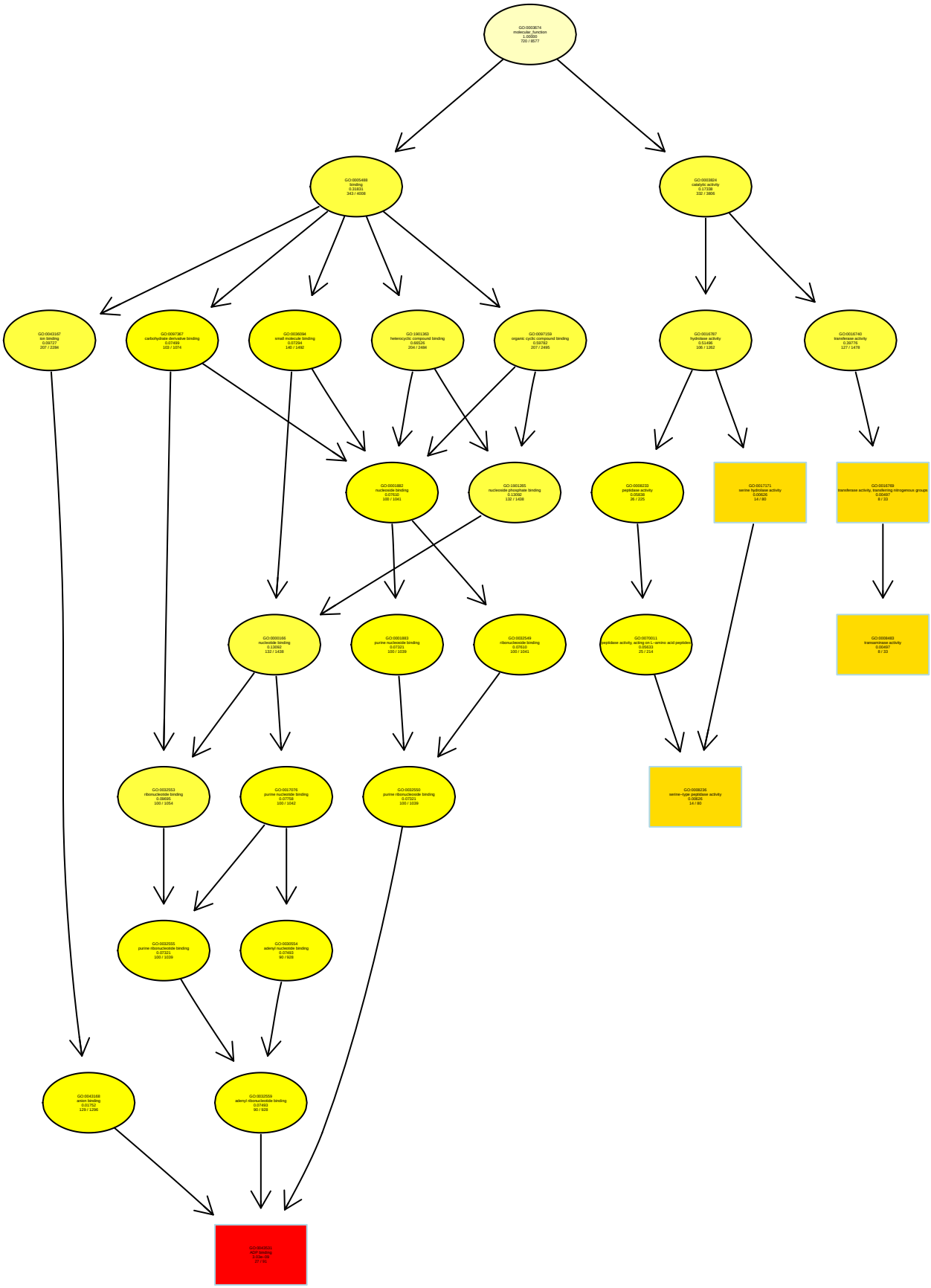
GO:0003674
molecular_function
1.00000
720 / 8577
GO:0005488
binding
0.31831
343 / 4008
GO:0003824
catalytic activity
0.17338
332 / 3806
GO:0043167
ion binding
0.08727
207 / 2284
GO:0097367
carbohydrate derivative binding
0.07489
153 / 1074
GO:0036094
small molecule binding
0.07246
148 / 1462
GO:1901363
heterocyclic compound binding
0.06538
204 / 2484
GO:0097159
organic cyclic compound binding
0.06782
207 / 2495
GO:0016787
hydrolase activity
0.51498
106 / 1262
GO:0016740
transferase activity
0.39775
127 / 1478
GO:0001882
nucleoside binding
0.07610
100 / 1041
GO:1901265
nucleoside phosphate binding
0.13092
132 / 1438
GO:0008233
peptidase activity
0.05639
26 / 225
GO:0017171
serine hydrolase activity
0.00626
14 / 80
GO:0016769
transferase activity, transferring nitrogenous groups
0.00497
8 / 33
GO:0001166
nucleotide binding
0.13092
132 / 1438
GO:0001883
purine nucleoside binding
0.07321
100 / 1039
GO:0032549
ribonucleoside binding
0.07610
100 / 1041
GO:0070011
peptidase activity, acting on L-amino acid peptides
0.05633
25 / 214
GO:0008468
transaminase activity
0.00497
8 / 33
GO:0032553
ribonucleotide binding
0.08695
100 / 1054
GO:0017076
purine nucleotide binding
0.07758
100 / 1042
GO:0032555
purine ribonucleotide binding
0.07321
100 / 1039
GO:0008236
serine-type peptidase activity
0.00626
14 / 80
GO:0032555
purine ribonucleotide binding
0.07321
100 / 1039
GO:0030554
adenyl nucleotide binding
0.07492
90 / 928
GO:0043168
anion binding
0.01752
129 / 1296
GO:0032559
adenyl ribonucleotide binding
0.07492
90 / 928
GO:0043531
ADP binding
3.03e-09
27 / 43