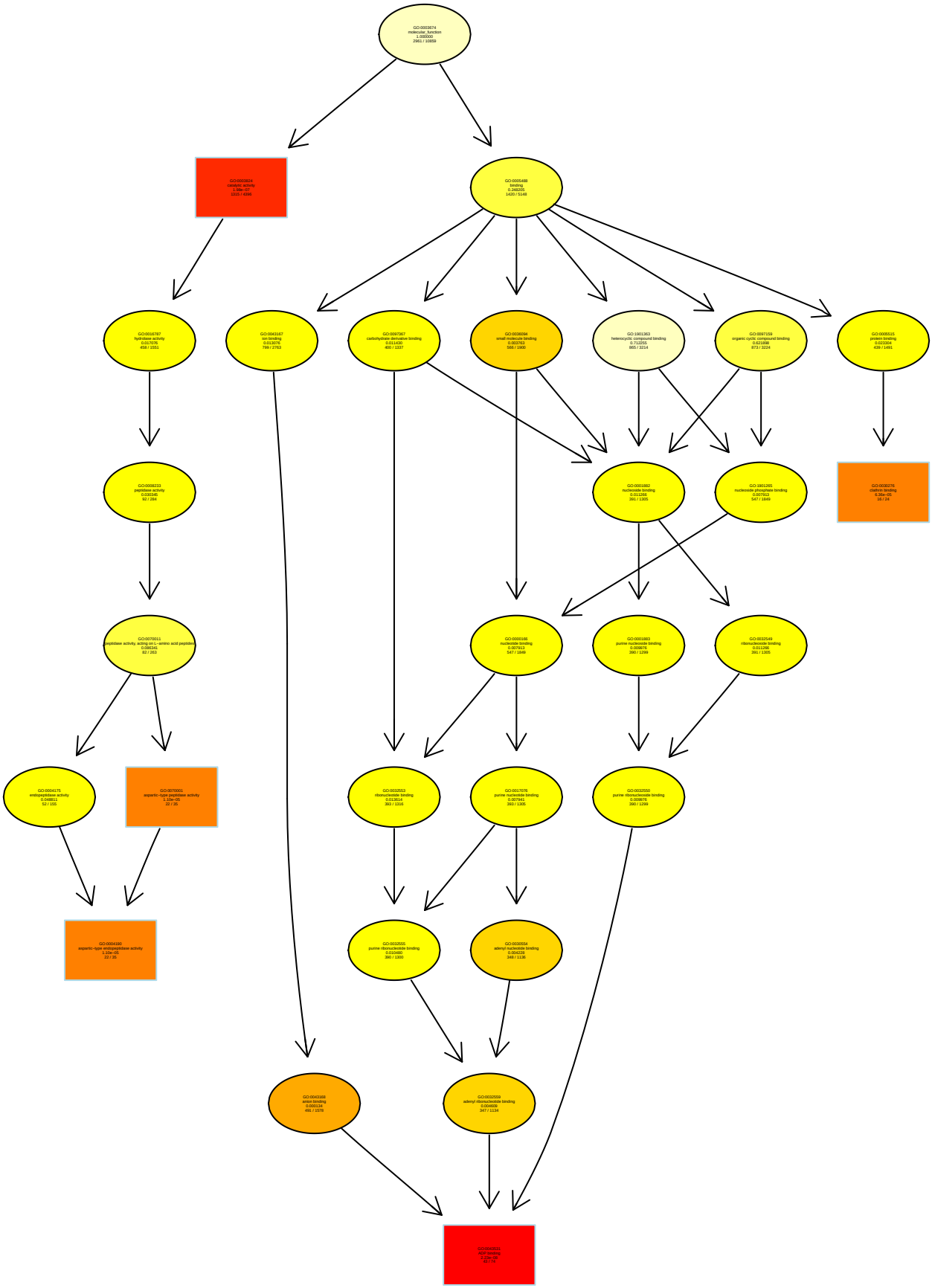
GO:0003674
molecular_function
1.000000
2961 / 10859
GO:0003824
catalytic activity
1.16e-07
1315 / 4396
GO:0005488
binding
0.248205
1420 / 5148
GO:0016787
hydrolase activity
0.017076
458 / 1551
GO:0043167
ion binding
0.012076
799 / 2763
GO:0097367
carbohydrate derivative binding
0.014450
400 / 1237
GO:0036094
small molecule binding
0.002763
966 / 1966
GO:1901363
heterocyclic compound binding
0.712255
865 / 3214
GO:0097159
organic cyclic compound binding
0.621084
873 / 3224
GO:0005515
protein binding
0.023054
439 / 1491
GO:0008233
peptidase activity
0.030345
92 / 284
GO:0001882
nucleoside binding
0.011266
391 / 1305
GO:1901265
nucleoside phosphate binding
0.007613
547 / 1849
GO:0003276
clathrin binding
6.36e-05
16 / 24
GO:0070011
peptidase activity, acting on L-amino acid peptides
0.086341
82 / 263
GO:0000166
nucleotide binding
0.007613
547 / 1849
GO:0001883
purine nucleoside binding
0.009876
290 / 1299
GO:0032549
ribonucleoside binding
0.011266
391 / 1305
GO:0004175
endopeptidase activity
0.048811
52 / 155
GO:0070001
aspartic-type peptidase activity
1.10e-05
22 / 35
GO:0032553
ribonucleotide binding
0.013614
393 / 1316
GO:0017076
purine nucleotide binding
0.007941
393 / 1305
GO:0032550
purine ribonucleoside binding
0.009876
290 / 1299
GO:0004190
aspartic-type endopeptidase activity
1.10e-05
22 / 35
GO:0032555
purine ribonucleotide binding
0.010460
390 / 1300
GO:0030554
adenyl nucleotide binding
0.004228
348 / 1136
GO:0043168
anion binding
0.000134
491 / 1578
GO:0032559
adenyl ribonucleotide binding
0.004008
347 / 1134
GO:0043531
ADP binding
2.23e-05
42 / 74