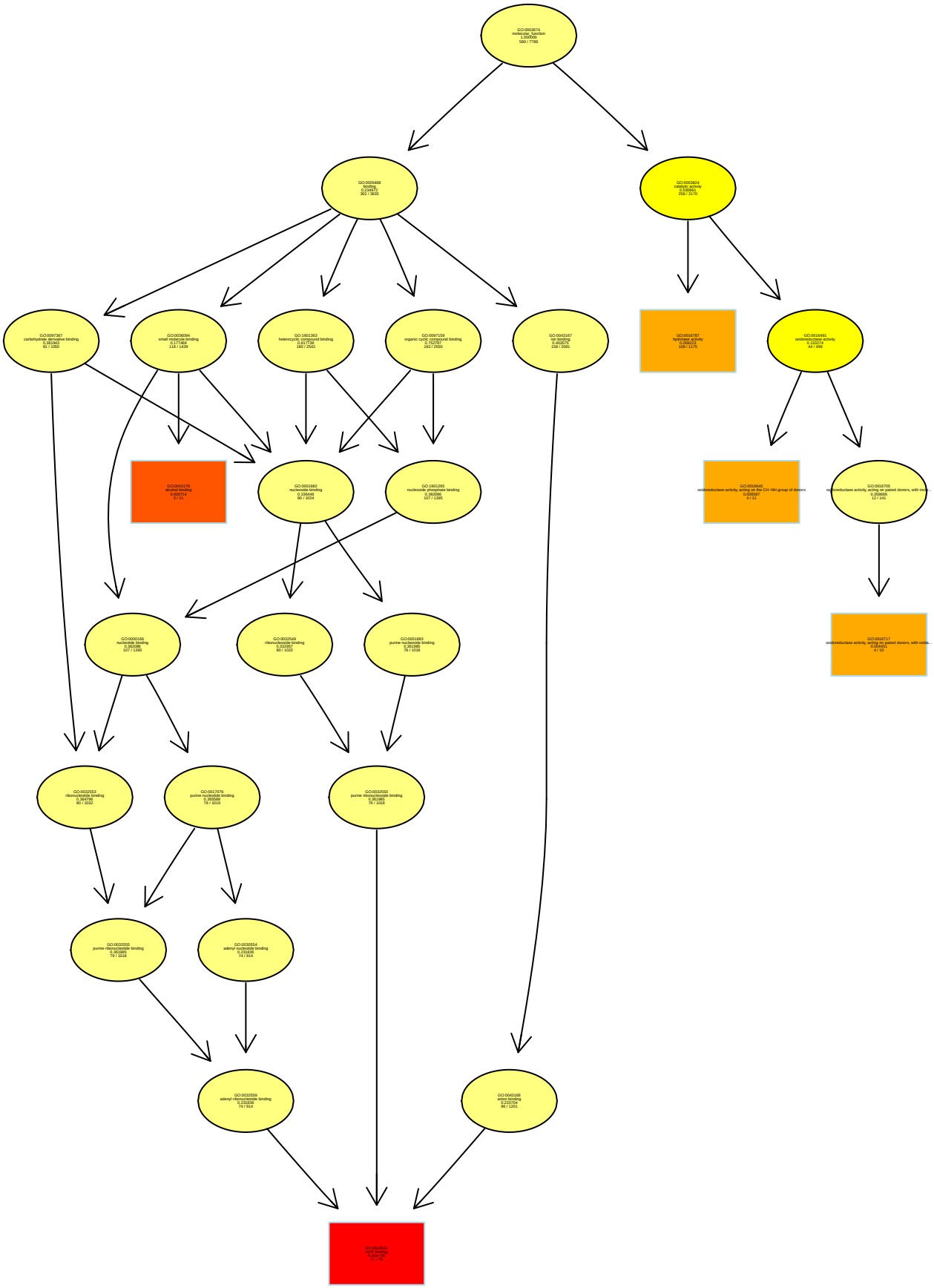
GO:0003674
molecular_function
1.000000
580 / 7798
GO:0005488
binding
0.216973
322 / 3635
GO:0003824
catalytic activity
0.030021
258 / 3170
GO:0097367
carbohydrate derivative binding
0.364796
82 / 1350
GO:0036094
small molecule binding
0.239836
116 / 1419
GO:1901363
heterocyclic compound binding
0.216973
180 / 2532
GO:0097159
organic cyclic compound binding
0.216973
183 / 2550
GO:0043167
ion binding
0.233754
158 / 2081
GO:0016787
hydrolase activity
0.030021
158 / 1175
GO:0016491
oxidoreductase activity
0.047453
44 / 499
GO:0043178
alcohol binding
0.000714
6 / 12
GO:0001882
nucleoside binding
0.339848
86 / 1024
GO:1901265
nucleoside phosphate binding
0.363598
107 / 1395
GO:0016645
oxidoreductase activity, acting on the CH–NH group of donors
0.000597
4 / 13
GO:0016705
oxidoreductase activity, acting on paired donors, with incor...
0.200565
12 / 141
GO:0000166
nucleotide binding
0.363598
107 / 1395
GO:0032549
ribonucleoside binding
0.323907
80 / 1023
GO:0001883
purine nucleoside binding
0.361985
79 / 1018
GO:0016717
oxidoreductase activity, acting on paired donors, with oxyg...
0.000453
4 / 10
GO:0032553
ribonucleotide binding
0.364796
82 / 1032
GO:0017076
purine nucleotide binding
0.365508
79 / 1019
GO:0032550
purine ribonucleoside binding
0.361985
79 / 1018
GO:0032555
purine ribonucleotide binding
0.361985
79 / 1018
GO:0030554
adenyl nucleotide binding
0.239836
74 / 914
GO:0032559
adenyl ribonucleotide binding
0.239836
74 / 914
GO:0043168
anion binding
0.233754
96 / 1201
GO:0043531
ADP binding
3.43e-05
77 / 79