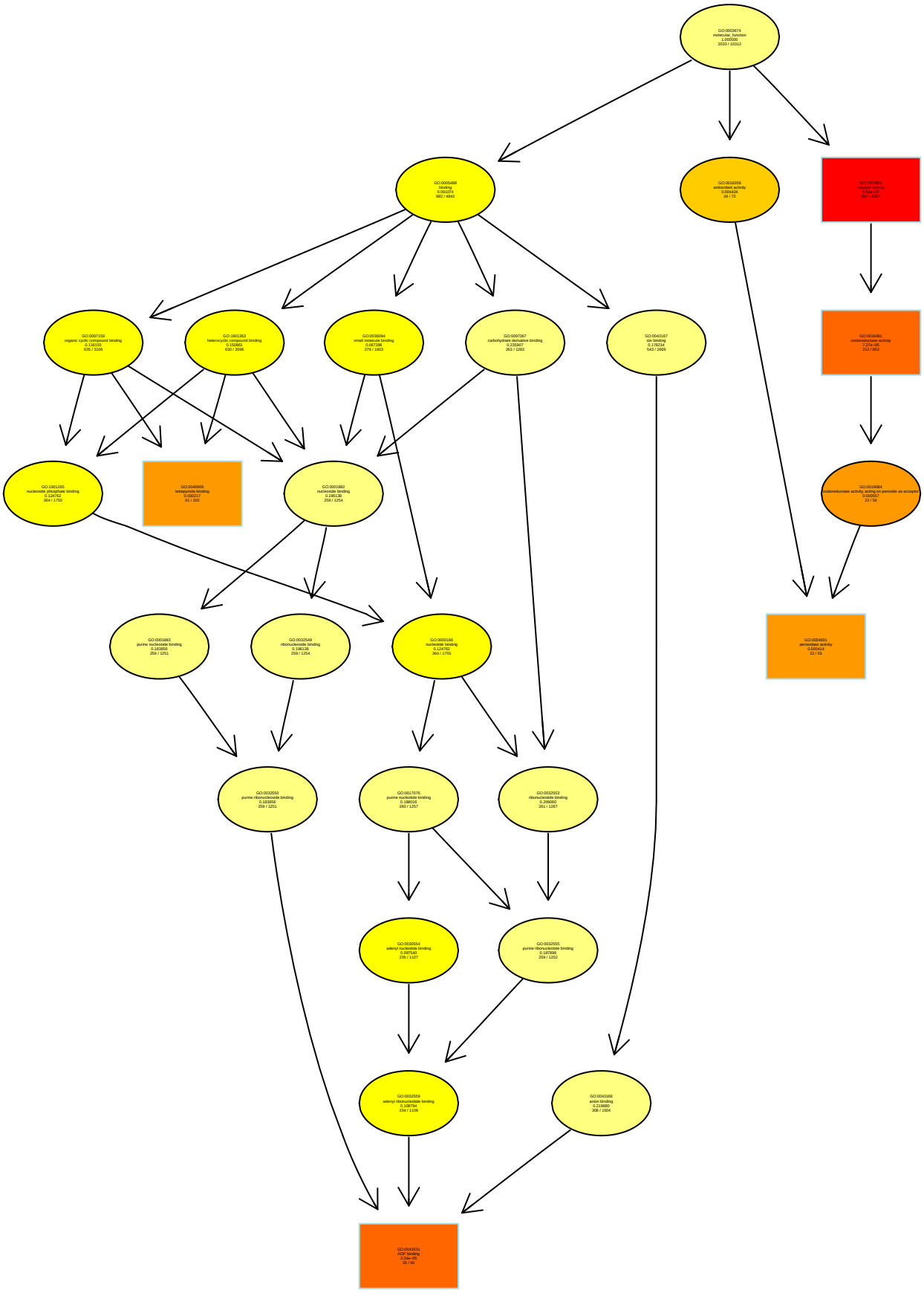
GO:0003674
molecular_function
1.000000
2033 / 10312
GO:0005488
binding
0.091074
982 / 4842
GO:0016209
antioxidant activity
0.004428
24 / 72
GO:0003824
catalytic activity
5.54e-07
947 / 4307
GO:0097159
organic cyclic compound binding
0.116155
635 / 3166
GO:1901363
heterocyclic compound binding
0.116093
630 / 3096
GO:0036094
small molecule binding
0.087288
379 / 1853
GO:0097367
carbohydrate derivative binding
0.235607
263 / 1283
GO:0043167
ion binding
0.178218
543 / 2669
GO:0016491
oxidoreductase activity
7.27e-05
212 / 653
GO:1901265
nucleoside phosphate binding
0.124762
364 / 1755
GO:0046906
tetrapyrrole binding
0.002217
61 / 252
GO:0001882
nucleoside binding
0.196138
259 / 1254
GO:0016684
oxidoreductase activity, acting on peroxide as acceptor
0.000557
22 / 56
GO:0001883
purine nucleoside binding
0.193816
259 / 1251
GO:0032549
ribonucleoside binding
0.196138
259 / 1254
GO:0000166
nucleotide binding
0.124762
364 / 1755
GO:0004601
peroxidase activity
0.000414
22 / 55
GO:0032550
purine ribonucleoside binding
0.193816
259 / 1251
GO:0017076
purine nucleotide binding
0.188516
260 / 1257
GO:0032553
ribonucleotide binding
0.200002
261 / 1267
GO:0030554
adenyl nucleotide binding
0.097540
235 / 1107
GO:0032555
purine ribonucleotide binding
0.187998
259 / 1252
GO:0032559
adenyl ribonucleotide binding
0.108794
234 / 1106
GO:0043168
anion binding
0.219882
208 / 1504
GO:0043531
ADP binding
3.34e-05
35 / 92