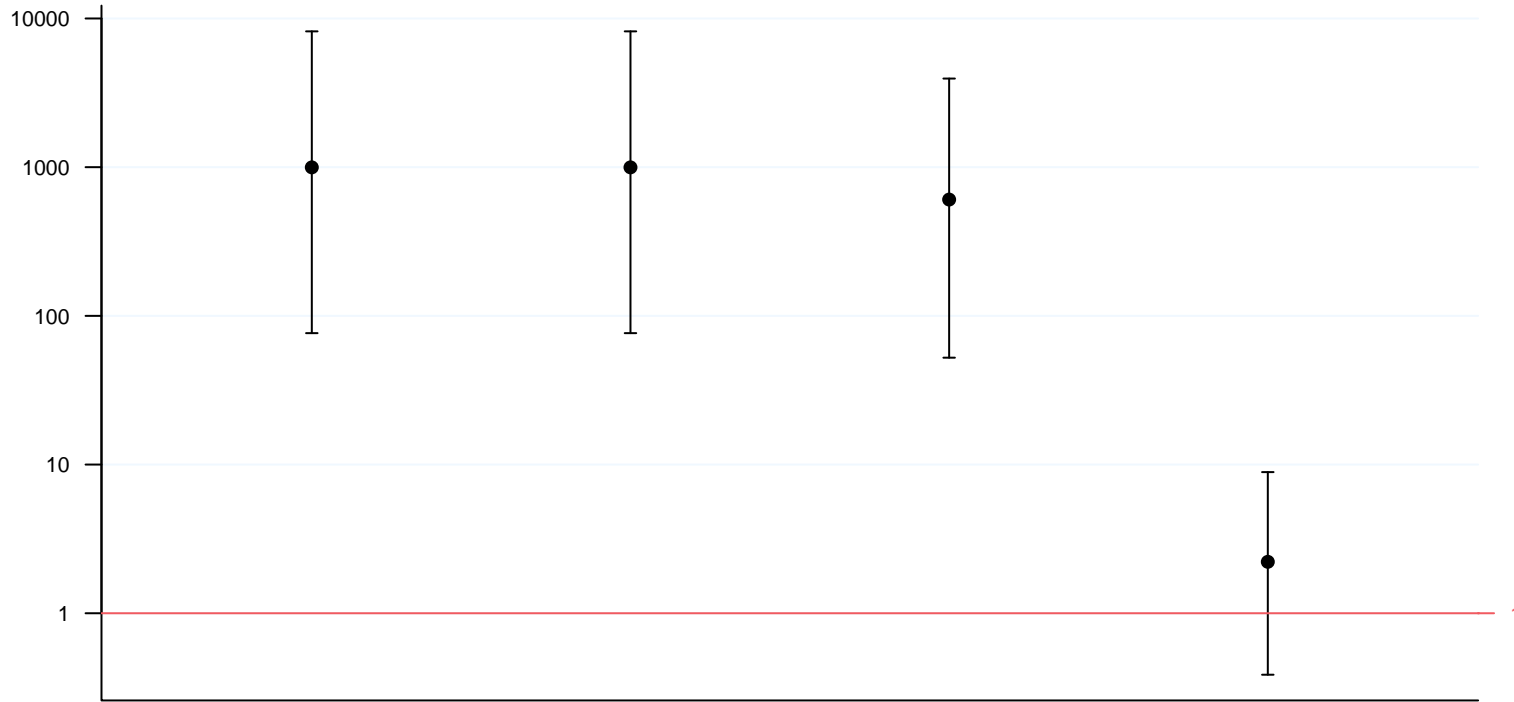
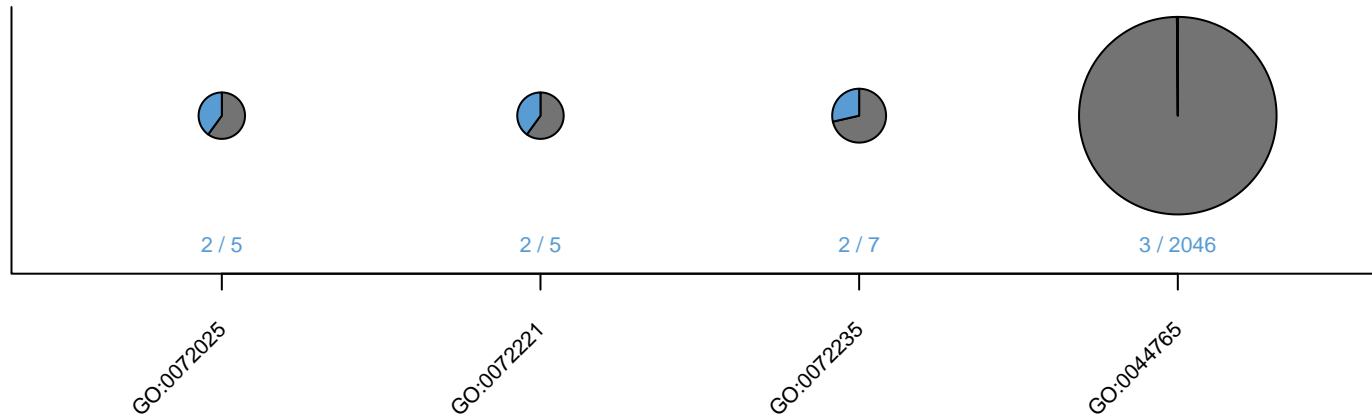


