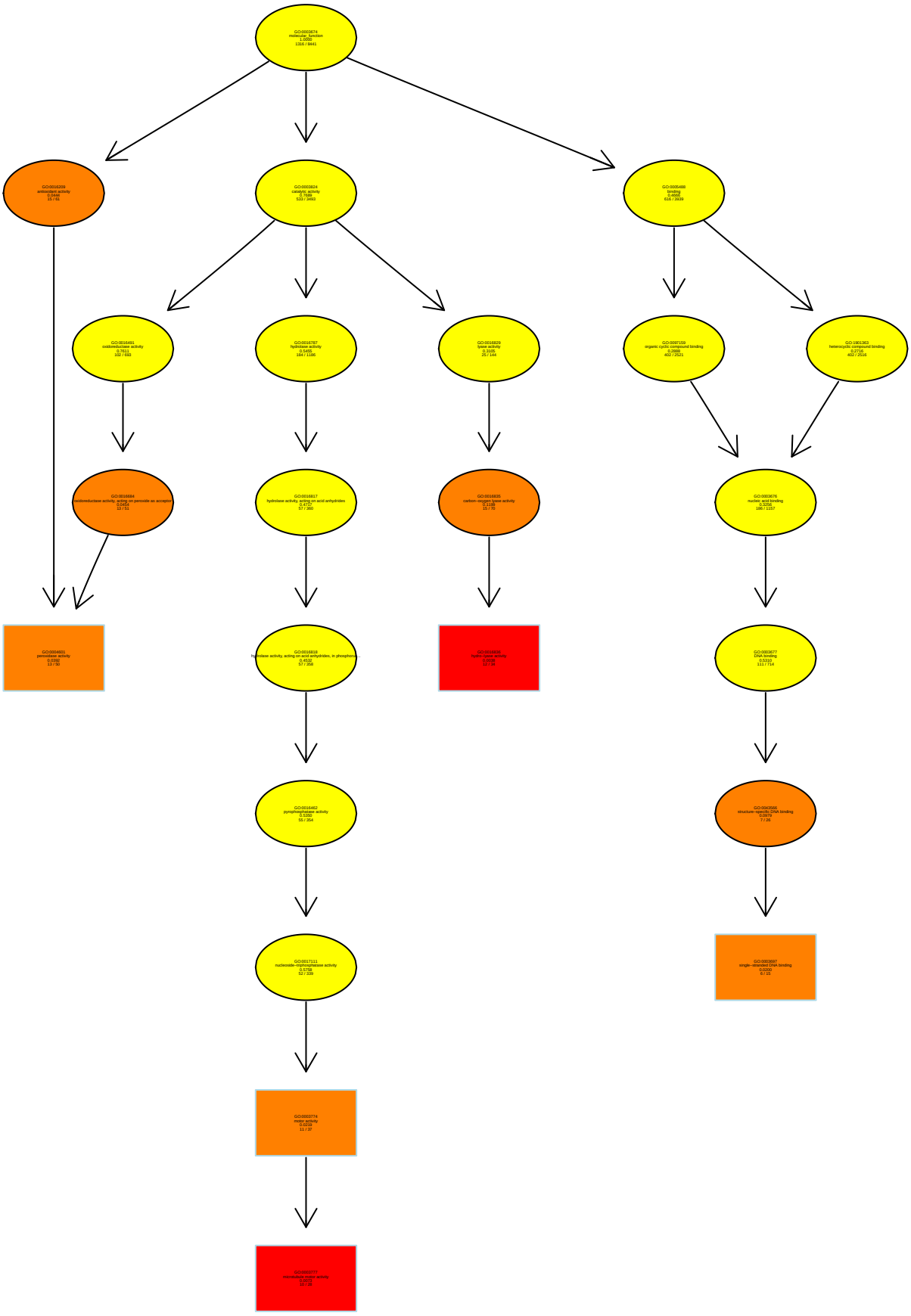
GO:0003674
molecular_function
1.0000
1316 / 8441
GO:0016209
antioxidant activity
0.0444
15 / 61
GO:0003824
catalytic activity
0.7430
533 / 3493
GO:0005488
binding
0.4652
616 / 3939
GO:0016491
oxidoreductase activity
0.7611
102 / 693
GO:0016787
hydrolase activity
0.5495
384 / 1186
GO:0016829
lyase activity
0.3105
25 / 144
GO:0097159
organic cyclic compound binding
0.2885
402 / 2521
GO:1901363
heterocyclic compound binding
0.2716
402 / 2516
GO:0016684
oxidoreductase activity, acting on peroxide as acceptor
0.0454
12 / 51
GO:0016817
hydrolase activity, acting on acid anhydrides
0.4737
57 / 360
GO:0016835
carbon-oxygen lyase activity
0.1148
10 / 70
GO:0003676
nucleic acid binding
0.3256
166 / 1157
GO:0004601
peroxidase activity
0.0392
12 / 50
GO:0016818
hydrolase activity, acting on acid anhydrides, in phosphorus-c...
0.4532
57 / 358
GO:0016836
hydro-lyase activity
0.0588
12 / 34
GO:0003677
DNA binding
0.3310
111 / 714
GO:0016462
pyrophosphatase activity
0.4496
55 / 354
GO:0043566
structure-specific DNA binding
0.0229
7 / 26
GO:0017111
nucleoside-triphosphatase activity
0.5736
52 / 339
GO:0003697
single-stranded DNA binding
0.0208
6 / 15
GO:0003774
motor activity
0.0219
11 / 37
GO:0003777
microtubule motor activity
0.0184
10 / 28