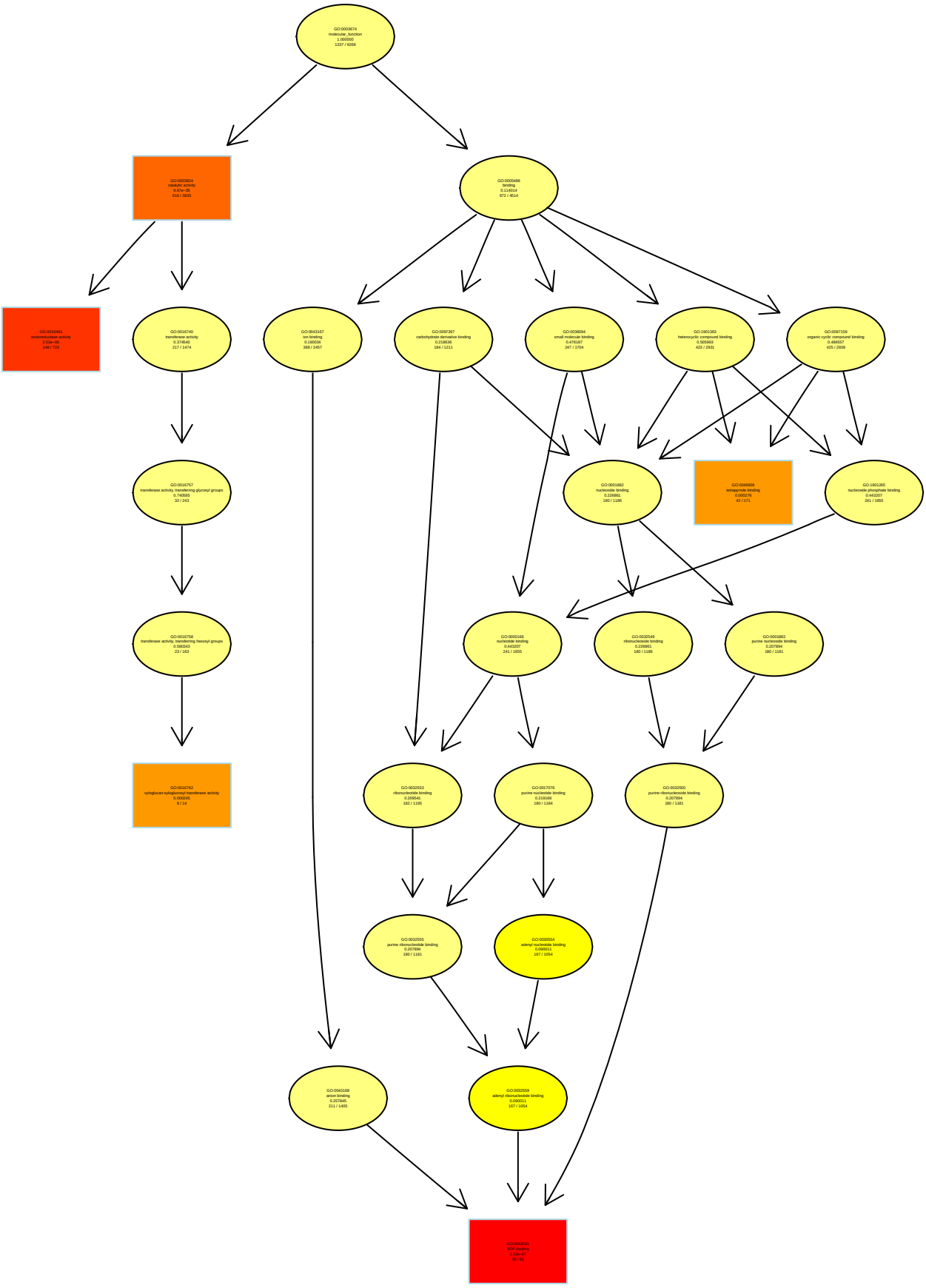
GO:0003674
molecular_function
1.000000
1327 / 9269
GO:0003824
catalytic activity
9.87e-05
616 / 3835
GO:0005488
binding
0.114014
672 / 4514
GO:0016491
oxidoreductase activity
2.52e-06
146 / 721
GO:0016740
transferase activity
0.378540
217 / 1474
GO:0043167
ion binding
0.190034
368 / 2457
GO:0097367
carbohydrate derivative binding
0.218836
184 / 1211
GO:0036094
small molecule binding
0.476187
247 / 1704
GO:1901363
heterocyclic compound binding
0.505993
423 / 2831
GO:0097159
organic cyclic compound binding
0.484657
425 / 2839
GO:0016757
transferase activity, transferring glycosyl groups
0.740565
32 / 243
GO:0001882
nucleoside binding
0.226861
180 / 1186
GO:0046906
tetrapyrrole binding
0.003276
42 / 171
GO:1901265
nucleoside phosphate binding
0.443207
241 / 1655
GO:0016758
transferase activity, transferring hexosyl groups
0.582043
23 / 163
GO:0000166
nucleotide binding
0.443207
241 / 1655
GO:0032549
ribonucleoside binding
0.226861
180 / 1186
GO:0001883
purine nucleoside binding
0.207894
180 / 1181
GO:0016762
xyloglucan:xyloglucosyl transferase activity
0.000245
8 / 14
GO:0032553
ribonucleotide binding
0.206541
182 / 1195
GO:0017076
purine nucleotide binding
0.219166
180 / 1184
GO:0032550
purine ribonucleoside binding
0.207894
180 / 1181
GO:0032555
purine ribonucleotide binding
0.207894
180 / 1181
GO:0030554
adenyl nucleotide binding
0.198011
167 / 1054
GO:0043168
anion binding
0.257845
211 / 1405
GO:0032559
adenyl ribonucleotide binding
0.198011
167 / 1054
GO:0043531
ADP binding
2.52e-07
30 / 61