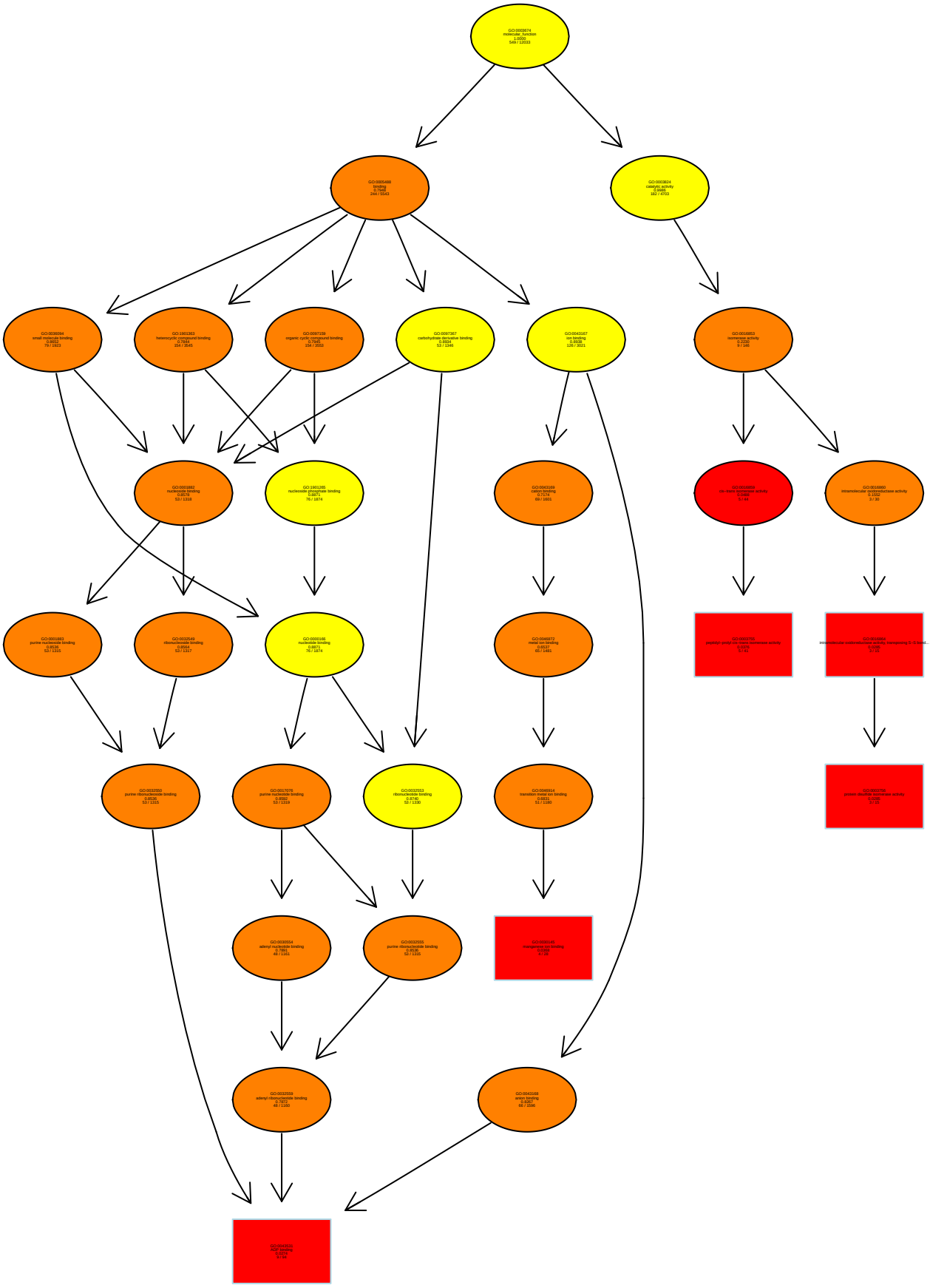
GO:0003674
molecular_function
0.0000
549 / 12933
GO:0005488
binding
0.7568
244 / 5543
GO:0003824
catalytic activity
0.3986
162 / 4703
GO:0036094
small molecule binding
0.9652
79 / 1923
GO:1901363
heterocyclic compound binding
0.7644
154 / 3545
GO:0097159
organic cyclic compound binding
0.7645
154 / 3553
GO:0097367
carbohydrate derivative binding
0.8684
53 / 1348
GO:0043167
ion binding
0.9538
126 / 3021
GO:0016853
isomerase activity
0.2250
9 / 145
GO:0001882
nucleoside binding
0.8539
53 / 1318
GO:1901265
nucleoside phosphate binding
0.8871
76 / 1874
GO:0043169
cation binding
0.7174
69 / 1601
GO:0016859
cis–trans isomerase activity
0.0454
5 / 44
GO:0016860
intramolecular oxidoreductase activity
0.1532
3 / 30
GO:0001883
purine nucleoside binding
0.8538
53 / 1315
GO:0032549
ribonucleoside binding
0.8564
53 / 1317
GO:0000166
nucleotide binding
0.8871
76 / 1874
GO:0046872
metal ion binding
0.6527
65 / 1481
GO:0003755
peptidyl–prolyl cis–trans isomerase activity
0.0276
3 / 41
GO:0016864
intramolecular oxidoreductase activity, transposing S–S bond...
0.0000
3 / 15
GO:0032550
purine ribonucleoside binding
0.8538
53 / 1315
GO:0017076
purine nucleotide binding
0.8532
53 / 1319
GO:0032553
ribonucleotide binding
0.8538
53 / 1316
GO:0046914
transition metal ion binding
0.6532
53 / 1318
GO:0003755
protein disulfide isomerase activity
0.0000
3 / 20
GO:0030554
adenyl nucleotide binding
0.7861
49 / 1161
GO:0032555
purine ribonucleotide binding
0.8538
53 / 1316
GO:0030145
manganese ion binding
0.0769
4 / 26
GO:0032559
adenyl ribonucleotide binding
0.7872
49 / 1160
GO:0043168
anion binding
0.4267
68 / 1596
GO:0043531
ADP binding
0.1000
3 / 34