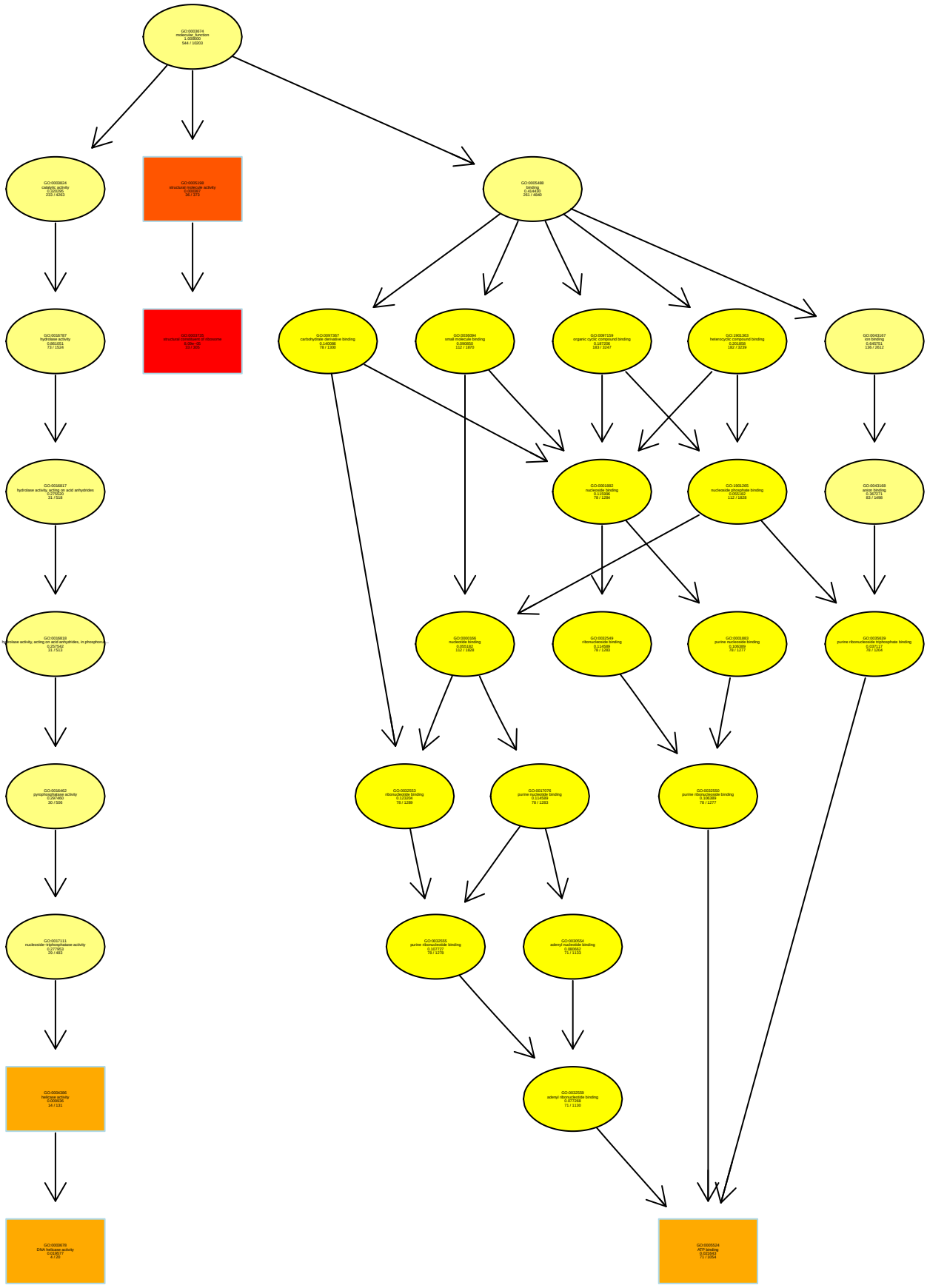
GO:0003674
molecular_function
1.000000
544 / 10203
GO:0003824
catalytic activity
0.000000
233 / 4263
GO:0005198
structural molecule activity
0.000000
36 / 373
GO:0005488
binding
0.000000
251 / 4840
GO:0016787
hydrolase activity
0.661051
73 / 1324
GO:0003735
structural constituent of ribosome
0.00e-00
37 / 372
GO:0097367
carbohydrate derivative binding
0.146598
78 / 1360
GO:0036094
small molecule binding
0.098993
112 / 1870
GO:0097159
organic cyclic compound binding
0.197256
163 / 2249
GO:1901363
heterocyclic compound binding
0.201858
163 / 2239
GO:0043167
ion binding
0.645701
136 / 2612
GO:0016817
hydrolase activity, acting on acid anhydrides
0.270520
31 / 518
GO:0001882
nucleoside binding
0.153946
78 / 1284
GO:1901265
nucleoside phosphate binding
0.200182
112 / 1828
GO:0043168
anion binding
0.367271
82 / 1468
GO:0016818
hydrolase activity, acting on acid anhydrides, in phosphorus-
0.257522
31 / 513
GO:0000166
nucleotide binding
0.200182
112 / 1828
GO:0032549
ribonucleoside binding
0.144598
78 / 1283
GO:0001883
purine nucleoside binding
0.196388
78 / 1277
GO:0035639
purine ribonucleoside triphosphate binding
0.027172
78 / 1204
GO:0016462
pyrophosphatase activity
0.241260
30 / 506
GO:0032553
ribonucleotide binding
0.196388
78 / 1288
GO:0017076
purine nucleotide binding
0.197256
78 / 1283
GO:0032550
purine ribonucleoside binding
0.196388
78 / 1277
GO:0017111
nucleoside-triphosphatase activity
0.277853
29 / 483
GO:0032555
purine ribonucleotide binding
0.197172
78 / 1278
GO:0030554
adenyl nucleotide binding
0.186882
71 / 1163
GO:0004386
helicase activity
0.036036
14 / 131
GO:0032559
adenyl ribonucleotide binding
0.077358
71 / 1130
GO:0003678
DNA helicase activity
0.036037
4 / 20
GO:0005524
ATP binding
0.077358
71 / 1204