odds ratio

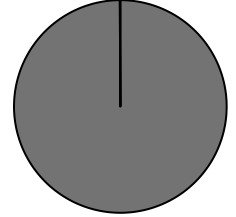


annotated genes



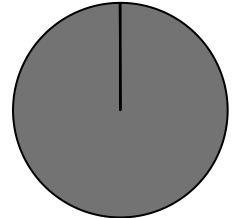
annotated genes root nodes

biological_process



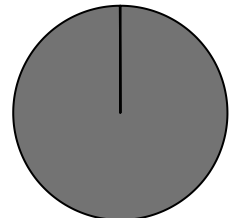
12 / 15655

cellular_component



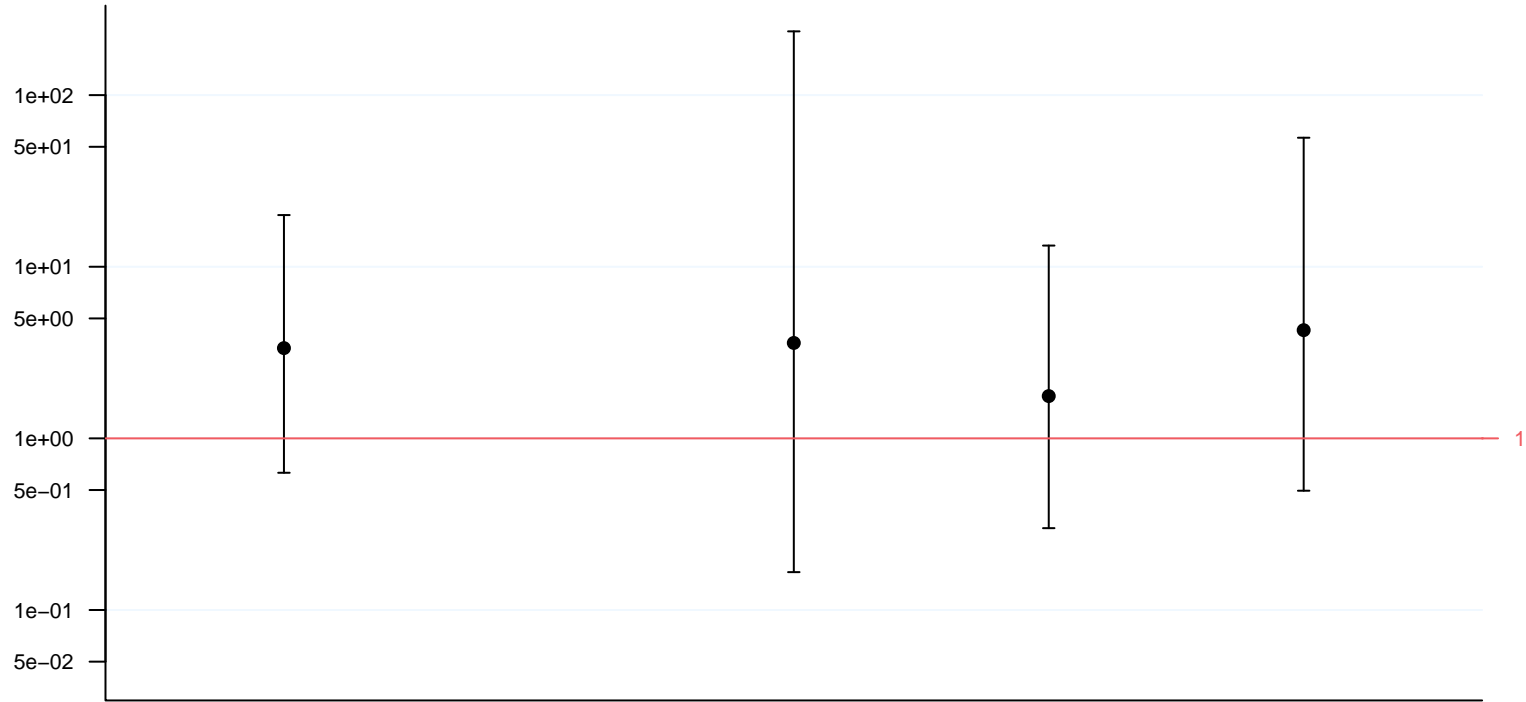
13 / 17106

molecular_function



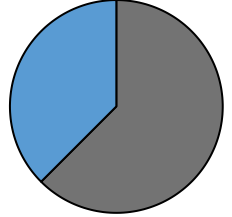
11 / 16809

odds ratio



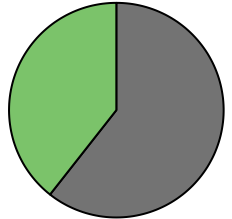
annotated genes root nodes

biological_process



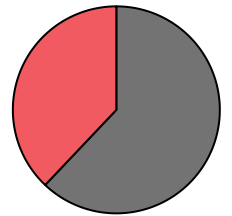
12 / 32

cellular_component



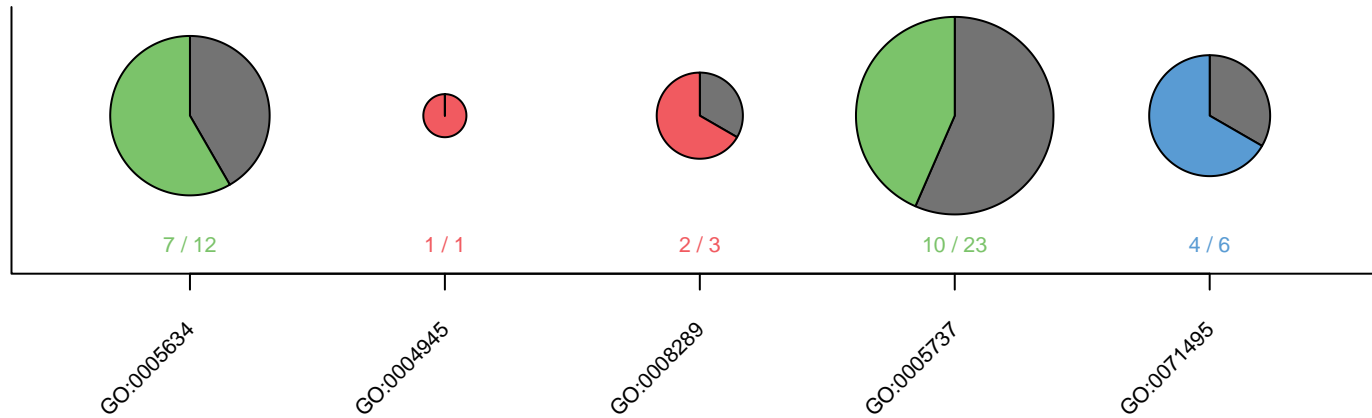
13 / 33

molecular_function

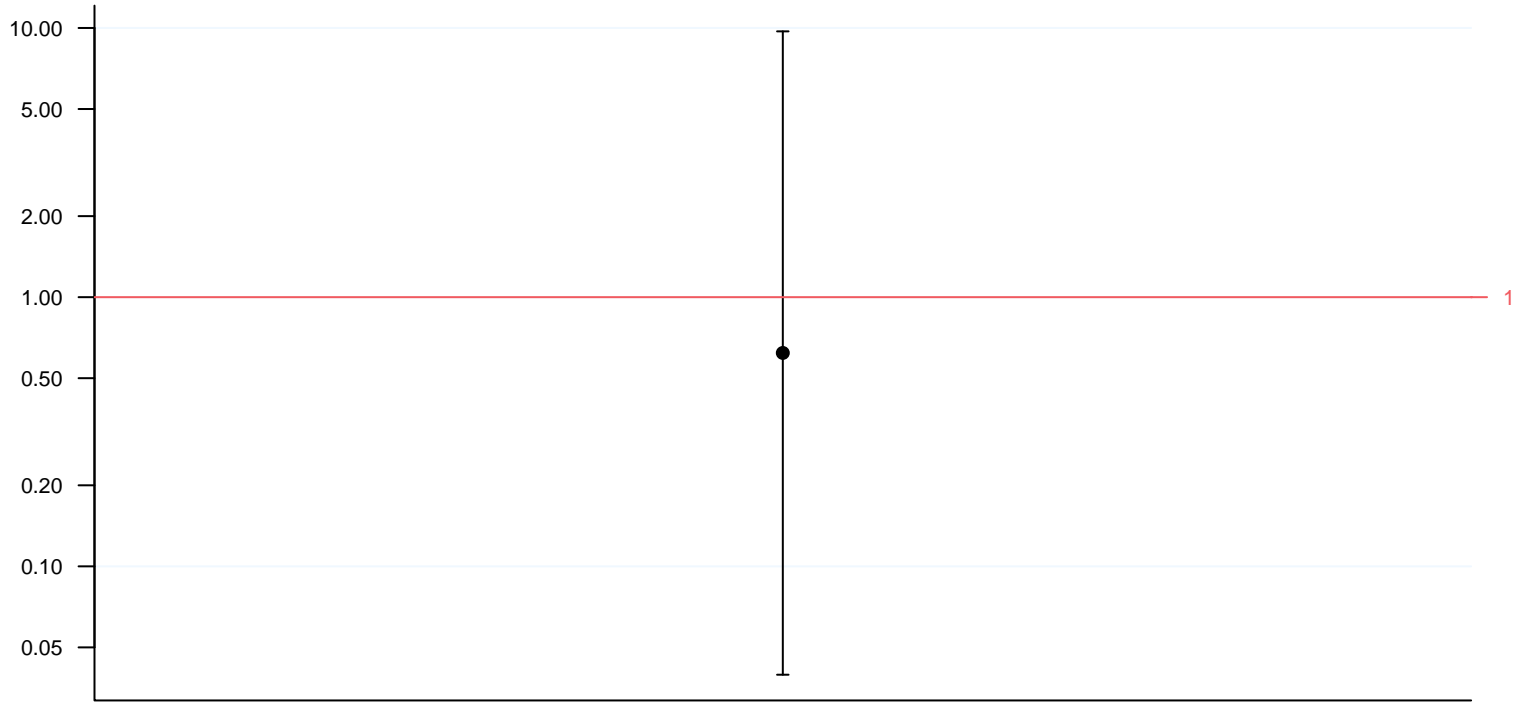


11 / 29

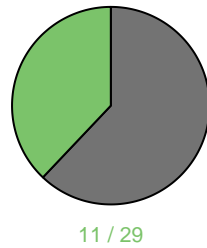
annotated genes



odds ratio



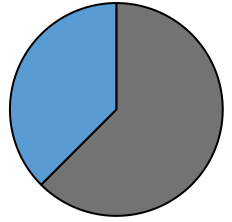
annotated genes



