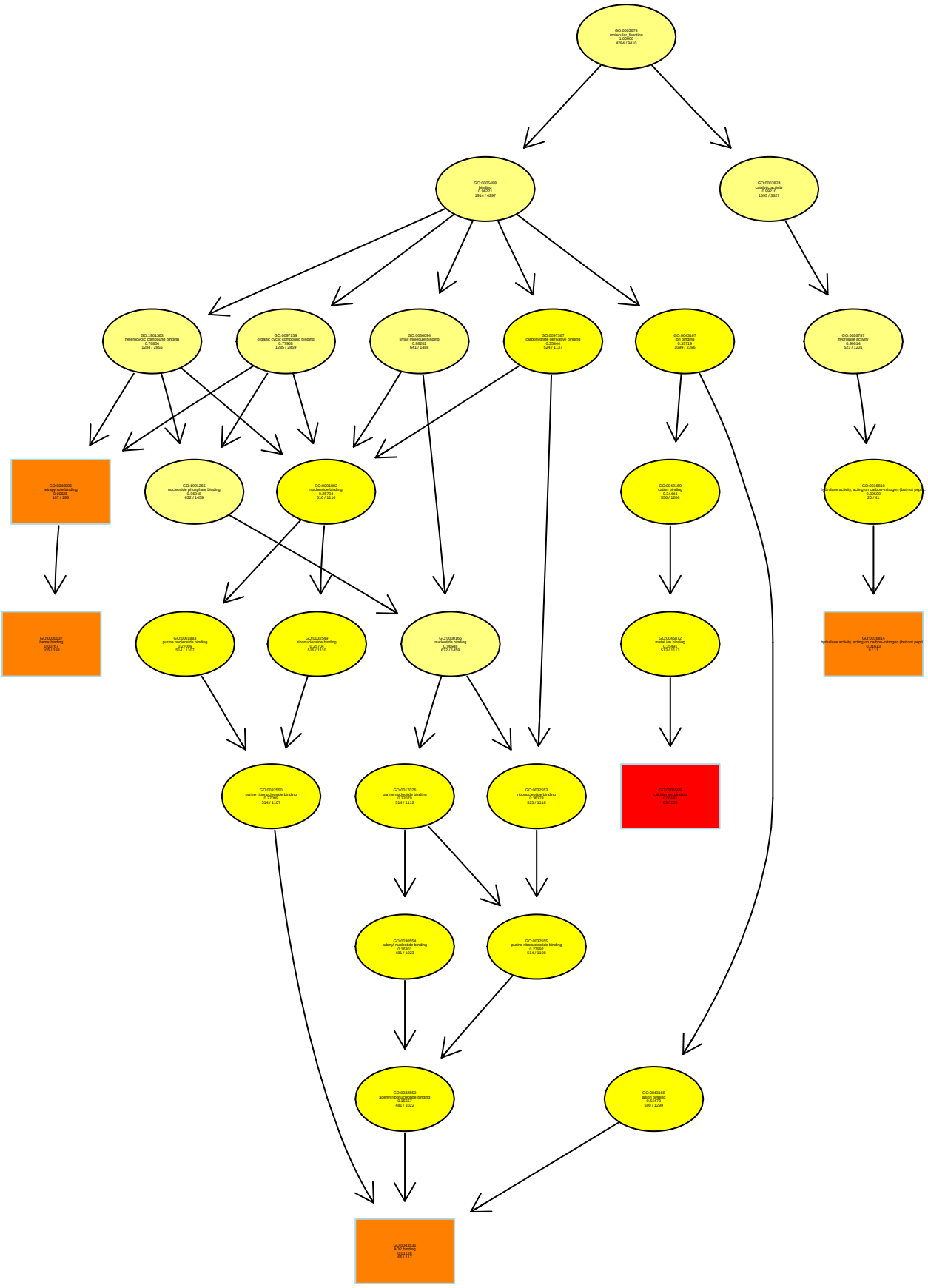
GO:0003674
molecular_function
1.00000
4268 / 9410
GO:0005488
binding
0.96223
1914 / 4297
GO:0003824
catalytic activity
0.95127
1595 / 3627
GO:1901363
heterocyclic compound binding
0.76804
1284 / 2859
GO:0097159
organic cyclic compound binding
0.77968
1285 / 2859
GO:0036094
small molecule binding
0.98283
641 / 1488
GO:0097367
carbohydrate derivative binding
0.35446
524 / 1137
GO:0043167
ion binding
0.57156
1099 / 2396
GO:0016787
hydrolase activity
0.98014
523 / 1231
GO:0046906
tetrapyrrole binding
0.00825
107 / 166
GO:1901265
nucleoside phosphate binding
0.98948
632 / 1459
GO:0001882
nucleoside binding
0.25704
516 / 1110
GO:0043169
cation binding
0.34444
556 / 1208
GO:0016810
hydrolase activity, acting on carbon-nitrogen (but not pep...
0.39059
20 / 41
GO:0020037
heme binding
0.00767
105 / 193
GO:0001883
purine nucleoside binding
0.27009
514 / 1107
GO:0032549
ribonucleoside binding
0.25704
516 / 1110
GO:0000166
nucleotide binding
0.98948
632 / 1459
GO:0046872
metal ion binding
0.25491
513 / 1113
GO:0016814
hydrolase activity, acting on carbon-nitrogen (but not pept...
0.01613
8 / 11
GO:0032550
purine ribonucleoside binding
0.27009
514 / 1107
GO:0017076
purine nucleotide binding
0.27992
514 / 1112
GO:0032553
ribonucleotide binding
0.25704
516 / 1110
GO:0005509
calcium ion binding
0.00108
2 / 185
GO:0030554
adenyl nucleotide binding
0.16557
491 / 1022
GO:0032555
purine ribonucleotide binding
0.27992
514 / 1108
GO:0032559
adenyl ribonucleotide binding
0.16557
491 / 1022
GO:0043168
anion binding
0.54479
590 / 1299
GO:0043531
ADP binding
0.01128
62 / 127