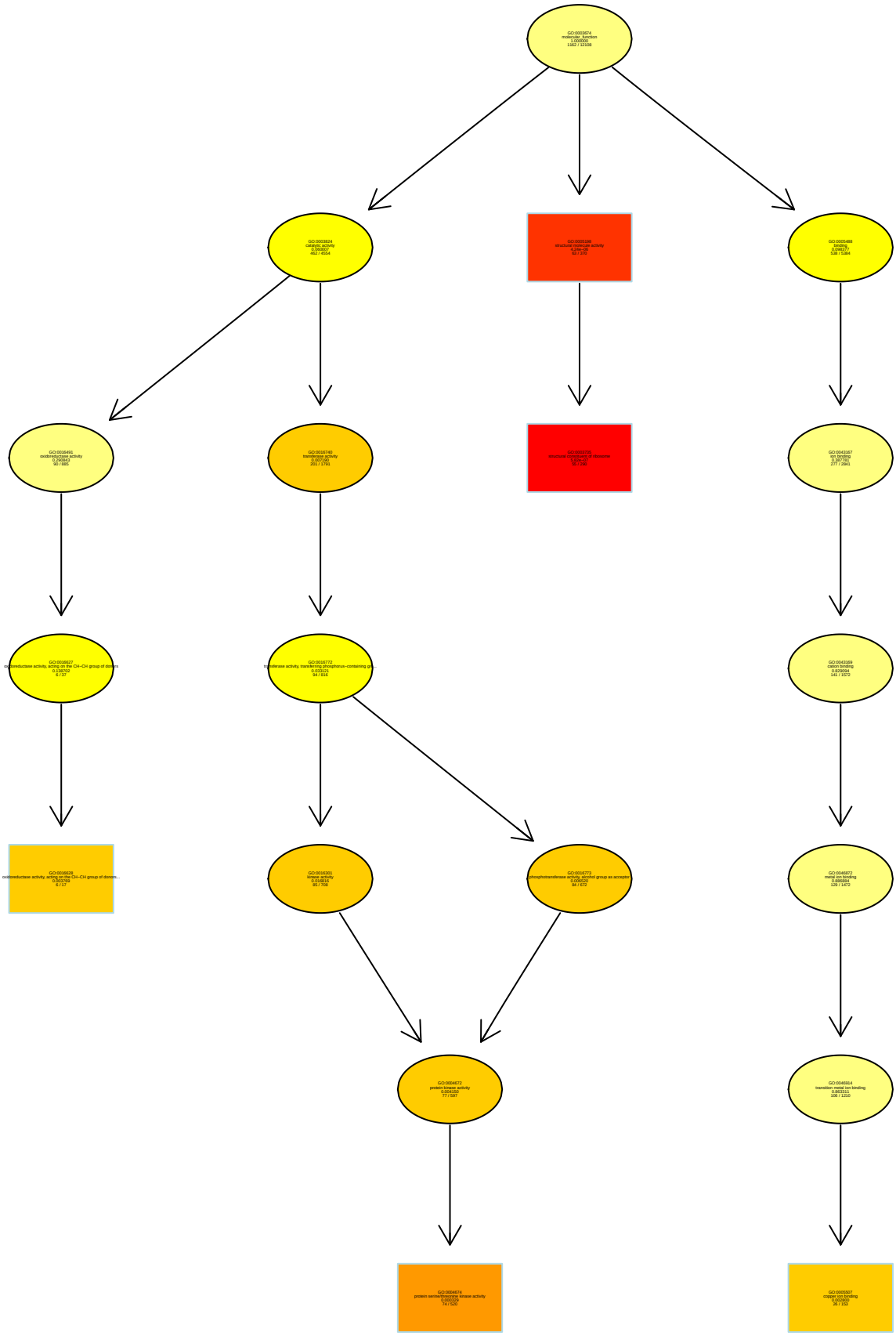
GO:0003674
molecular_function
0.000000
1357 / 12108
GO:0003824
catalytic activity
0.000000
462 / 4554
GO:0005198
structural molecule activity
0.000000
53 / 255
GO:0005488
binding
0.000000
539 / 5364
GO:0016491
oxidoreductase activity
0.250943
90 / 685
GO:0016740
transferase activity
0.007190
201 / 1791
GO:0003735
structural constituent of ribosome
0.00e-07
34 / 202
GO:0043167
ion binding
0.397791
277 / 2861
GO:0016627
oxidoreductase activity, acting on the CH–CH group of donors
0.138702
0 / 17
GO:0016772
transferase activity, transferring phosphorus-containing gr...
0.033121
94 / 816
GO:0043169
cation binding
0.329264
141 / 1572
GO:0016628
oxidoreductase activity, acting on the CH–CH group of donors...
0.138702
0 / 17
GO:0016301
kinase activity
0.016816
85 / 708
GO:0016773
phosphotransferase activity, alcohol group as acceptor
0.006000
94 / 672
GO:0046872
metal ion binding
0.466866
129 / 1472
GO:0004672
protein kinase activity
0.004150
77 / 597
GO:0046914
transition metal ion binding
0.463311
108 / 1210
GO:0004674
protein serine/threonine kinase activity
0.000329
74 / 520
GO:0005507
copper ion binding
0.003800
26 / 102