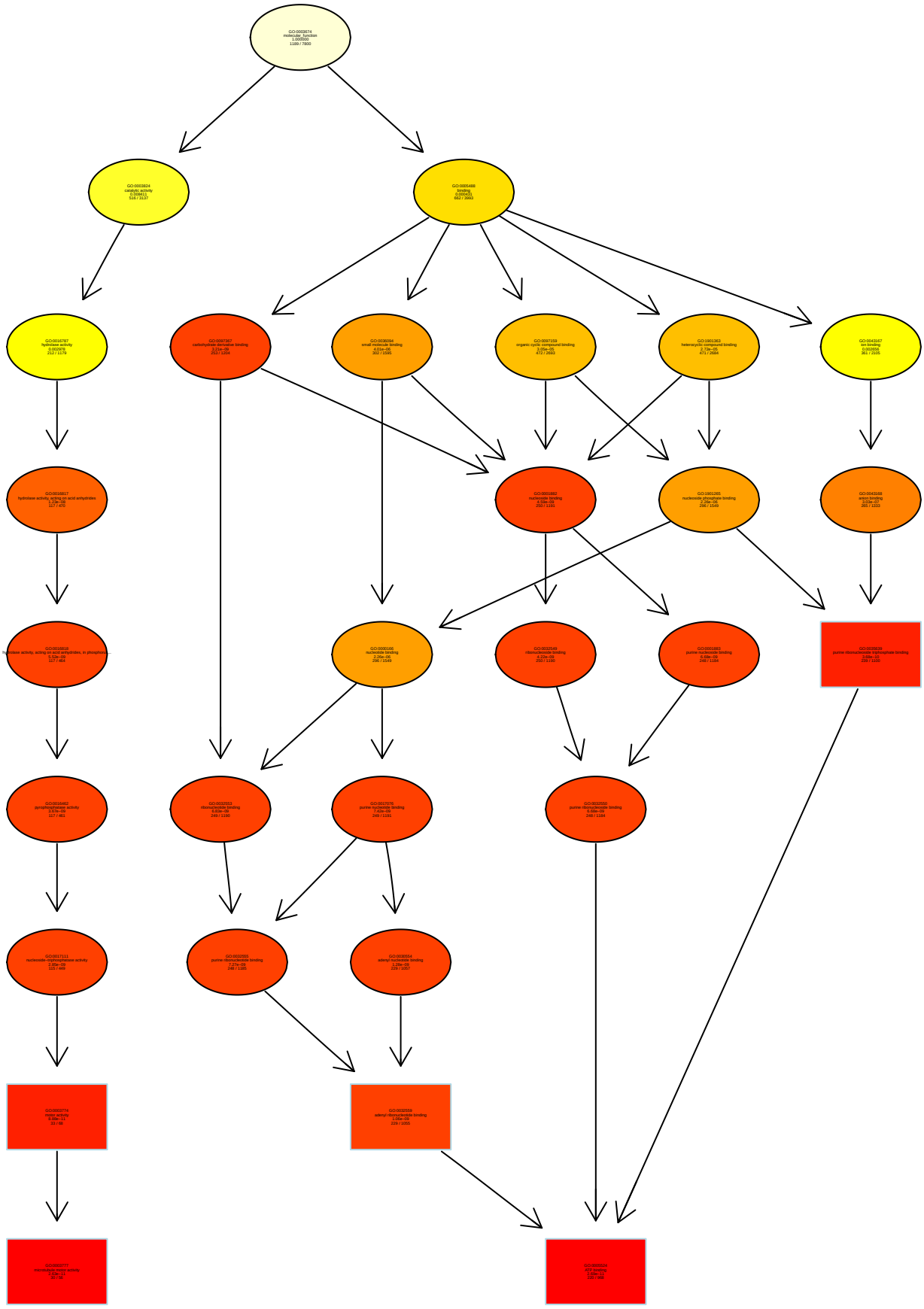
GO:0003824
catalytic activity
1.000000
516 / 3137
GO:0016787
hydrolase activity
0.000000
212 / 1179
GO:0016817
hydrolase activity, acting on acid anhydrides
1.23e-02
117 / 470
GO:0016818
hydrolase activity, acting on acid anhydrides, in phosphorus-containing anhydrides
5.52e-06
117 / 464
GO:0016462
pyrophosphatase activity
1.00e-02
117 / 461
GO:0017111
nucleoside-triphosphatase activity
1.00e-02
121 / 449
GO:0003774
motor activity
8.08e-11
23 / 68
GO:0003777
microtubule motor activity
3.47e-11
20 / 62
GO:0005488
binding
0.000000
662 / 3993
GO:0097367
carbohydrate derivative binding
2.21e-09
253 / 1204
GO:0036094
small molecule binding
4.61e-06
302 / 1595
GO:0097159
organic cyclic compound binding
3.55e-05
472 / 2993
GO:1901363
heterocyclic compound binding
2.74e-05
471 / 2984
GO:0043167
ion binding
0.000000
361 / 2335
GO:0043168
anion binding
3.03e-10
265 / 1333
GO:0001882
nucleoside binding
4.19e-09
250 / 1191
GO:1901265
nucleoside phosphate binding
2.26e-06
298 / 1549
GO:0000166
nucleotide binding
2.26e-06
298 / 1549
GO:0032549
ribonucleoside binding
4.22e-09
250 / 1190
GO:0001883
purine nucleoside binding
6.68e-09
248 / 1184
GO:0032553
ribonucleotide binding
2.21e-09
249 / 1190
GO:0017076
purine nucleotide binding
2.26e-06
298 / 1191
GO:0032550
purine ribonucleoside binding
1.68e-09
248 / 1184
GO:0032555
purine ribonucleotide binding
2.21e-09
248 / 1190
GO:0030554
adenyl nucleotide binding
3.39e-07
229 / 1037
GO:0032559
adenyl ribonucleotide binding
1.56e-09
229 / 1055
GO:0005524
ATP binding
3.39e-07
229 / 1037