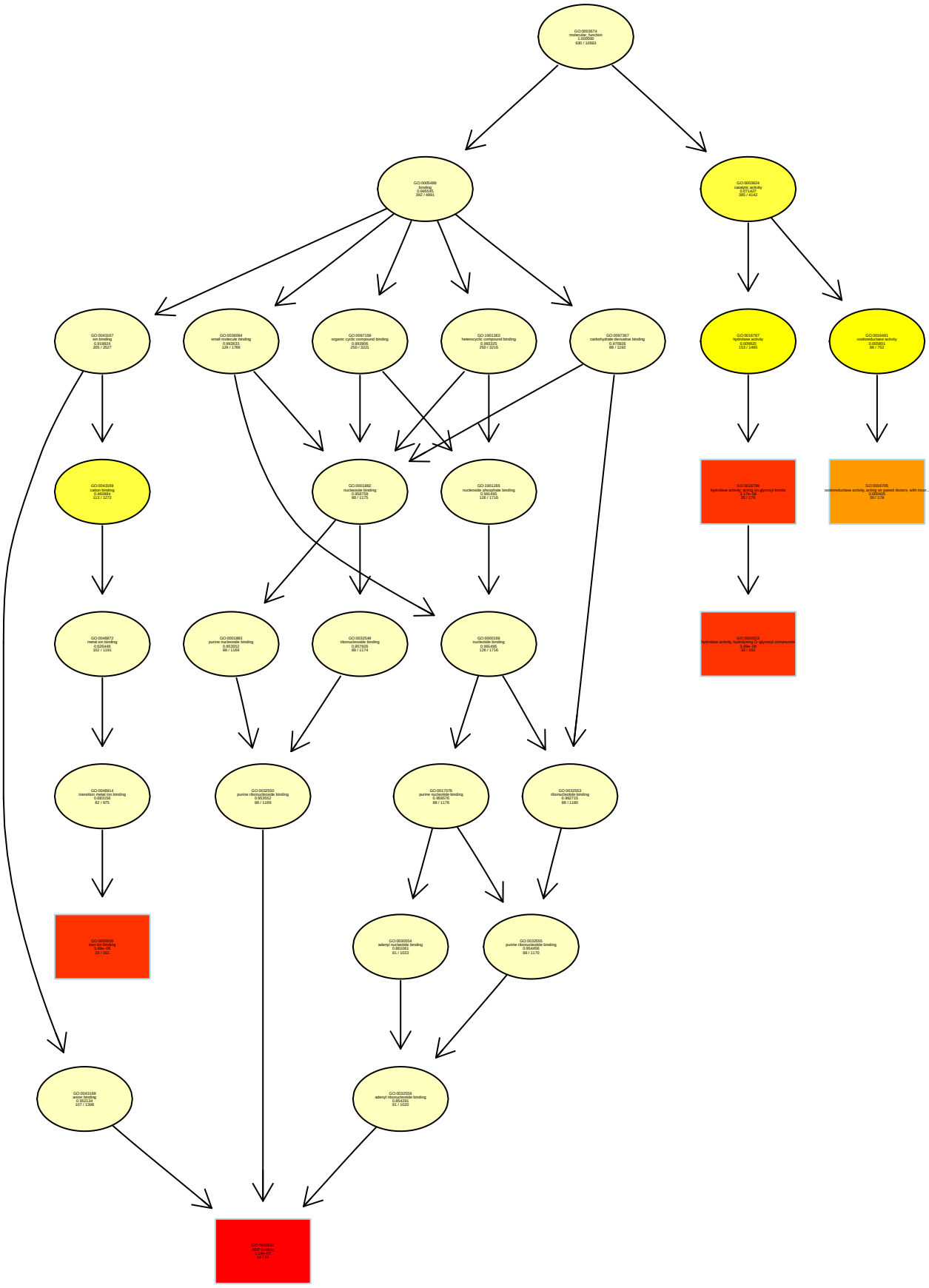
GO:0003674
molecular_function
1.000000
930 / 10593
GO:0005488
binding
0.940526
262 / 4891
GO:0003824
catalytic activity
0.931325
265 / 4142
GO:0043167
ion binding
0.918824
205 / 2527
GO:0036094
small molecule binding
0.909033
139 / 1766
GO:0097159
organic cyclic compound binding
0.900806
250 / 3221
GO:1901363
heterocyclic compound binding
0.893325
250 / 3216
GO:0097367
carbohydrate derivative binding
0.870826
88 / 1192
GO:0016787
hydrolase activity
0.906825
153 / 1985
GO:0016491
oxidoreductase activity
0.856801
66 / 752
GO:0043169
cation binding
0.890884
113 / 1472
GO:0001882
nucleoside binding
0.868758
88 / 1175
GO:1901265
nucleoside phosphate binding
0.866495
126 / 1716
GO:0016798
hydrolase activity, acting on glycosyl bonds
3.17e-36
35 / 179
GO:0016705
oxidoreductase activity, acting on paired donors, with incor...
0.036405
30 / 179
GO:0046872
metal ion binding
0.826448
322 / 1191
GO:0001883
purine nucleoside binding
0.953552
88 / 1169
GO:0032549
ribonucleoside binding
0.957526
88 / 1174
GO:0000166
nucleotide binding
0.866495
126 / 1716
GO:0004553
hydrolase activity, hydrolyzing O-glycosyl compounds
3.39e-32
33 / 162
GO:0046914
transition metal ion binding
0.826448
82 / 975
GO:0032550
purine ribonucleoside binding
0.953552
88 / 1169
GO:0017076
purine nucleotide binding
0.866495
88 / 1176
GO:0032553
ribonucleotide binding
0.866495
88 / 1180
GO:0005506
iron ion binding
0.826448
32 / 162
GO:0030554
adenyl nucleotide binding
0.881281
81 / 1023
GO:0032555
purine ribonucleotide binding
0.866495
88 / 1170
GO:0043168
anion binding
0.905234
107 / 1398
GO:0032559
adenyl ribonucleotide binding
0.854281
81 / 1020
GO:0005524
ATP binding
0.854281
77 / 1020