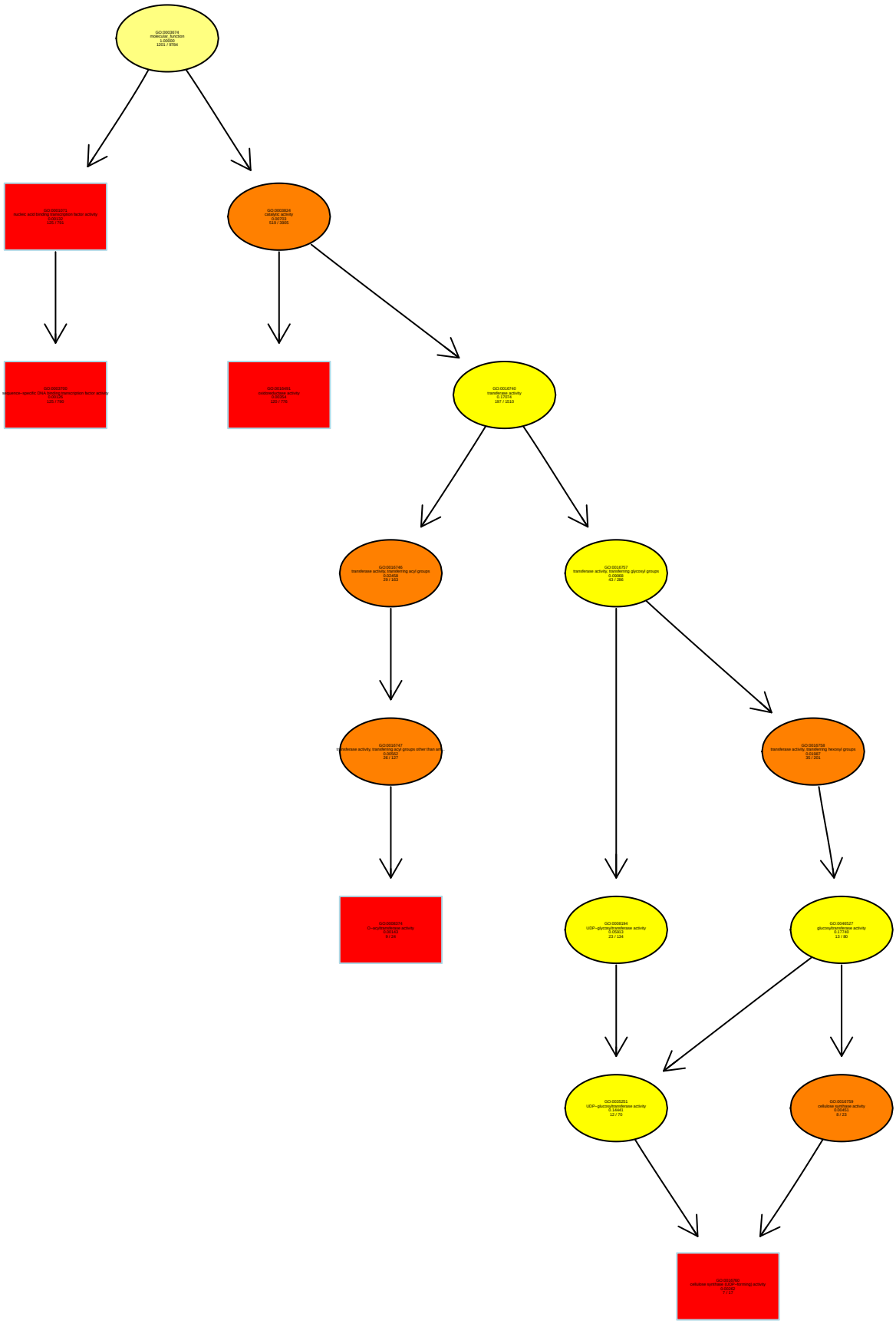
GO:0003674
molecular_function
1.00000
1201 / 9784
GO:0001071
nucleic acid binding transcription factor activity
0.00132
120 / 791
GO:0003824
catalytic activity
0.00703
519 / 3955
GO:0003700
sequence-specific DNA binding transcription factor activity
0.00132
120 / 790
GO:0016491
oxidoreductase activity
0.00204
120 / 776
GO:0016740
transferase activity
0.12674
197 / 1510
GO:0016746
transferase activity, transferring acyl groups
0.02092
29 / 163
GO:0016757
transferase activity, transferring glycosyl groups
0.07066
43 / 288
GO:0016747
transferase activity, transferring acyl groups other than amino-acyl groups
0.00652
26 / 127
GO:0016769
transferase activity, transferring hexosyl groups
0.01987
25 / 101
GO:0008274
C-acylltransferase activity
0.00149
9 / 24
GO:0008194
UDP-glycosyltransferase activity
0.05812
23 / 124
GO:0046527
glucosyltransferase activity
0.17730
13 / 80
GO:0035251
UDP-glucosyltransferase activity
0.14441
12 / 70
GO:0016759
cellulose synthase activity
0.04451
8 / 23
GO:0016760
cellulose synthase (UDP-forming) activity
0.02067
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