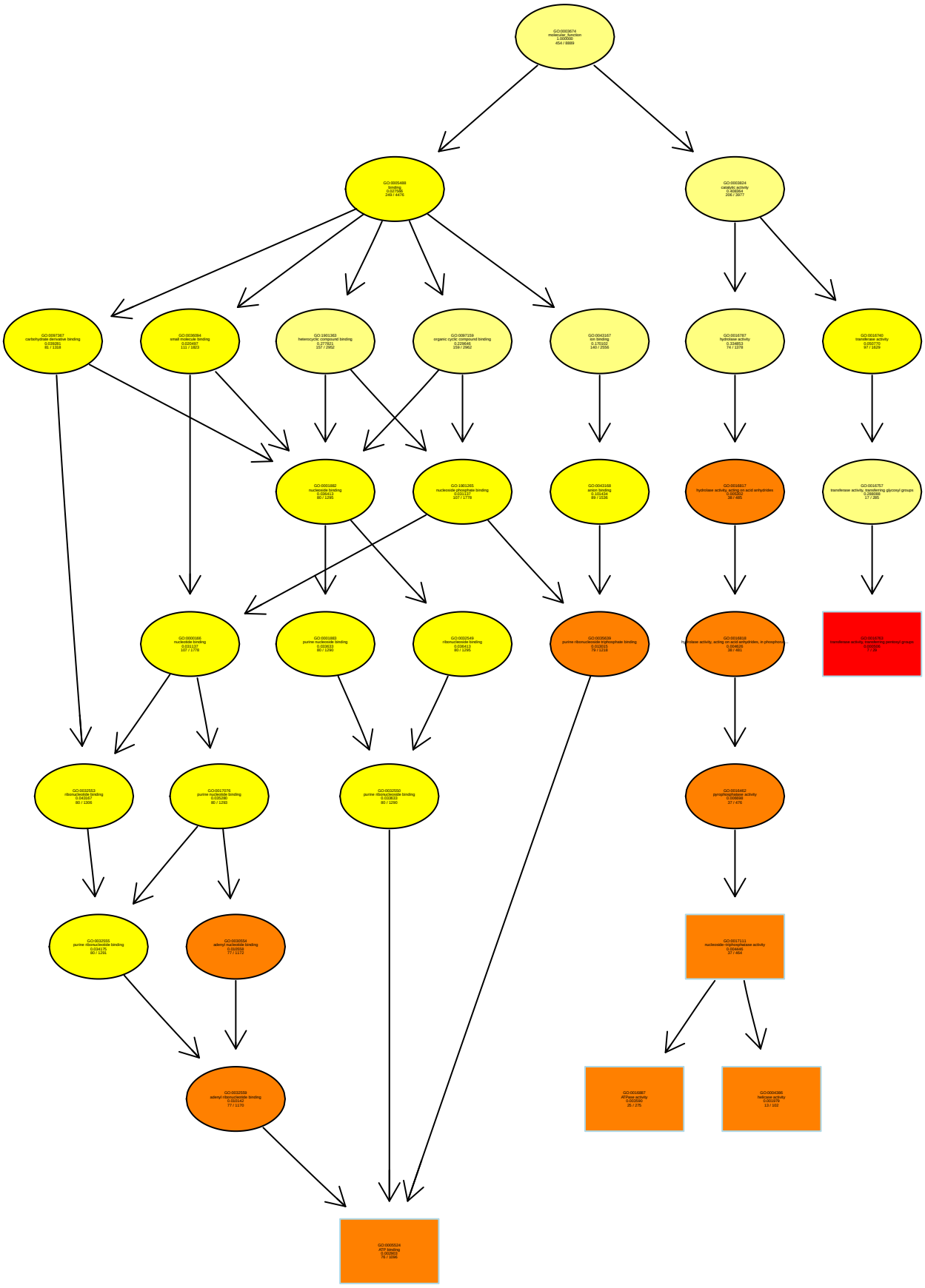
GO:0003674
molecular_function
1.000000
454 / 8889
GO:0005488
binding
0.537756
249 / 4476
GO:0003824
catalytic activity
0.456264
206 / 3977
GO:0097367
carbohydrate derivative binding
0.239281
81 / 1318
GO:0036094
small molecule binding
0.320457
111 / 1853
GO:1901363
heterocyclic compound binding
0.277821
157 / 2562
GO:0097159
organic cyclic compound binding
0.229645
159 / 2562
GO:0043167
ion binding
0.176102
140 / 2556
GO:0016787
hydrolase activity
0.334853
74 / 1378
GO:0016740
transferase activity
0.292776
67 / 1629
GO:0001882
nucleoside binding
0.099413
80 / 1295
GO:1901265
nucleoside phosphate binding
0.101137
107 / 1778
GO:0043169
anion binding
0.101434
89 / 1536
GO:0016817
hydrolase activity, acting on acid anhydrides
0.395392
38 / 480
GO:0016757
transferase activity, transferring glycosyl groups
0.288088
17 / 365
GO:0000166
nucleotide binding
0.101137
107 / 1778
GO:0001883
purine nucleoside binding
0.098413
80 / 1295
GO:0032549
ribonucleoside binding
0.098413
80 / 1295
GO:0035639
purine ribonucleoside triphosphate binding
0.093015
79 / 1218
GO:0016818
hydrolase activity, acting on acid anhydrides, in phosphorus-containing anhydrides
0.394826
38 / 481
GO:0016763
transferase activity, transferring pentosyl groups
0.000000
7 / 29
GO:0032553
ribonucleotide binding
0.098413
80 / 1295
GO:0017076
purine nucleotide binding
0.098413
80 / 1295
GO:0032550
purine ribonucleoside binding
0.098413
80 / 1295
GO:0016462
pyrophosphatase activity
0.394826
37 / 476
GO:0032555
purine ribonucleotide binding
0.098175
80 / 1291
GO:0030554
adenyl nucleotide binding
0.092558
77 / 1172
GO:0017111
nucleoside-triphosphatase activity
0.394648
37 / 464
GO:0032559
adenyl ribonucleotide binding
0.092142
77 / 1170
GO:0016887
ATPase activity
0.192330
25 / 275
GO:0004386
helicase activity
0.101379
13 / 102
GO:0005524
ATP binding
0.092558
76 / 1156