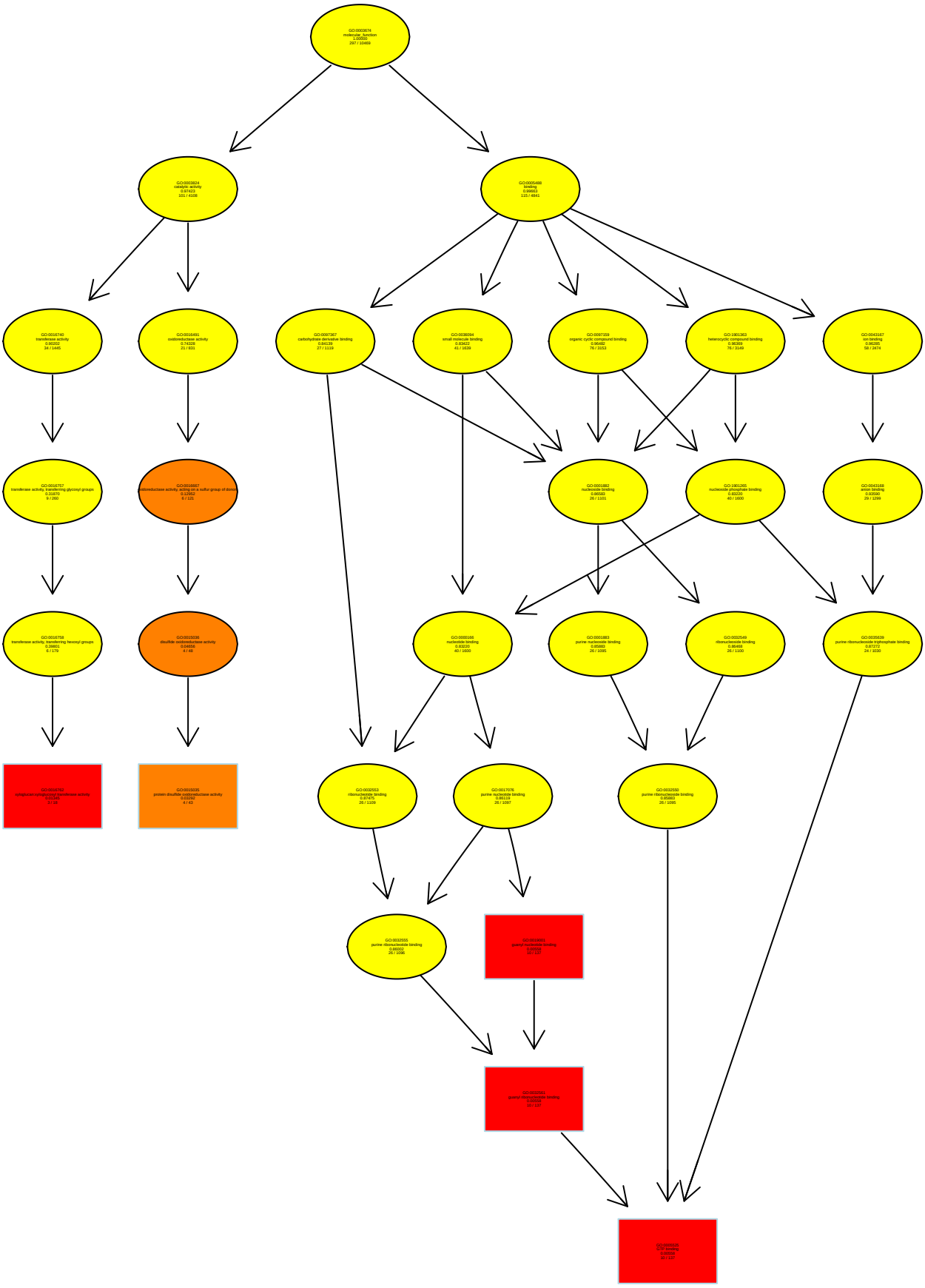
GO:0003674
molecular_function
1.00000
297 / 10469
GO:0003824
catalytic activity
0.67423
101 / 4108
GO:0005488
binding
0.99663
115 / 4841
GO:0016740
transferase activity
0.80202
34 / 1445
GO:0016491
oxidoreductase activity
0.74328
21 / 831
GO:0097367
carbohydrate derivative binding
0.94139
27 / 1119
GO:0036094
small molecule binding
0.83822
41 / 1639
GO:0097159
organic cyclic compound binding
0.96482
76 / 3153
GO:1901363
heterocyclic compound binding
0.96384
76 / 3149
GO:0043167
ion binding
0.96285
58 / 2474
GO:0016757
transferase activity, transferring glycosyl groups
0.31870
9 / 260
GO:0016667
oxidoreductase activity, acting on a sulfur group of donors
0.12852
6 / 121
GO:0001882
nucleoside binding
0.96583
26 / 1101
GO:1901265
nucleoside phosphate binding
0.83220
40 / 1600
GO:0043168
anion binding
0.93590
29 / 1299
GO:0016758
transferase activity, transferring hexosyl groups
0.28601
6 / 179
GO:0015036
disulfide oxidoreductase activity
0.04656
4 / 48
GO:0000166
nucleotide binding
0.83220
40 / 1600
GO:0001883
purine nucleoside binding
0.85883
26 / 1095
GO:0032549
ribonucleoside binding
0.86468
26 / 1100
GO:0035639
purine ribonucleoside triphosphate binding
0.87272
24 / 1030
GO:0016762
xyloglucan:xyloglucosyl transferase activity
0.00135
2 / 15
GO:0015035
protein disulfide oxidoreductase activity
0.04656
4 / 48
GO:0032553
ribonucleotide binding
0.86468
26 / 1100
GO:0017076
purine nucleotide binding
0.86468
26 / 1100
GO:0032550
purine ribonucleoside binding
0.85883
26 / 1095
GO:0032555
purine ribonucleotide binding
0.86202
26 / 1096
GO:0019001
guanyl nucleotide binding
0.00258
10 / 137
GO:0032561
guanyl ribonucleotide binding
0.00258
10 / 137
GO:0005525
GTP binding
0.00258
10 / 137