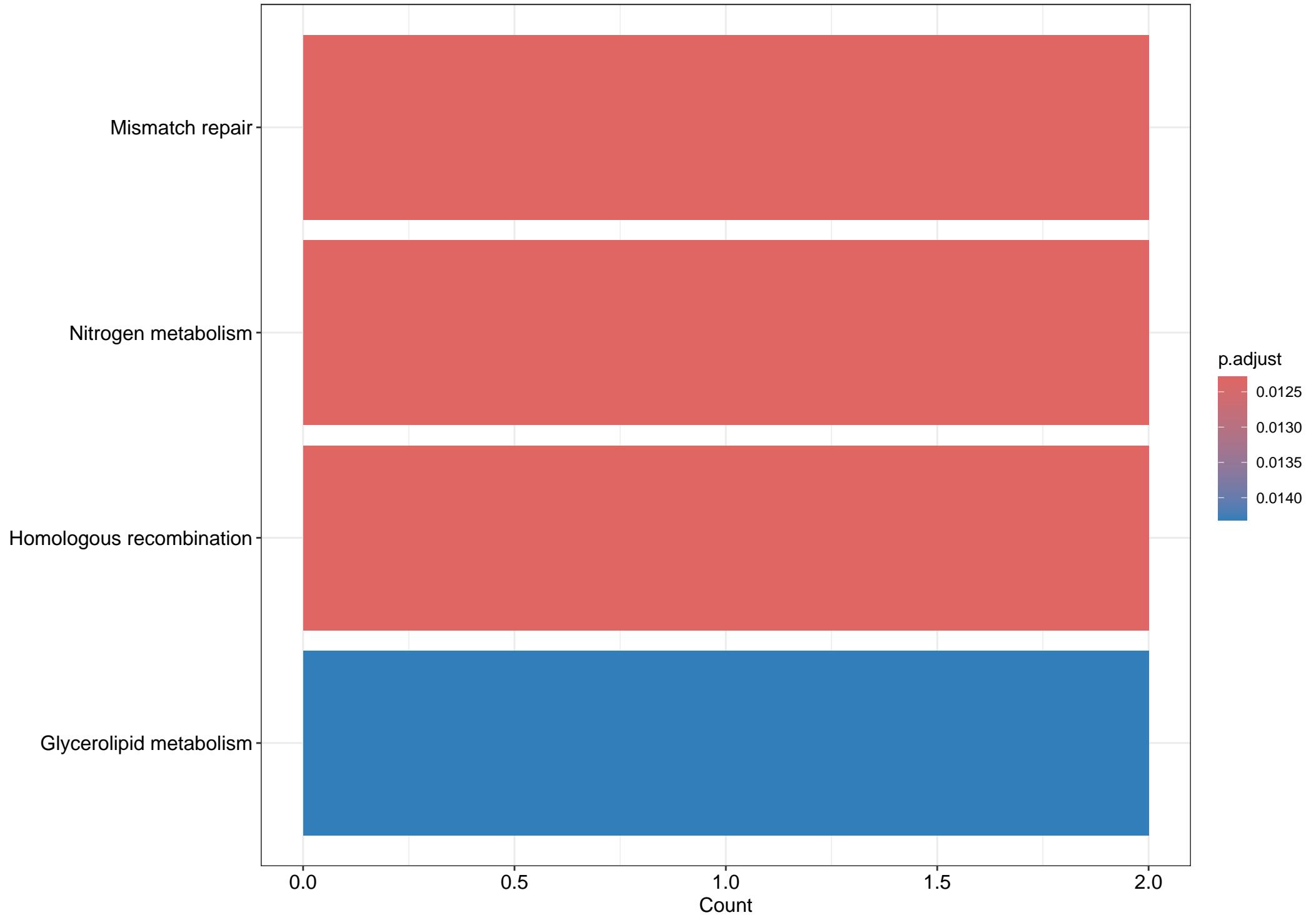


D geom_path

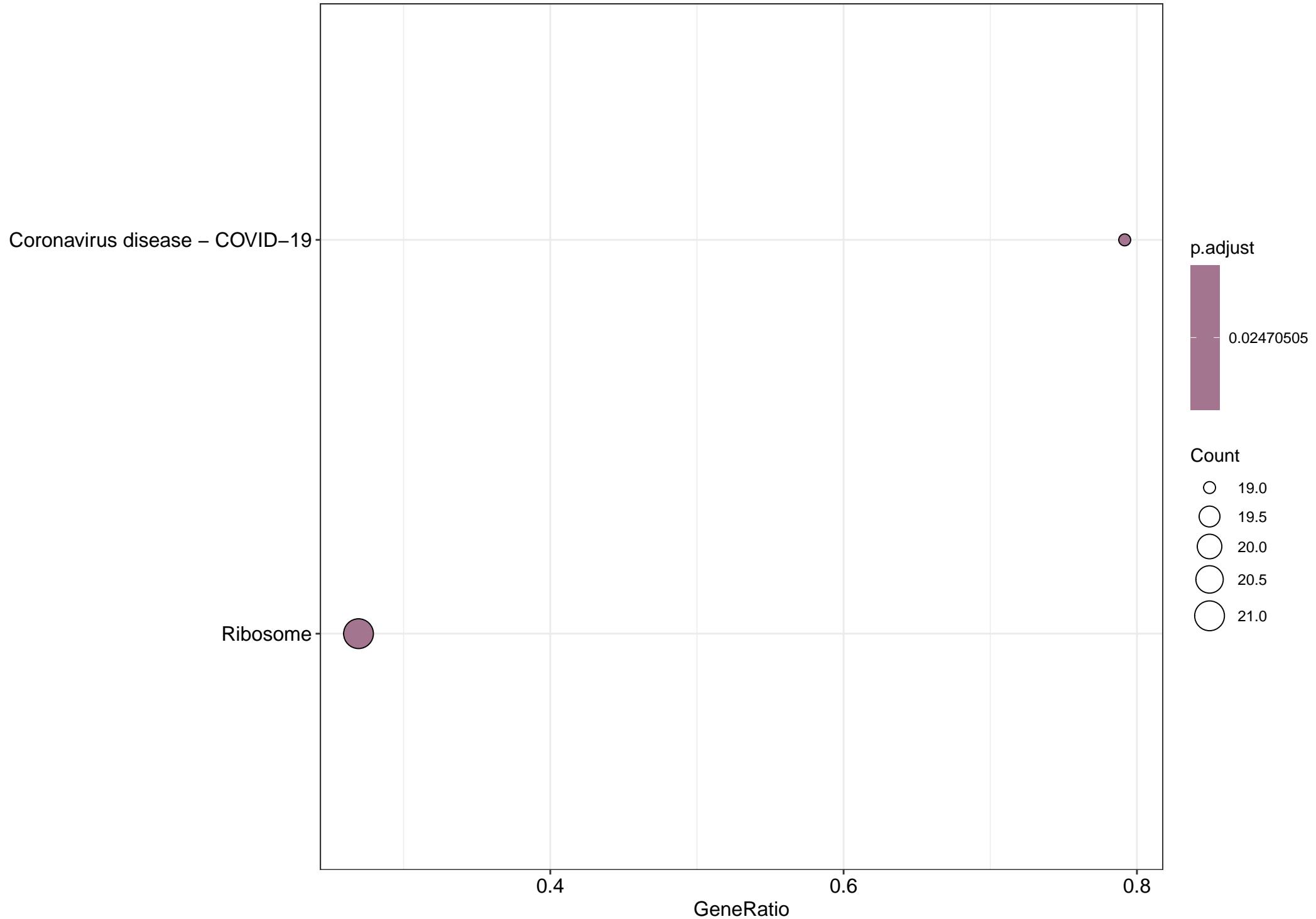




