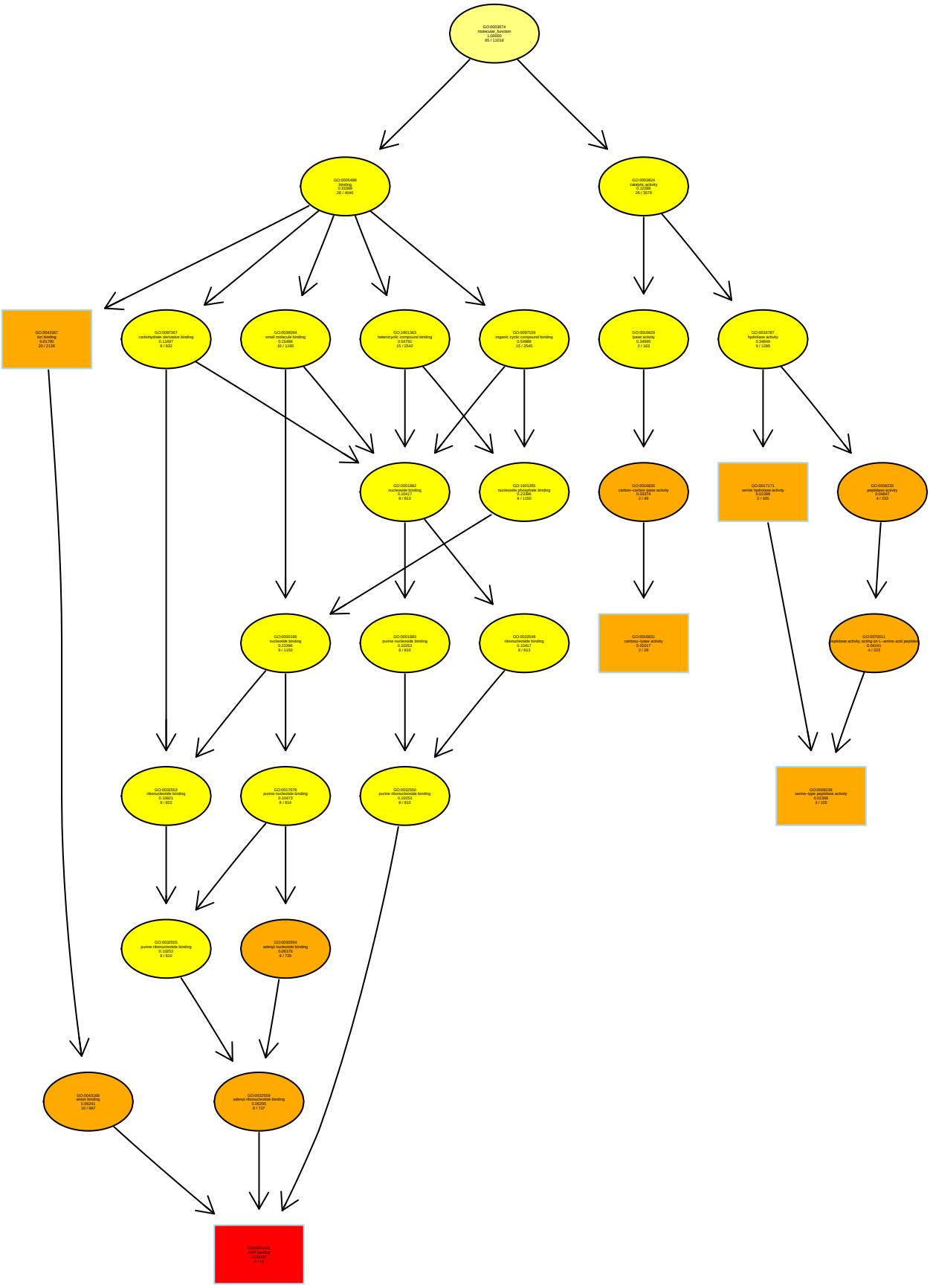
GO:0003824
catalytic activity
0.12288
26 / 3579
GO:0043167
ion binding
0.01795
26 / 3126
GO:0097367
carbohydrate derivative binding
0.11497
8 / 832
GO:0036094
small molecule binding
0.15488
10 / 1183
GO:1901363
heterocyclic compound binding
0.54791
15 / 2542
GO:0097159
organic cyclic compound binding
0.54968
15 / 2545
GO:0016829
lyase activity
0.24995
2 / 163
GO:0016787
hydrolase activity
0.34849
9 / 1289
GO:0001882
nucleoside binding
0.10417
8 / 813
GO:1901265
nucleoside phosphate binding
0.23356
9 / 1150
GO:0016830
carbon-carbon lyase activity
0.01374
2 / 49
GO:0017171
serine hydrolase activity
0.02098
3 / 105
GO:0008233
peptidase activity
0.04847
4 / 223
GO:0000166
nucleotide binding
0.23356
9 / 1150
GO:0001883
purine nucleoside binding
0.10253
8 / 810
GO:0032549
ribonucleoside binding
0.10417
8 / 813
GO:0016831
carbon-lyase activity
0.01017
2 / 26
GO:0070011
peptidase activity, acting on L-amino acid peptides
0.04241
4 / 223
GO:0032553
ribonucleotide binding
0.10921
8 / 822
GO:0017076
purine nucleotide binding
0.10473
8 / 814
GO:0032550
purine ribonucleoside binding
0.10253
8 / 810
GO:0008236
serine-type peptidase activity
0.02098
3 / 105
GO:0032555
purine ribonucleotide binding
0.10253
8 / 810
GO:0030554
adenyl nucleotide binding
0.08376
8 / 709
GO:0043168
anion binding
0.06241
10 / 987
GO:0032559
adenyl ribonucleotide binding
0.08256
8 / 727
GO:0043531
ADP binding
0.06256
8 / 727