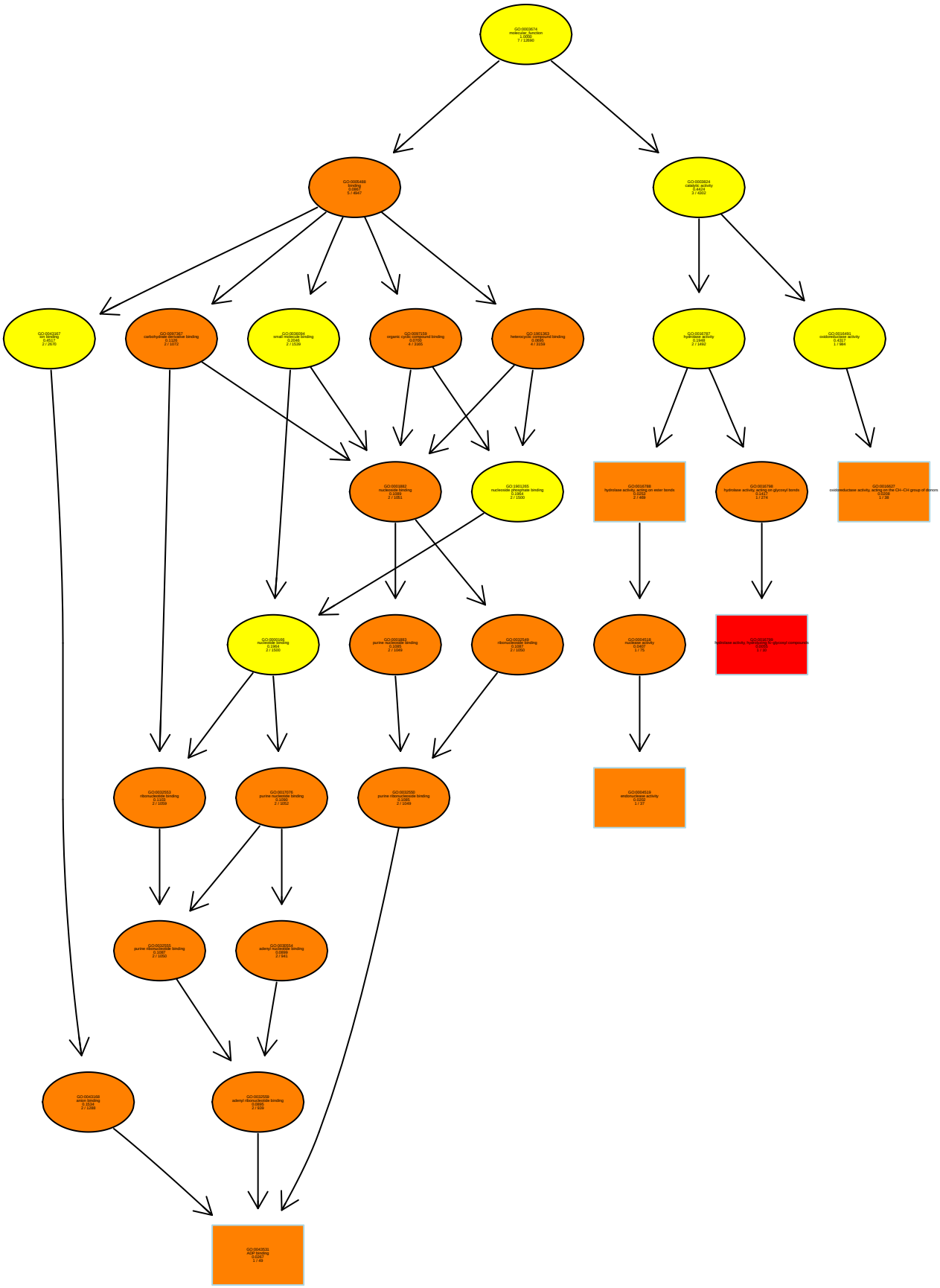
GO:0003674
molecular_function
1.0000
7 / 10690
GO:0005488
binding
0.9267
5 / 4847
GO:0003824
catalytic activity
0.4147
3 / 4302
GO:0043167
ion binding
0.1534
2 / 2670
GO:0097367
carbohydrate derivative binding
0.1534
2 / 1573
GO:0036094
small molecule binding
0.2046
2 / 1535
GO:0097159
organic cyclic compound binding
0.3858
4 / 3365
GO:1901363
heterocyclic compound binding
0.3858
4 / 3359
GO:0016787
hydrolase activity
0.3446
2 / 1452
GO:0016491
oxidoreductase activity
0.4202
1 / 864
GO:0001882
nucleoside binding
0.1858
2 / 1051
GO:1901265
nucleoside phosphate binding
0.1864
2 / 1500
GO:0016788
hydrolase activity, acting on ester bonds
0.0202
2 / 469
GO:0016798
hydrolase activity, acting on glycosyl bonds
0.0437
1 / 274
GO:0016620
oxidoreductase activity, acting on the CH-CH group of donors
0.0158
1 / 28
GO:0001883
purine nucleoside binding
0.1864
2 / 1500
GO:0001883
purine nucleoside binding
0.1585
2 / 1069
GO:0032549
ribonucleoside binding
0.1587
2 / 1050
GO:0004518
nuclease activity
0.0485
1 / 75
GO:0016798
hydrolase activity, acting on glycosyl compounds
0.0025
1 / 10
GO:0032553
ribonucleotide binding
0.1585
2 / 1059
GO:0017076
purine nucleotide binding
0.1585
2 / 1052
GO:0032555
purine ribonucleotide binding
0.1585
2 / 1049
GO:0004519
endonuclease activity
0.0202
1 / 237
GO:0032555
purine ribonucleotide binding
0.1587
2 / 1050
GO:0030554
adenyl nucleotide binding
0.0846
2 / 941
GO:0043168
anion binding
0.1534
2 / 1288
GO:0032559
adenyl ribonucleotide binding
0.0846
2 / 939
GO:0043531
ADP binding
0.0267
1 / 49