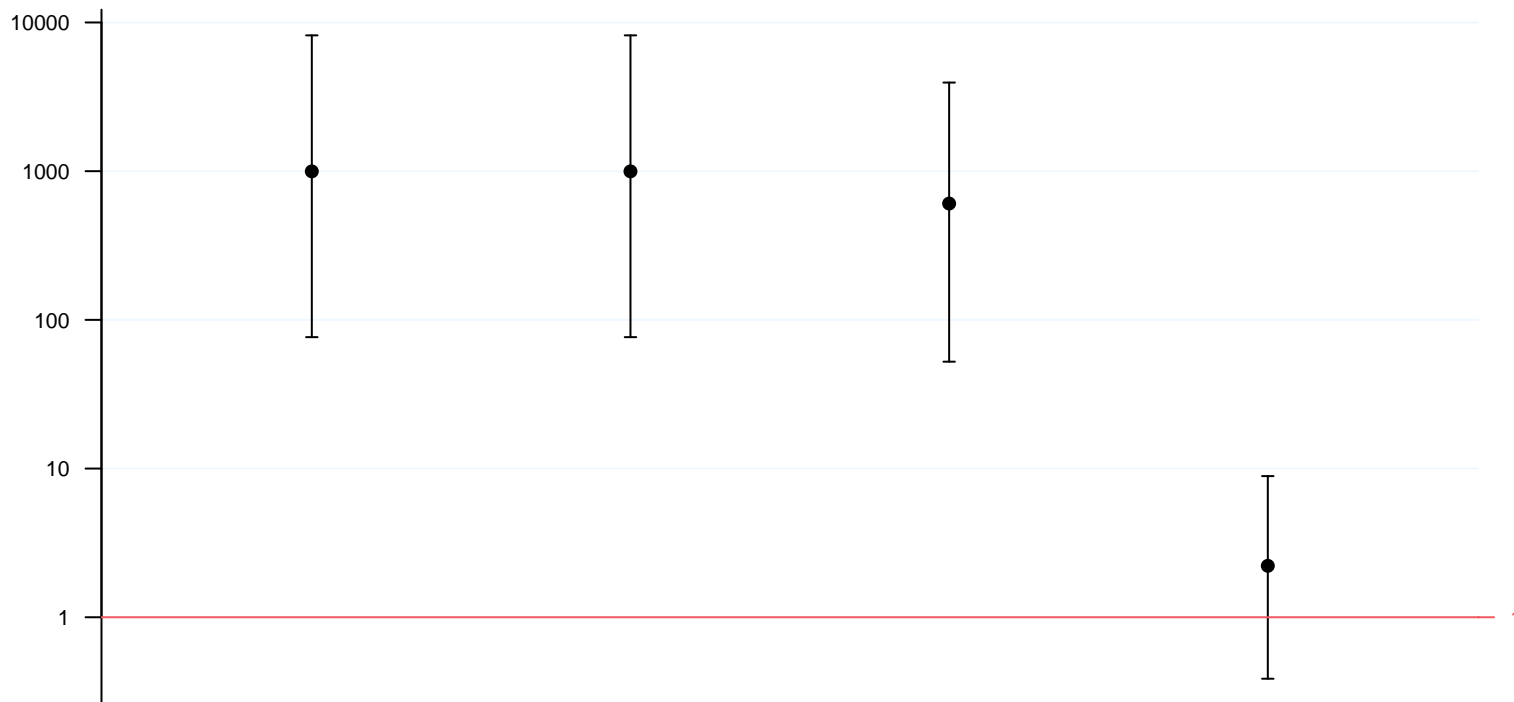
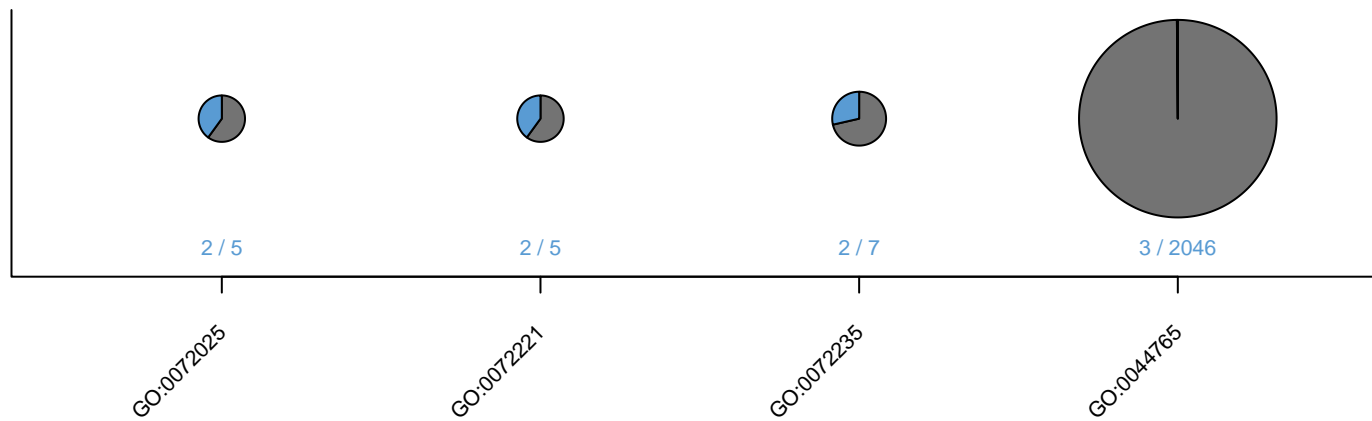