GO:0005623

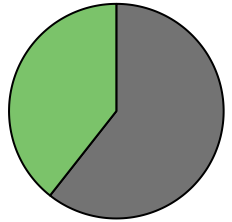
annotated genes root nodes

biological_process



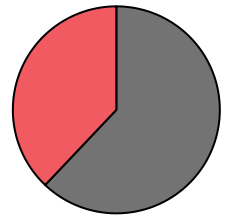
12 / 32

cellular_component



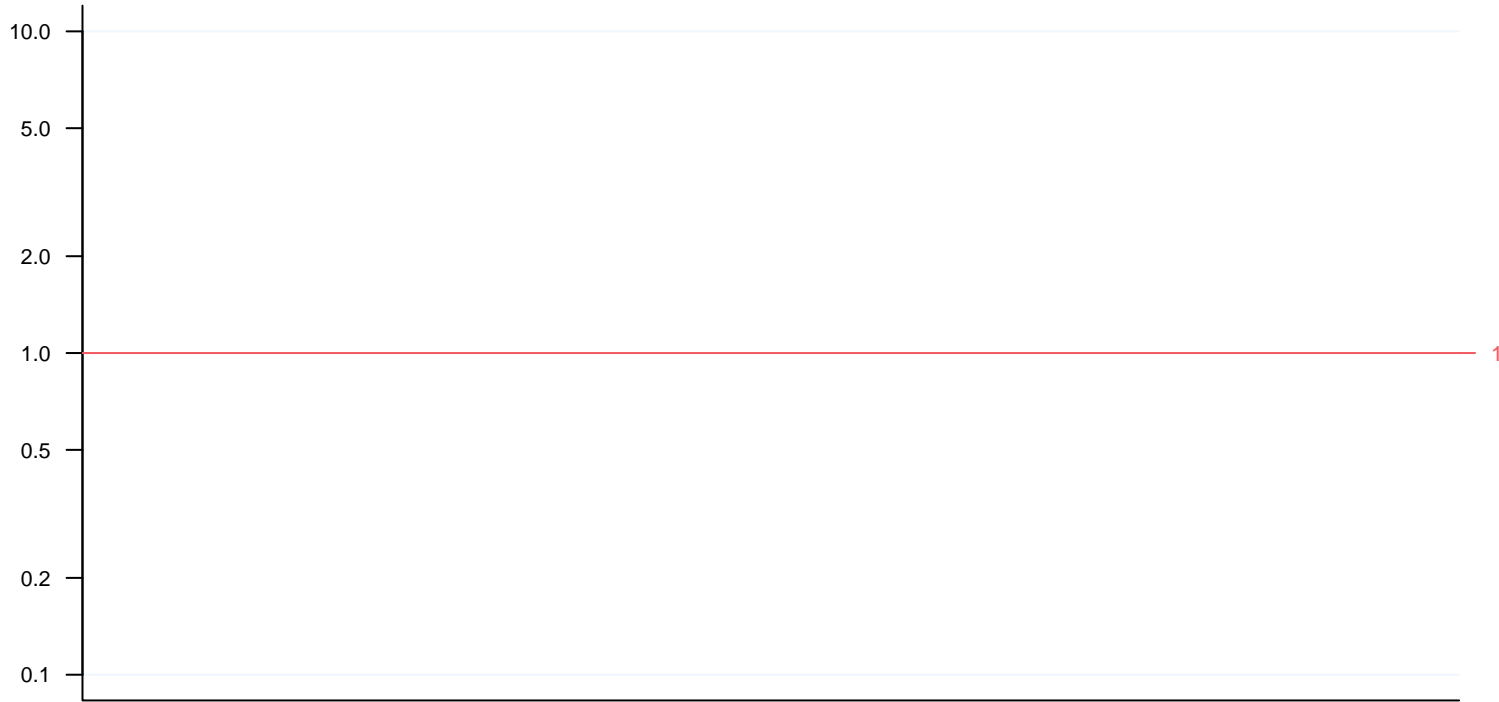
13 / 33

molecular_function

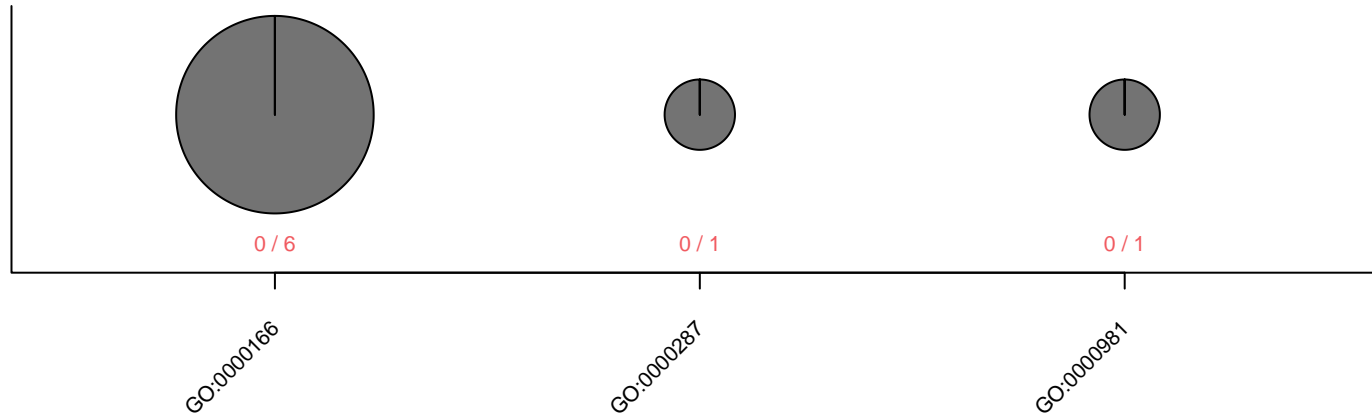


11 / 29

odds ratio

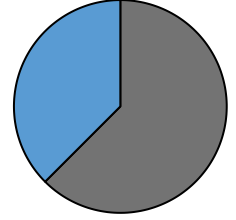


annotated genes



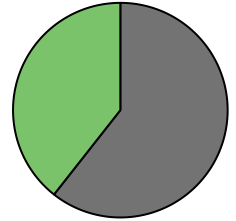
annotated genes root nodes

biological_process



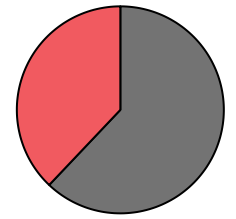
12 / 32

cellular_component



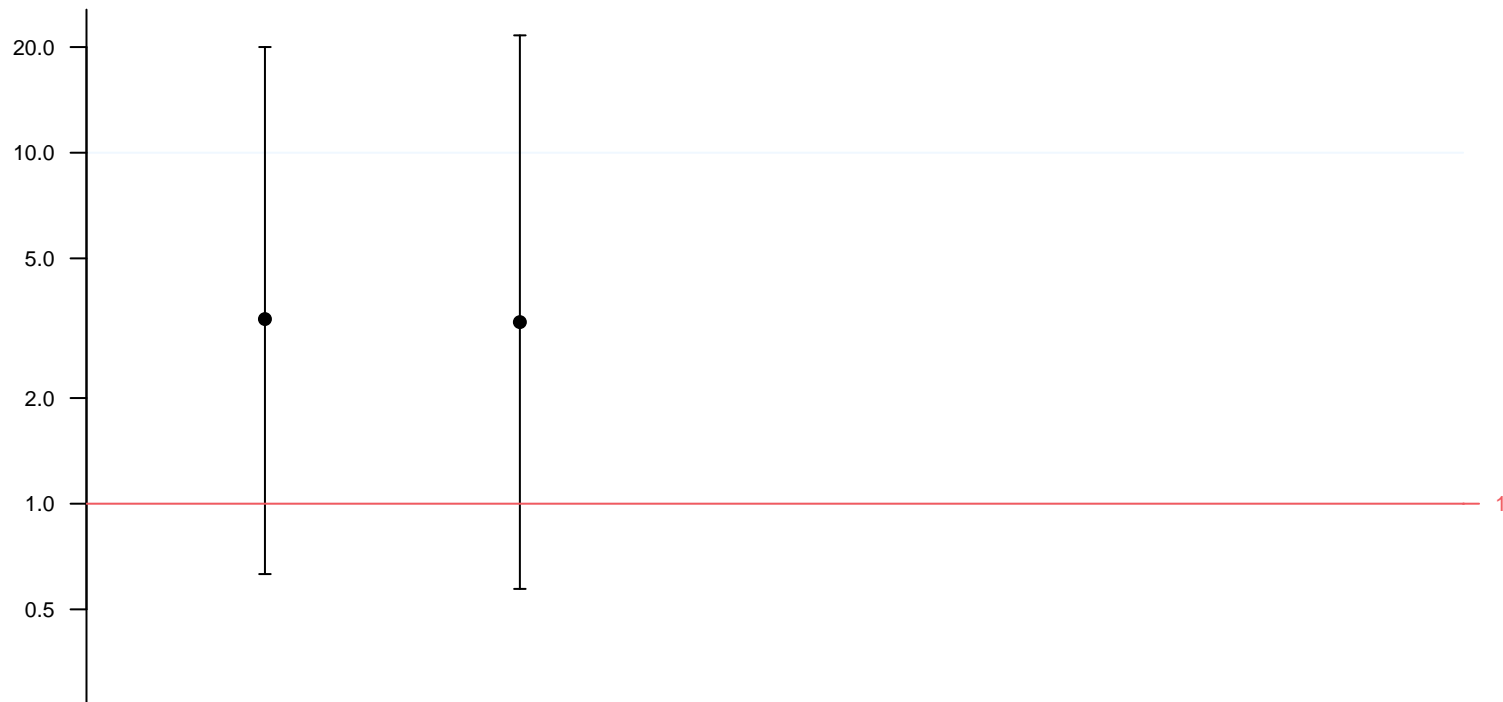
13 / 33

molecular_function

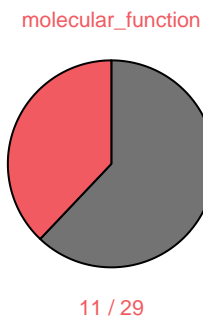
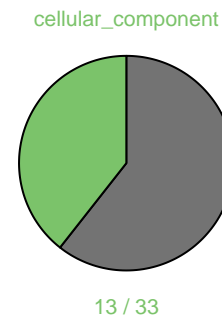


11 / 29

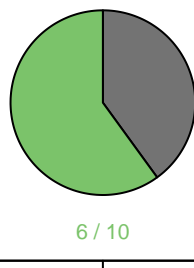
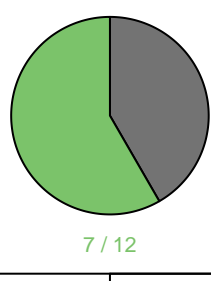
odds ratio



