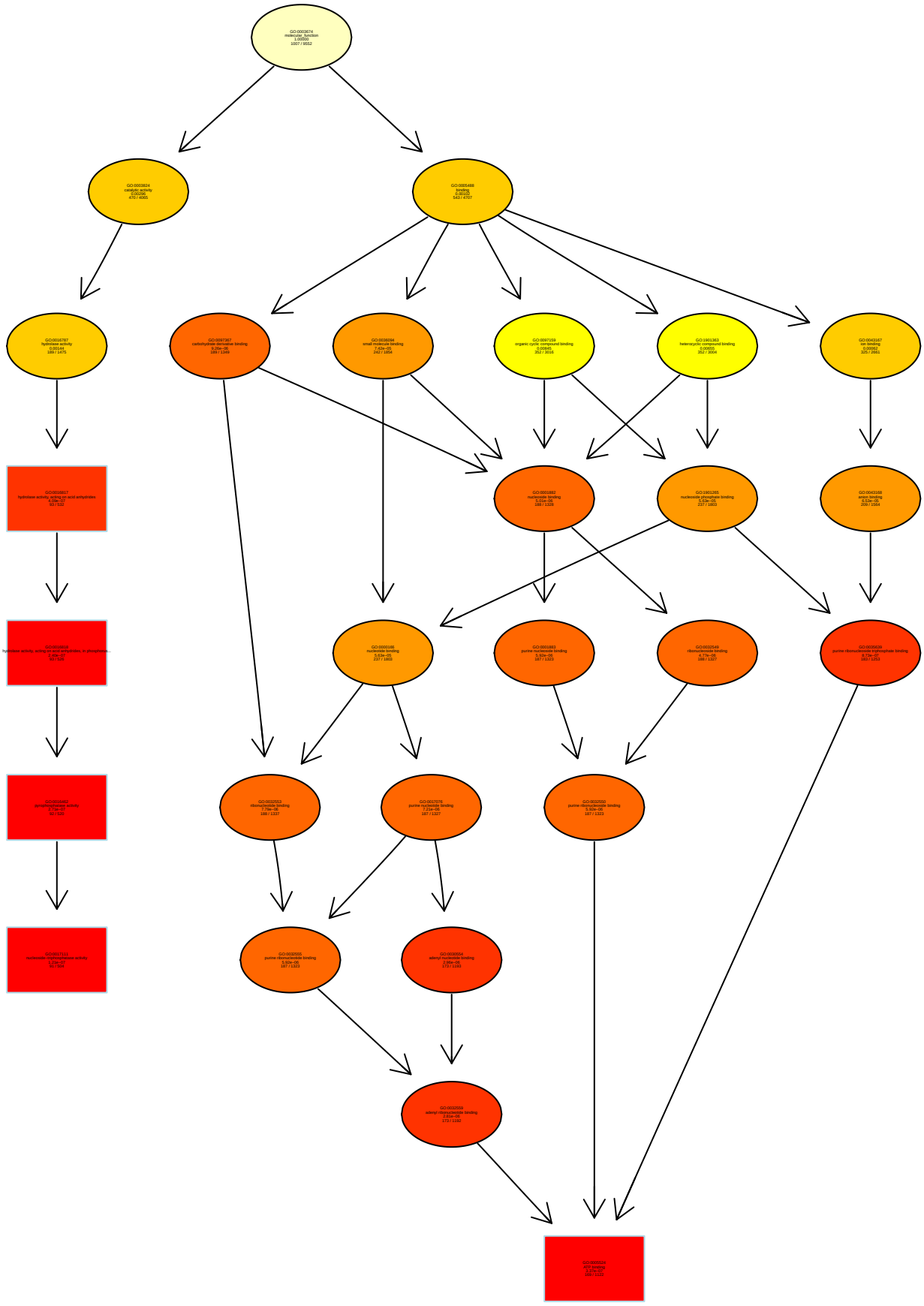
GO:0003674
molecular_function
1.00000
1027 / 9552
GO:0003824
catalytic activity
0.00000
670 / 4055
GO:0005488
binding
0.00000
543 / 4707
GO:0016787
hydrolase activity
0.00144
189 / 1476
GO:0097367
carbohydrate derivative binding
8.35e-05
199 / 1349
GO:0036094
small molecule binding
7.41e-05
242 / 1864
GO:0097159
organic cyclic compound binding
0.00845
352 / 3016
GO:1901363
heterocyclic compound binding
0.00800
352 / 3004
GO:0043167
ion binding
0.00582
325 / 2861
GO:0016817
hydrolase activity, acting on acid anhydrides
4.19e-07
93 / 532
GO:0001882
nucleoside binding
5.01e-05
188 / 1328
GO:1901265
nucleoside phosphate binding
5.63e-05
237 / 1803
GO:0043168
anion binding
6.53e-05
269 / 2164
GO:0016818
hydrolase activity, acting on acid anhydrides, in phosphorus-c...
4.49e-07
93 / 526
GO:0000166
nucleotide binding
5.62e-05
237 / 1803
GO:0001883
purine nucleoside binding
5.93e-05
187 / 1323
GO:0032549
ribonucleoside binding
6.77e-05
188 / 1327
GO:0035639
purine ribonucleoside triphosphate binding
6.73e-07
193 / 1253
GO:0016462
pyrophosphatase activity
4.49e-07
93 / 520
GO:0032553
ribonucleotide binding
5.93e-05
188 / 1327
GO:0017076
purine nucleotide binding
5.93e-05
187 / 1327
GO:0032550
purine ribonucleoside binding
5.93e-05
187 / 1323
GO:0017111
nucleoside-triphosphatase activity
5.27e-07
91 / 504
GO:0032555
purine ribonucleotide binding
5.93e-05
187 / 1323
GO:0005524
adenyl nucleotide binding
7.93e-05
173 / 1193
GO:0032559
adenyl ribonucleotide binding
7.93e-05
173 / 1192
GO:0005524
ATP binding
5.93e-05
187 / 1323