odds ratio

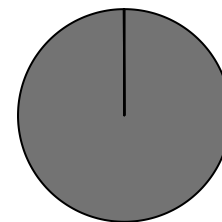


annotated genes



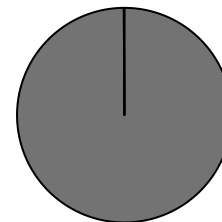
annotated genes
root nodes

biological_process



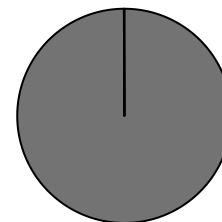
12 / 15655

cellular_component



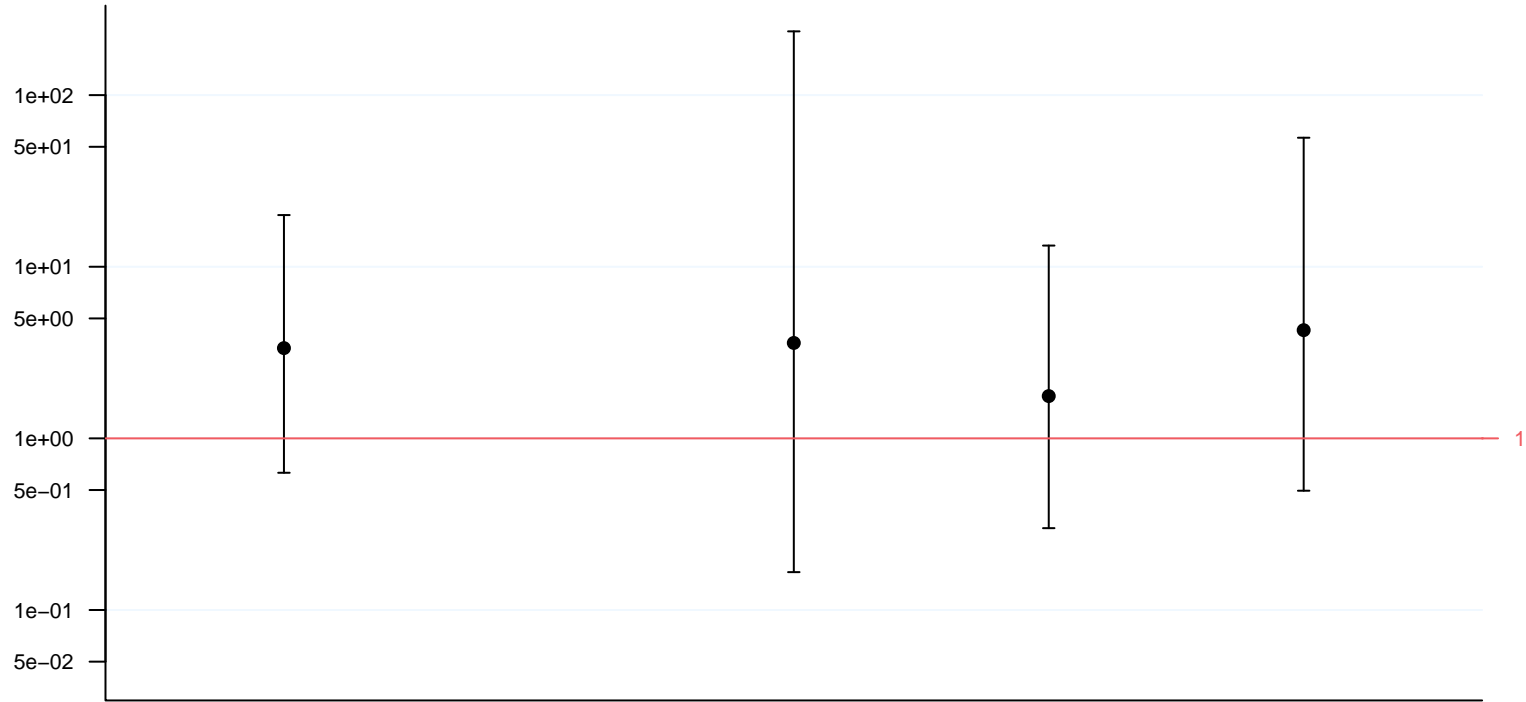
13 / 17106

molecular_function

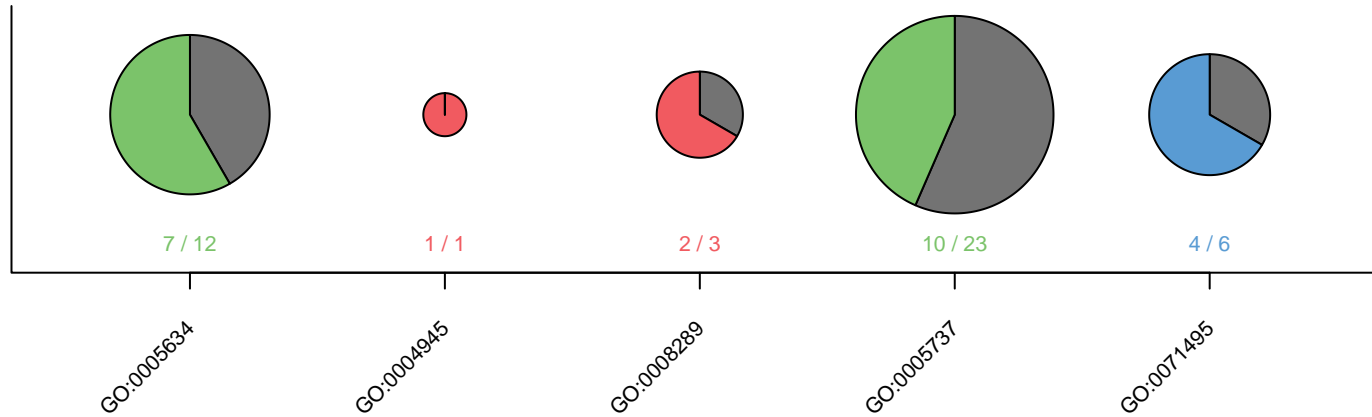


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odds ratio

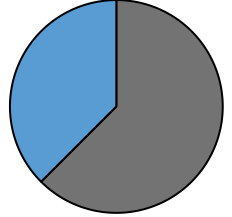


annotated genes



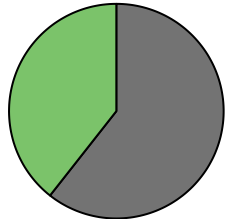
annotated genes root nodes

biological_process



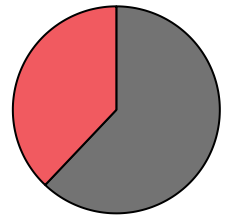
12 / 32

cellular_component



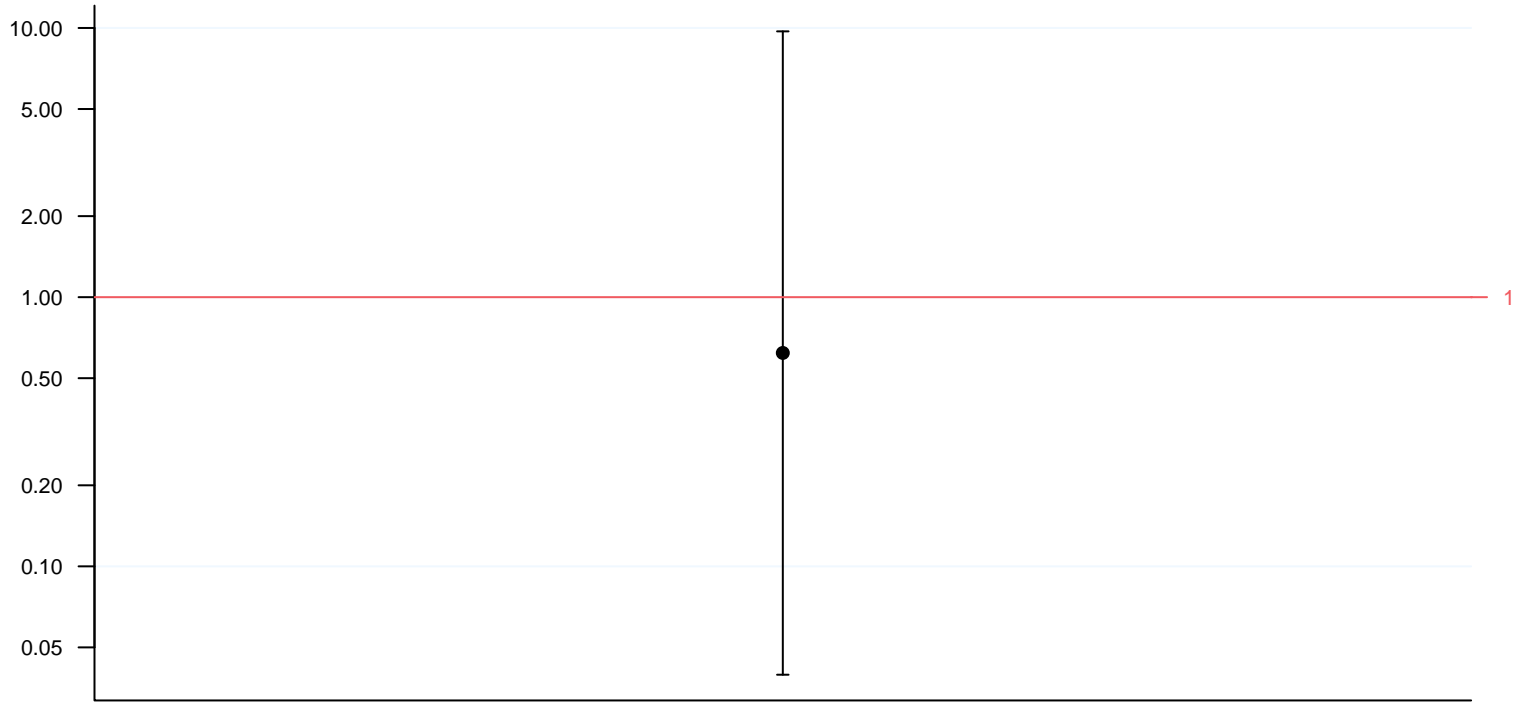
13 / 33

molecular_function



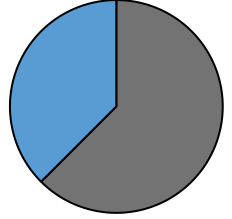
11 / 29

odds ratio



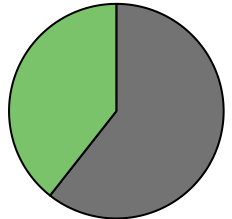
annotated genes root nodes

biological_process



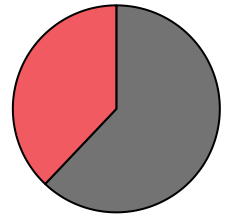
12 / 32

cellular_component



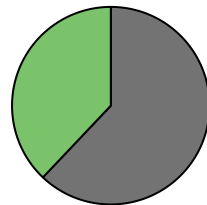
13 / 33

molecular_function



11 / 29

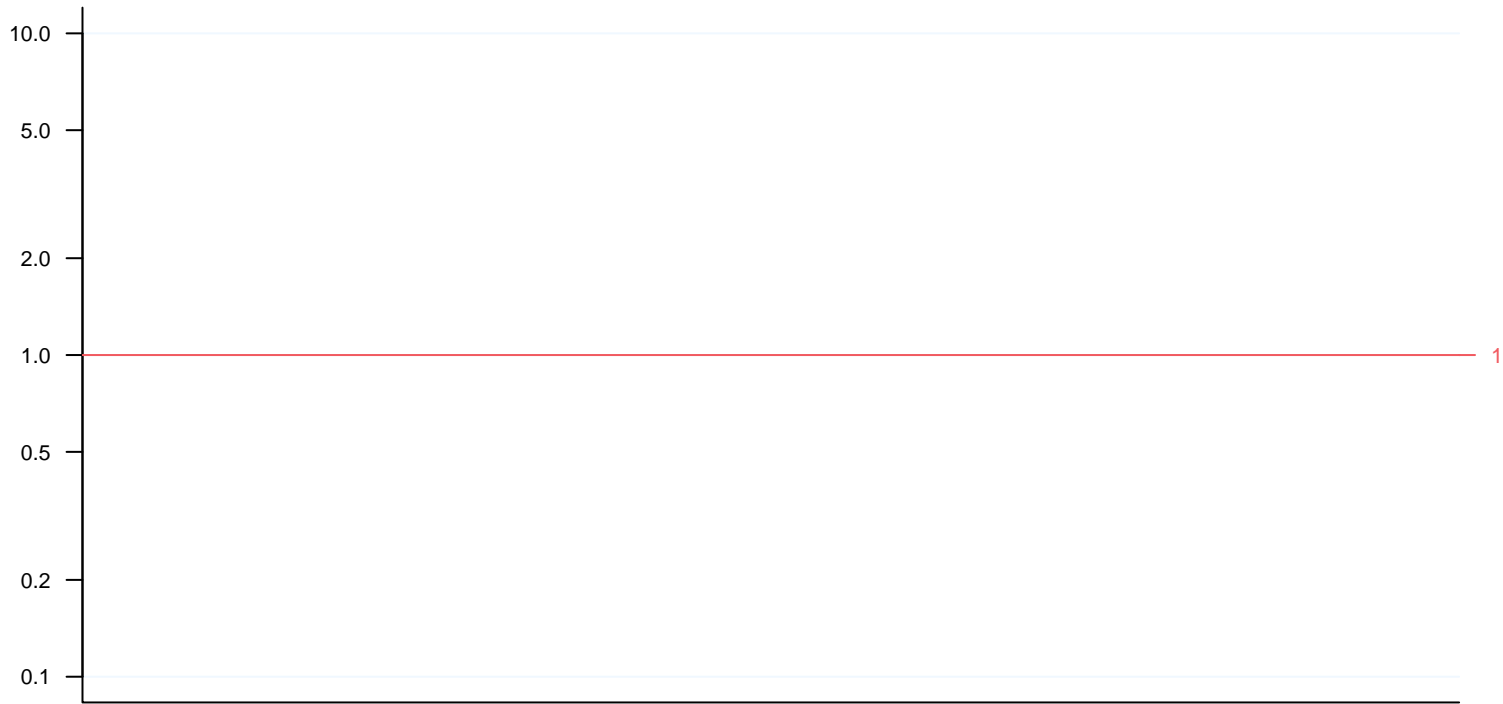
annotated genes



11 / 29

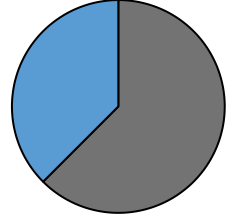
GO:0005623

odds ratio



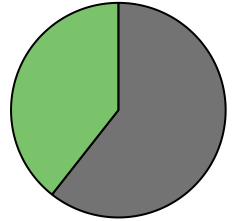
annotated genes root nodes

biological_process



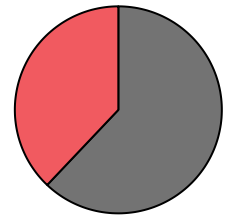
12 / 32

cellular_component



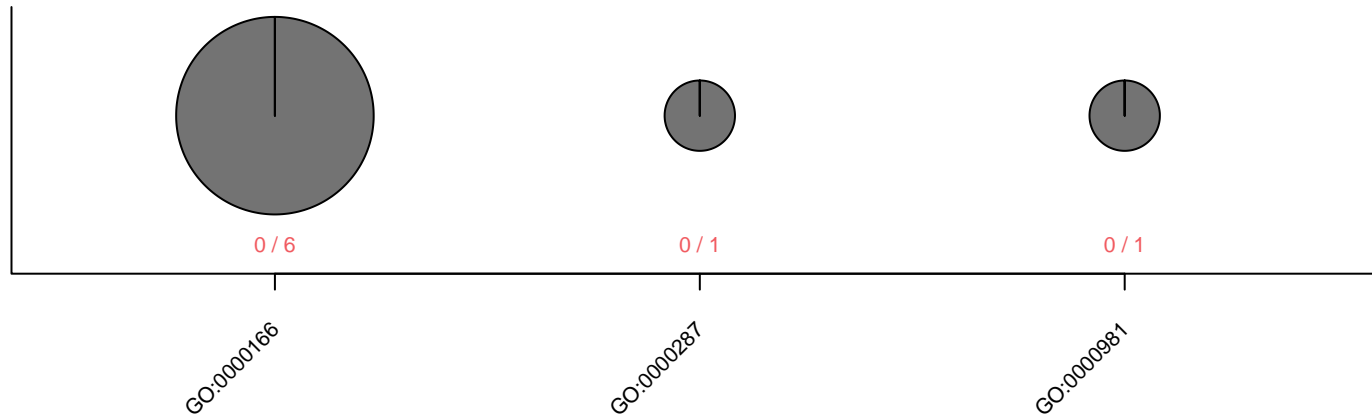
13 / 33

molecular_function

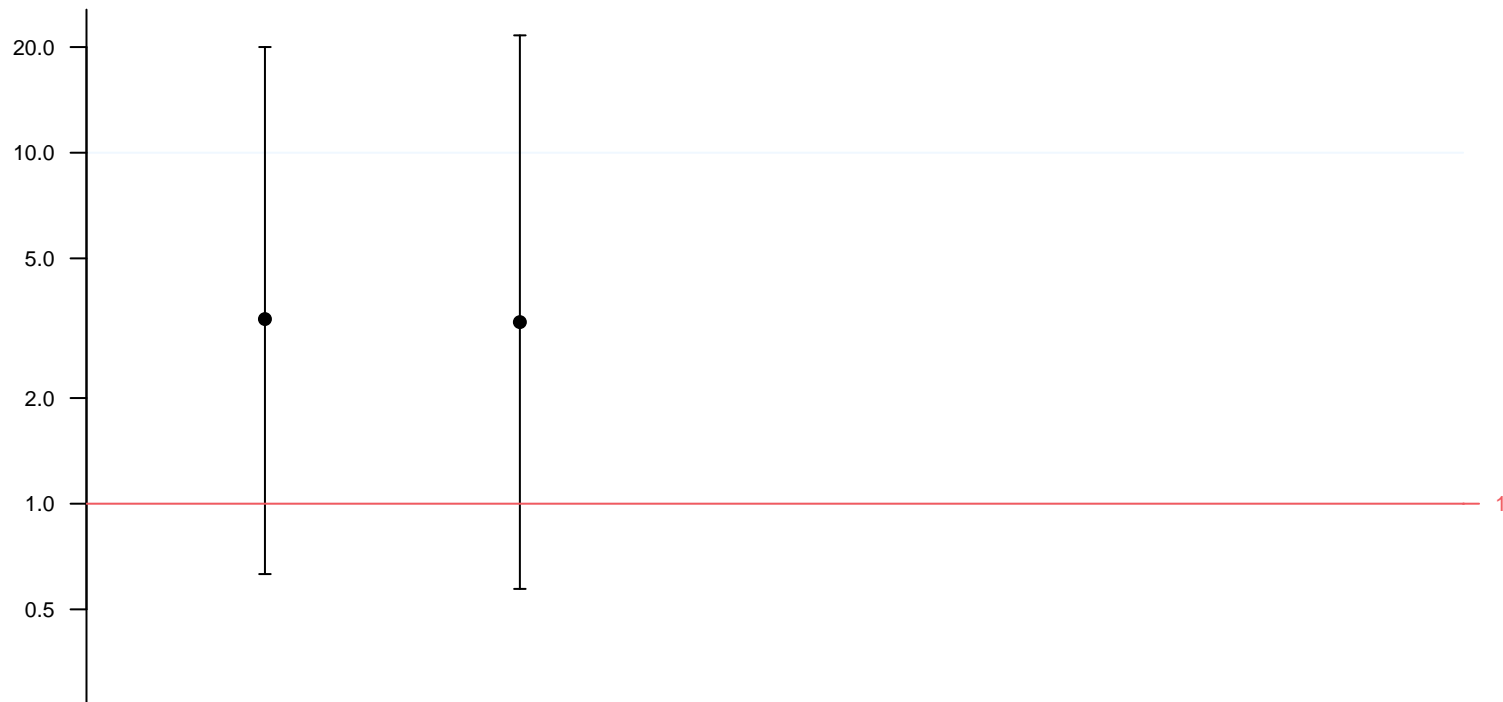


11 / 29

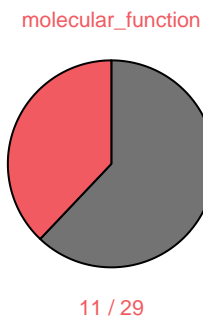
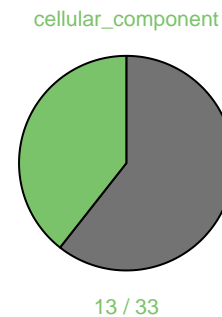
annotated genes