annotated genes root nodes



annotated genes



GO:0005634

GO:0005829

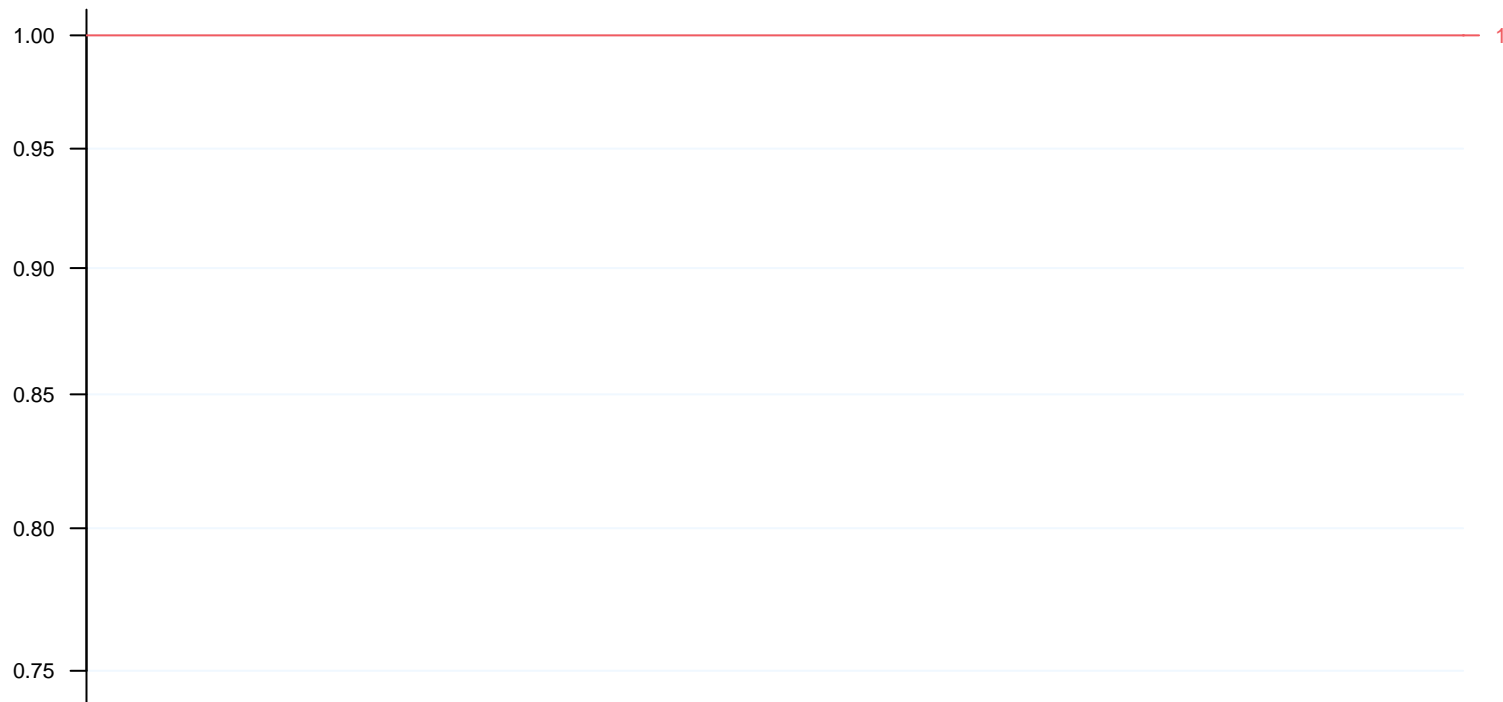
GO:0005216

GO:0015075

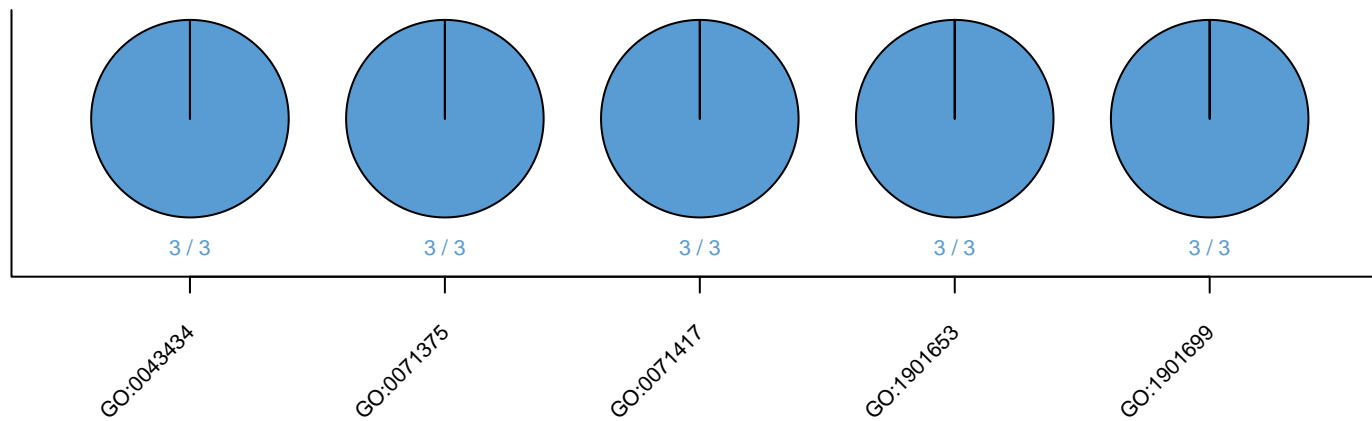
GO:0015267

odds ratio

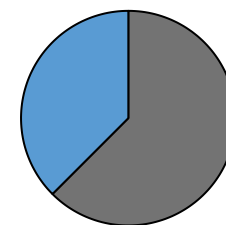
annotated genes
root nodes



annotated genes

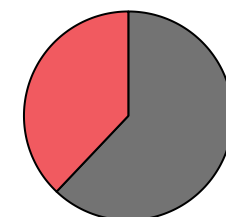


biological_process



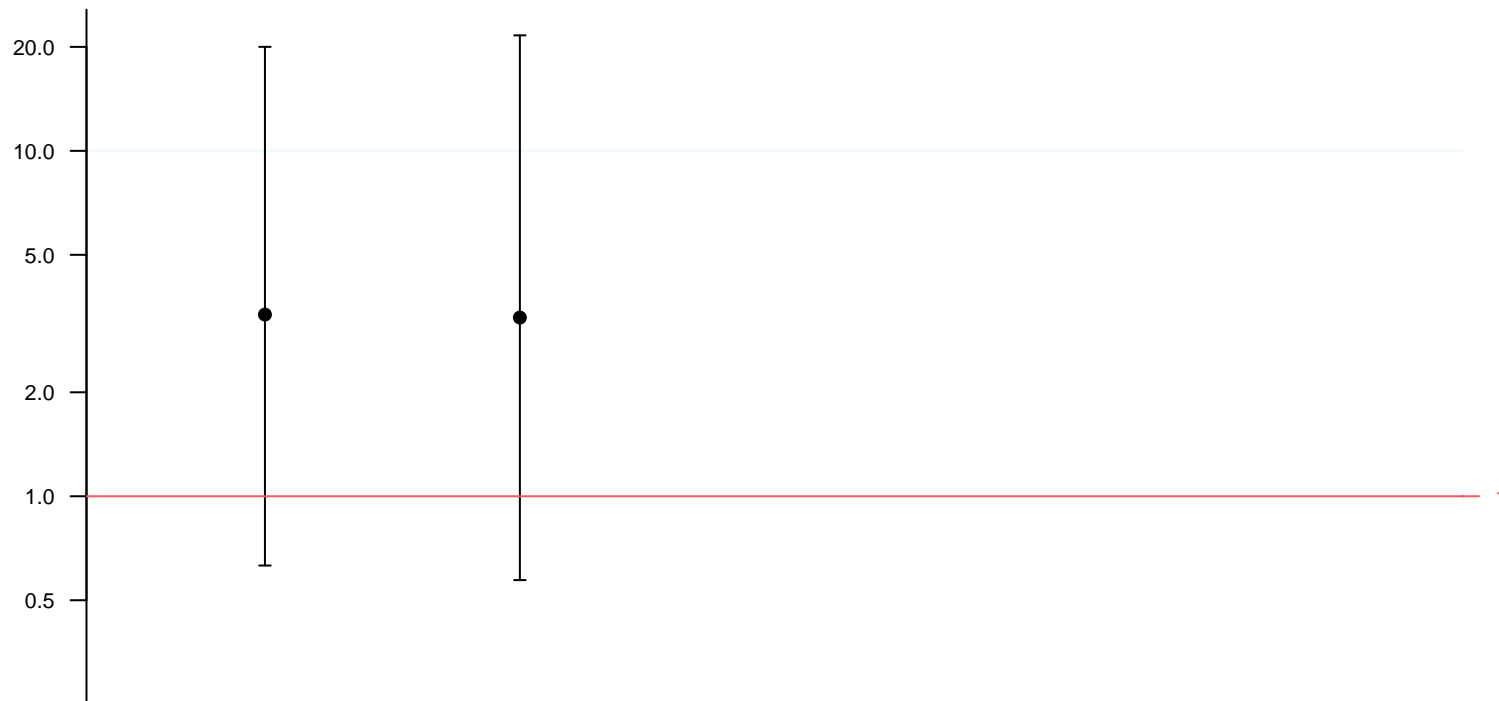
12 / 32

molecular_function



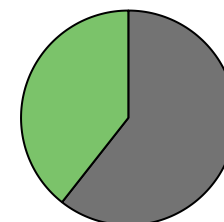
11 / 29

odds ratio



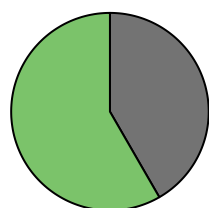
annotated genes
root nodes

