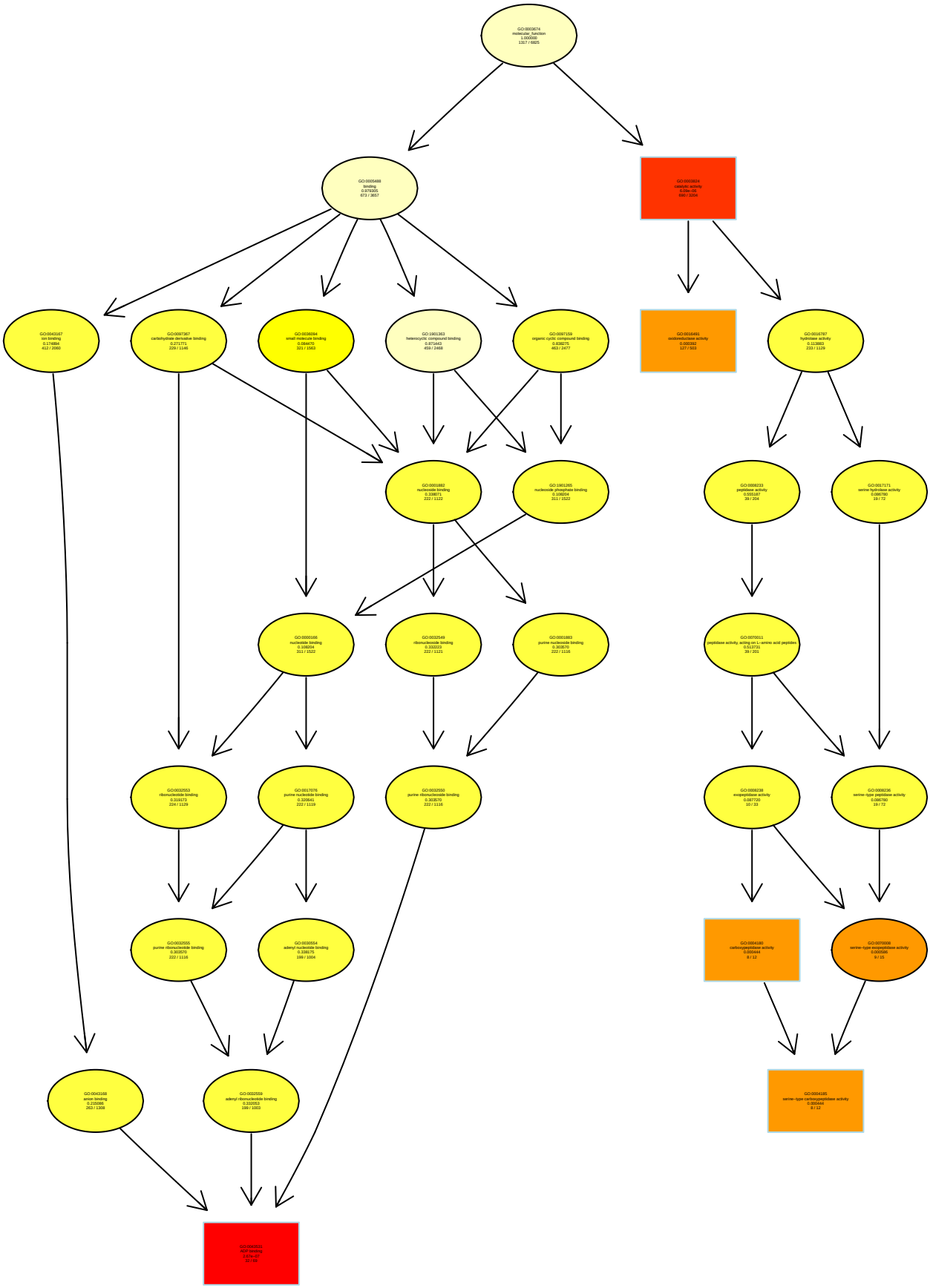
GO:0003674
molecular_function
1.000000
1357 / 6825
GO:0005488
binding
0.970305
673 / 3657
GO:0003824
catalytic activity
0.08e-02
690 / 3204
GO:0043167
ion binding
0.174894
412 / 2060
GO:0097367
carbohydrate derivative binding
0.071771
229 / 1146
GO:0036094
small molecule binding
0.064405
321 / 1563
GO:1901363
heterocyclic compound binding
0.874443
459 / 2458
GO:0097159
organic cyclic compound binding
0.838275
463 / 2477
GO:0016491
oxidoreductase activity
0.000382
127 / 593
GO:0016787
hydrolase activity
0.113803
233 / 1129
GO:0001882
nucleoside binding
0.338071
222 / 1122
GO:1901265
nucleoside phosphate binding
0.108204
311 / 1522
GO:0008233
peptidase activity
0.555187
39 / 204
GO:0017171
serine hydrolase activity
0.086780
19 / 72
GO:0000166
nucleotide binding
0.108204
311 / 1522
GO:0032549
ribonucleoside binding
0.332223
222 / 1121
GO:0001883
purine nucleoside binding
0.303670
222 / 1116
GO:0070011
peptidase activity, acting on L-amino acid peptides
0.513731
39 / 201
GO:0032553
ribonucleotide binding
0.319173
224 / 1129
GO:0017076
purine nucleotide binding
0.320641
222 / 1119
GO:0032555
purine ribonucleotide binding
0.303670
222 / 1116
GO:0008238
exopeptidase activity
0.087720
10 / 23
GO:0008236
serine-type peptidase activity
0.086780
19 / 72
GO:0032555
purine ribonucleotide binding
0.303670
222 / 1116
GO:0030554
adenyl nucleotide binding
0.338175
199 / 1004
GO:0004180
carboxypeptidase activity
0.000444
8 / 12
GO:0070008
serine-type exopeptidase activity
0.000586
8 / 15
GO:0043168
anion binding
0.218286
263 / 1308
GO:0032559
adenyl ribonucleotide binding
0.322053
199 / 1003
GO:0004185
serine-type carboxypeptidase activity
0.000444
8 / 12
GO:0043531
ADP binding
2.67e-07
22 / 88