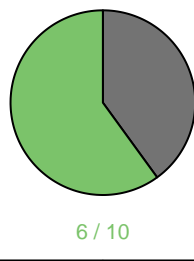
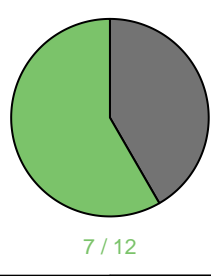
odds ratio



annotated genes root nodes



annotated genes



GO:0005634

GO:0005829

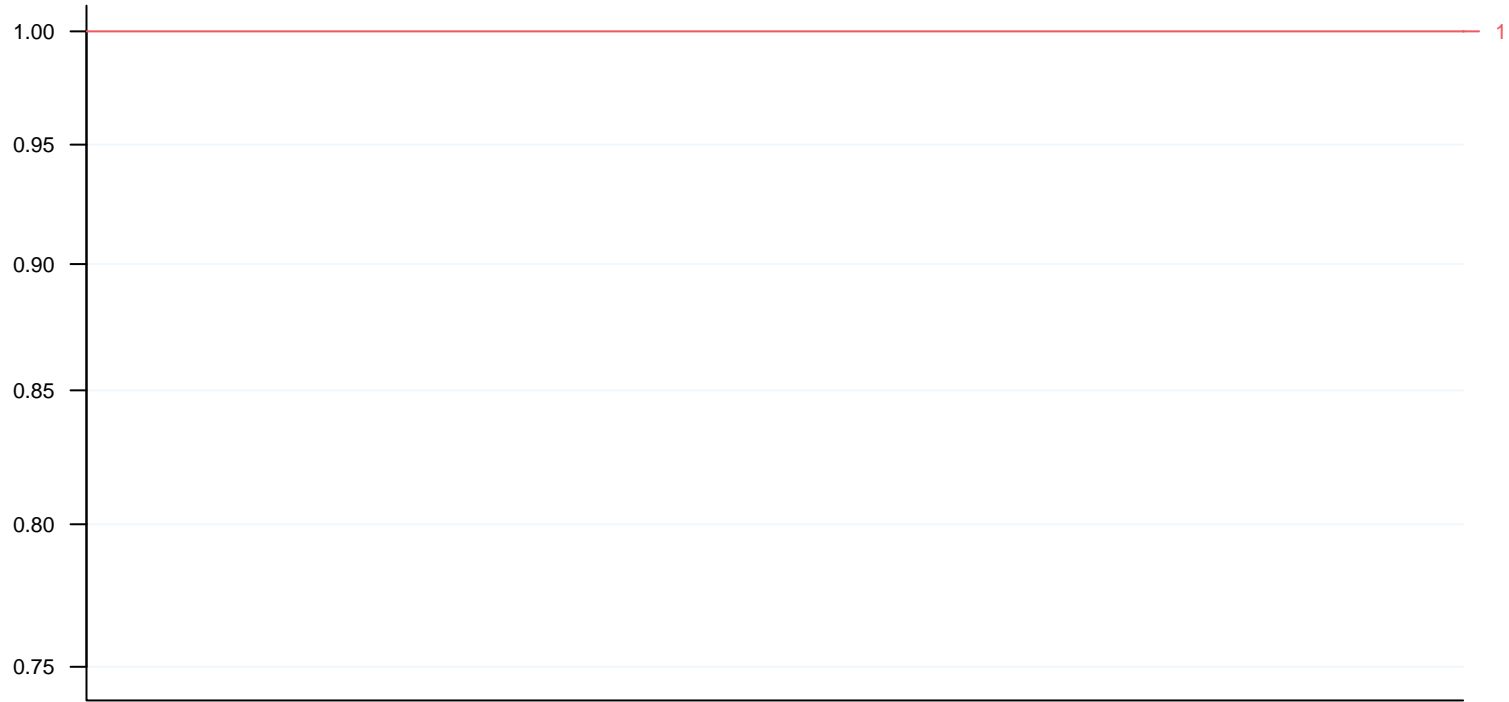
GO:0005216

GO:0015075

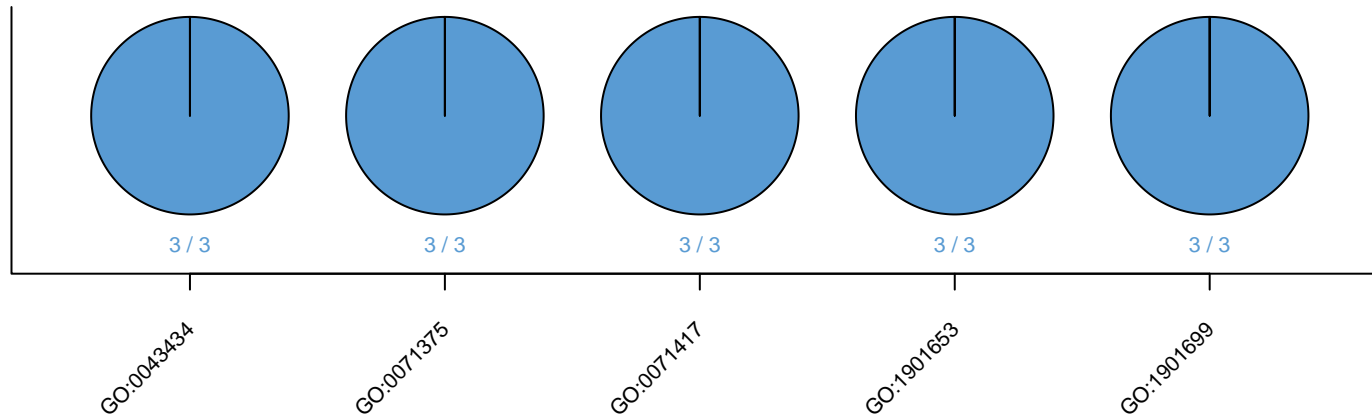
GO:0015267

odds ratio

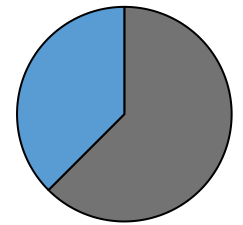
annotated genes
root nodes



annotated genes

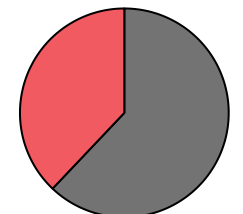


biological_process



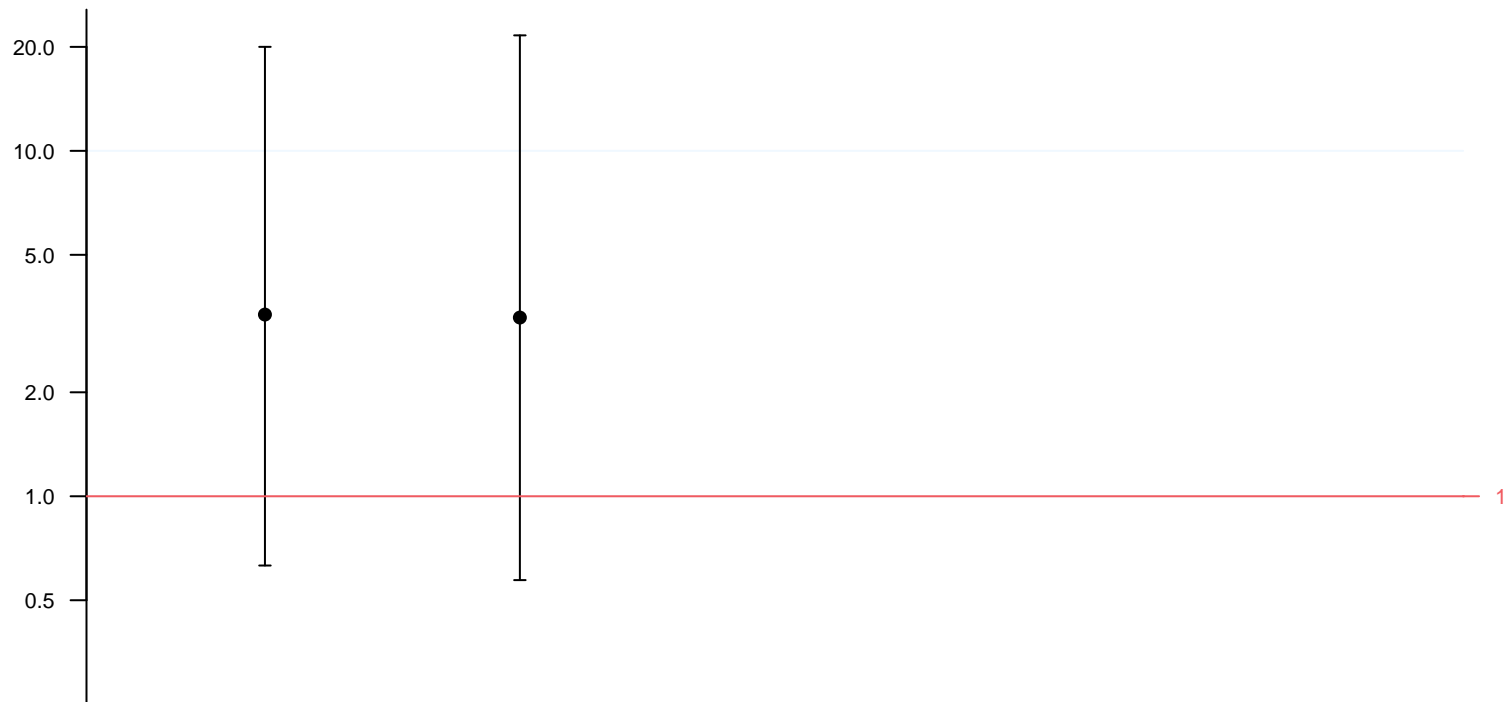
12 / 32

