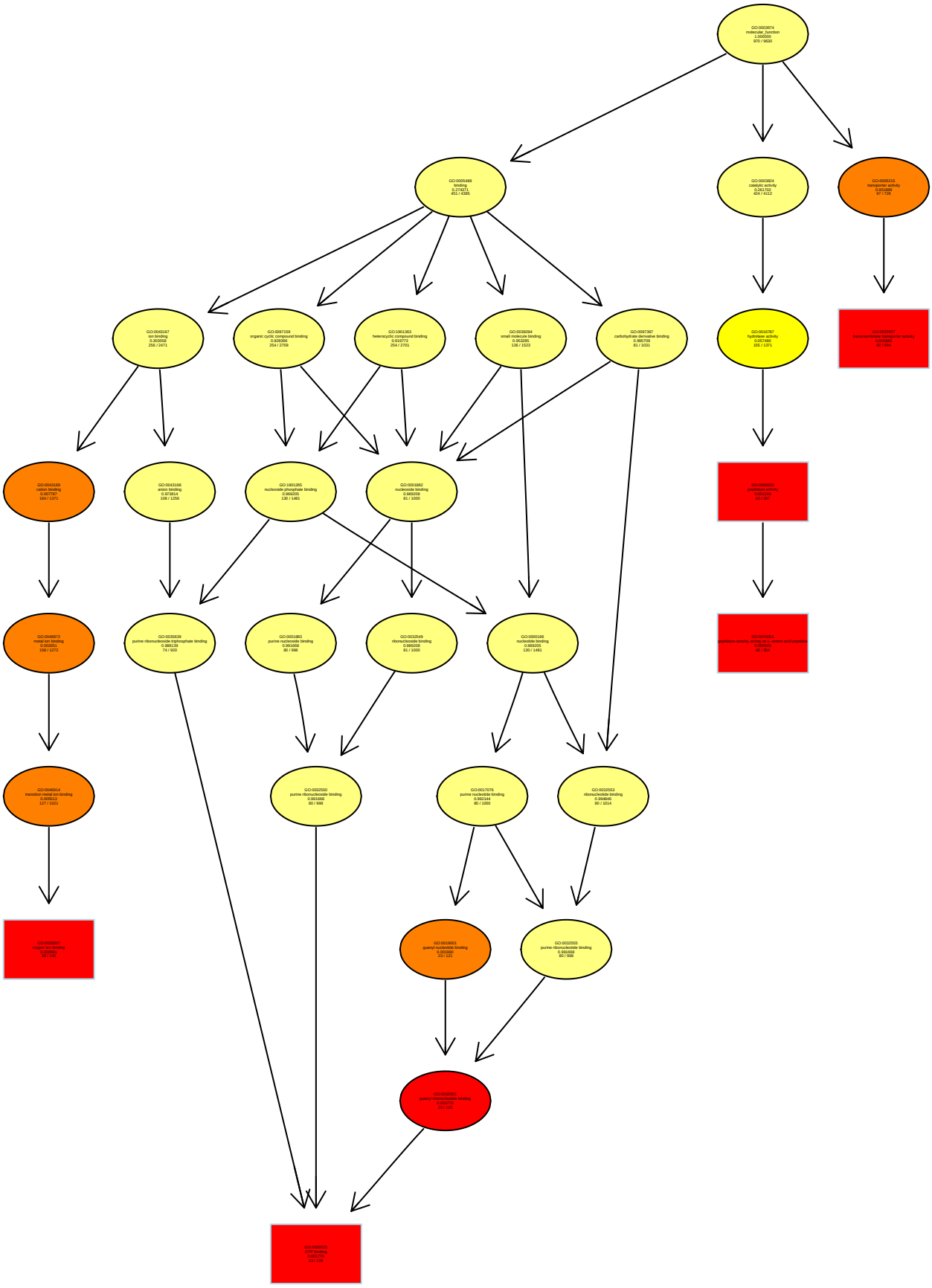
GO:0003674
molecular_function
1.000000
970 / 9630
GO:0005488
binding
0.274371
451 / 4365
GO:0003824
catalytic activity
0.261703
424 / 4112
GO:0005215
transporter activity
0.070588
97 / 726
GO:0043167
ion binding
0.163055
258 / 2471
GO:0097159
organic cyclic compound binding
0.928369
254 / 2709
GO:1901363
heterocyclic compound binding
0.919753
254 / 2701
GO:0036094
small molecule binding
0.902895
136 / 1523
GO:0097367
carbohydrate derivative binding
0.965708
81 / 1031
GO:0016787
hydrolase activity
0.263264
155 / 1371
GO:0022857
transmembrane transporter activity
0.021962
10 / 594
GO:0043169
cation binding
0.007787
156 / 1271
GO:0043168
anion binding
0.973814
108 / 1256
GO:1901265
nucleoside phosphate binding
0.969205
130 / 1461
GO:0001882
nucleoside binding
0.969208
81 / 1000
GO:0008233
peptidase activity
0.001244
43 / 357
GO:0046872
metal ion binding
0.002551
158 / 1272
GO:0035639
purine ribonucleoside triphosphate binding
0.989139
74 / 920
GO:0001883
purine nucleoside binding
0.991668
80 / 998
GO:0032549
ribonucleoside binding
0.989208
81 / 1000
GO:0000166
nucleotide binding
0.969205
130 / 1461
GO:0070011
peptidase activity, acting on L-amino acid peptides
0.000026
42 / 254
GO:0046914
transition metal ion binding
0.000551
127 / 1221
GO:0032550
purine ribonucleoside binding
0.991668
80 / 998
GO:0017076
purine nucleotide binding
0.992144
80 / 1000
GO:0032553
ribonucleotide binding
0.994846
80 / 1014
GO:0005507
copper ion binding
0.000591
29 / 145
GO:0019001
guanyl nucleotide binding
0.001968
23 / 121
GO:0032555
purine ribonucleotide binding
0.991668
80 / 998
GO:0032561
guanyl ribonucleotide binding
0.001776
23 / 120
GO:0005525
GTP binding
0.001776
23 / 120