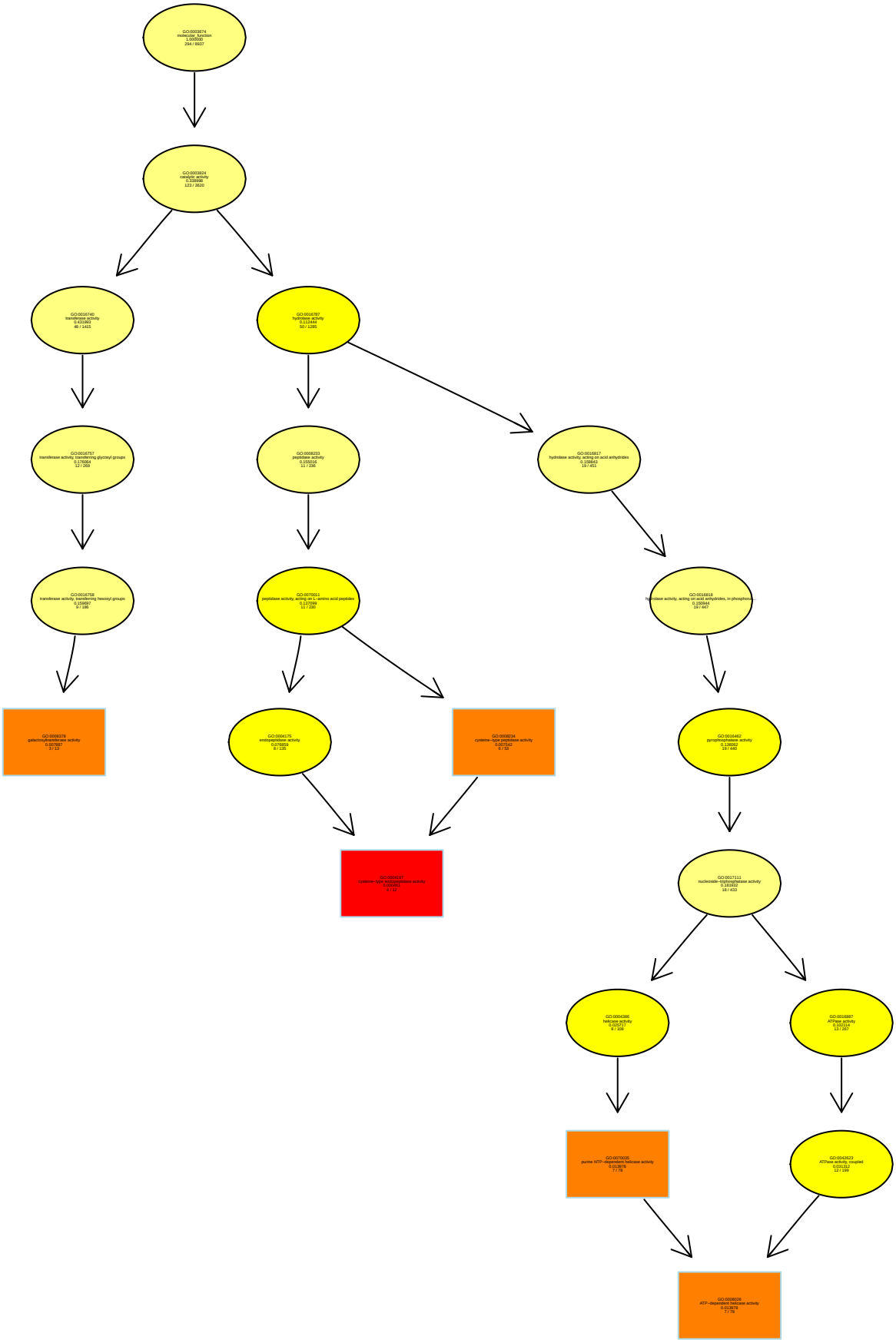
GO:0003674
molecular_function
1.000000
294 / 8937
GO:0003824
catalytic activity
0.938808
123 / 3620
GO:0016740
transferase activity
0.431983
48 / 1415
GO:0016787
hydrolase activity
0.112444
50 / 1285
GO:0016757
transferase activity, transferring glycosyl groups
0.176064
12 / 269
GO:0008233
peptidase activity
0.155016
11 / 226
GO:0016817
hydrolase activity, acting on acid anhydrides
0.159843
19 / 451
GO:0016758
transferase activity, transferring hexosyl groups
0.156657
9 / 188
GO:0070011
peptidase activity, acting on L-amino acid peptides
0.137099
11 / 220
GO:0016818
hydrolase activity, acting on acid anhydrides, in phosphorus-
0.150944
18 / 447
GO:0008378
galactosyltransferase activity
0.007667
2 / 13
GO:0004175
endopeptidase activity
0.076859
6 / 135
GO:0008234
cysteine-type peptidase activity
0.067542
6 / 53
GO:0016462
pyrophosphatase activity
0.136062
18 / 440
GO:0004197
cysteine-type endopeptidase activity
0.044001
2 / 11
GO:0017111
nucleoside-triphosphatase activity
0.136062
18 / 432
GO:0004386
helicase activity
0.013878
6 / 108
GO:0016887
ATPase activity
0.151152
12 / 267
GO:0070035
purine NTP-dependent helicase activity
0.013878
7 / 78
GO:0042623
ATPase activity, coupled
0.151152
12 / 198
GO:0008026
ATP-dependent helicase activity
0.013878
7 / 78