molecular_function



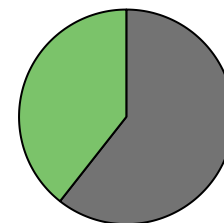
11 / 29

odds ratio



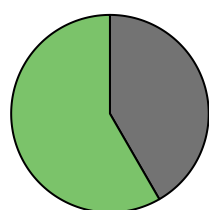
annotated genes
root nodes

cellular_component

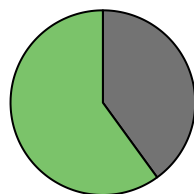


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annotated genes



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2 / 2



2 / 2



2 / 2

GO:0005634

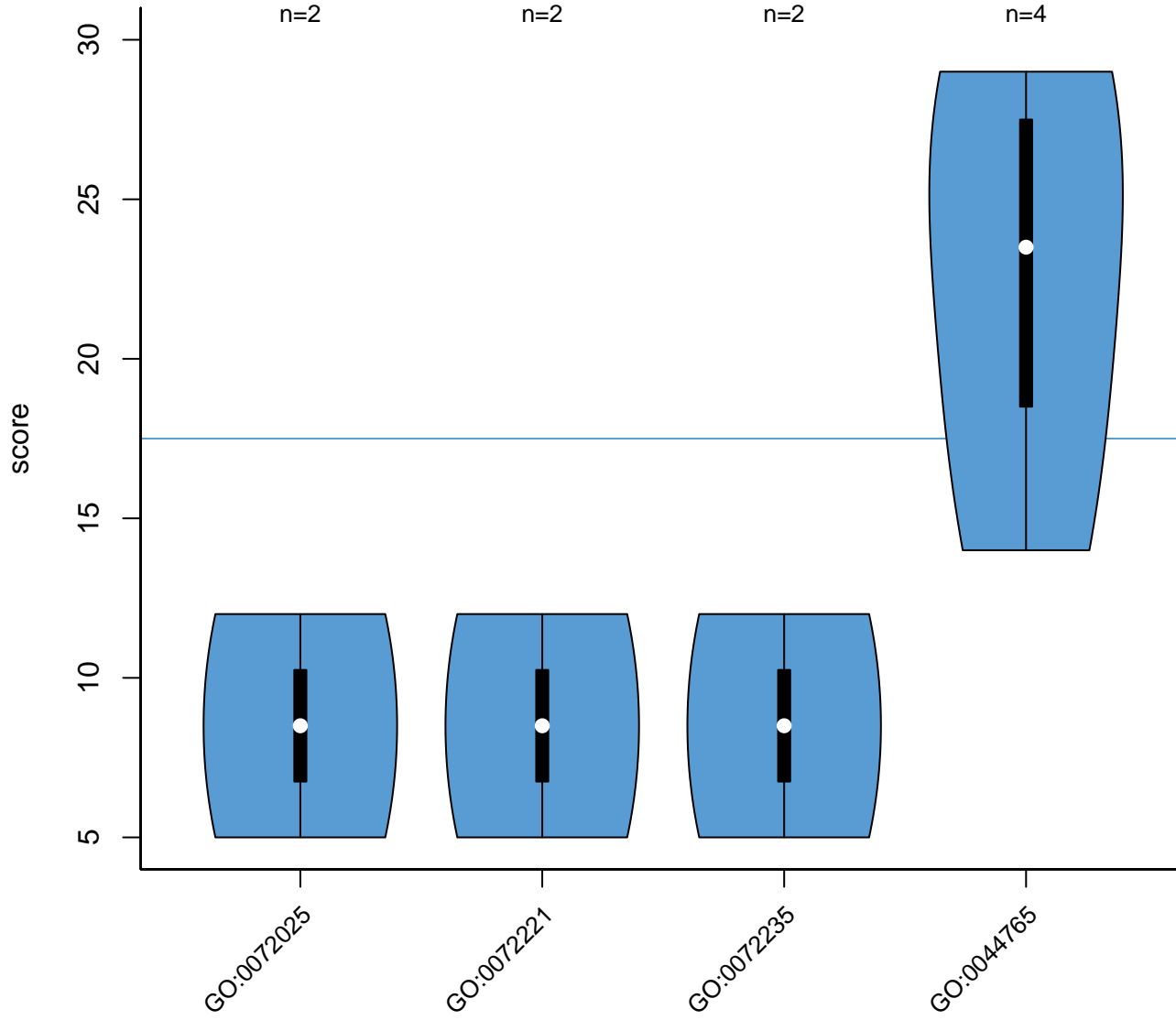
GO:0005829

GO:0034702

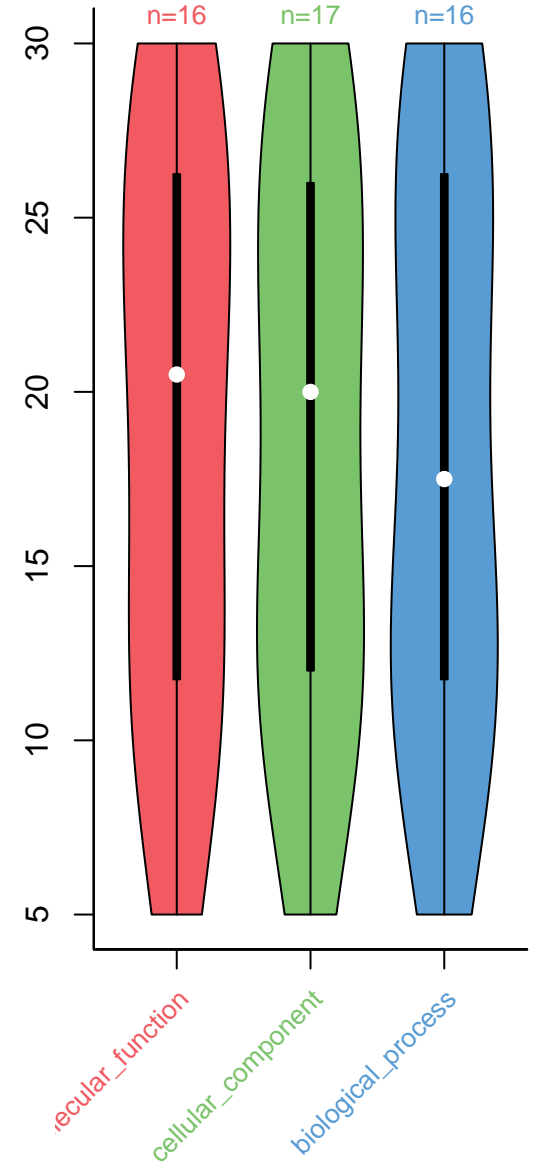
GO:1902495

GO:1990351