cellular_component

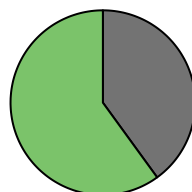


13 / 33

annotated genes



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6 / 10



2 / 2



2 / 2



2 / 2

GO:0005634

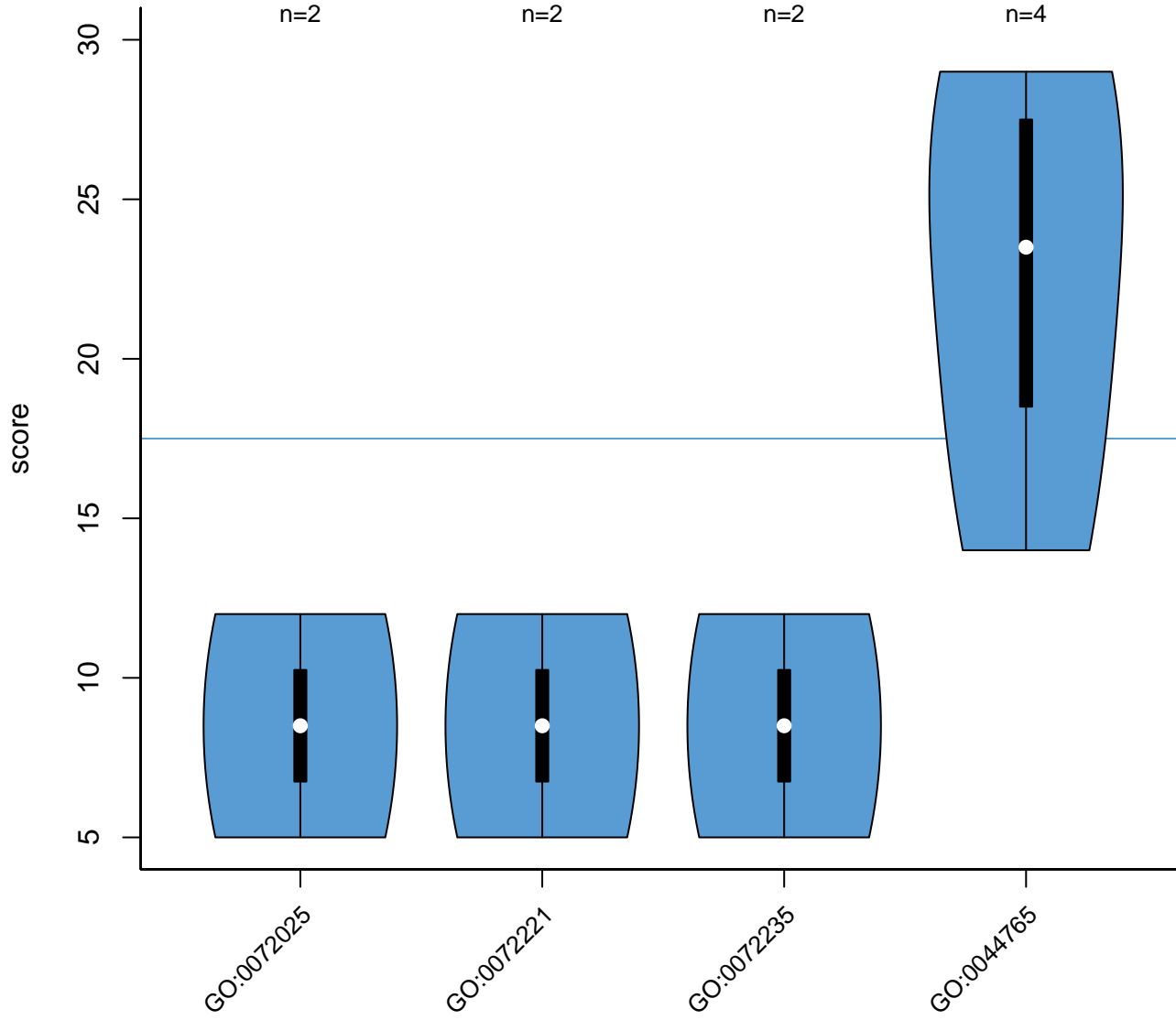
GO:0005829

GO:0034702

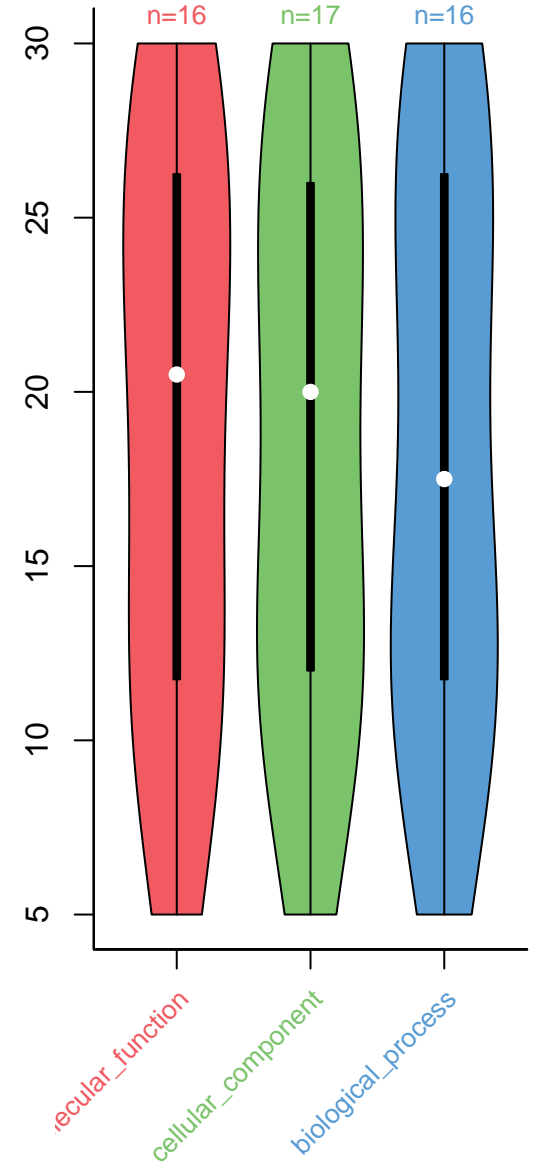
GO:1902495

GO:1990351

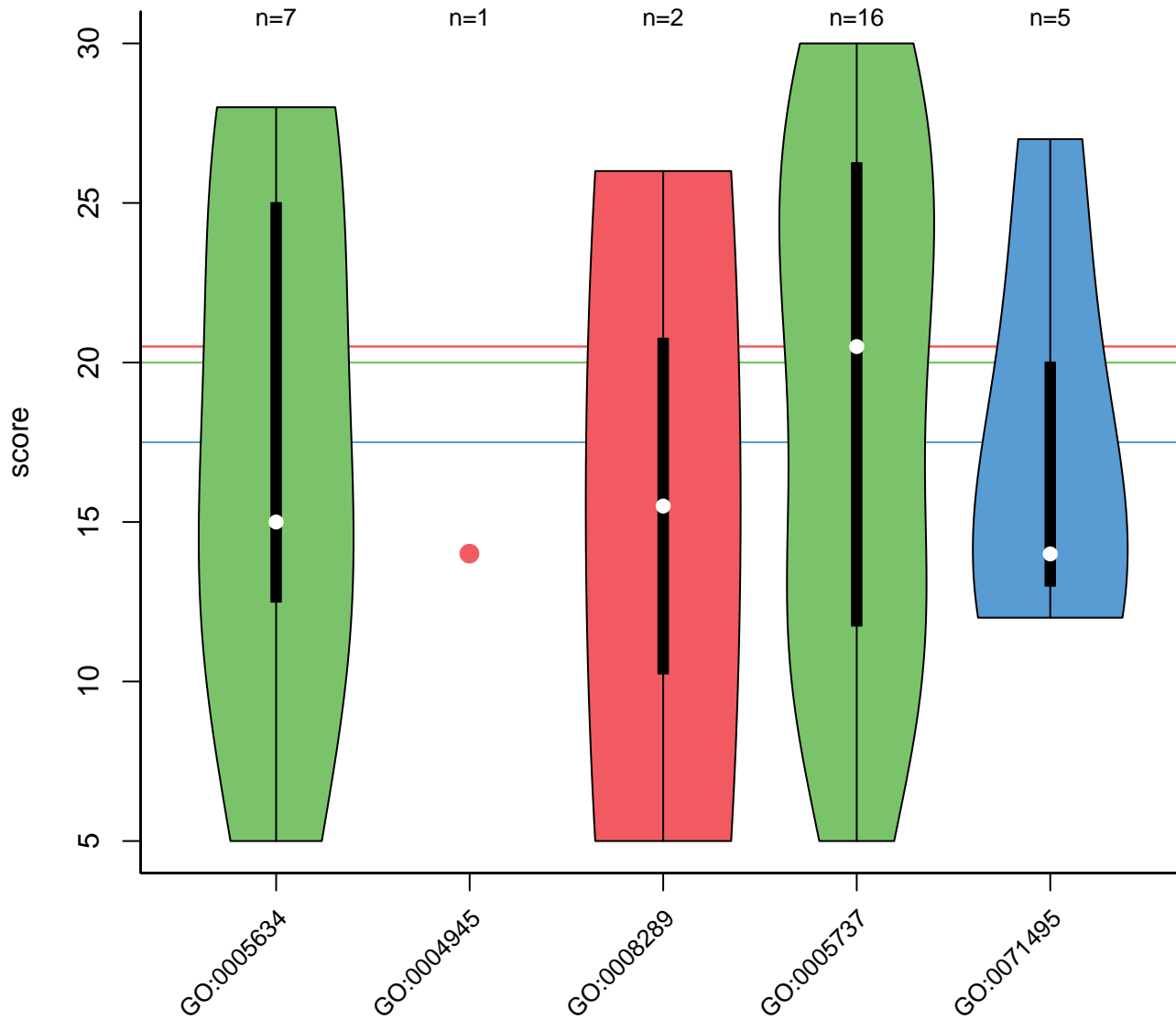
scores of annotated genes



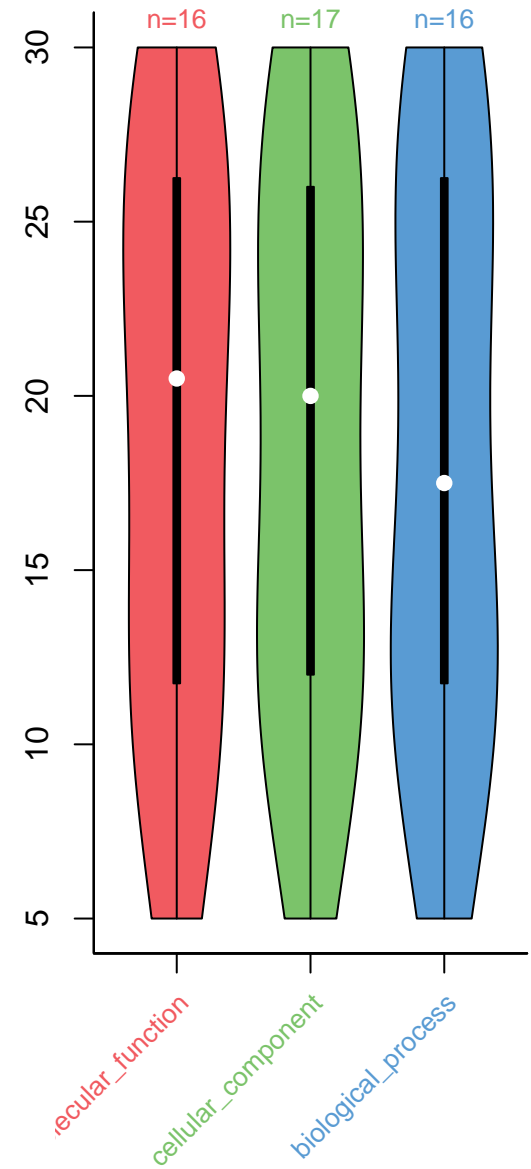
root nodes



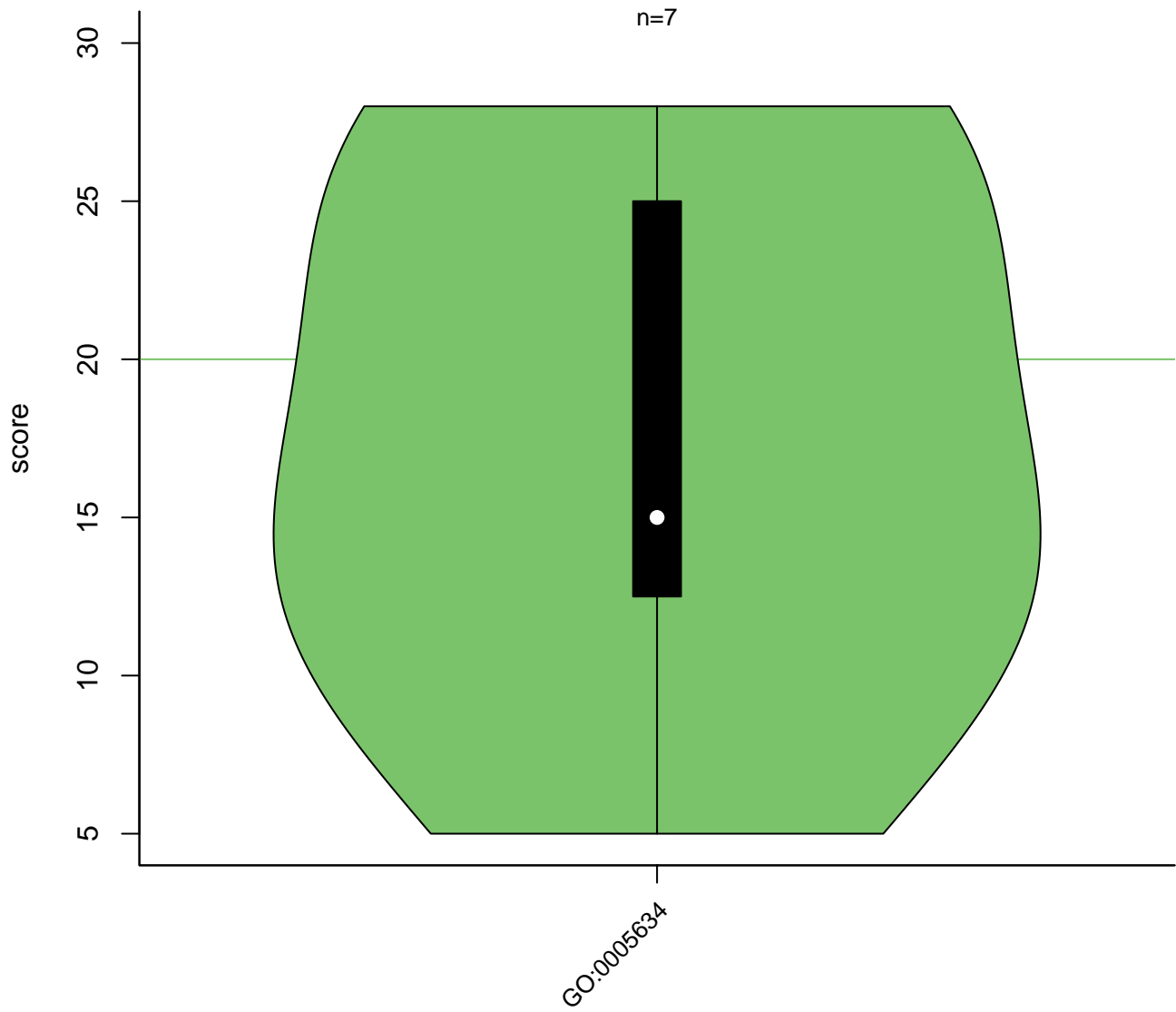
scores of annotated genes



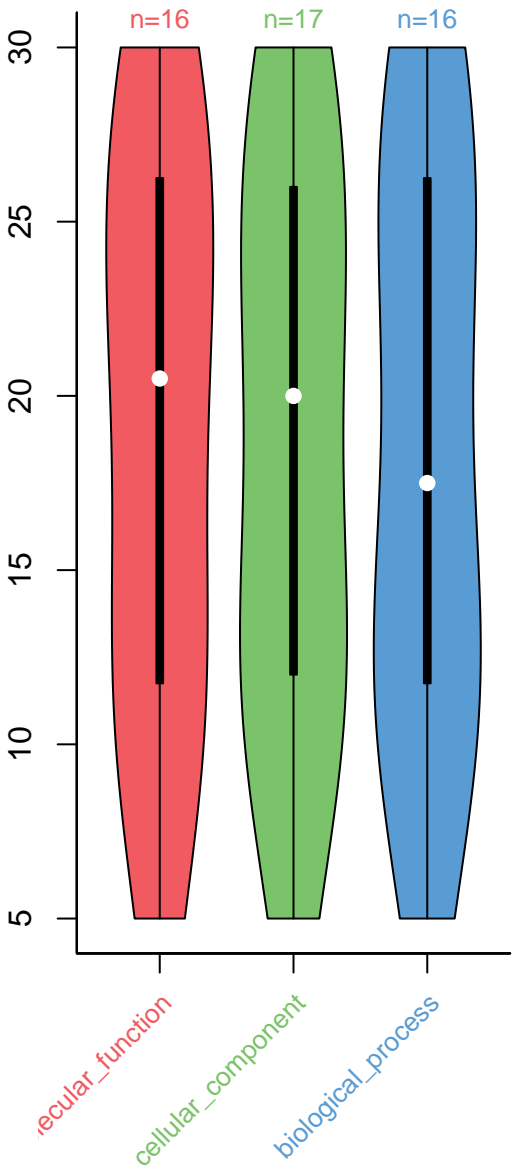
root nodes



