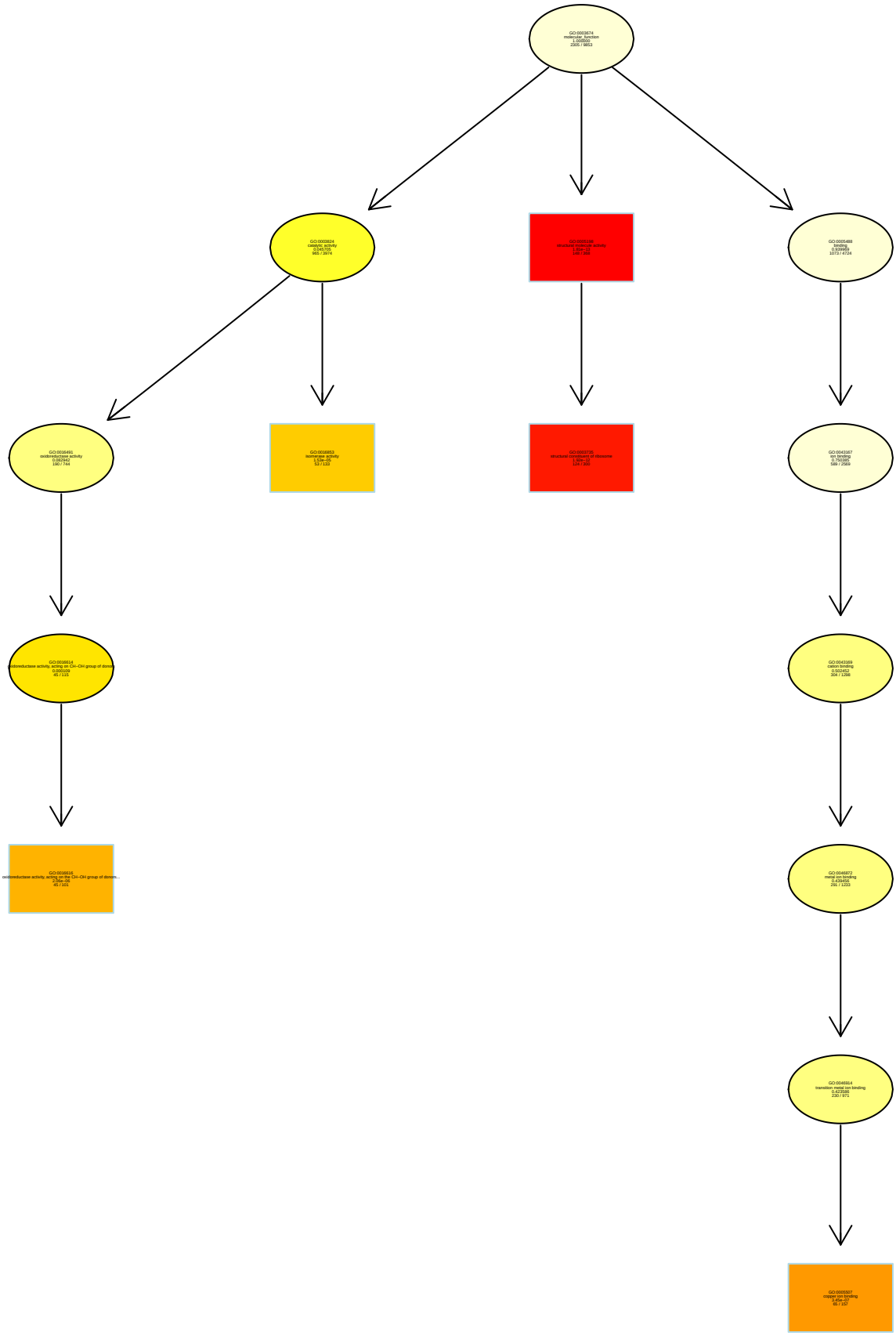
GO:0003674
molecular_function
1.000000
2305 / 9853
GO:0003824
catalytic activity
0.404000
965 / 3974
GO:0005198
structural molecule activity
1.60e-10
148 / 388
GO:0005488
binding
0.339689
1072 / 4724
GO:0016491
oxidoreductase activity
0.025942
190 / 744
GO:0016853
isomerase activity
1.51e-05
53 / 133
GO:0003735
structural constituent of ribosome
1.30e-12
124 / 300
GO:0043167
ion binding
0.705385
588 / 2589
GO:0016614
oxidoreductase activity, acting on CH–OH group of donors
0.000109
45 / 115
GO:0043169
cation binding
0.010452
304 / 1298
GO:0016616
oxidoreductase activity, acting on the CH–OH group of donors, ...
1.90e-05
45 / 103
GO:0046872
metal ion binding
0.438453
291 / 1233
GO:0046914
transition metal ion binding
0.423586
281 / 971
GO:0005507
copper ion binding
3.45e-07
65 / 107