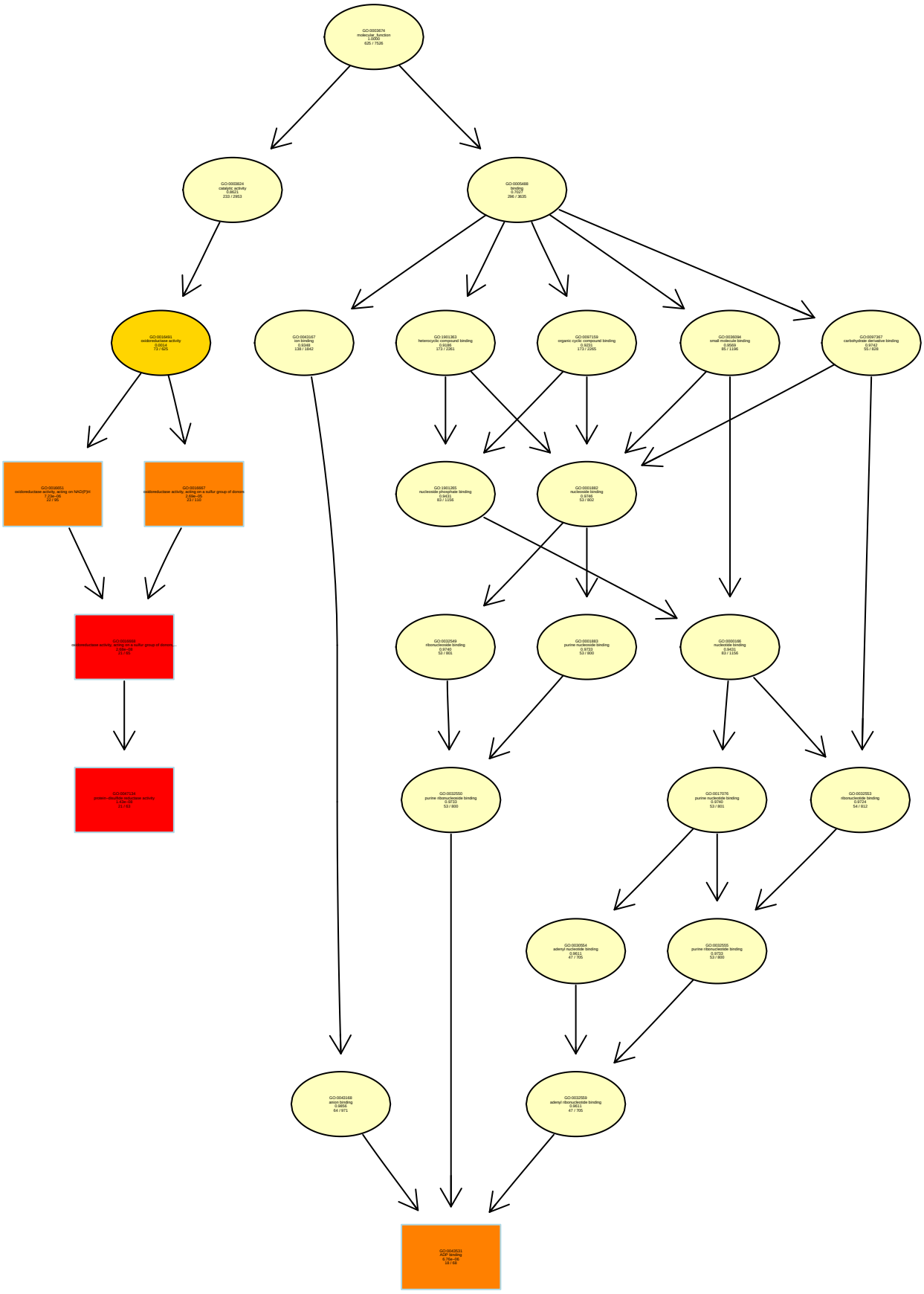
GO:0003674
molecular_function
1.0000
625 / 7526
GO:0003824
catalytic activity
0.9621
233 / 2953
GO:0005488
binding
0.7027
296 / 3635
GO:0016491
oxidoreductase activity
0.0014
73 / 825
GO:0043167
ion binding
0.9148
138 / 1842
GO:1901363
heterocyclic compound binding
0.9168
173 / 2281
GO:0097159
organic cyclic compound binding
0.9025
173 / 2265
GO:0036094
small molecule binding
0.9569
85 / 1156
GO:0097367
carbohydrate derivative binding
0.9742
55 / 828
GO:0016651
oxidoreductase activity, acting on NAD(P)H
7.22e-06
22 / 55
GO:0016667
oxidoreductase activity, acting on a sulfur group of donors
2.69e-05
23 / 110
GO:1901265
nucleoside phosphate binding
0.9431
83 / 1156
GO:0001882
nucleoside binding
0.9746
53 / 802
GO:0016668
oxidoreductase activity, acting on a sulfur group of donors, ...
2.69e-05
21 / 63
GO:0032549
ribonucleoside binding
0.9740
53 / 801
GO:0001883
purine nucleoside binding
0.9733
53 / 800
GO:0000166
nucleotide binding
0.9421
83 / 1156
GO:0047134
protein-disulfide reductase activity
1.34e-05
21 / 63
GO:0032550
purine ribonucleoside binding
0.9740
53 / 800
GO:0017076
purine nucleotide binding
0.9739
53 / 801
GO:0032553
ribonucleotide binding
0.9733
54 / 812
GO:0030554
adenyl nucleotide binding
0.9611
47 / 705
GO:0032555
purine ribonucleotide binding
0.9733
53 / 800
GO:0043168
anion binding
0.9656
64 / 971
GO:0032559
adenyl ribonucleotide binding
0.9611
47 / 705
GO:0043531
ADP binding
6.76e-06
18 / 68