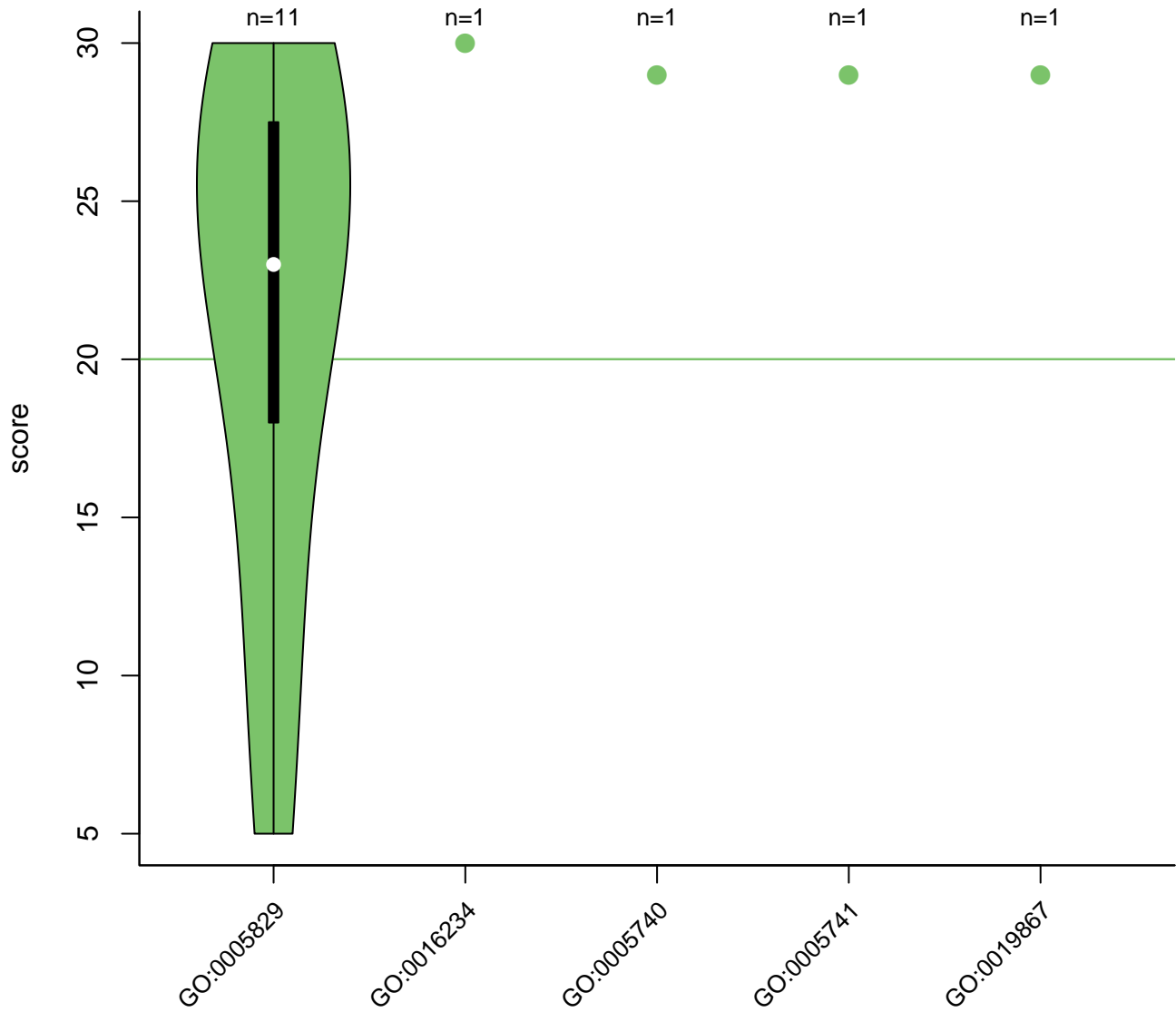
scores of annotated genes



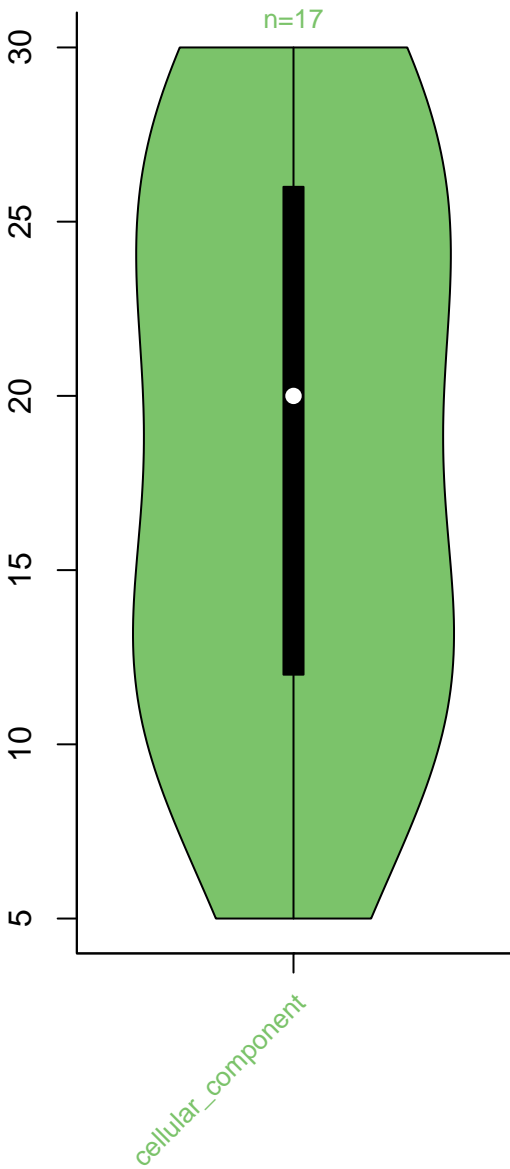
root nodes



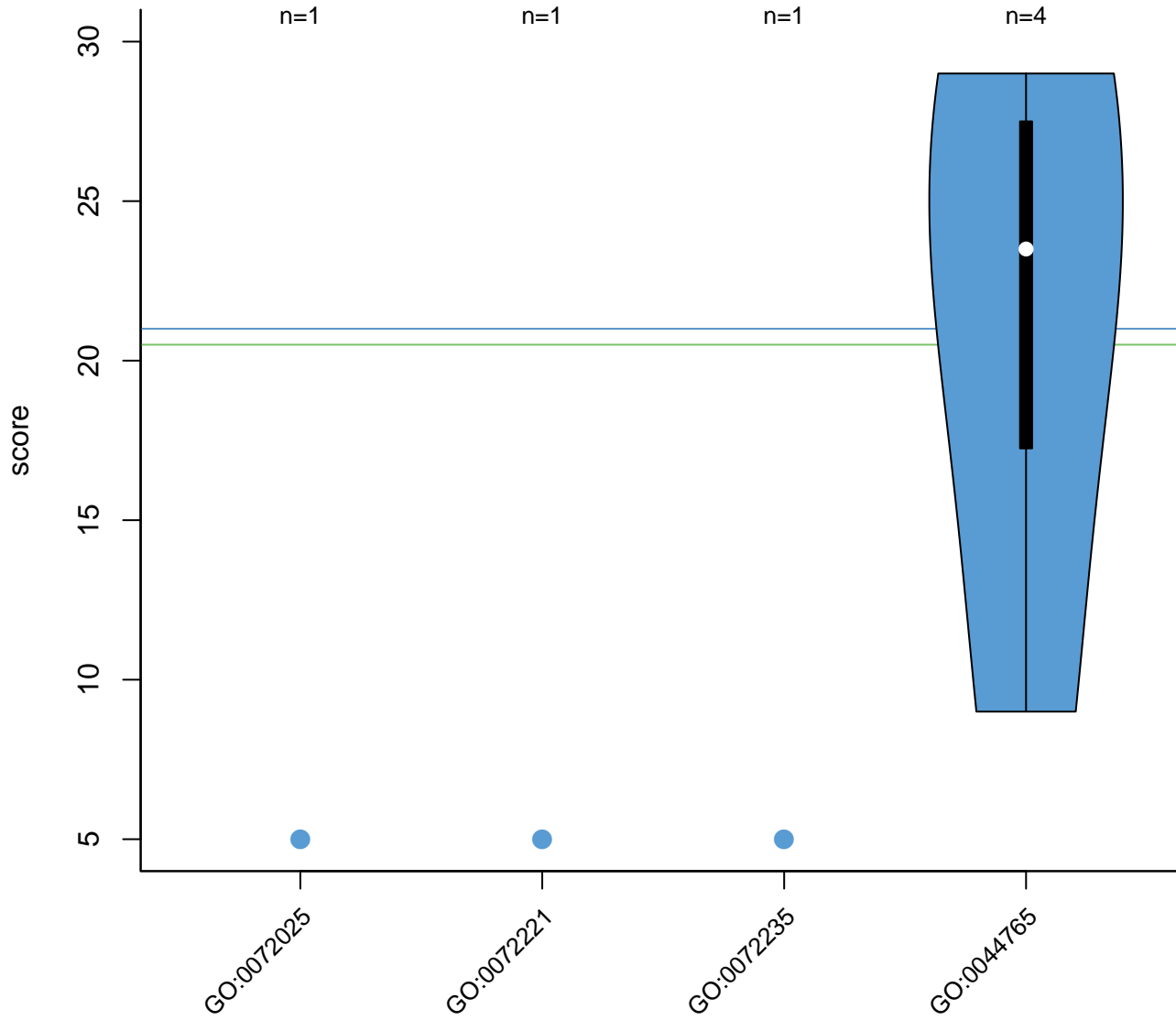
scores of annotated genes



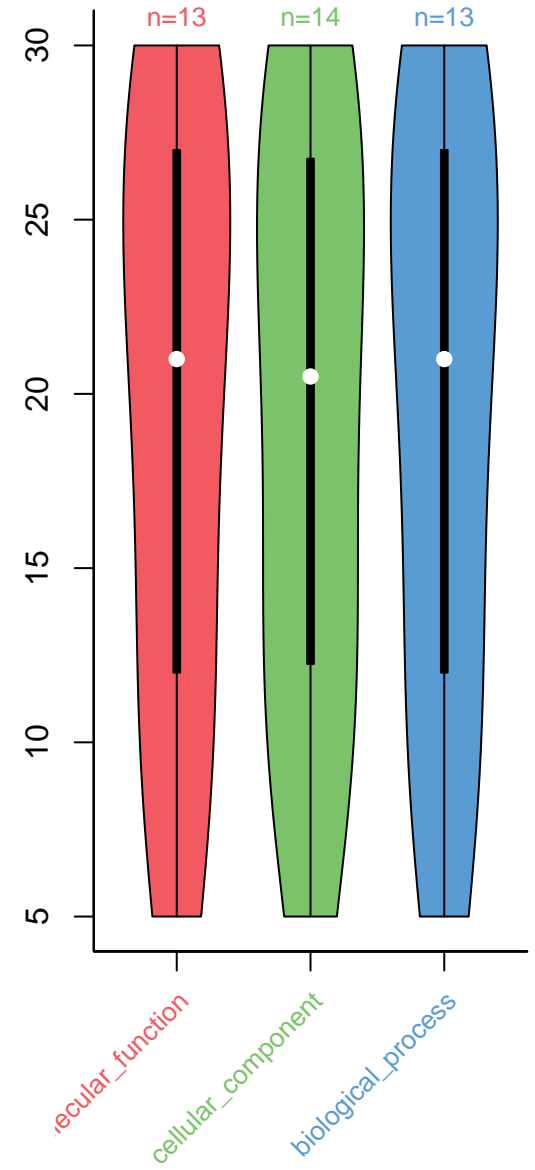
root nodes



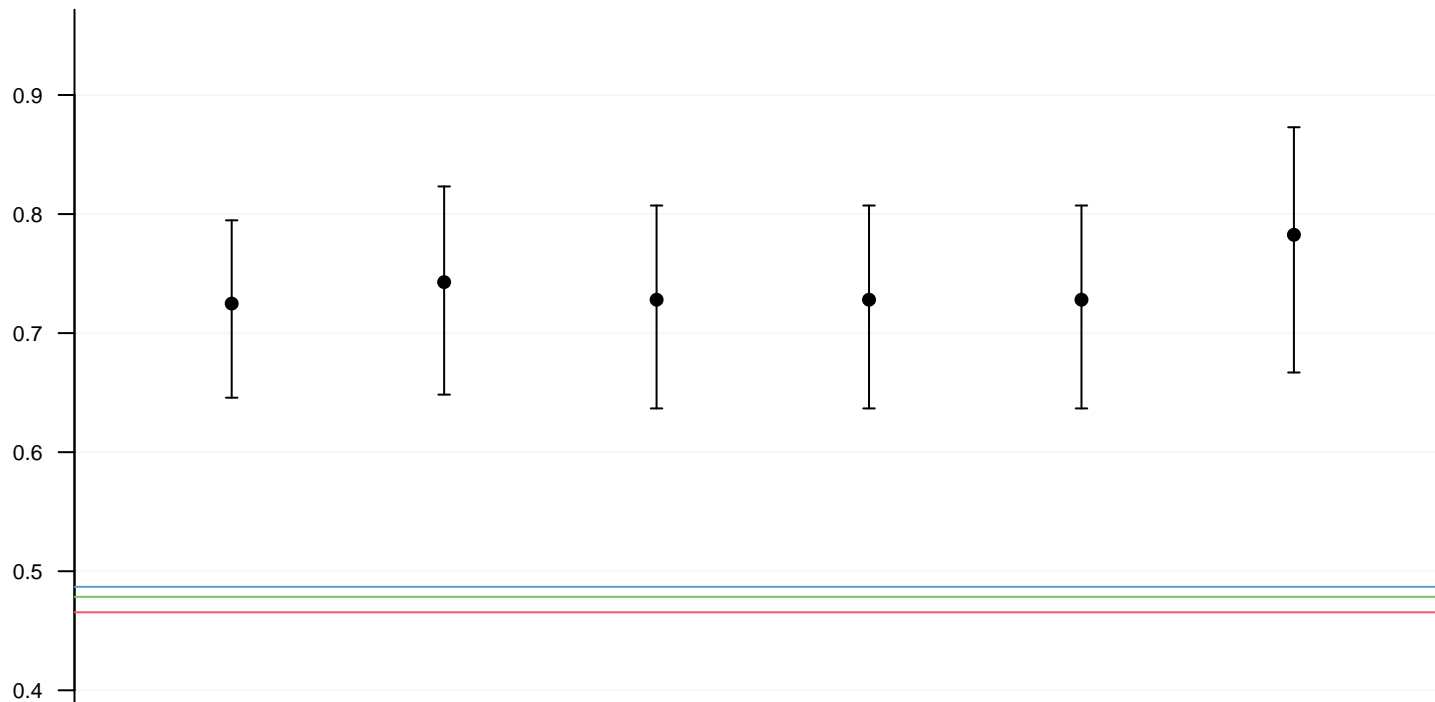
scores of annotated genes



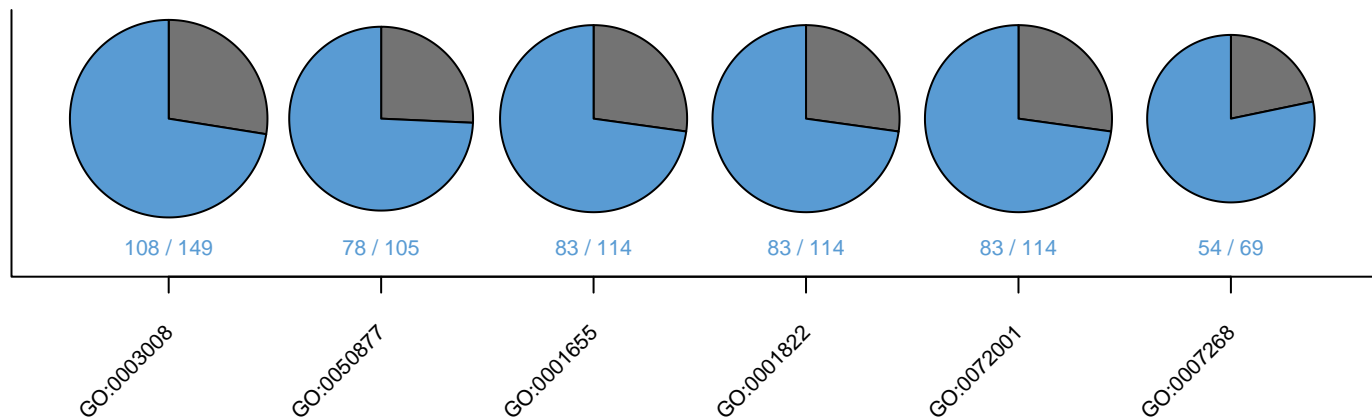
root nodes



p(A) in node

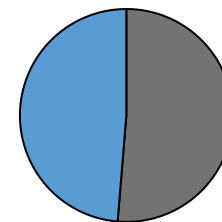


proportion of A



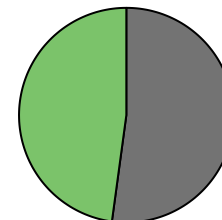
proportion of A
root nodes

biological_process



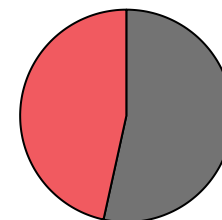
