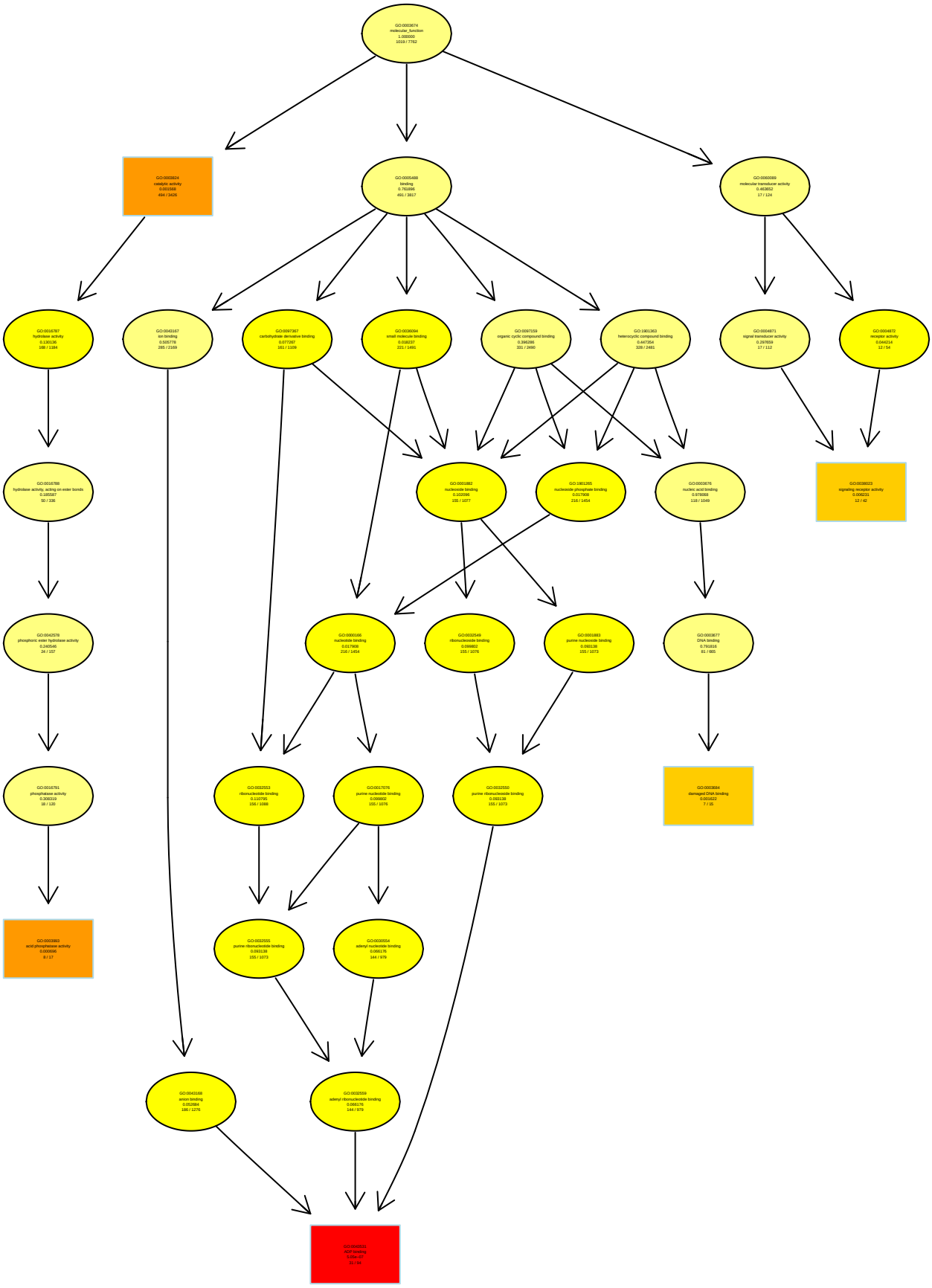
GO:0003674
molecular_function
1.000000
1010 / 7752
GO:0003824
catalytic activity
0.501568
494 / 3426
GO:0005488
binding
0.761896
491 / 3817
GO:0060089
molecular transducer activity
0.453652
17 / 124
GO:0016787
hydrolase activity
0.130136
168 / 1184
GO:0043167
ion binding
0.555778
285 / 2169
GO:0097367
carbohydrate derivative binding
0.077587
161 / 1109
GO:0036094
small molecule binding
0.018237
221 / 1491
GO:0097159
organic cyclic compound binding
0.396266
331 / 2490
GO:1901363
heterocyclic compound binding
0.447354
328 / 2481
GO:0004871
signal transducer activity
0.267659
17 / 112
GO:0004872
receptor activity
0.044214
12 / 54
GO:0016788
hydrolase activity, acting on ester bonds
0.185587
50 / 336
GO:0001882
nucleoside binding
0.102096
155 / 1077
GO:1901265
nucleoside phosphate binding
0.017908
216 / 1454
GO:0003676
nucleic acid binding
0.978868
118 / 1049
GO:0038023
signaling receptor activity
0.006231
12 / 42
GO:0042578
phosphoric ester hydrolase activity
0.240546
24 / 157
GO:0000166
nucleotide binding
0.017908
216 / 1454
GO:0032549
ribonucleoside binding
0.099802
155 / 1076
GO:0001883
purine nucleoside binding
0.098138
155 / 1073
GO:0003677
DNA binding
0.781816
81 / 665
GO:0016791
phosphatase activity
0.308319
18 / 120
GO:0032553
ribonucleotide binding
0.110795
156 / 1088
GO:0017076
purine nucleotide binding
0.099802
155 / 1076
GO:0032550
purine ribonucleoside binding
0.098138
155 / 1073
GO:0003684
damaged DNA binding
0.001622
7 / 15
GO:0003993
acid phosphatase activity
0.000696
8 / 17
GO:0032555
purine ribonucleotide binding
0.098138
155 / 1073
GO:0030554
adenyl nucleotide binding
0.066176
144 / 979
GO:0043168
anion binding
0.052084
166 / 1276
GO:0032559
adenyl ribonucleotide binding
0.066176
144 / 979
GO:0000166
ADP binding
5.05e-07
31 / 64