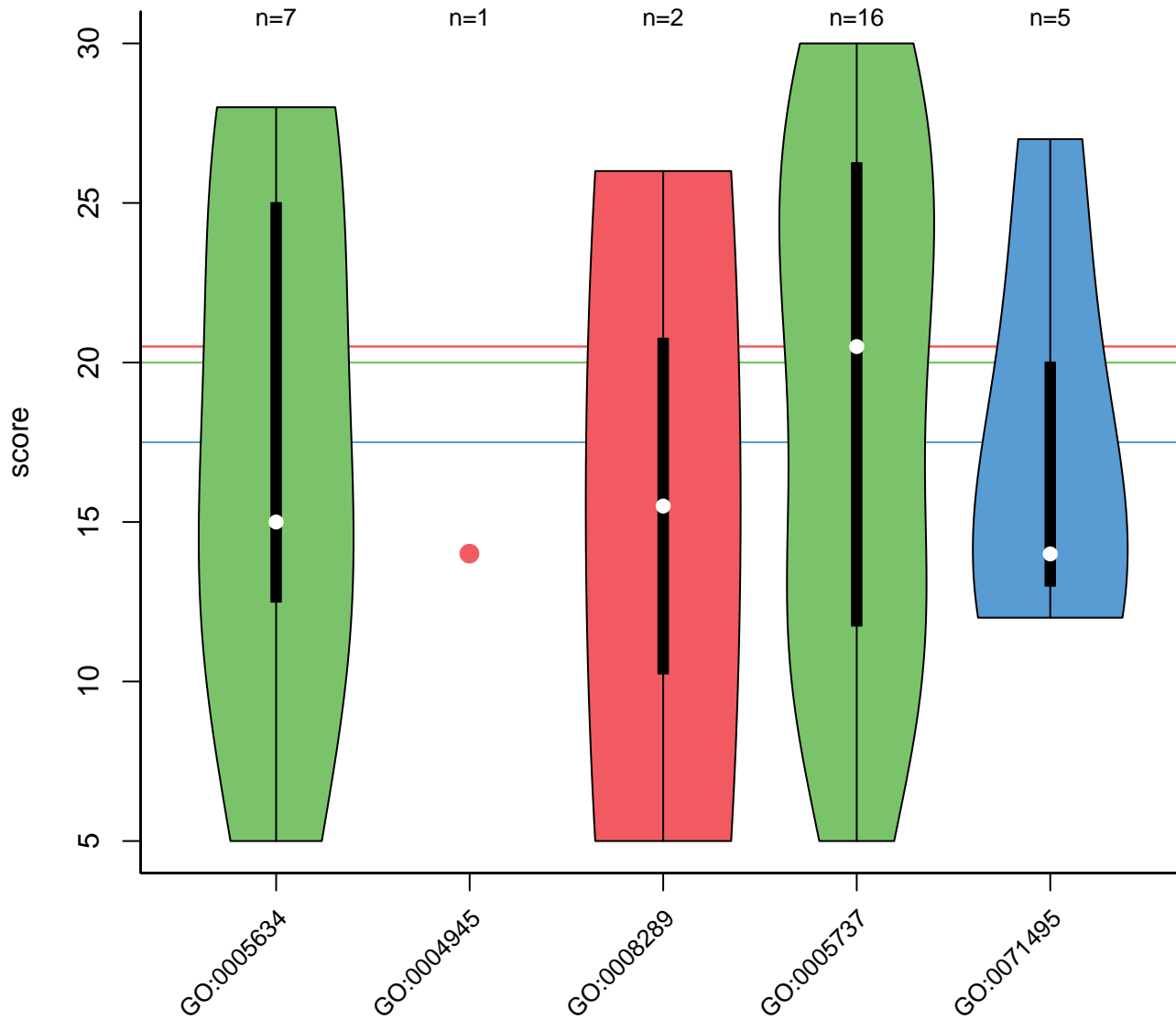
scores of annotated genes



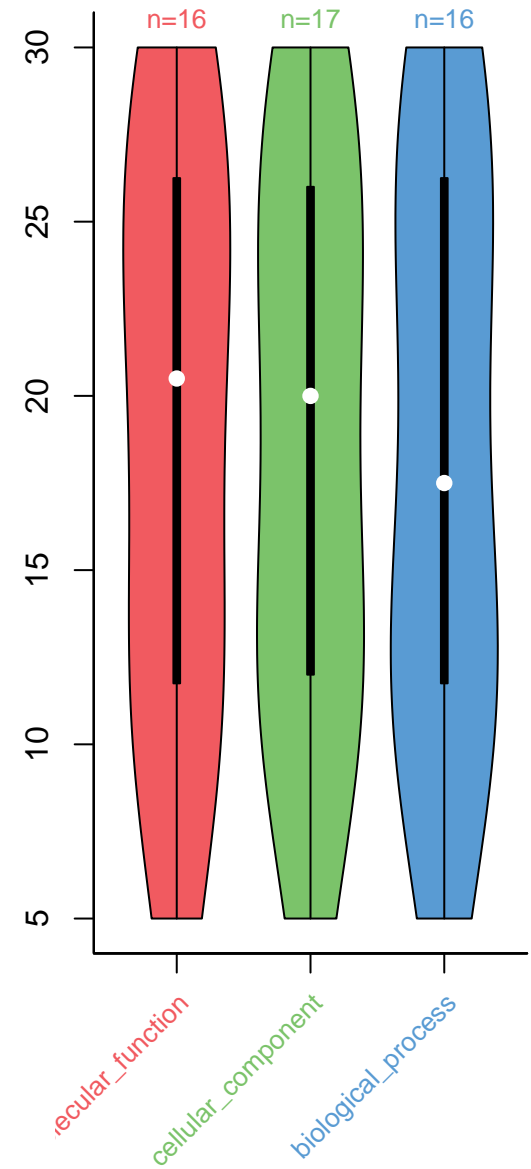
root nodes



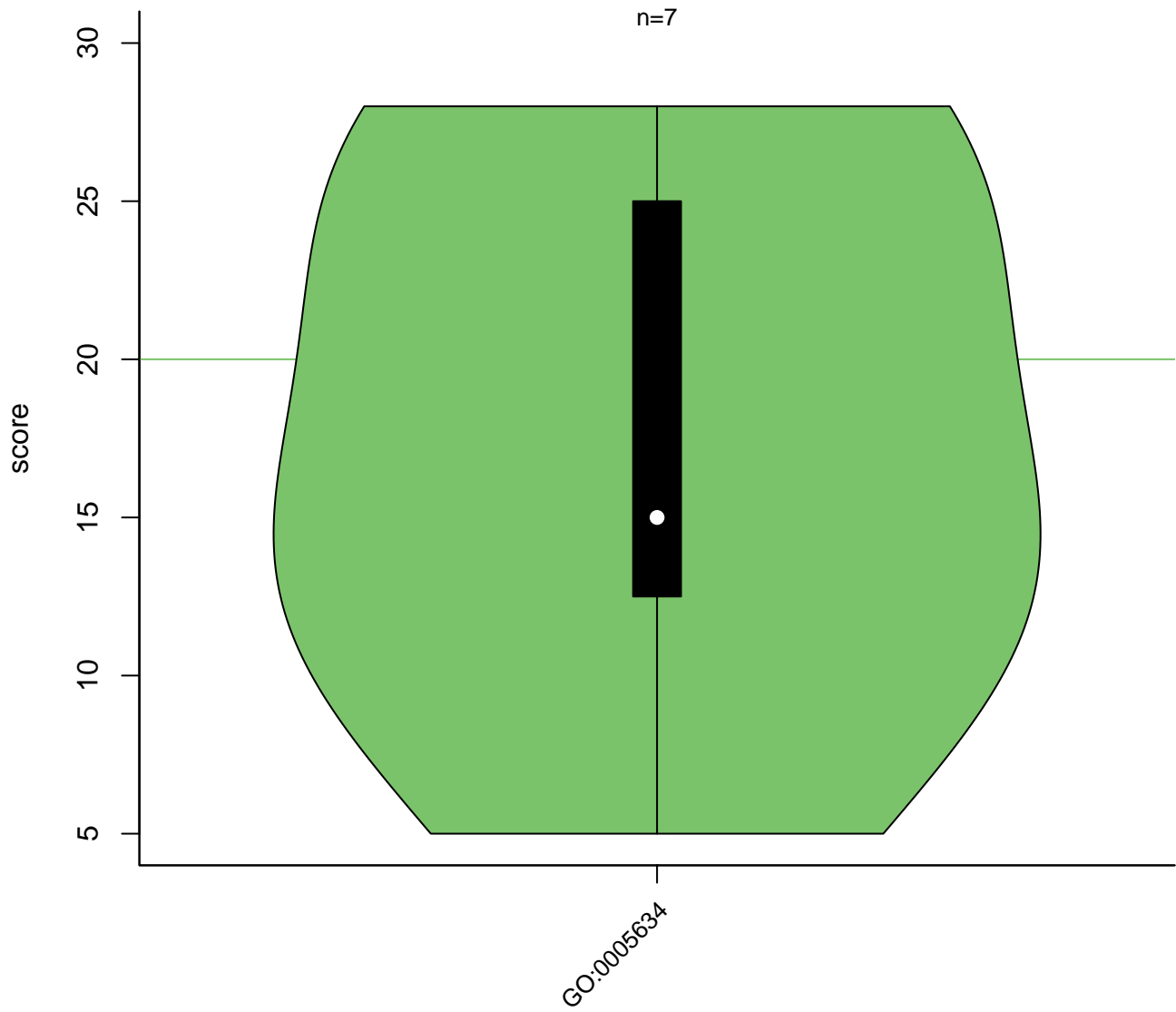
scores of annotated genes



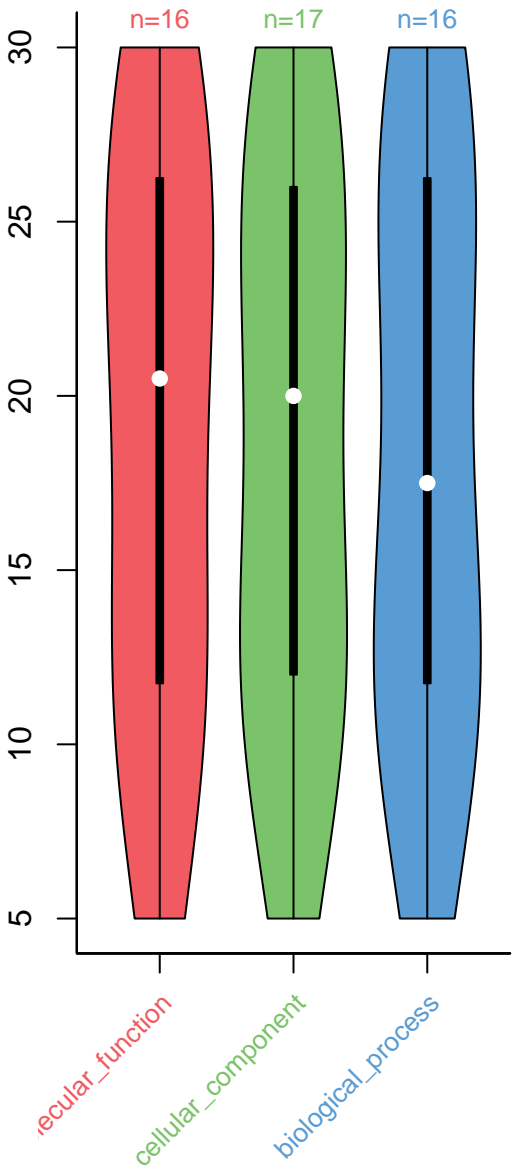
root nodes



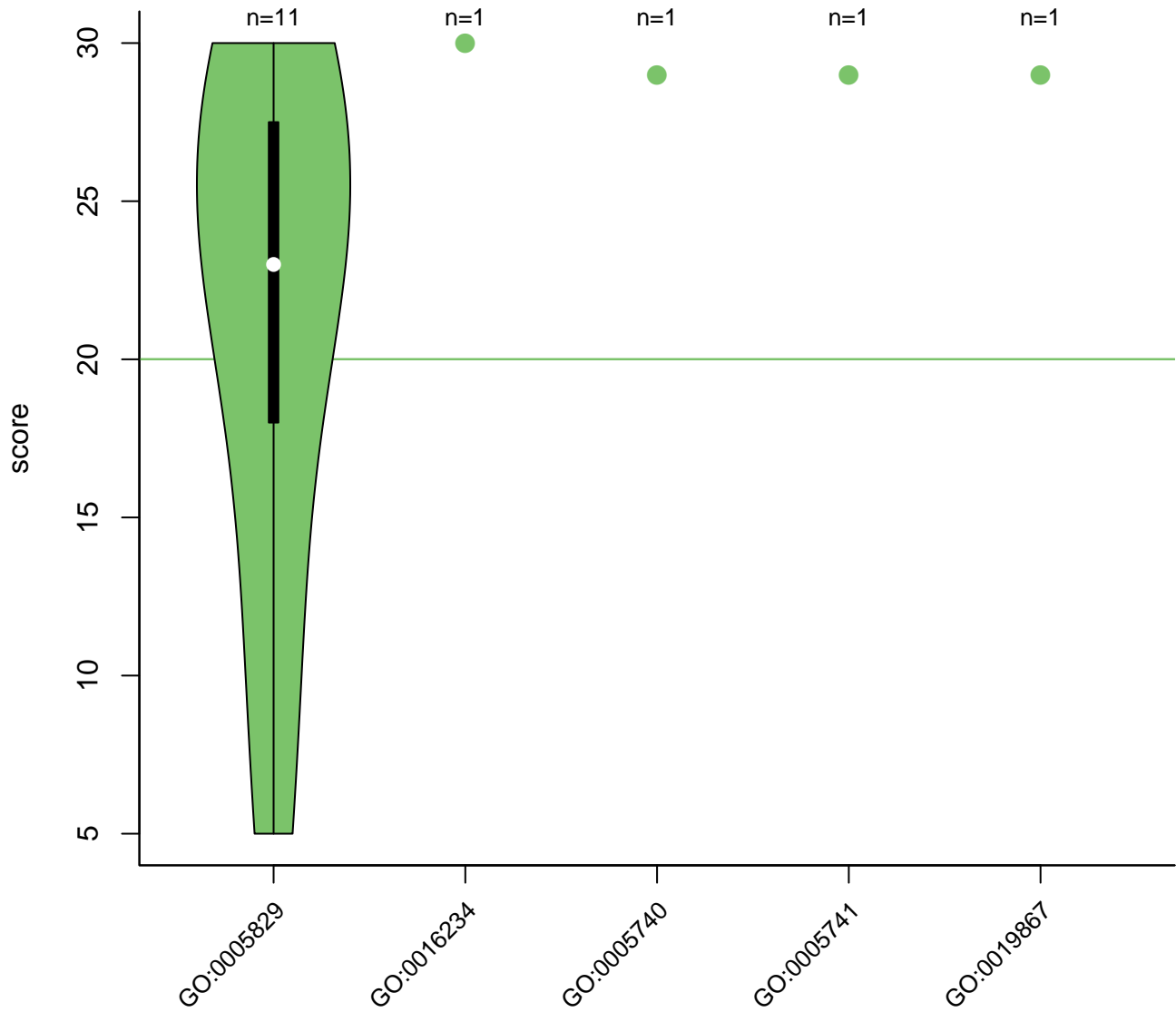
scores of annotated genes



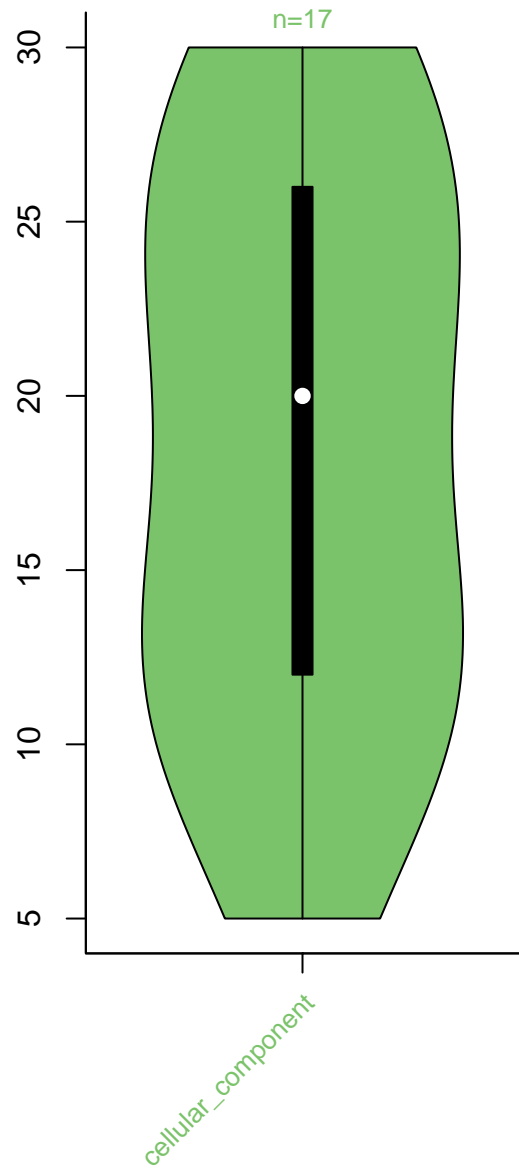
root nodes



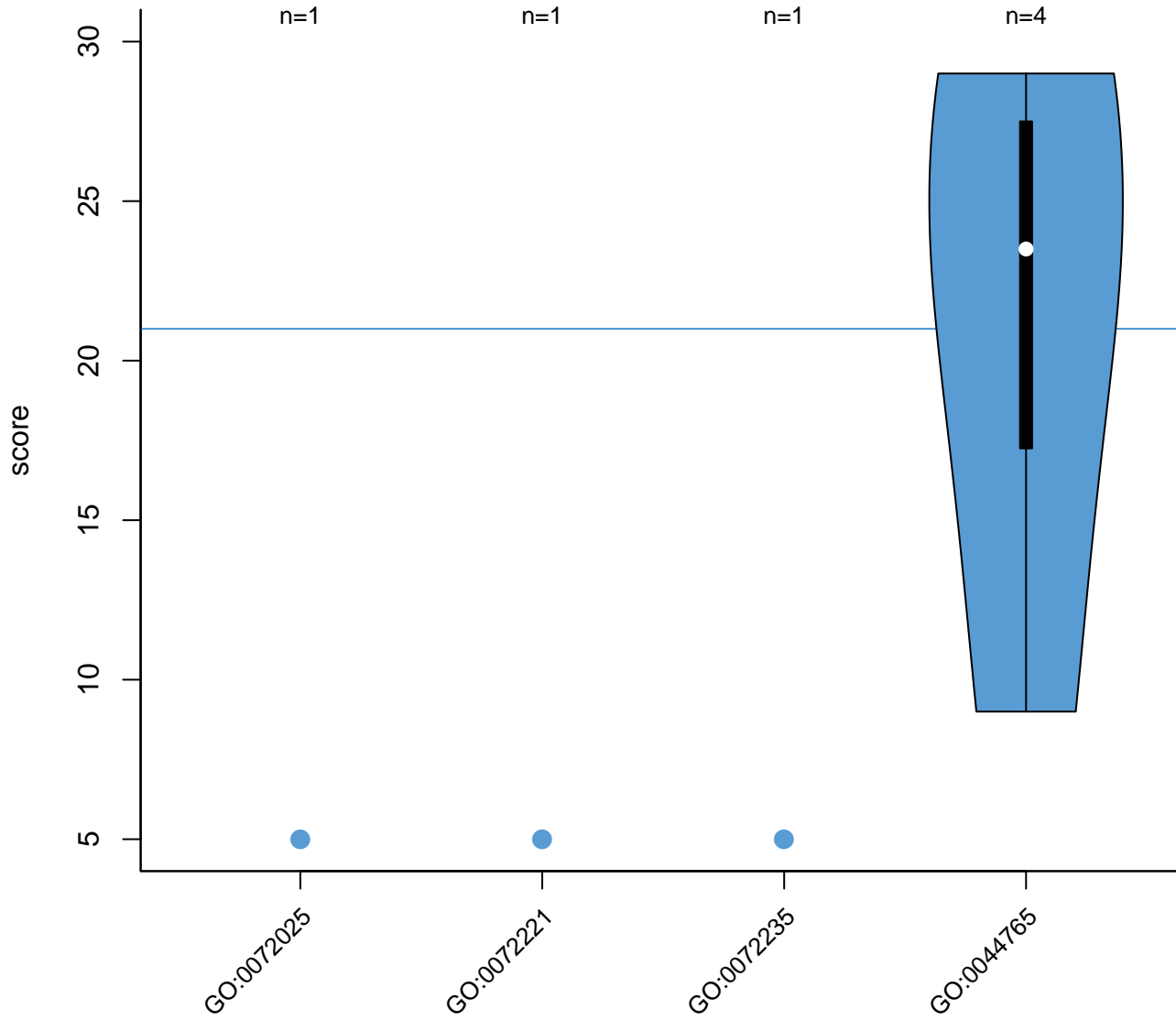
scores of annotated genes



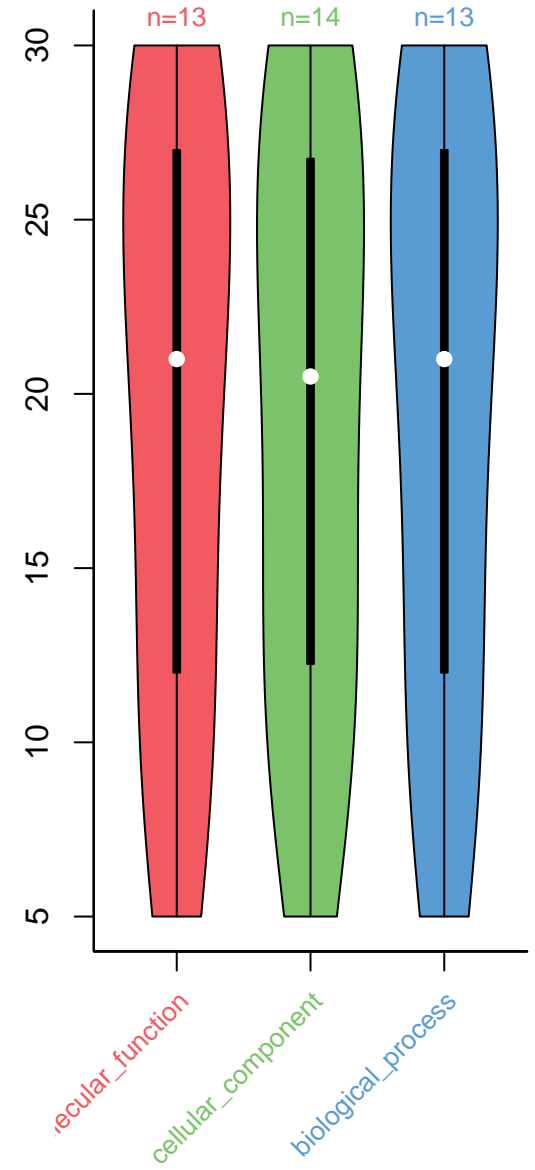
root nodes