279 / 573

cellular_component



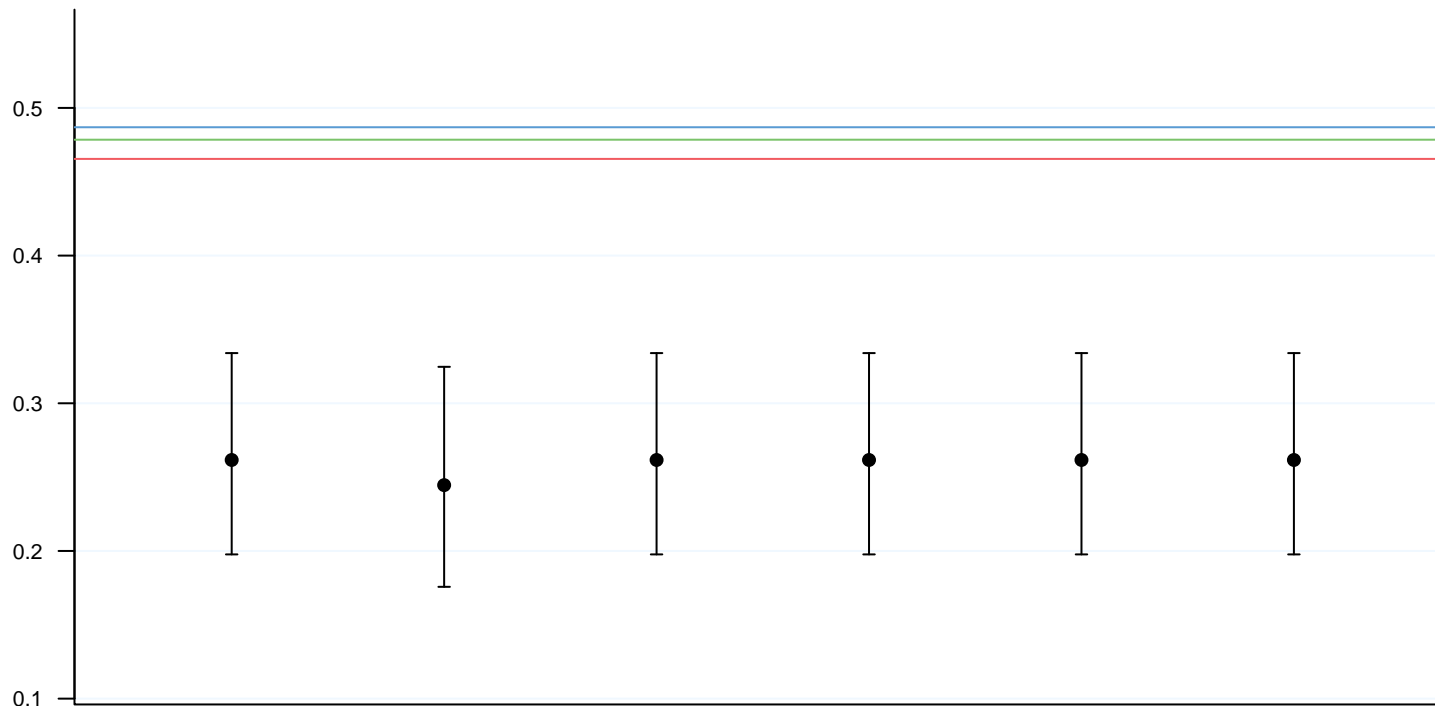
289 / 604

molecular_function

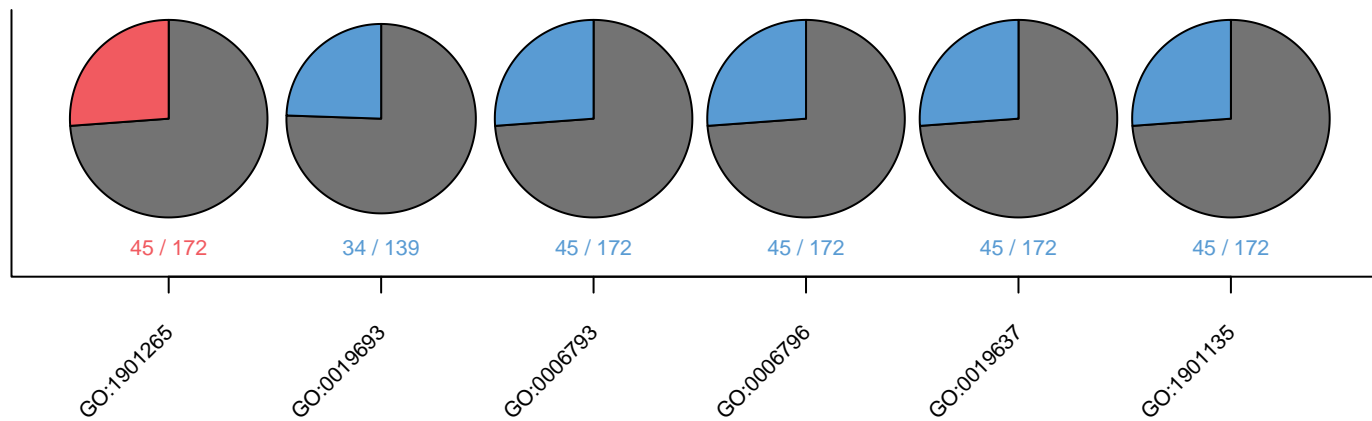


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p(A) in node

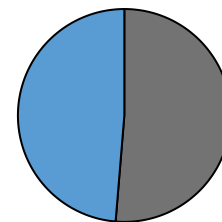


proportion of A



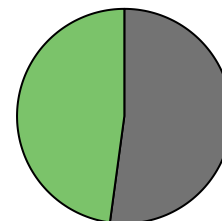
proportion of A
root nodes

biological_process



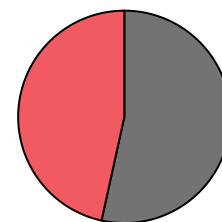
279 / 573

cellular_component



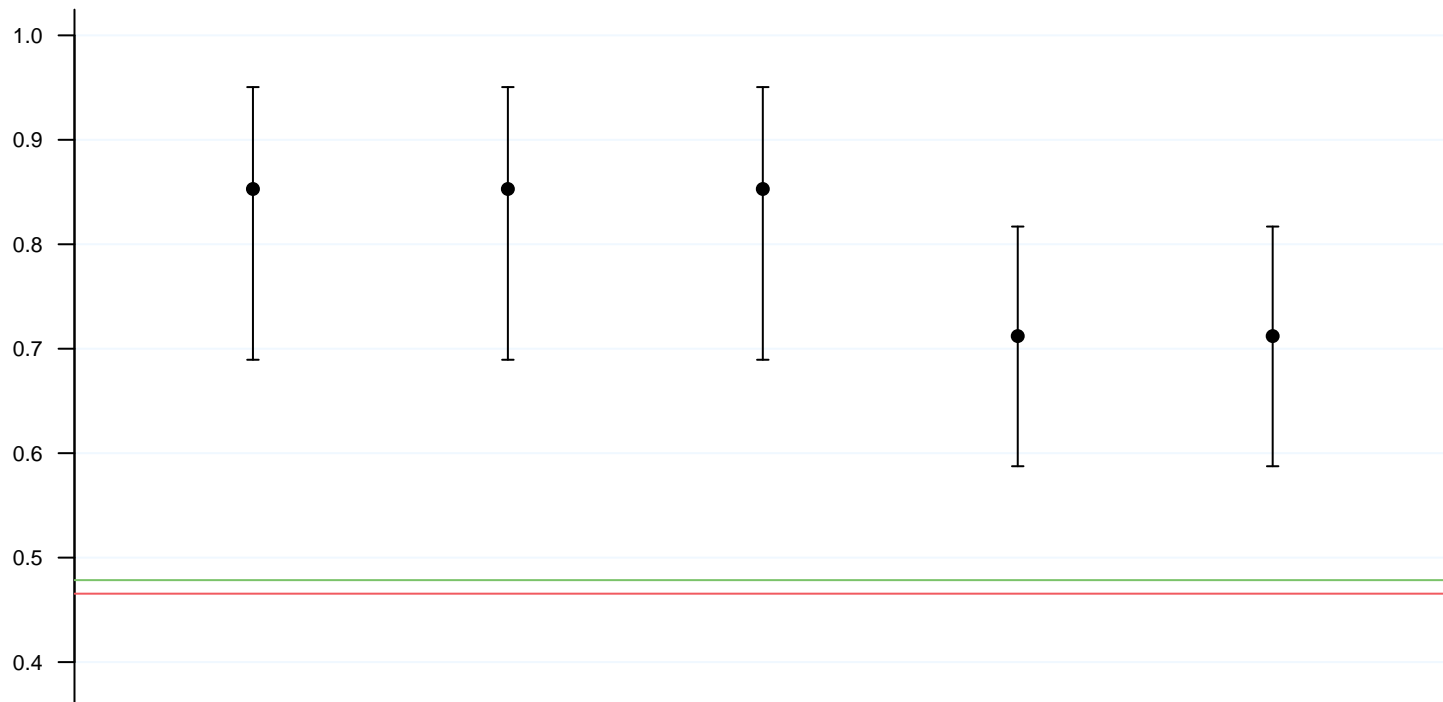
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molecular_function

