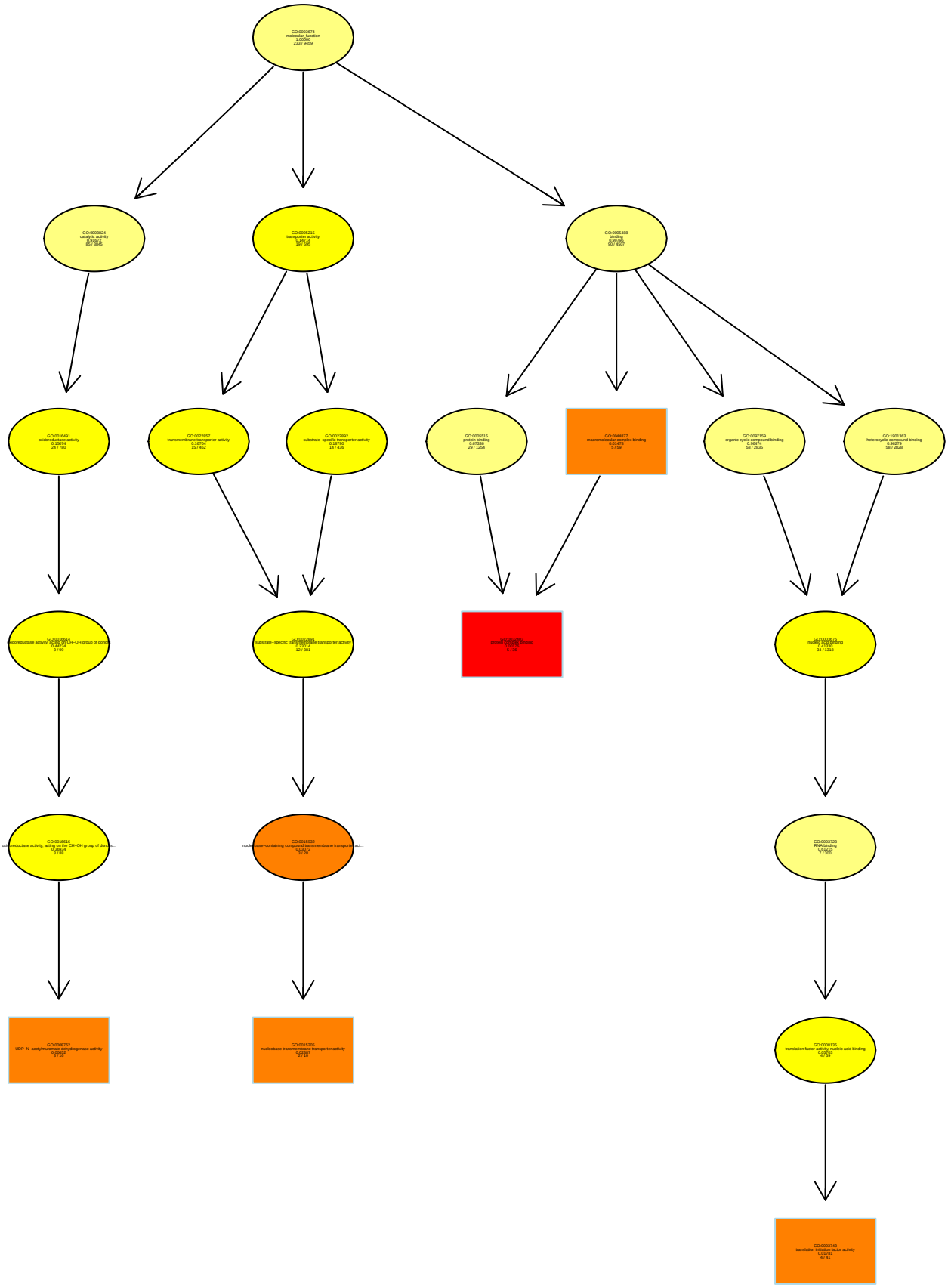
GO:0003674
molecular_function
1.00000
232 / 9959
GO:0003824
catalytic activity
0.61215
85 / 3845
GO:0005215
transporter activity
0.07000
14 / 705
GO:0005488
binding
0.68258
90 / 4027
GO:0016491
oxidoreductase activity
0.19270
24 / 760
GO:0022857
transmembrane transporter activity
0.14162
20 / 482
GO:0022892
substrate-specific transporter activity
0.13730
14 / 436
GO:0005515
protein binding
0.67156
29 / 1254
GO:0044877
macromolecular complex binding
0.01478
5 / 59
GO:0097159
organic cyclic compound binding
0.60474
58 / 2635
GO:1901363
heterocyclic compound binding
0.60275
58 / 2628
GO:0016614
oxidoreductase activity, acting on CH-OH group of donors
0.48239
3 / 99
GO:0022891
substrate-specific transmembrane transporter activity
0.29014
12 / 381
GO:0003676
nucleic acid binding
0.41230
34 / 1318
GO:0016616
oxidoreductase activity, acting on the CH-OH group of donors...
0.38018
3 / 88
GO:0015932
nucleobase-containing compound transmembrane transporter act...
0.05010
2 / 28
GO:0003723
RNA binding
0.21215
7 / 300
GO:0008762
UDP-N-acetylmuramate dehydrogenase activity
0.09652
2 / 19
GO:0015205
nucleobase transmembrane transporter activity
0.09397
2 / 18
GO:0008135
translation factor activity, nucleic acid binding
0.02304
4 / 59
GO:0003743
translation initiation factor activity
0.01781
4 / 41