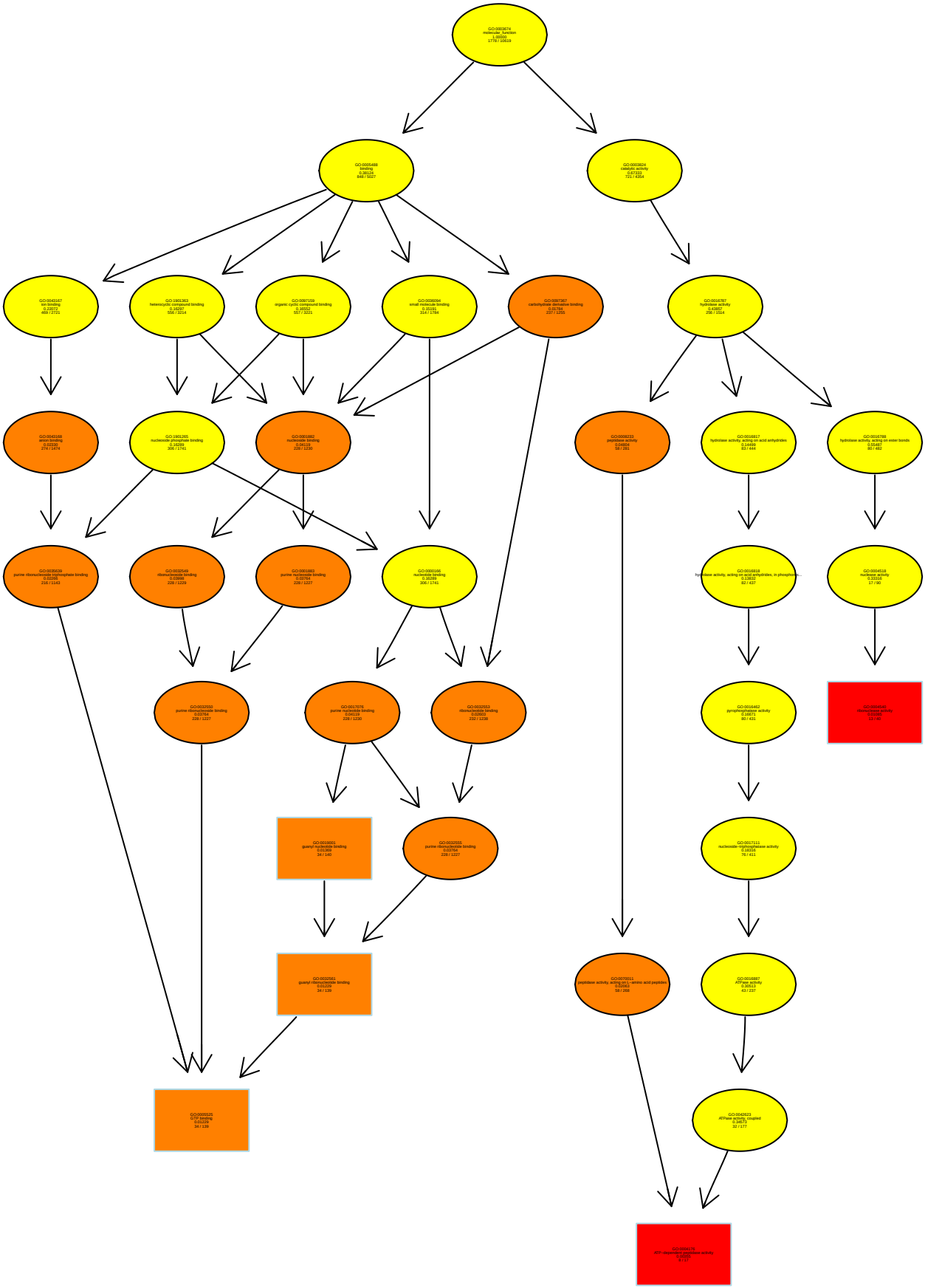
GO:0003674
molecular_function
1.00000
1770 / 10619
GO:0005488
binding
0.38115
848 / 5027
GO:0003824
catalytic activity
0.67123
721 / 4354
GO:0043167
ion binding
0.23311
468 / 2721
GO:1901363
heterocyclic compound binding
0.18307
958 / 5214
GO:0097159
organic cyclic compound binding
0.18312
957 / 5211
GO:0036094
small molecule binding
0.25151
316 / 1794
GO:0097367
carbohydrate derivative binding
0.01784
237 / 1355
GO:0016787
hydrolase activity
0.24811
256 / 1514
GO:0043169
anion binding
0.22230
274 / 1474
GO:1901265
nucleoside phosphate binding
0.18284
306 / 1741
GO:0001882
nucleoside binding
0.04119
228 / 1230
GO:0008233
peptidase activity
0.04804
59 / 291
GO:0016817
hydrolase activity, acting on acid anhydrides
0.14439
82 / 444
GO:0016788
hydrolase activity, acting on ester bonds
0.55487
87 / 402
GO:0035639
purine ribonucleoside triphosphate binding
0.32266
216 / 1143
GO:0032549
ribonucleoside binding
0.03968
228 / 1229
GO:0001883
purine nucleoside binding
0.03754
228 / 1227
GO:0000166
nucleotide binding
0.18269
306 / 1741
GO:0016818
hydrolase activity, acting on acid anhydrides, in phosphorus-
0.13832
82 / 437
GO:0004518
nuclease activity
0.13316
17 / 90
GO:0032550
purine ribonucleoside binding
0.03764
228 / 1227
GO:0017076
purine nucleotide binding
0.04119
228 / 1230
GO:0032553
ribonucleotide binding
0.02603
222 / 1238
GO:0016462
pyrophosphatase activity
0.16671
80 / 431
GO:0004540
ribonuclease activity
0.01065
13 / 60
GO:0019001
guanyl nucleotide binding
0.01369
34 / 140
GO:0032555
purine ribonucleotide binding
0.03764
228 / 1227
GO:0017111
nucleoside-triphosphatase activity
0.18257
76 / 411
GO:0032561
guanyl ribonucleotide binding
0.01229
34 / 139
GO:0070011
peptidase activity, acting on L-amino acid peptides
0.01966
58 / 266
GO:0016887
ATPase activity
0.30537
63 / 237
GO:0005525
GTP binding
0.01229
34 / 139
GO:0042623
ATPase activity, coupled
0.34573
32 / 177
GO:0004175
ATP-dependent peptidase activity
0.00355
0 / 17