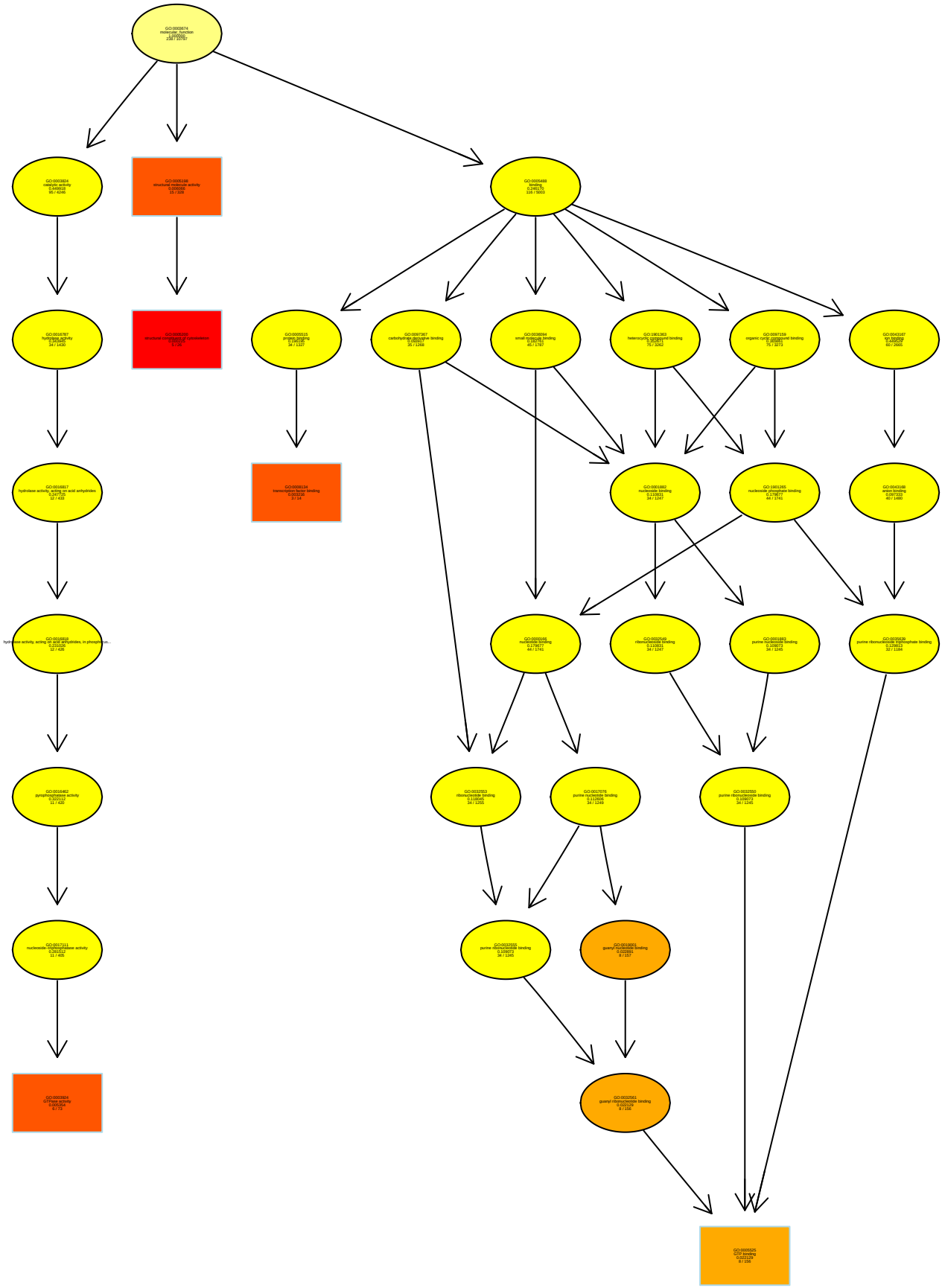
GO:0003824
catalytic activity
0.546819
95 / 4246
GO:0005198
structural molecule activity
0.008049
15 / 328
GO:0005488
binding
0.348175
116 / 5003
GO:0016787
hydrolase activity
0.247725
34 / 1430
GO:0005200
structural constituent of cytoskeleton
0.000204
7 / 226
GO:0005515
protein binding
0.179677
34 / 1327
GO:0097367
carbohydrate derivative binding
0.110831
35 / 1268
GO:0036094
small molecule binding
0.179677
60 / 1787
GO:1901363
heterocyclic compound binding
0.179677
76 / 2262
GO:0097159
organic cyclic compound binding
0.179677
76 / 2273
GO:0043167
ion binding
0.179677
60 / 2665
GO:0016817
hydrolase activity, acting on acid anhydrides
0.247725
12 / 433
GO:0008134
transcription factor binding
0.003236
3 / 14
GO:0001882
nucleoside binding
0.110831
34 / 1247
GO:1901265
nucleoside phosphate binding
0.179677
44 / 1741
GO:0043168
anion binding
0.097333
40 / 1460
GO:0016818
hydrolase activity, acting on acid anhydrides, in phosphorus-...
0.251026
12 / 435
GO:0000166
nucleotide binding
0.179677
44 / 1741
GO:0002549
ribonucleotide binding
0.110831
34 / 1247
GO:0001883
purine nucleoside binding
0.109072
34 / 1245
GO:0032559
purine ribonucleotide triphosphate binding
0.128873
32 / 1184
GO:0016462
pyrophosphatase activity
0.251026
11 / 420
GO:0032553
ribonucleotide binding
0.109072
34 / 1255
GO:0017076
purine nucleotide binding
0.109072
34 / 1249
GO:0032550
purine ribonucleoside binding
0.109072
34 / 1245
GO:0017111
nucleoside-triphosphatase activity
0.281012
11 / 405
GO:0032555
purine ribonucleotide binding
0.109072
34 / 1245
GO:0019001
guanyl nucleotide binding
0.032884
8 / 157
GO:0003924
GTPase activity
0.000204
6 / 73
GO:0032561
guanyl ribonucleotide binding
0.032122
8 / 156
GO:0005525
GTP binding
0.032122
8 / 154