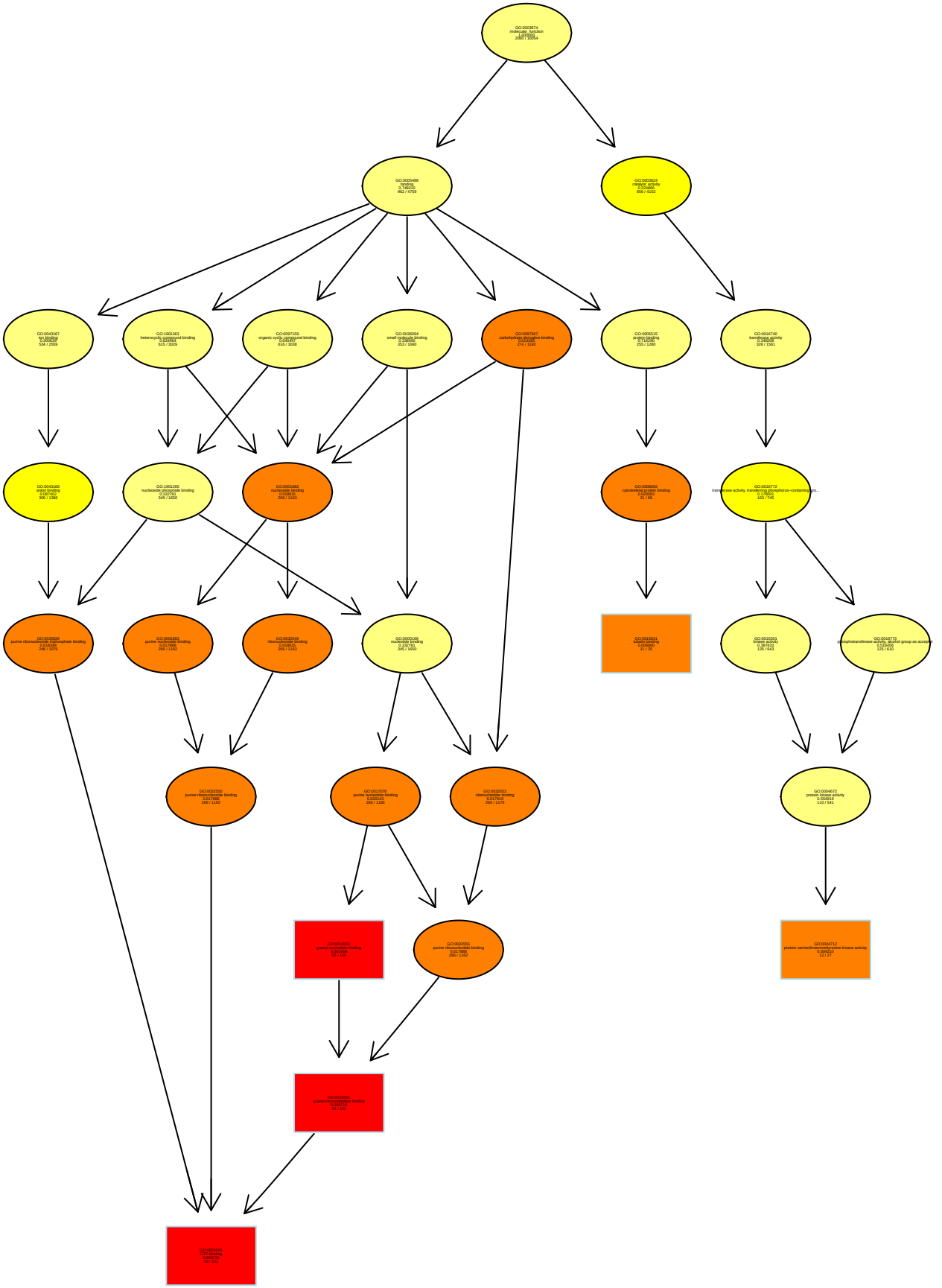
GO:0003674
molecular_function
1.000000
2667 / 39264
GO:0005488
binding
0.748255
962 / 4759
GO:0003824
catalytic activity
0.534868
856 / 4102
GO:0043167
ion binding
0.396897
538 / 2559
GO:1901363
heterocyclic compound binding
0.377358
619 / 3035
GO:0097159
organic cyclic compound binding
0.377358
619 / 3038
GO:0036094
small molecule binding
0.219512
353 / 1696
GO:0097367
carbohydrate derivative binding
0.165829
274 / 1192
GO:0005515
protein binding
0.159204
255 / 1286
GO:0016740
transferase activity
0.135135
326 / 1561
GO:0043168
anion binding
0.067402
308 / 1398
GO:1901265
nucleoside phosphate binding
0.332761
345 / 1650
GO:0032182
nucleoside binding
0.198455
266 / 1163
GO:0008092
cytoskeletal protein binding
0.030303
21 / 66
GO:0016772
transferase activity, transferring phosphorus-containing gro...
0.176501
143 / 745
GO:0032555
purine ribonucleotide binding
0.197183
249 / 1079
GO:0001883
purine nucleoside binding
0.197666
266 / 1162
GO:0032549
ribonucleoside binding
0.198455
266 / 1163
GO:0000166
nucleotide binding
0.332761
345 / 1650
GO:0015631
tubulin binding
0.066667
11 / 25
GO:0016301
kinase activity
0.187419
135 / 643
GO:0016773
phosphotransferase activity, alcohol group as acceptor
0.151456
125 / 610
GO:0032550
purine ribonucleoside binding
0.197666
266 / 1162
GO:0017076
purine nucleotide binding
0.329341
266 / 1165
GO:0032553
ribonucleoside binding
0.197183
269 / 1176
GO:0004672
protein kinase activity
0.164491
115 / 541
GO:0019001
guanyl nucleotide binding
0.035294
43 / 124
GO:0032555
purine ribonucleotide binding
0.017889
266 / 1162
GO:0004712
protein serine/threonine/tyrosine kinase activity
0.026230
12 / 27
GO:0032561
guanyl ribonucleotide binding
0.035715
43 / 120
GO:0005525
GTP binding
0.035294
43 / 124