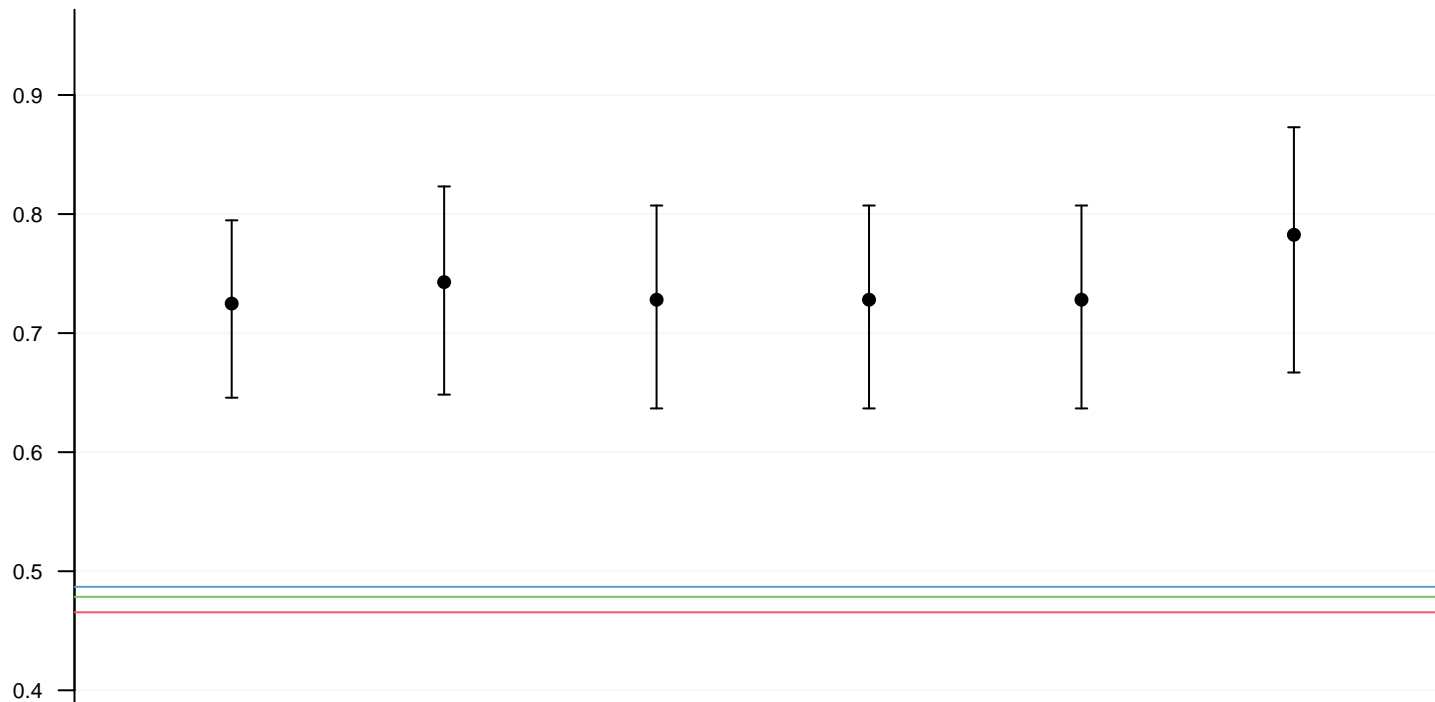
scores of annotated genes



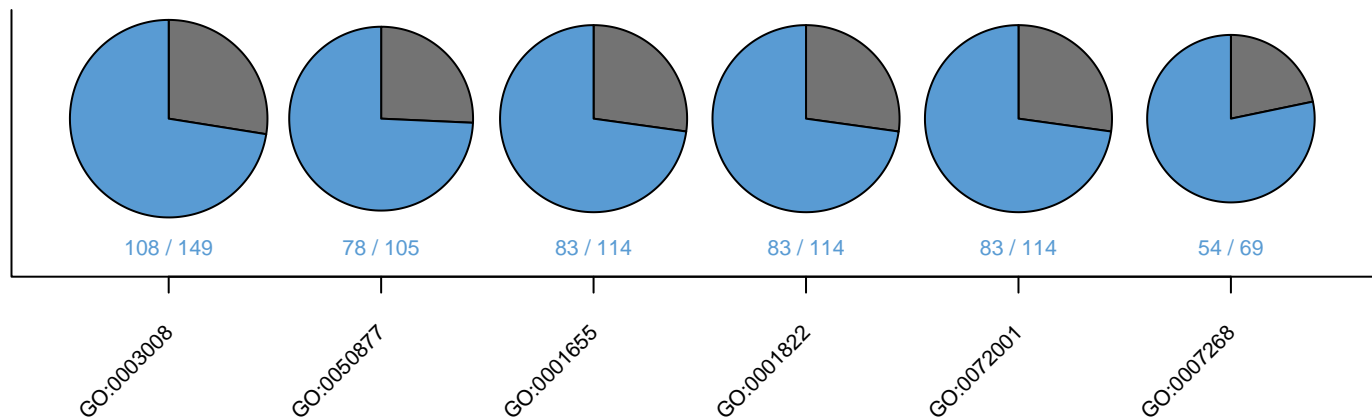
root nodes



p(A) in node

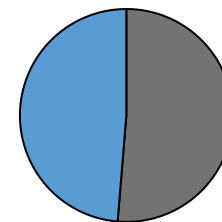


proportion of A



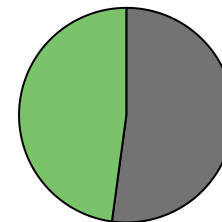
proportion of A
root nodes

biological_process



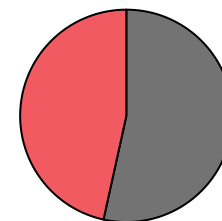
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cellular_component



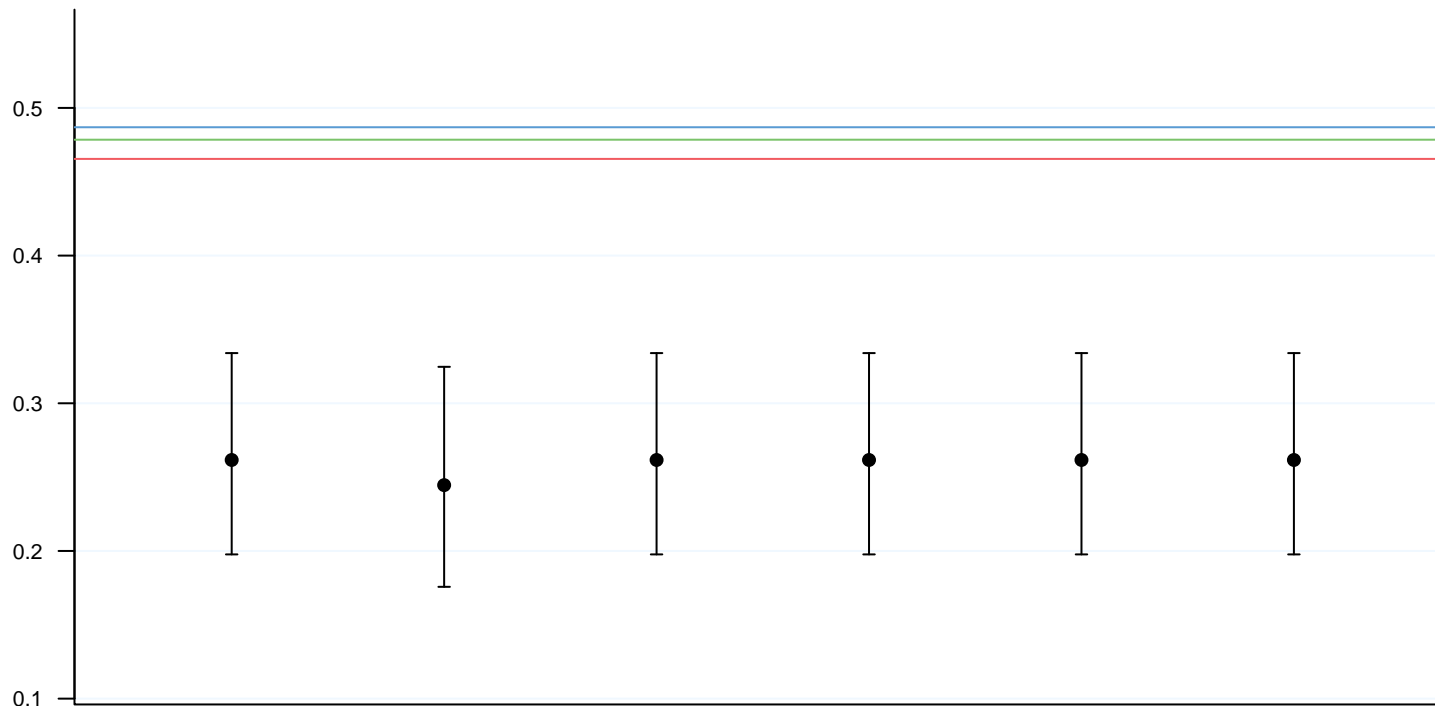
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molecular_function

