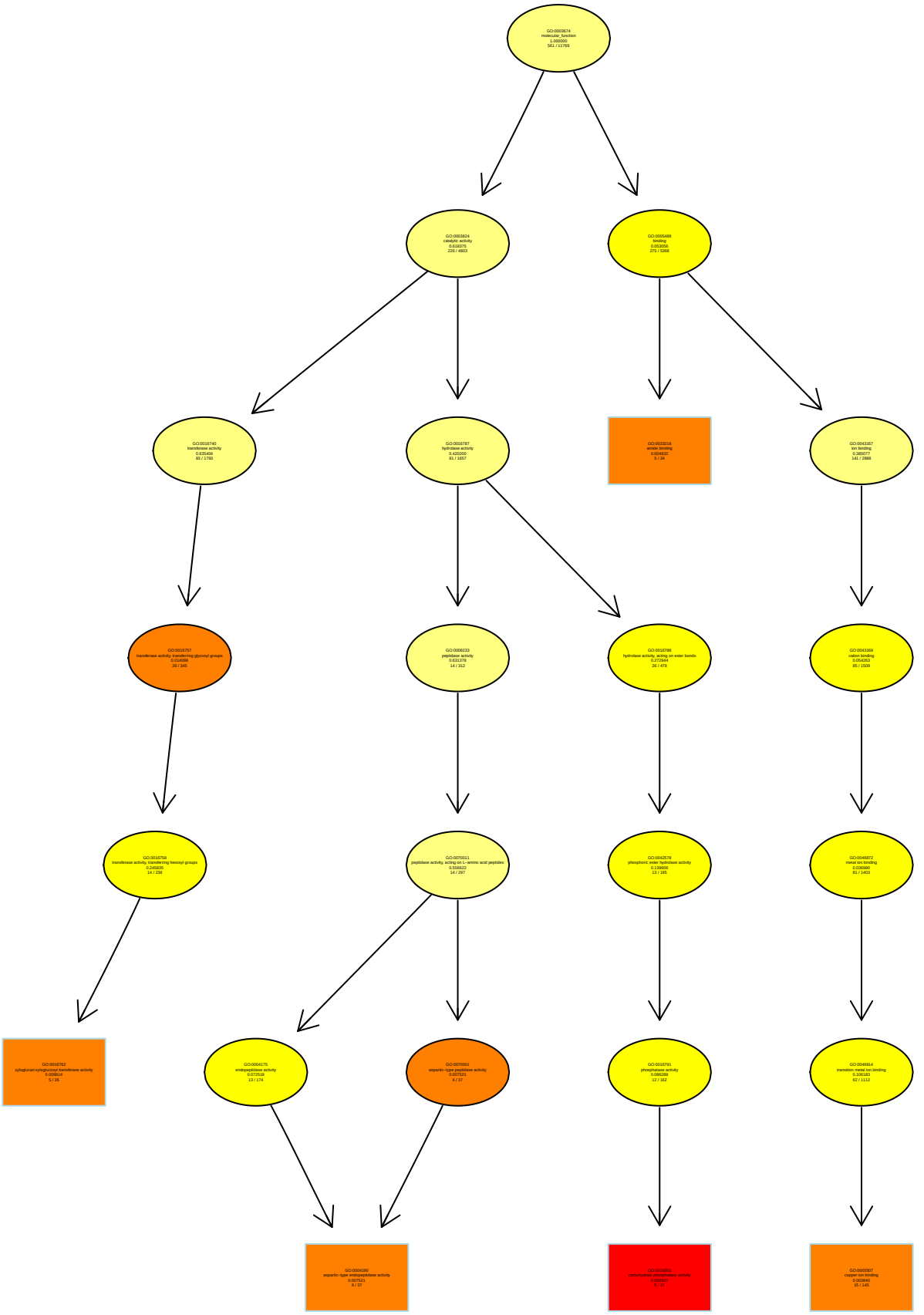
GO:0003674
molecular_function
1.000000
561 / 11769
GO:0003824
catalytic activity
0.618375
226 / 4663
GO:0005488
binding
0.053056
275 / 5368
GO:0016740
transferase activity
0.535494
83 / 1793
GO:0016787
hydrolase activity
0.435200
81 / 1657
GO:0033218
amide binding
0.004822
5 / 24
GO:0043167
ion binding
0.365077
141 / 2868
GO:0016757
transferase activity, transferring glycosyl groups
0.104088
26 / 345
GO:0008233
peptidase activity
0.631378
14 / 312
GO:0016788
hydrolase activity, acting on ester bonds
0.272944
26 / 479
GO:0043169
cation binding
0.054263
85 / 1509
GO:0016758
transferase activity, transferring hexosyl groups
0.240205
14 / 238
GO:0070011
peptidase activity, acting on L-amino acid peptides
0.156822
14 / 297
GO:0042578
phosphoric ester hydrolase activity
0.139606
13 / 265
GO:0046872
metal ion binding
0.039560
81 / 1403
GO:0016762
xyloglucan:xyloglucosyl transferase activity
0.009914
5 / 26
GO:0004175
endopeptidase activity
0.072516
13 / 174
GO:0070001
aspartic-type peptidase activity
0.007521
6 / 37
GO:0016791
phosphatase activity
0.088288
12 / 182
GO:0046914
transition metal ion binding
0.106193
62 / 1132
GO:0004190
aspartic-type endopeptidase activity
0.007521
6 / 37
GO:0016820
carbohydrate phosphatase activity
0.000927
5 / 37
GO:0005507
copper ion binding
0.003840
10 / 145