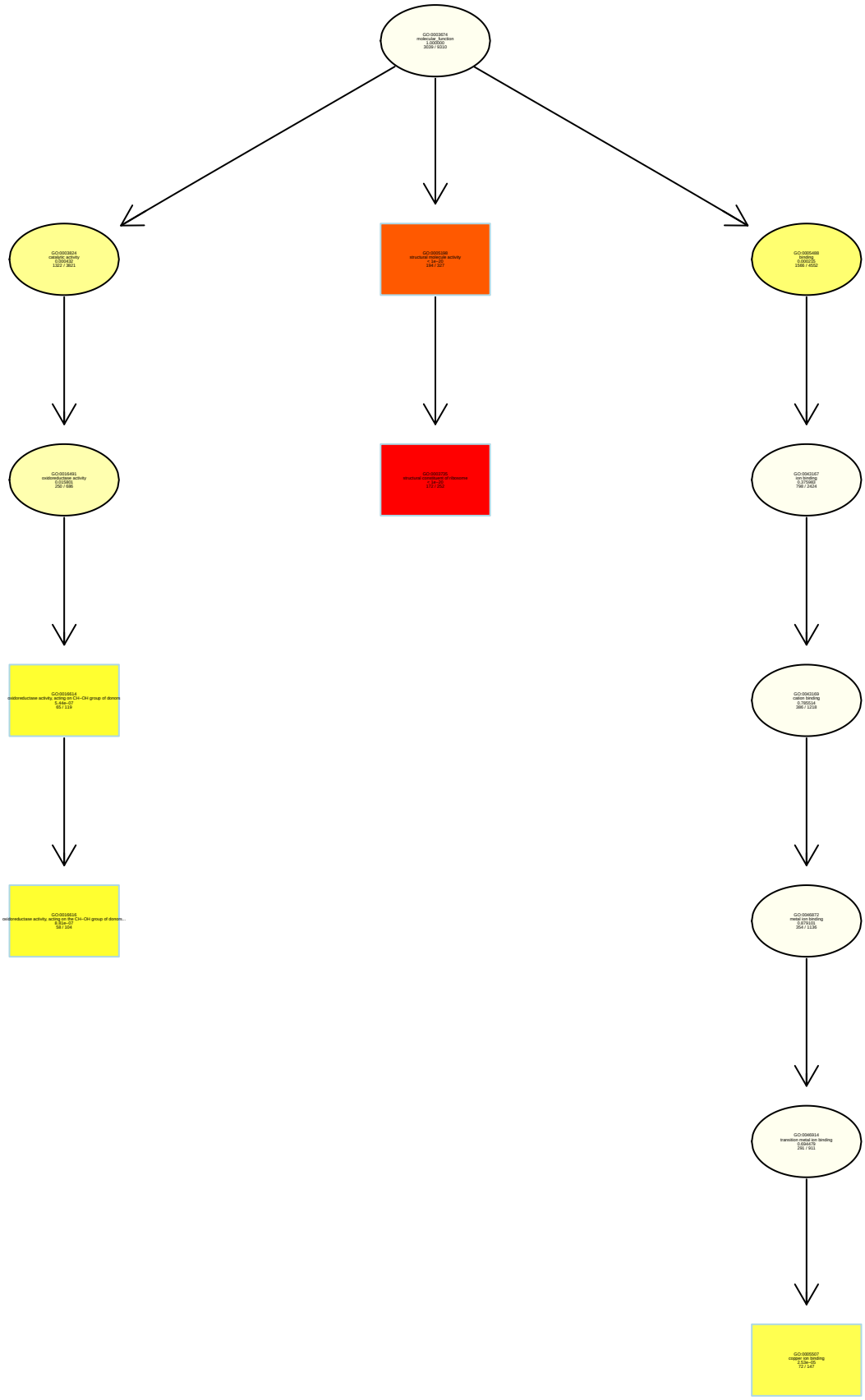
GO:0003674
molecular_function
1.000000
3519 / 9310
GO:0003824
catalytic activity
0.000000
1327 / 3621
GO:0005198
structural molecule activity
0.000000
194 / 227
GO:0005488
binding
0.000000
3168 / 4552
GO:0016491
oxidoreductase activity
0.000000
250 / 686
GO:0003735
structural constituent of ribosome
1.11e-25
172 / 203
GO:0043167
ion binding
0.370982
798 / 2424
GO:0016614
oxidoreductase activity, acting on CH–OH group of donors
5.44e-07
65 / 119
GO:0043169
cation binding
0.765114
386 / 1218
GO:0016616
oxidoreductase activity, acting on the CH–OH group of donors...
8.33e-07
58 / 104
GO:0046872
metal ion binding
0.376102
354 / 1136
GO:0046914
transition metal ion binding
0.694478
291 / 911
GO:0005507
copper ion binding
2.53e-05
72 / 147