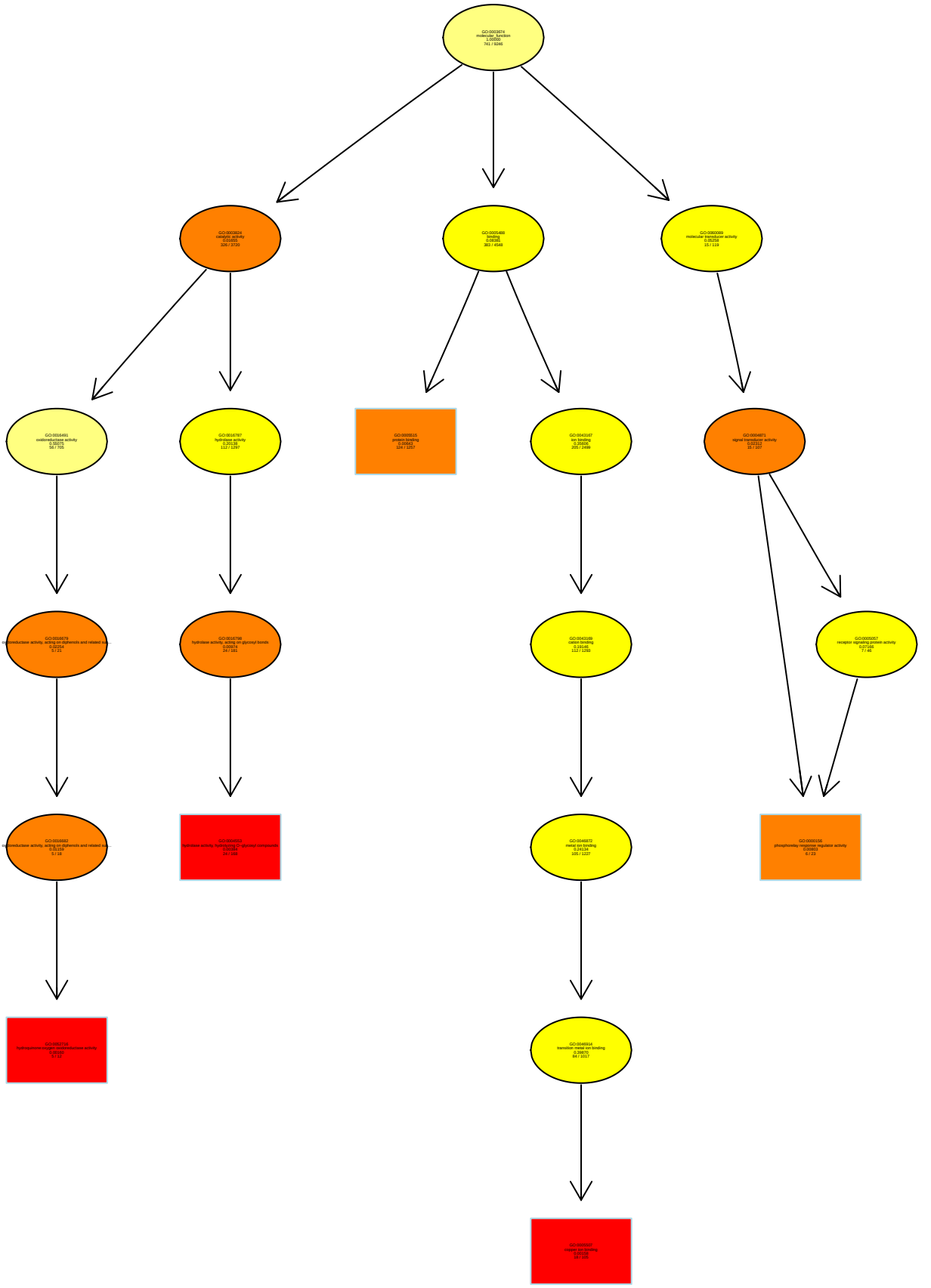
GO:0003874
molecular function
1.00000
741 / 9246
GO:0003824
catalytic activity
0.02655
306 / 3720
GO:0005488
binding
0.06361
363 / 4548
GO:0060089
molecular transducer activity
0.05298
15 / 119
GO:0016491
oxidoreductase activity
0.05075
56 / 705
GO:0016787
hydrolase activity
0.20139
112 / 1297
GO:0005515
protein binding
0.00643
124 / 1257
GO:0043167
ion binding
0.35606
205 / 2499
GO:0004871
signal transducer activity
0.02317
15 / 107
GO:0016679
oxidoreductase activity, acting on diphenols and related sub...
0.02254
5 / 21
GO:0016798
hydrolase activity, acting on glycosyl bonds
0.00974
24 / 181
GO:0043169
cation binding
0.19146
112 / 1293
GO:0005057
receptor signaling protein activity
0.07166
7 / 46
GO:0016682
oxidoreductase activity, acting on diphenols and related sub...
0.00159
5 / 18
GO:0004553
hydrolase activity, hydrolyzing O-glycosyl compounds
0.00384
24 / 168
GO:0046872
metal ion binding
0.24134
105 / 1227
GO:0000156
phosphorelay response regulator activity
0.00803
6 / 23
GO:0052716
hydroquinone:oxygen oxidoreductase activity
0.00165
5 / 12
GO:0046914
transition metal ion binding
0.39870
84 / 1017
GO:0005507
copper ion binding
0.00118
12 / 105