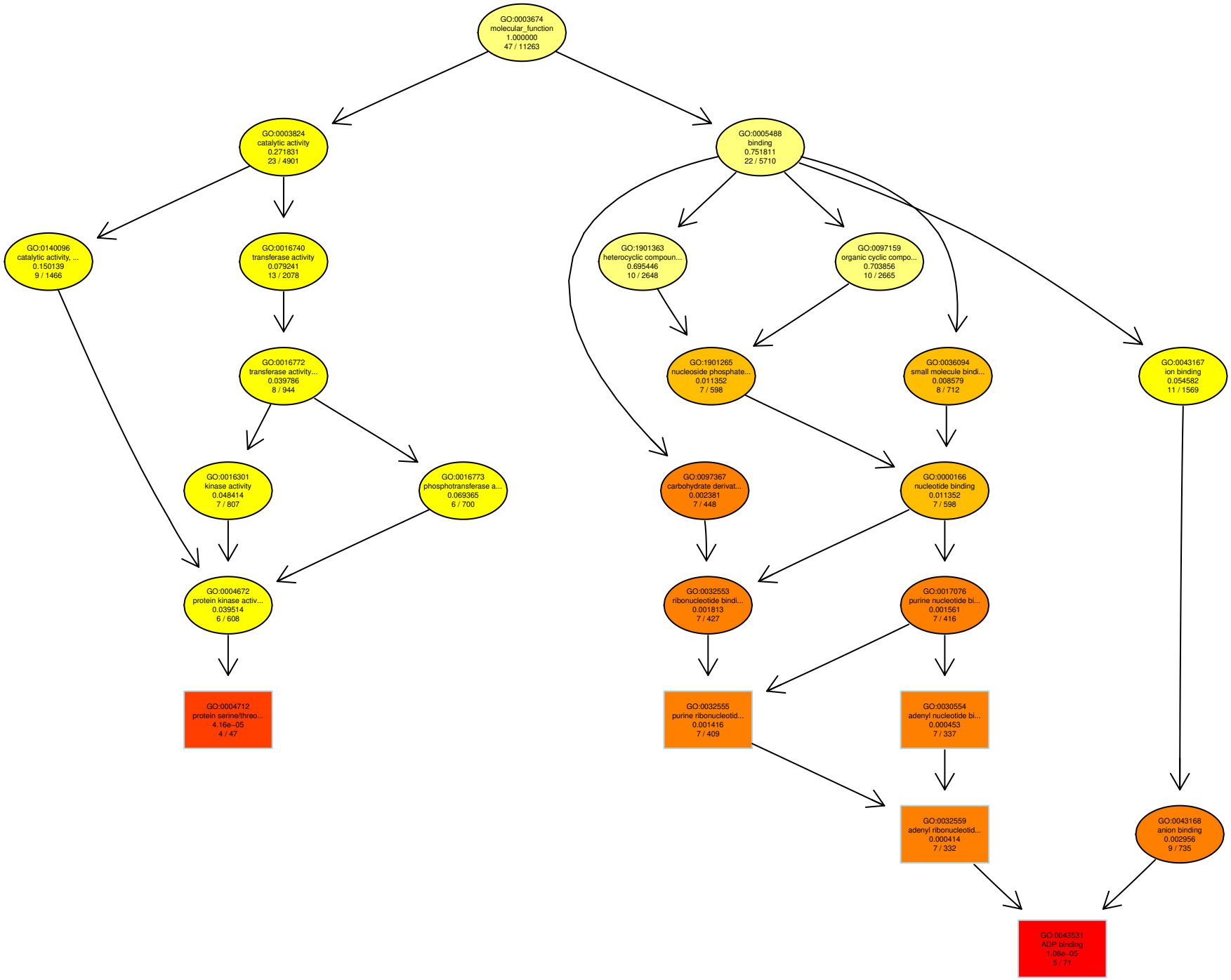
GO:0003674
molecular_function
1.000000
47 / 11263
GO:0003824
catalytic activity
0.271831
23 / 4901
GO:0005488
binding
0.751811
22 / 5710
GO:0140096
catalytic activity, ...
0.150139
9 / 1466
GO:0016740
transferase activity
0.079241
13 / 2078
GO:1901363
heterocyclic compoun...
0.695446
10 / 2648
GO:0097159
organic cyclic compo...
0.703856
10 / 2665
GO:0016772
transferase activity...
0.039786
8 / 944
GO:1901265
nucleoside phosphate...
0.011352
7 / 598
GO:0036094
small molecule bindi...
0.008579
8 / 712
GO:0043167
ion binding
0.054582
11 / 1569
GO:0016301
kinase activity
0.048414
7 / 807
GO:0016773
phosphotransferase a...
0.069365
6 / 700
GO:0097367
carbohydrate derivat...
0.002381
7 / 448
GO:0000166
nucleotide binding
0.011352
7 / 598
GO:0004672
protein kinase activ...
0.039514
6 / 608
GO:0032553
ribonucleotide bindi...
0.001813
7 / 427
GO:0017076
purine nucleotide bi...
0.001561
7 / 416
GO:0004712
protein serine/threo...
4.16e−05
4 / 47
GO:0032555
purine ribonucleotid...
0.001416
7 / 409
GO:0030554
adenyl nucleotide bi...
0.000453
7 / 337
GO:0032559
adenyl ribonucleotid...
0.000414
7 / 332
GO:0043168
anion binding
0.002956
9 / 735
GO:0043531
ADP binding
1.08e−05
5 / 71