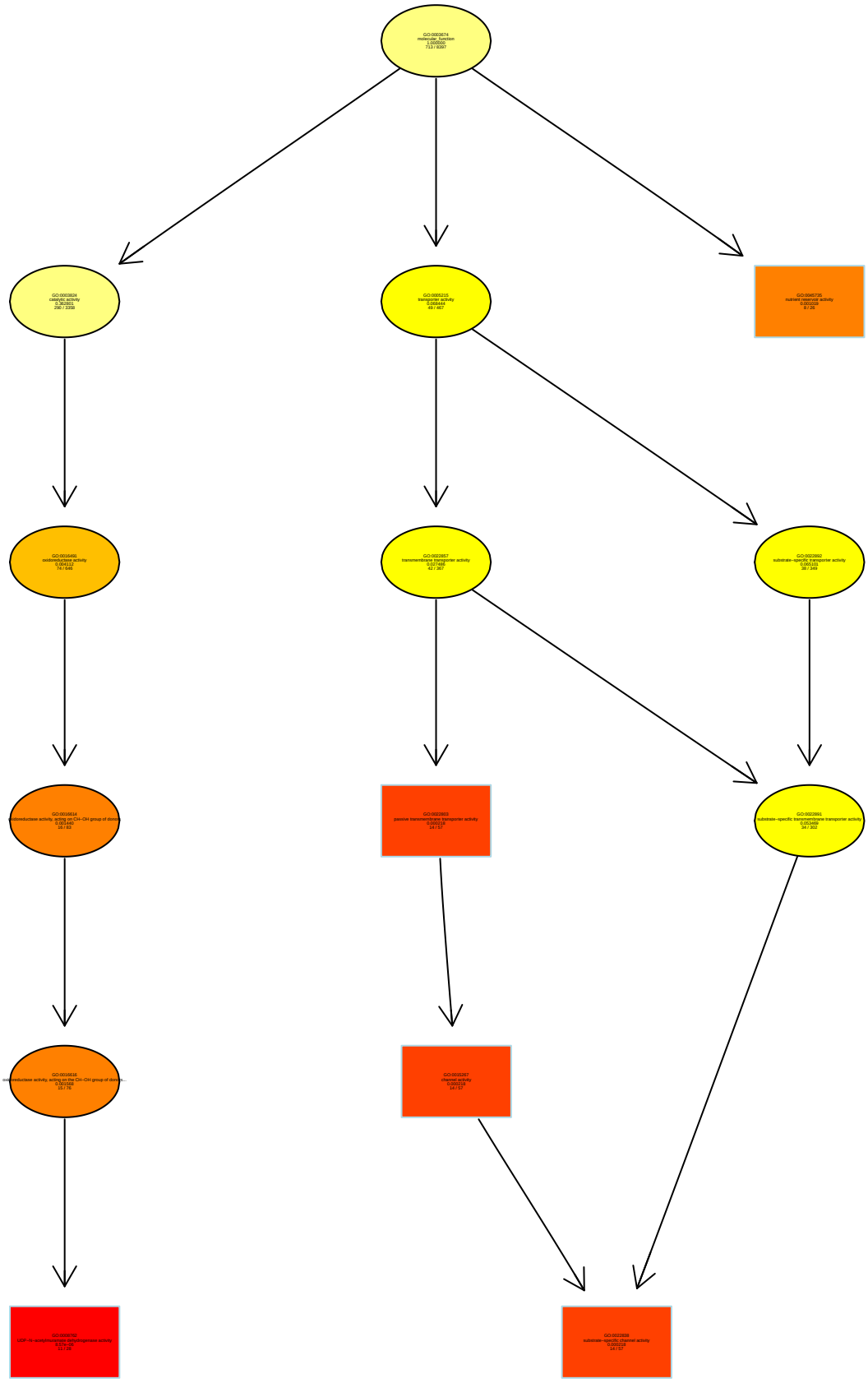
GO:0003674
molecular_function
1.000000
7131 / 8287
GO:0003824
catalytic activity
0.362000
290 / 3358
GO:0005215
transporter activity
0.058400
49 / 667
GO:0045735
nutrient reservoir activity
0.000218
8 / 26
GO:0016491
oxidoreductase activity
0.124000
79 / 648
GO:0022857
transmembrane transporter activity
0.058400
42 / 667
GO:0022892
substrate-specific transporter activity
0.053400
38 / 349
GO:0016614
oxidoreductase activity, acting on CH-OH group of donors
0.053400
16 / 63
GO:0022803
passive transmembrane transporter activity
0.000218
14 / 57
GO:0022891
substrate-specific transmembrane transporter activity
0.053400
34 / 302
GO:0016616
oxidoreductase activity, acting on the CH-OH group of donors, ...
0.024200
15 / 76
GO:0015267
channel activity
0.000218
14 / 57
GO:0008792
UDP-N-acetylmuramate dehydrogenase activity
0.E2e-50
5 / 7 26
GO:0022838
substrate-specific channel activity
0.000218
14 / 57