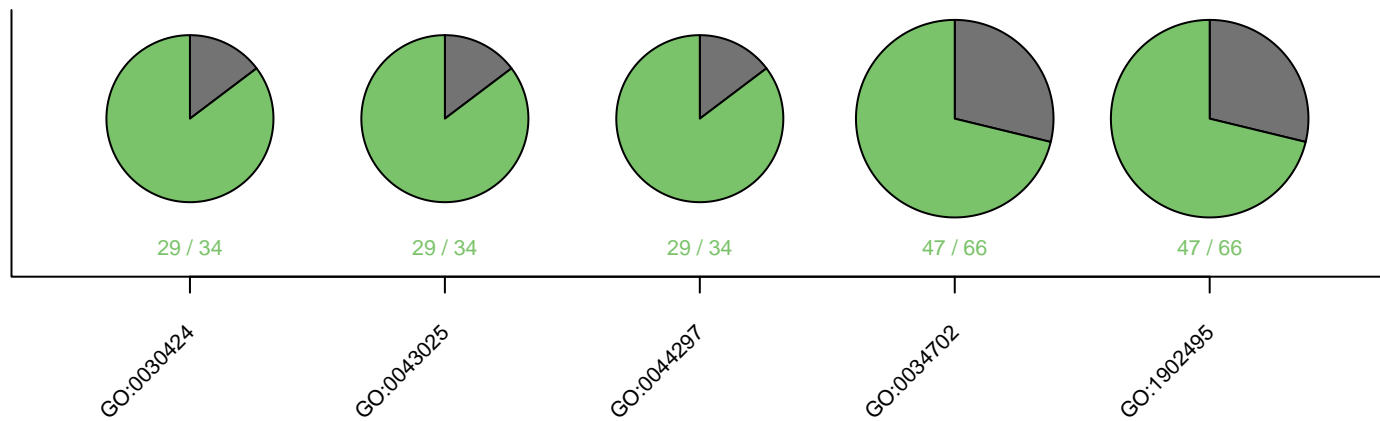


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p(A) in node

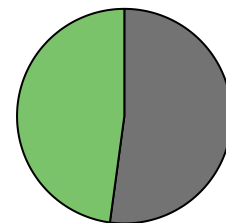


proportion of A



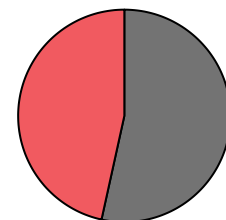
proportion of A
root nodes

cellular_component



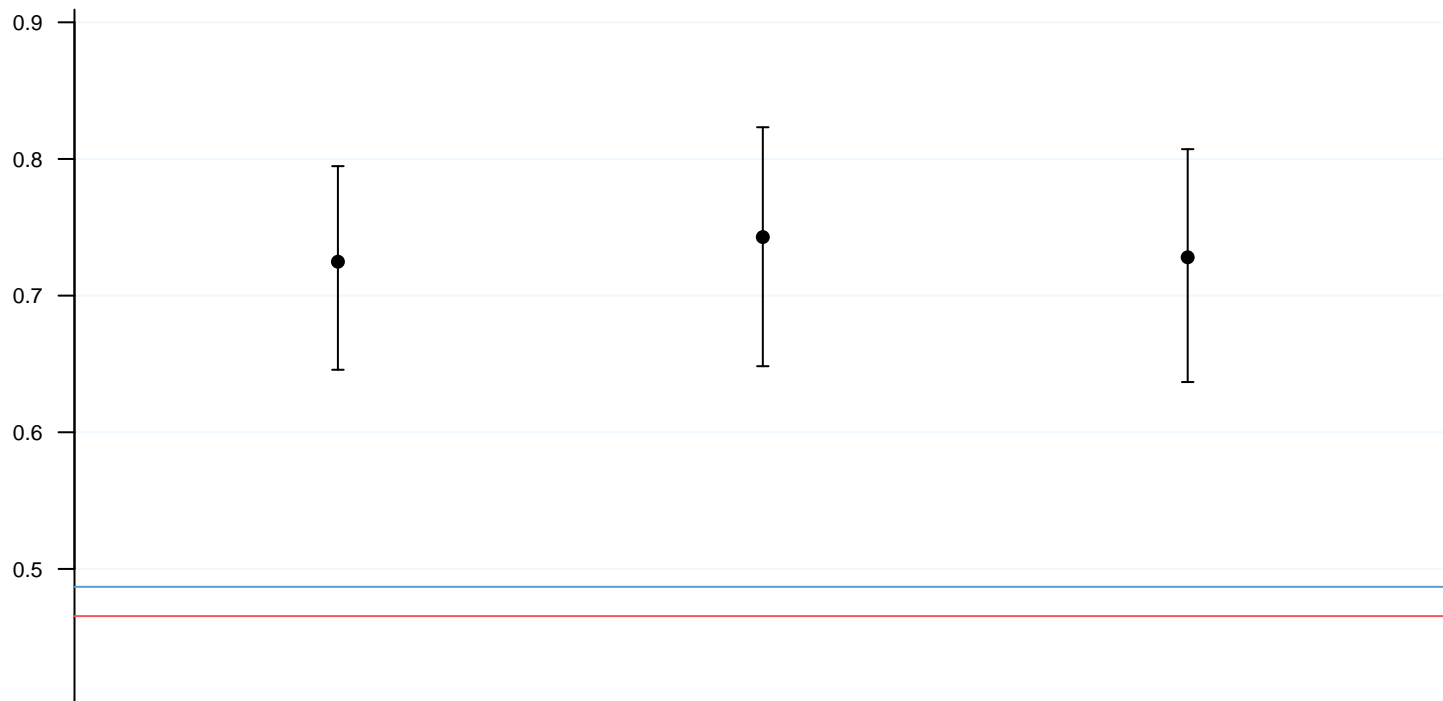
289 / 604

molecular_function



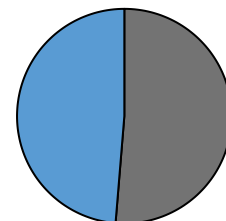
263 / 565

p(A) in node



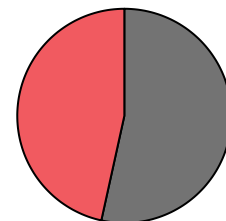
proportion of A
root nodes

biological_process



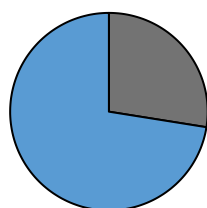
279 / 573

molecular_function

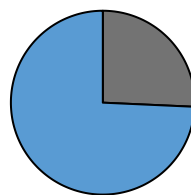


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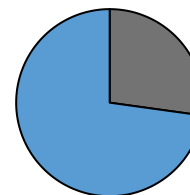
proportion of A



108 / 149



78 / 105



83 / 114

GO:0003008

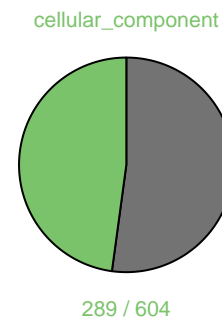
GO:0050877

GO:0001655

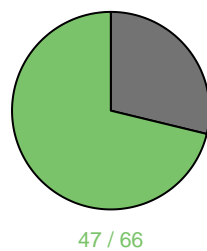
p(A) in node



proportion of A
root nodes

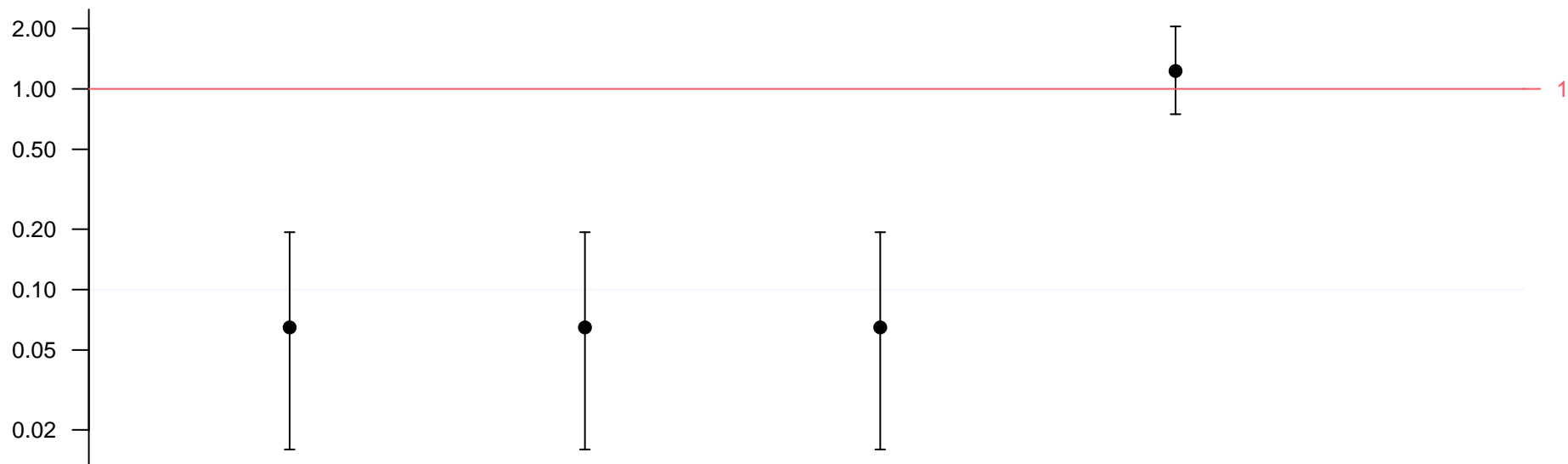


proportion of A

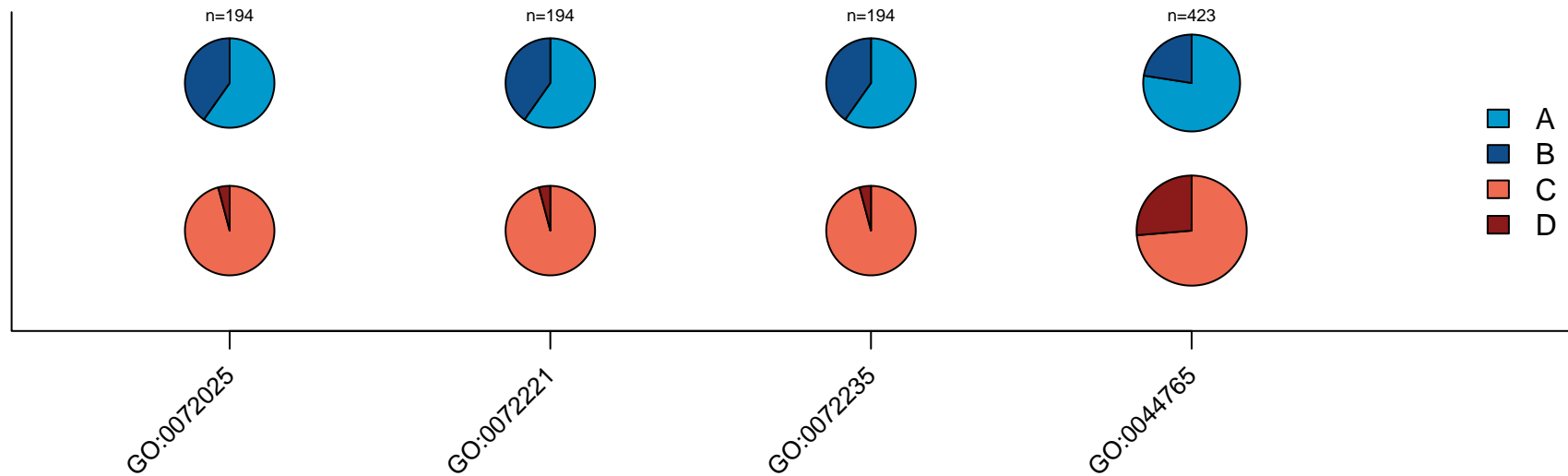


GO:0034702

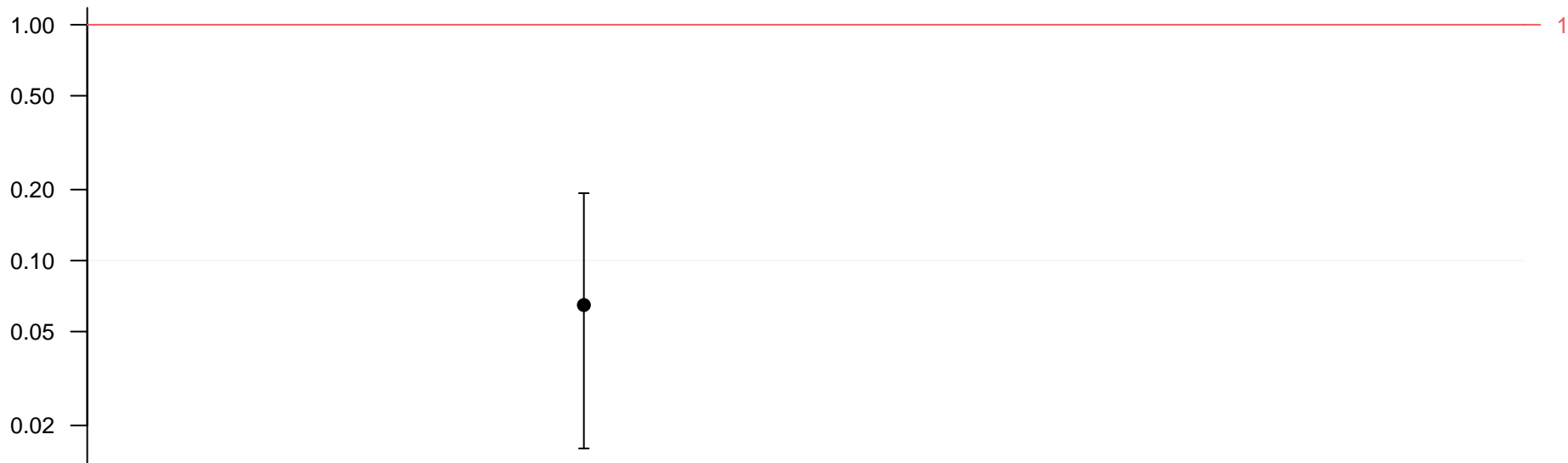
odds ratio (A/B) / (C/D)



annotated genes



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



annotated genes

