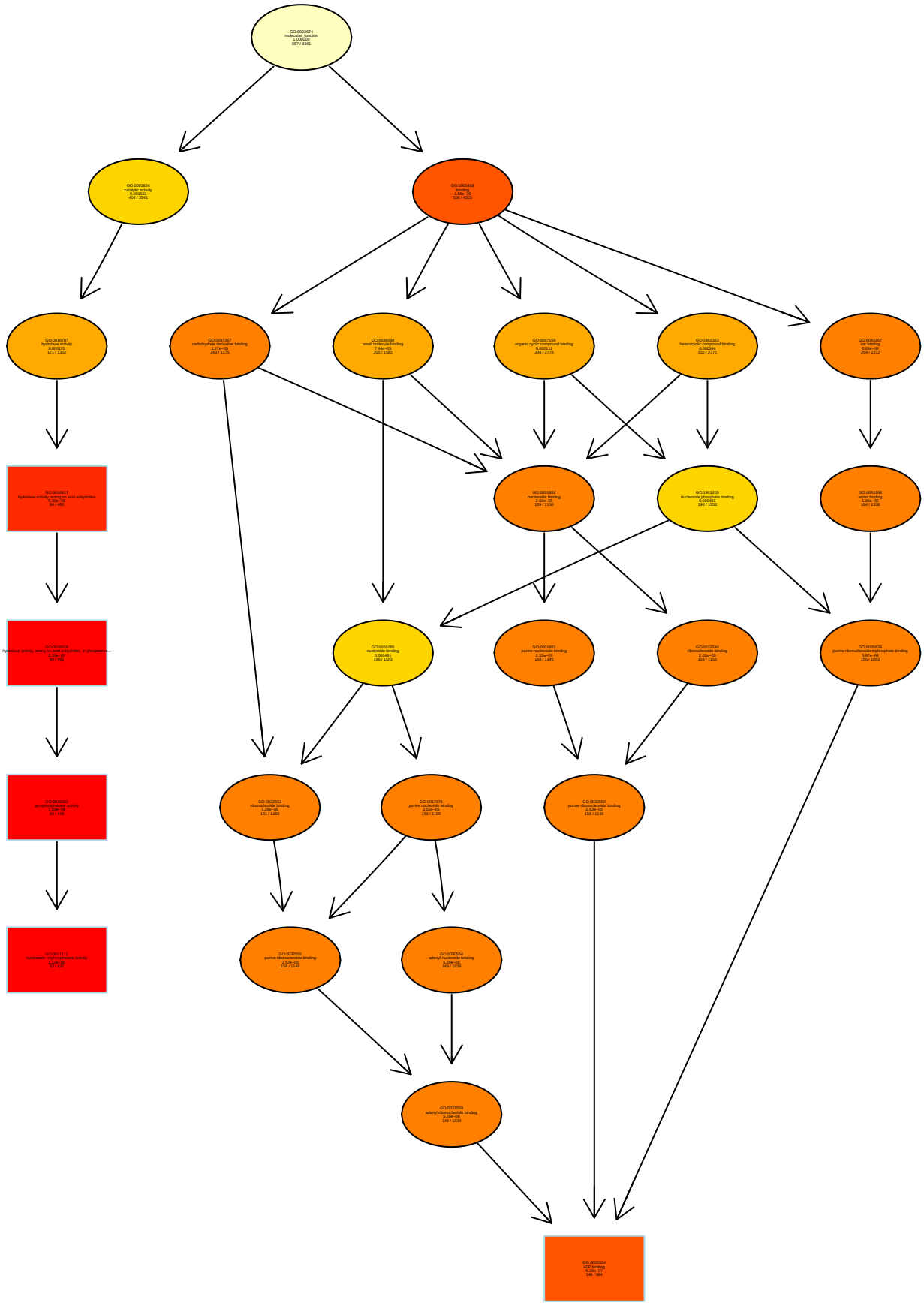
GO:0003674
molecular_function
1.000000
857 / 8281
GO:0003824
catalytic activity
0.000000
404 / 3541
GO:0005488
binding
1.00e-06
506 / 4305
GO:0016787
hydrolase activity
0.000000
171 / 1302
GO:0097367
carbohydrate derivative binding
1.27e-05
163 / 1175
GO:0036094
small molecule binding
7.44e-05
205 / 1588
GO:0097159
organic cyclic compound binding
0.000111
334 / 2776
GO:1901363
heterocyclic compound binding
0.000164
332 / 2772
GO:0043167
ion binding
6.69e-05
269 / 2272
GO:0016817
hydrolase activity, acting on acid anhydrides
5.39e-18
94 / 450
GO:0001882
nucleoside binding
2.03e-05
159 / 1150
GO:1901265
nucleoside phosphate binding
0.000491
196 / 1553
GO:0043168
anion binding
1.36e-05
199 / 1358
GO:0016818
hydrolase activity, acting on acid anhydrides, in phosphorus-...
2.39e-08
94 / 451
GO:0000166
nucleotide binding
0.000491
196 / 1553
GO:0001883
purine nucleoside binding
2.53e-05
158 / 1146
GO:0032549
ribonucleoside binding
2.03e-05
159 / 1150
GO:0035639
purine ribonucleoside triphosphate binding
5.87e-06
155 / 1092
GO:0016462
pyrophosphatase activity
0.000000
94 / 448
GO:0032553
ribonucleotide binding
2.03e-05
161 / 1158
GO:0017076
purine nucleotide binding
2.03e-05
159 / 1150
GO:0032550
purine ribonucleoside binding
2.53e-05
158 / 1146
GO:0017111
nucleoside-triphosphatase activity
1.17e-08
87 / 427
GO:0032555
purine ribonucleotide binding
2.53e-05
158 / 1146
GO:0000554
adenyl nucleotide binding
2.39e-05
149 / 1039
GO:0032559
adenyl ribonucleotide binding
5.28e-05
149 / 1039
GO:0005524
ATP binding
2.39e-05
149 / 1039