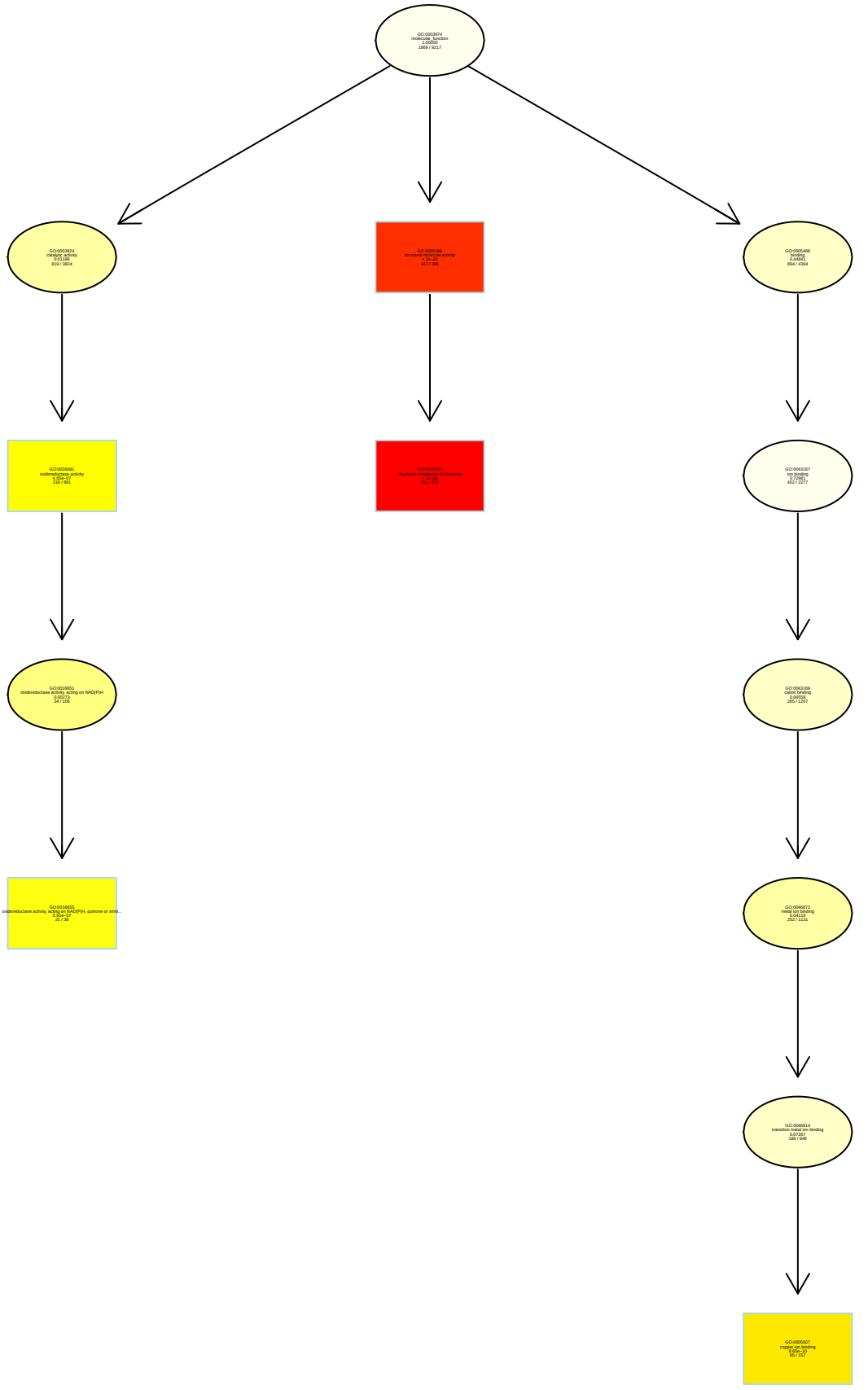
GO:0003674
molecular_function
1.00e-50
1869 / 9217
GO:0003824
catalytic activity
0.01165
819 / 3824
GO:0005198
structural molecule activity
1.71e-25
167 / 385
GO:0005488
binding
0.44845
894 / 4394
GO:0016491
oxidoreductase activity
8.05e-07
218 / 801
GO:0003735
structural constituent of ribosome
1.71e-25
167 / 373
GO:0043167
ion binding
0.72951
452 / 2277
GO:0016651
oxidoreductase activity, acting on NAD(P)H
0.00272
34 / 106
GO:0043169
cation binding
0.06058
265 / 1207
GO:0016655
oxidoreductase activity, acting on NAD(P)H, quinone or simil...
0.00272
21 / 36
GO:0046872
metal ion binding
0.00001
252 / 1131
GO:0046914
transition metal ion binding
0.07267
169 / 866
GO:0005507
copper ion binding
6.65e-10
65 / 107