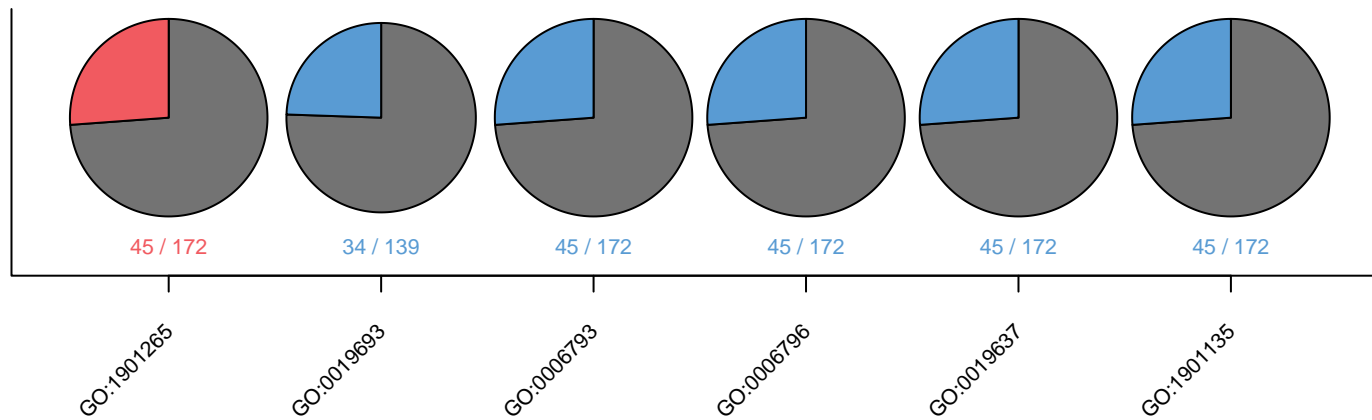


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p(A) in node

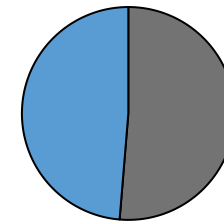


proportion of A



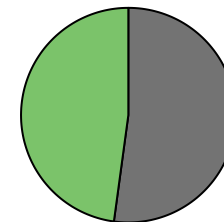
proportion of A
root nodes

biological_process



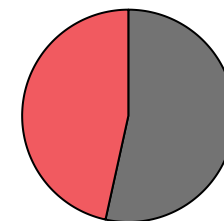
279 / 573

cellular_component



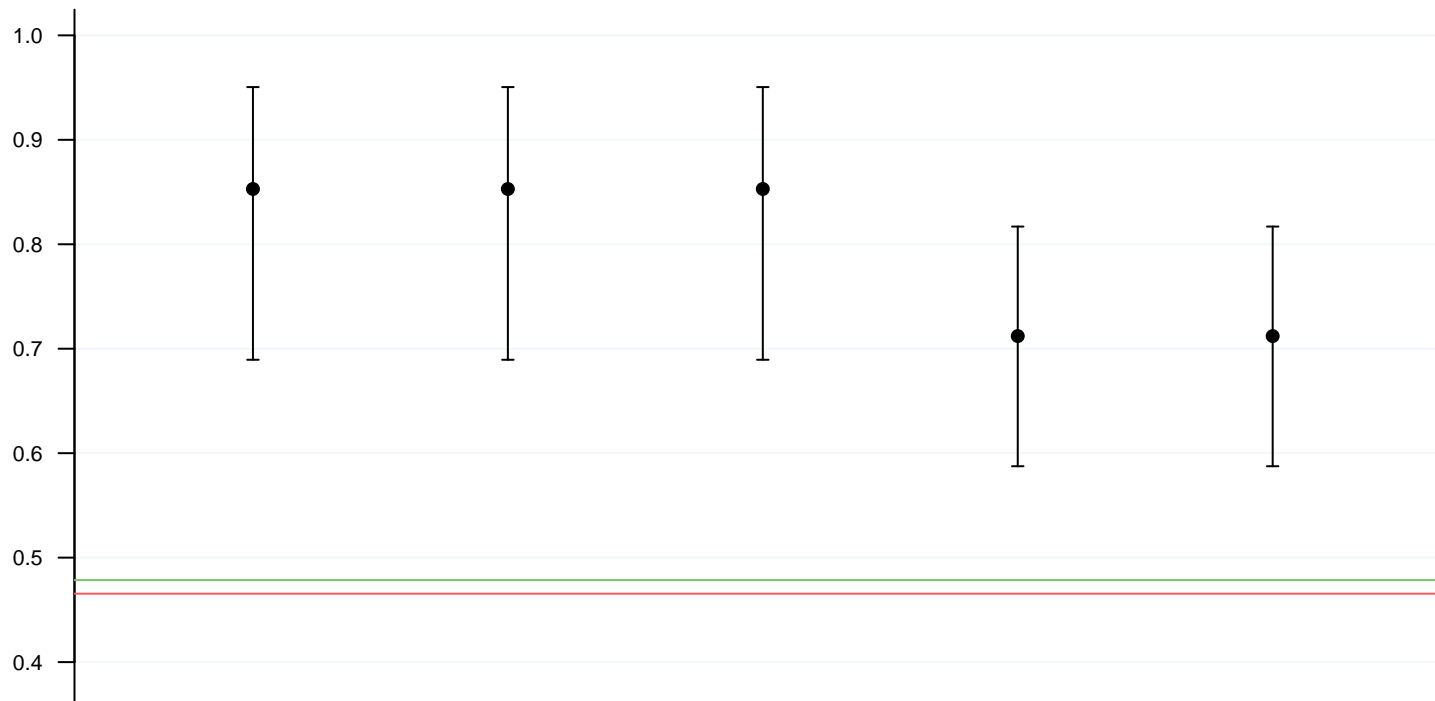
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molecular_function

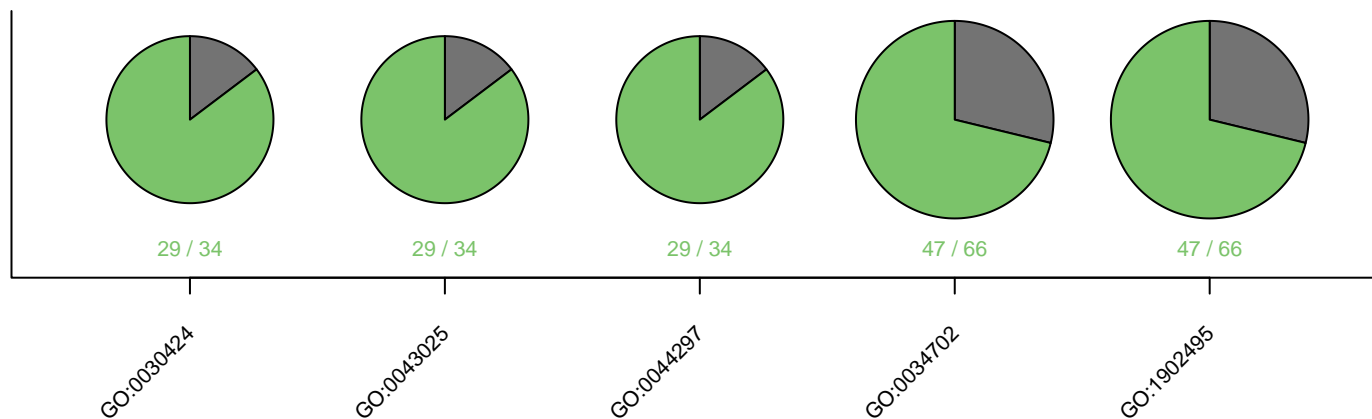


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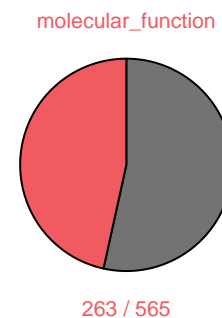
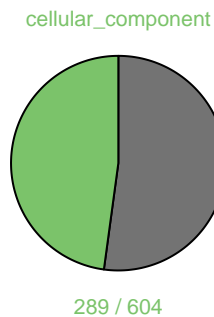
p(A) in node



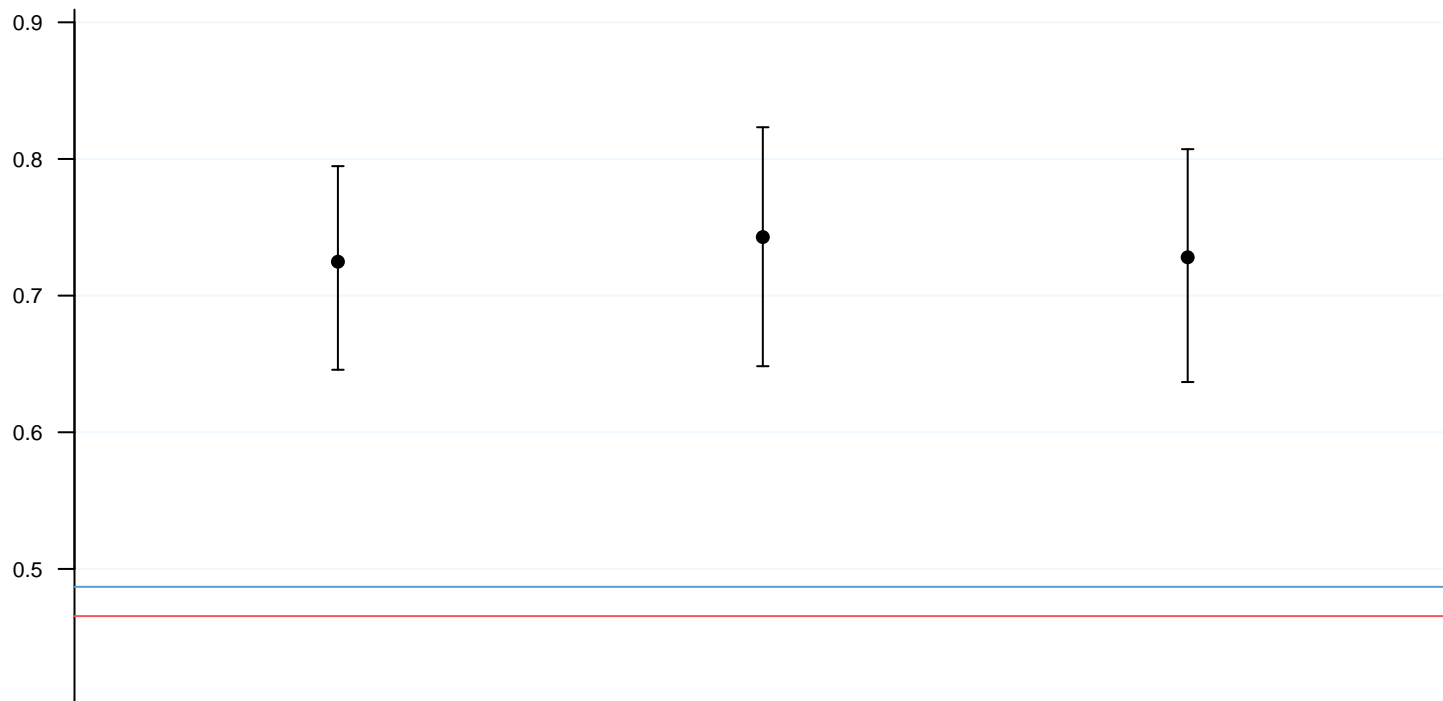
proportion of A



proportion of A
root nodes

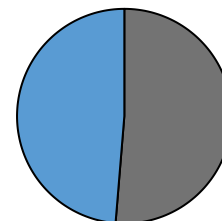


p(A) in node



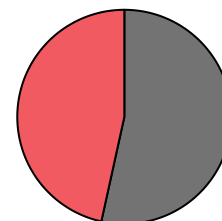
proportion of A
root nodes

biological_process



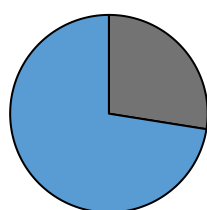
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molecular_function

