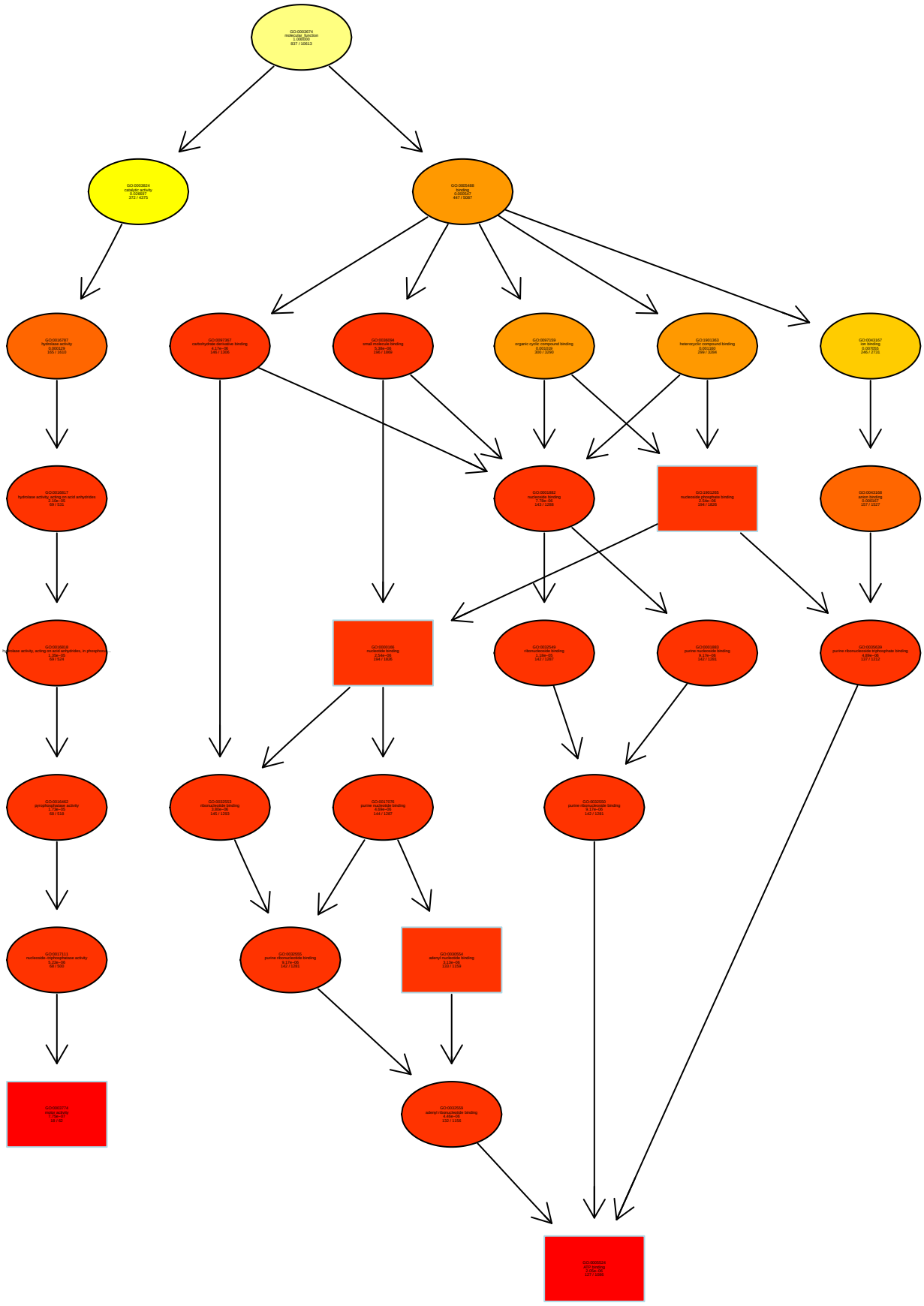
GO:0003674
molecular_function
1.000000
837 / 16613
GO:0003824
catalytic activity
0.525837
372 / 4375
GO:0005488
binding
0.500000
447 / 5087
GO:0016787
hydrolase activity
0.000191
157 / 1616
GO:0097367
carbohydrate derivative binding
4.17e-05
148 / 1306
GO:0036094
small molecule binding
5.29e-05
158 / 1386
GO:0097159
organic cyclic compound binding
0.001155
300 / 3290
GO:1901363
heterocyclic compound binding
0.001180
298 / 3284
GO:0043167
ion binding
0.007055
246 / 2731
GO:0016817
hydrolase activity, acting on acid anhydrides
2.19e-10
69 / 521
GO:0001882
nucleoside binding
7.79e-06
143 / 1288
GO:1901265
nucleoside phosphate binding
2.54e-05
194 / 1826
GO:0043168
anion binding
0.000157
157 / 1527
GO:0016818
hydrolase activity, acting on acid anhydrides, in phosphorus-containing anhydrides
1.26e-09
69 / 524
GO:0000166
nucleotide binding
2.64e-05
194 / 1826
GO:0032549
ribonucleoside binding
1.18e-05
142 / 1287
GO:0001883
purine nucleoside binding
9.17e-06
142 / 1281
GO:0035639
purine ribonucleoside triphosphate binding
4.69e-06
137 / 1212
GO:0016462
pyrophosphatase activity
1.00e-09
69 / 518
GO:0032553
ribonucleotide binding
2.64e-05
142 / 1293
GO:0017076
purine nucleotide binding
2.64e-05
144 / 1300
GO:0032550
purine ribonucleoside binding
9.17e-06
142 / 1281
GO:0017111
nucleoside-triphosphatase activity
2.19e-10
69 / 350
GO:0032555
purine ribonucleotide binding
9.17e-06
142 / 1281
GO:0005524
adenyl nucleotide binding
3.19e-05
132 / 1159
GO:0003774
motor activity
7.79e-07
18 / 62
GO:0032559
adenyl ribonucleotide binding
4.46e-05
132 / 1156
GO:0005524
ATP binding
3.19e-05
127 / 1085