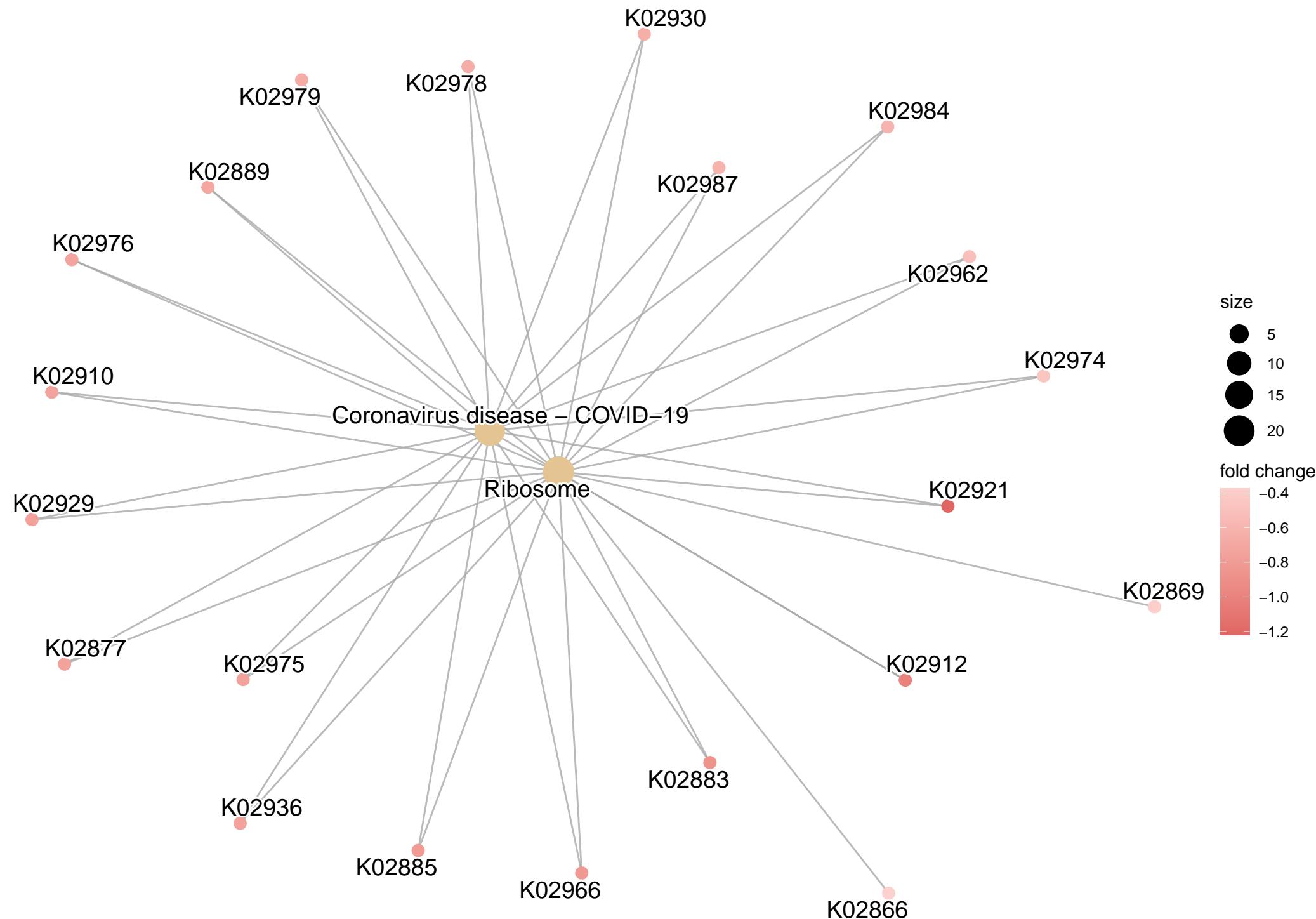
Nitrogen metabolism
Glycerolipid metabolism
Homologous recombination
Mismatch repair



D gsea_path



D gsea_path



D gsea_path

