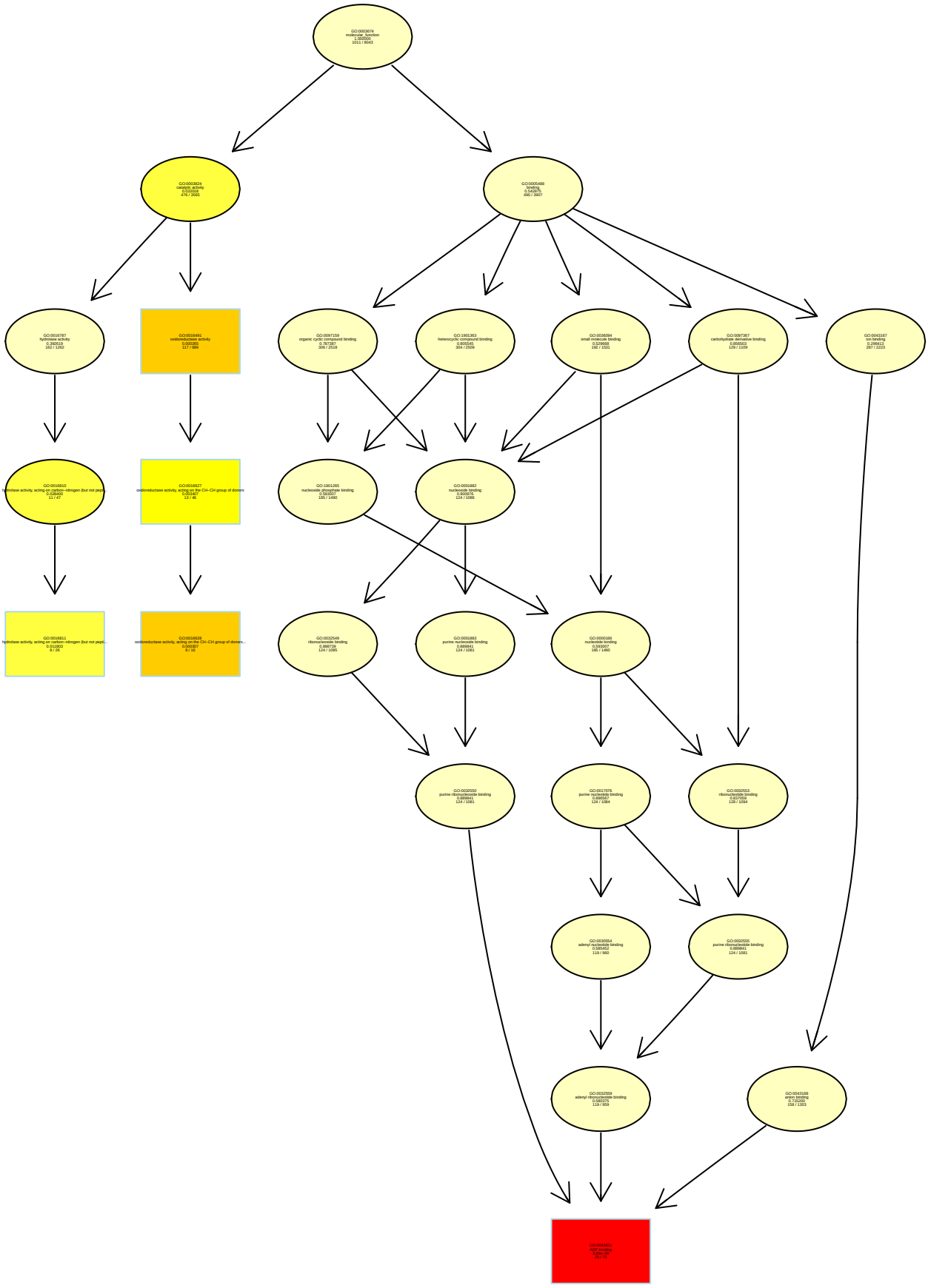
GO:0003674
molecular_function
1.000000
1011 / 8043
GO:0003824
catalytic activity
0.470588
476 / 3565
GO:0005488
binding
0.542857
490 / 3907
GO:0016787
hydrolase activity
0.360015
167 / 1262
GO:0016491
oxidoreductase activity
0.050265
117 / 889
GO:0097159
organic cyclic compound binding
0.787387
306 / 2516
GO:1901363
heterocyclic compound binding
0.805545
304 / 2509
GO:0036094
small molecule binding
0.529658
192 / 1521
GO:0097367
carbohydrate derivative binding
0.565501
129 / 1109
GO:0043167
ion binding
0.296413
287 / 2223
GO:0016810
hydrolase activity, acting on carbon–nitrogen (but not pept...
0.038400
31 / 47
GO:0016627
oxidoreductase activity, acting on the CH–CH group of donors
0.003407
3 / 46
GO:1901265
nucleoside phosphate binding
0.593007
185 / 1490
GO:0000166
nucleotide binding
0.800876
124 / 1085
GO:0016811
hydrolase activity, acting on carbon–nitrogen (but not pept...
0.011803
8 / 26
GO:0016628
oxidoreductase activity, acting on the CH–CH group of donors...
0.000307
8 / 16
GO:0032549
ribonucleoside binding
0.984739
124 / 1085
GO:0001883
purine nucleoside binding
0.888841
124 / 1081
GO:0000166
nucleotide binding
0.593007
185 / 1490
GO:0032550
purine ribonucleoside binding
0.888841
124 / 1081
GO:0017076
purine nucleotide binding
0.845188
128 / 1084
GO:0032553
ribonucleotide binding
0.845188
128 / 1084
GO:0030554
adenyl nucleotide binding
0.595453
119 / 960
GO:0032555
purine ribonucleotide binding
0.888841
124 / 1081
GO:0032559
adenyl ribonucleotide binding
0.595275
119 / 959
GO:0043169
anion binding
0.745200
158 / 1303
GO:0043531
ADP binding
0.199000
20 / 79