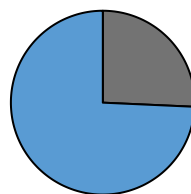


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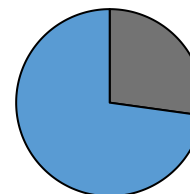
proportion of A



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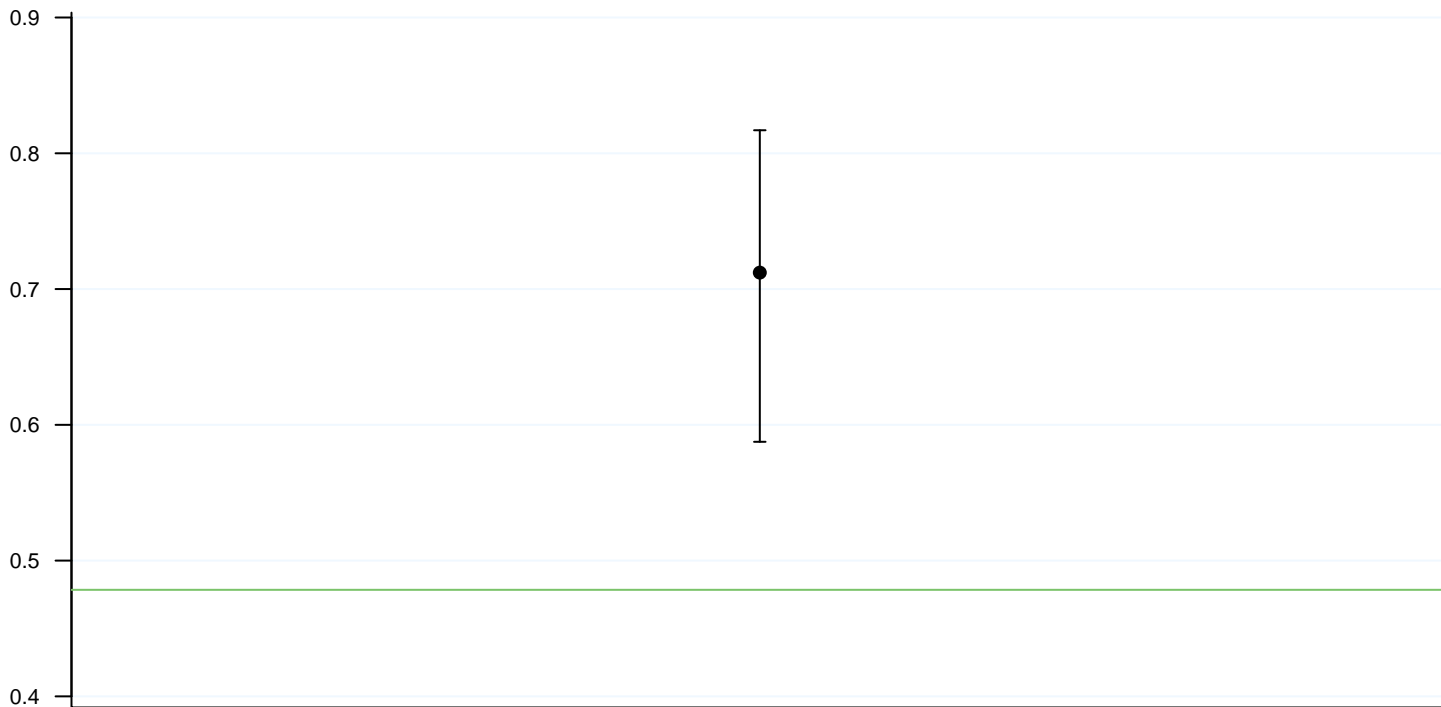
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GO:0003008

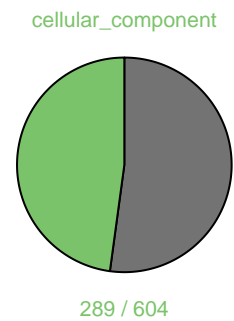
GO:0050877

GO:0001655

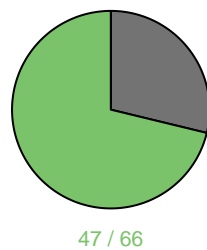
p(A) in node



proportion of A
root nodes

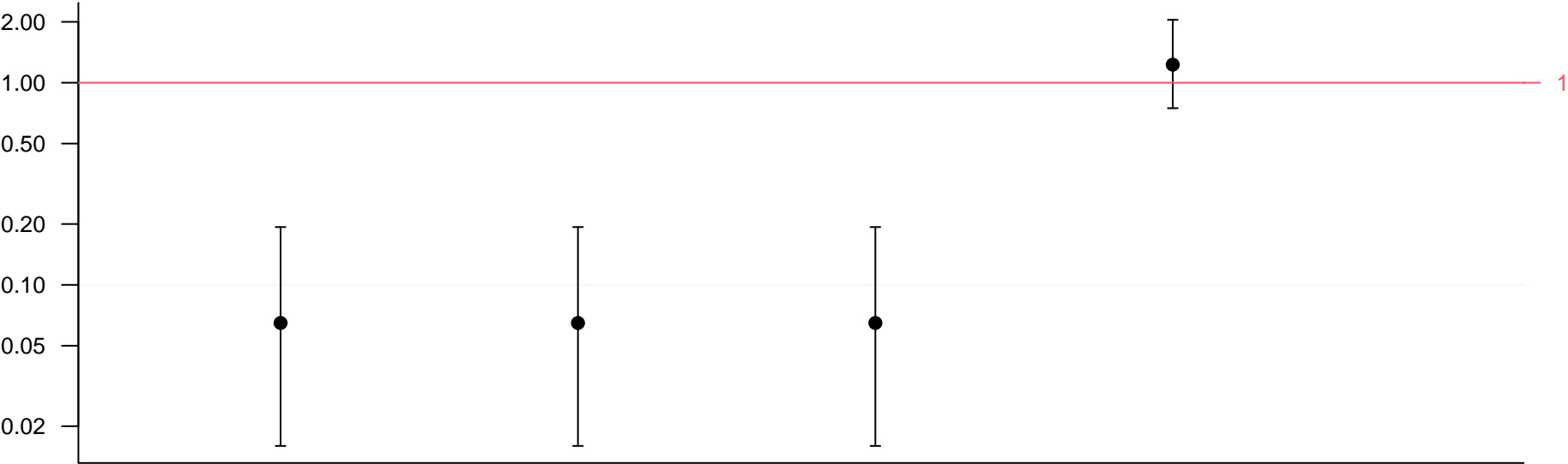


proportion of A

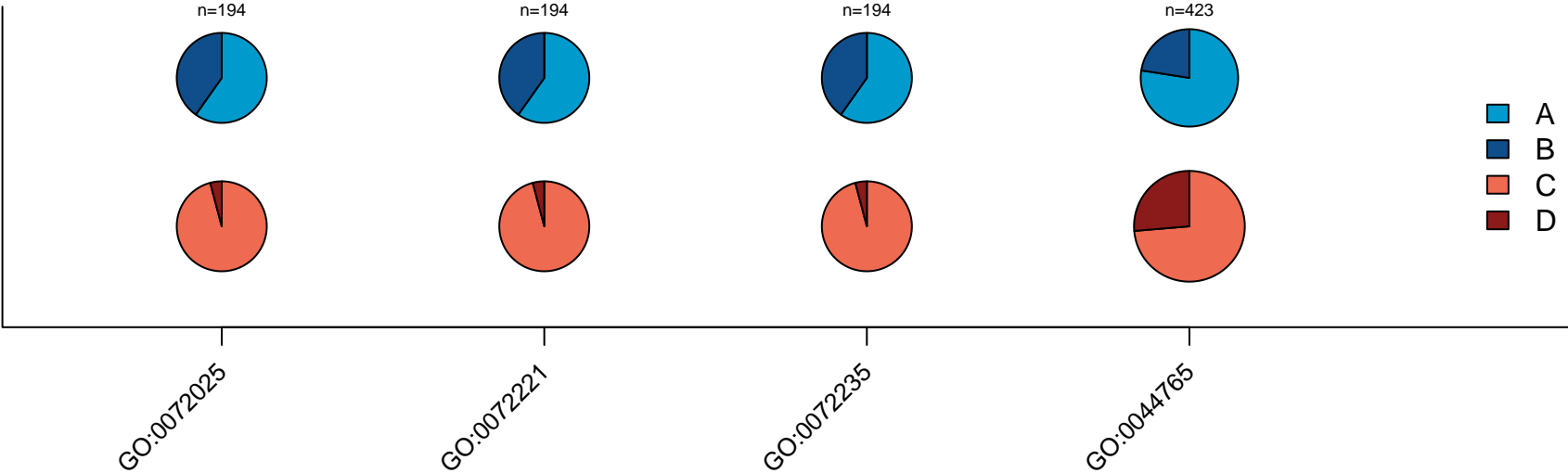


GO:0034702

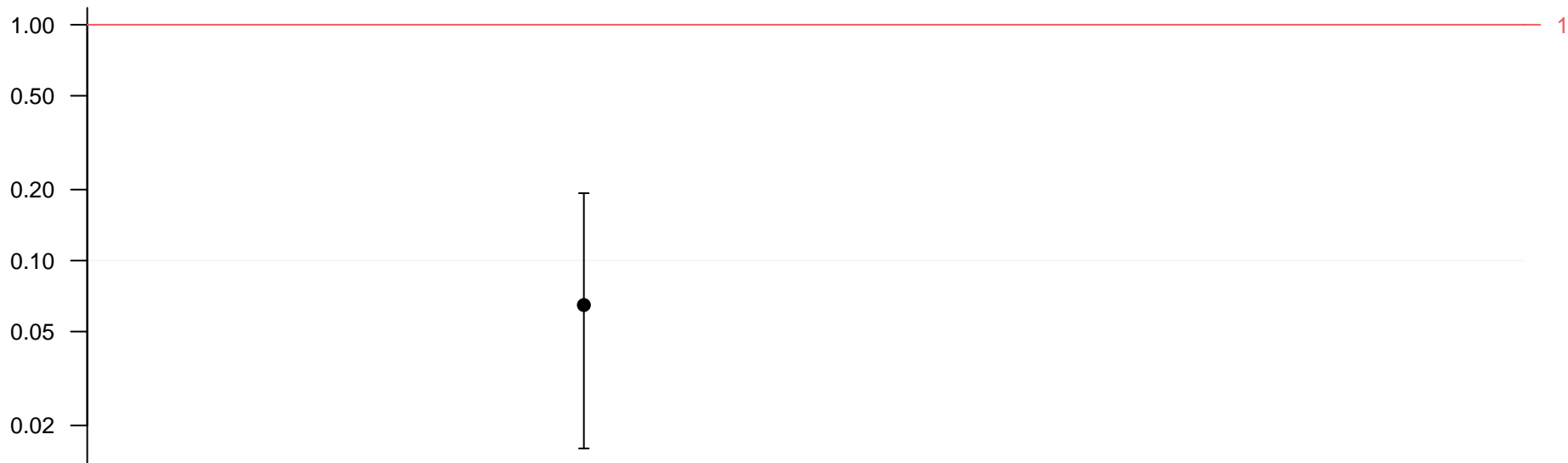
odds ratio (A/B) / (C/D)



annotated genes



odds ratio (A/B) / (C/D)



annotated genes

