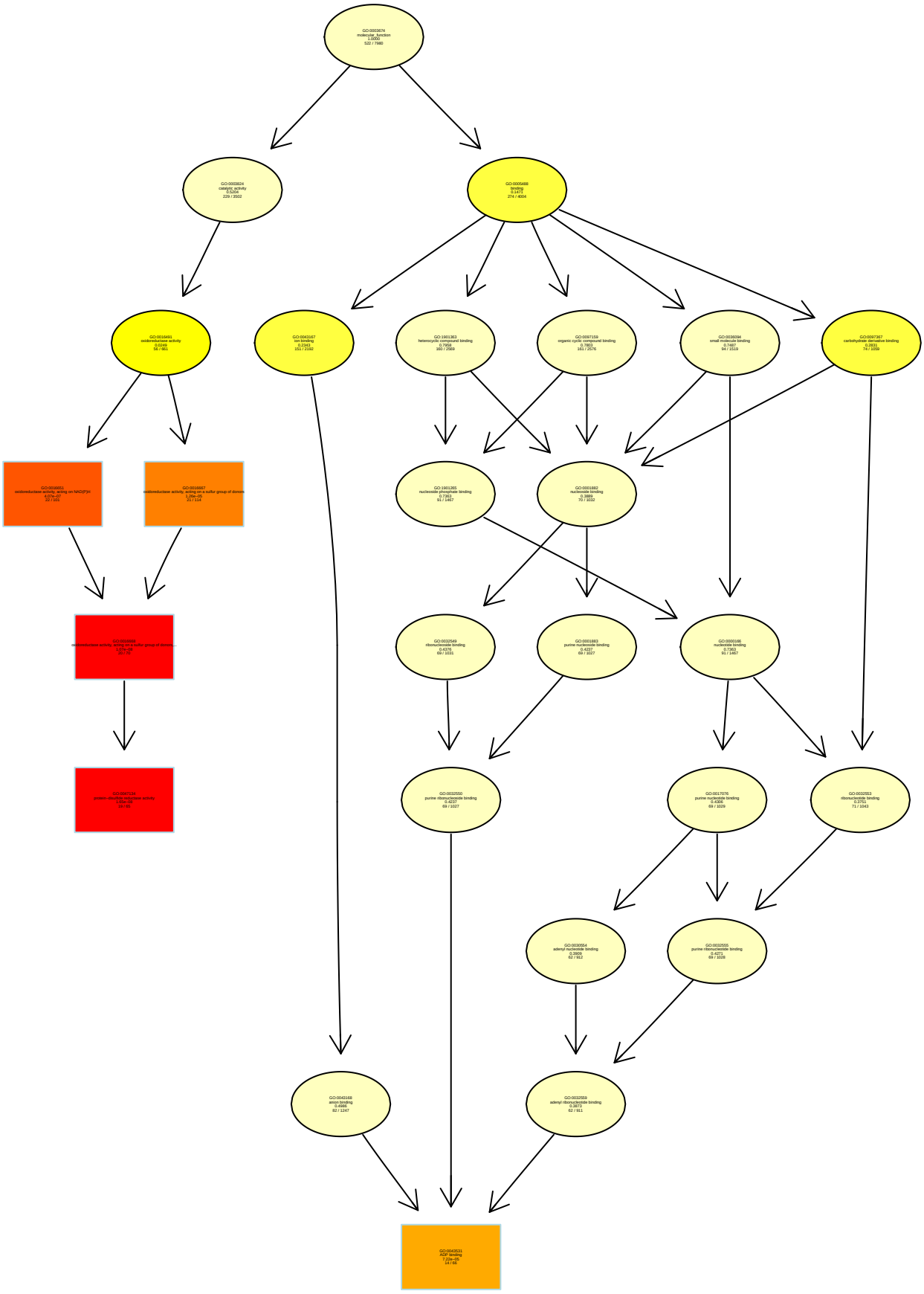
GO:0003674
molecular_function
1.0000
522 / 7980
GO:0003824
catalytic activity
0.5268
229 / 3502
GO:0005488
binding
0.5472
274 / 4004
GO:0016491
oxidoreductase activity
0.5248
56 / 861
GO:0043167
ion binding
0.5161
153 / 2392
GO:1901363
heterocyclic compound binding
0.7958
160 / 2589
GO:0097159
organic cyclic compound binding
0.7923
161 / 2576
GO:0036094
small molecule binding
0.7687
94 / 1519
GO:0097367
carbohydrate derivative binding
0.8931
74 / 1259
GO:0016651
oxidoreductase activity, acting on NADH/NH
4.07e-07
22 / 101
GO:0016667
oxidoreductase activity, acting on a sulfur group of donors
1.26e-05
21 / 114
GO:1901265
nucleoside phosphate binding
0.7363
91 / 1467
GO:0001882
nucleoside binding
0.3889
70 / 1032
GO:0016668
oxidoreductase activity, acting on a sulfur group of donors,...
1.07e-05
20 / 70
GO:0032549
ribonucleoside binding
0.4276
69 / 1031
GO:0001883
purine nucleoside binding
0.4237
69 / 1027
GO:0000166
nucleotide binding
0.7363
91 / 1467
GO:0047134
protein-disulfide reductase activity
1.88e-05
17 / 63
GO:0032550
purine ribonucleoside binding
0.4237
69 / 1027
GO:0017076
purine nucleotide binding
0.4276
69 / 1029
GO:0032553
ribonucleotide binding
0.4884
71 / 1043
GO:0030554
adenyl nucleotide binding
0.3865
62 / 913
GO:0032555
purine ribonucleotide binding
0.4271
69 / 1028
GO:0043168
anion binding
0.4886
82 / 1247
GO:0032559
adenyl ribonucleotide binding
0.3873
62 / 911
GO:0043531
ADP binding
7.23e-05
14 / 68