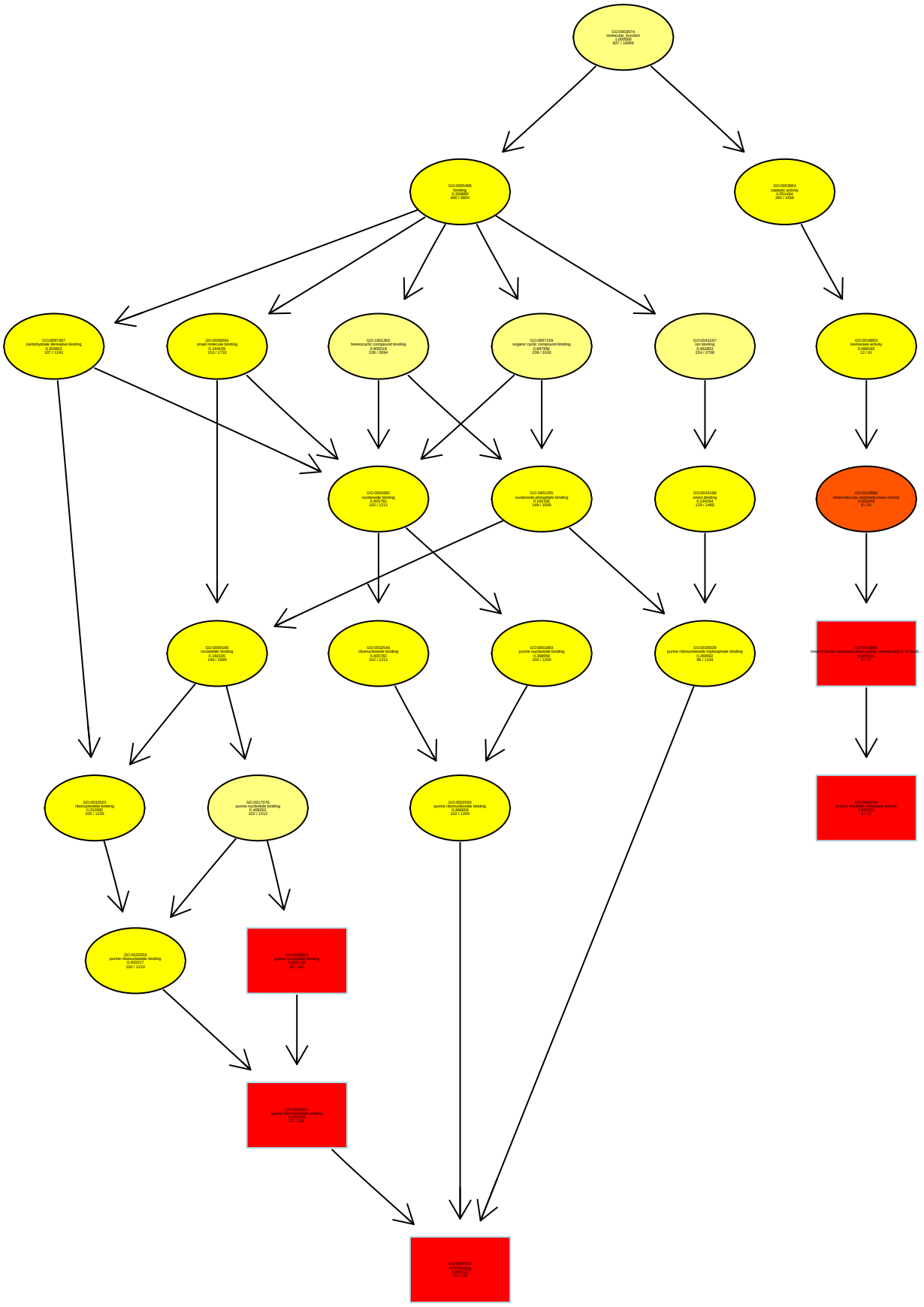
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molecular_function
1.000000
827 / 10069
GO:0005488
binding
0.205859
400 / 4804
GO:0003824
catalytic activity
0.551494
364 / 4156
GO:0097367
carbohydrate derivative binding
0.303815
107 / 1241
GO:0036094
small molecule binding
0.164109
153 / 1733
GO:1901363
heterocyclic compound binding
0.905019
238 / 3094
GO:0097159
organic cyclic compound binding
0.897892
239 / 3100
GO:0043167
ion binding
0.462603
224 / 2708
GO:0016853
isomerase activity
0.068160
12 / 91
GO:0001882
nucleoside binding
0.405782
102 / 1211
GO:1901265
nucleoside phosphate binding
0.192105
149 / 1699
GO:0043168
anion binding
0.199194
129 / 1465
GO:0016860
intramolecular oxidoreductase activity
0.000449
8 / 24
GO:0000166
nucleotide binding
0.192105
149 / 1699
GO:0032549
ribonucleoside binding
0.405782
102 / 1211
GO:0001883
purine nucleoside binding
0.398659
102 / 1209
GO:0035639
purine ribonucleoside triphosphate binding
0.266592
99 / 1134
GO:0016864
intramolecular oxidoreductase activity, transposing S–S bond...
0.000181
6 / 12
GO:0032553
ribonucleoside binding
0.332930
105 / 1226
GO:0017076
purine nucleotide binding
0.408253
102 / 1212
GO:0032550
purine ribonucleoside binding
0.398659
102 / 1209
GO:0003756
protein disulfide isomerase activity
0.000181
6 / 12
GO:0032555
purine ribonucleotide binding
0.402217
102 / 1210
GO:0019001
guanyl nucleotide binding
0.000144
25 / 140
GO:0032561
guanyl ribonucleotide binding
0.000129
25 / 138
GO:0005525
GTP binding
0.000129
25 / 138