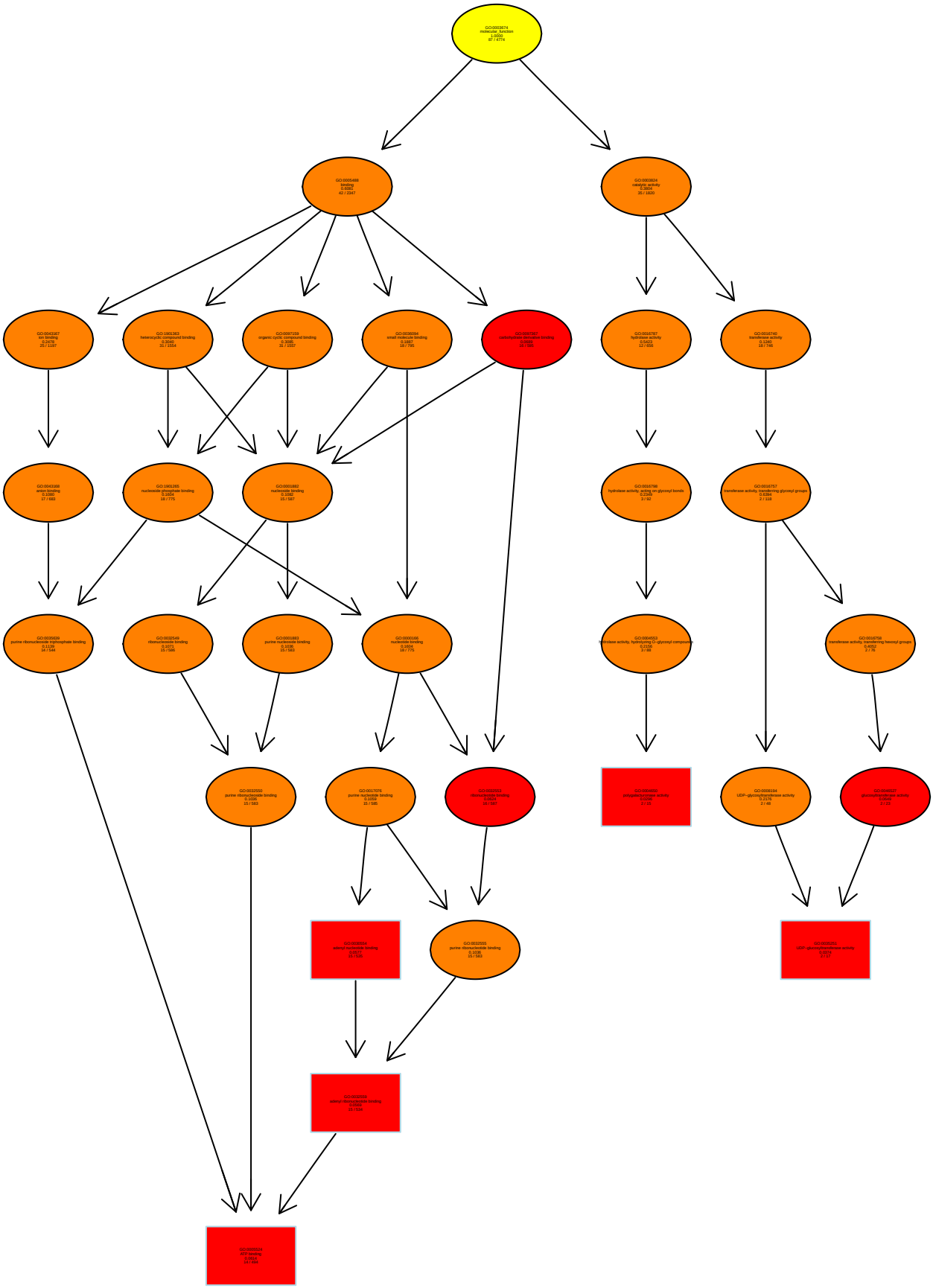
GO:0003674
molecular_function
1.0000
87 / 4774
GO:0005488
binding
0.6897
42 / 2347
GO:0003824
catalytic activity
0.3804
35 / 1820
GO:0043167
ion binding
0.1980
25 / 1197
GO:1901363
heterocyclic compound binding
0.3723
21 / 1004
GO:0097159
organic cyclic compound binding
0.3723
21 / 1007
GO:0036094
small molecule binding
0.3298
18 / 796
GO:0097367
carbohydrate derivative binding
0.2989
17 / 722
GO:0016787
hydrolase activity
0.3723
21 / 956
GO:0016740
transferase activity
0.3298
18 / 765
GO:0043168
anion binding
0.1980
17 / 683
GO:1901265
nucleoside phosphate binding
0.1604
18 / 775
GO:0001882
nucleoside binding
0.1582
15 / 587
GO:0016798
hydrolase activity, acting on glycosyl bonds
0.2349
3 / 92
GO:0016757
transferase activity, transferring glycosyl groups
0.6394
2 / 118
GO:0032559
purine ribonucleoside triphosphate binding
0.1129
14 / 544
GO:0032549
ribonucleoside binding
0.1071
15 / 586
GO:0001883
purine nucleoside binding
0.1036
15 / 583
GO:0000166
nucleotide binding
0.1604
18 / 775
GO:0004553
hydrolase activity, hydrolyzing O-glycosyl compounds
0.2155
3 / 88
GO:0016758
transferase activity, transferring hexosyl groups
0.4652
2 / 76
GO:0032550
purine ribonucleoside binding
0.1036
15 / 583
GO:0017076
purine nucleotide binding
0.1036
15 / 586
GO:0032553
ribonucleotide binding
0.1029
14 / 587
GO:0004650
polygalacturonase activity
0.0000
2 / 53
GO:0008194
UDP-glycosyltransferase activity
0.0000
2 / 48
GO:0046527
glucosyltransferase activity
0.0000
2 / 22
GO:0003954
adenyl nucleotide binding
0.0577
15 / 526
GO:0032555
purine ribonucleotide binding
0.1036
15 / 583
GO:0005351
UDP-glucose/galactose transferase activity
0.0274
2 / 17
GO:0032559
adenyl ribonucleotide binding
0.0569
15 / 524
GO:0005524
ATP binding
0.1013
14 / 584