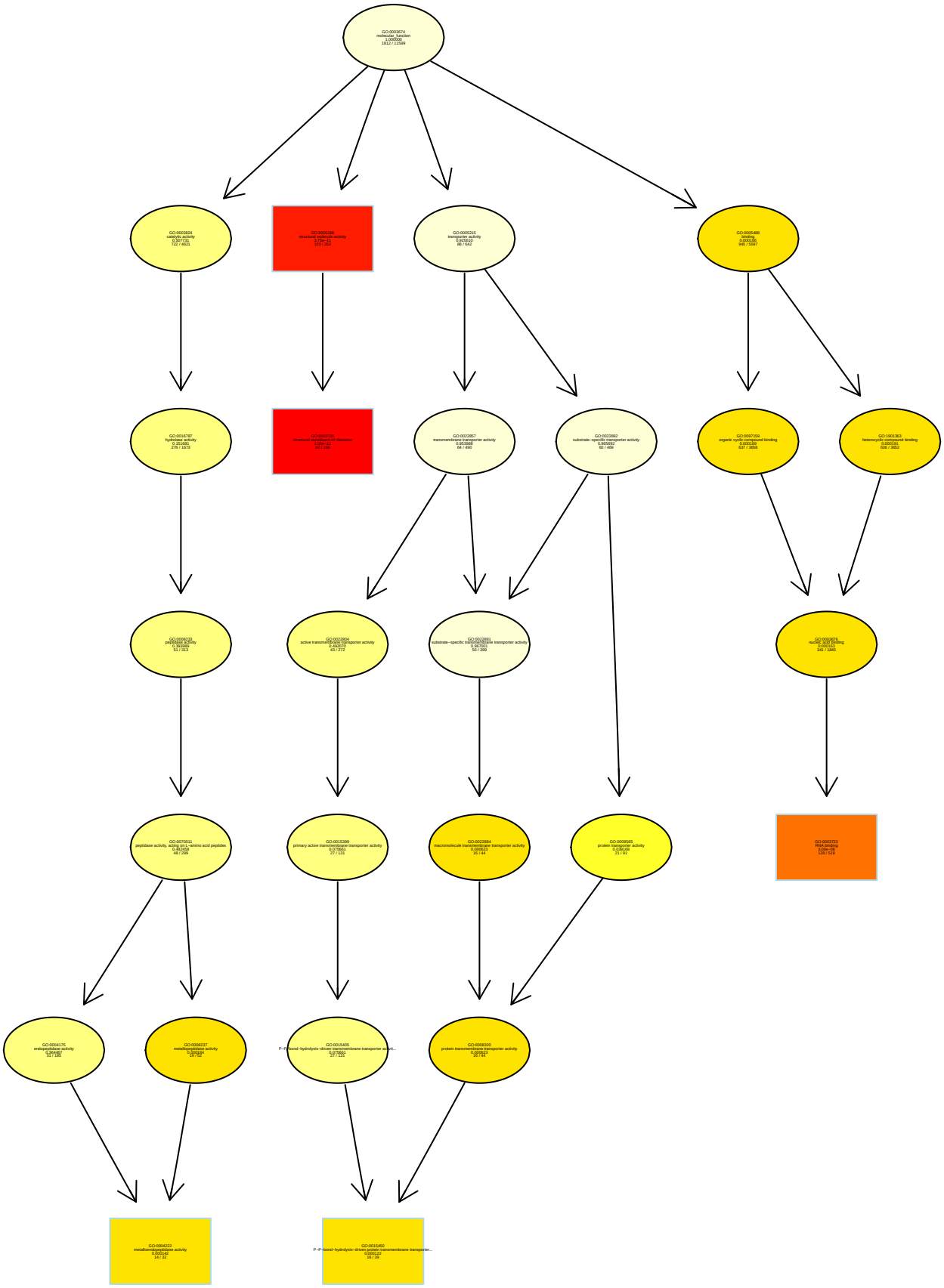
GO:0003674
molecular_function
1.000000
1612 / 11599
GO:0003824
catalytic activity
0.000000
722 / 4621
GO:0005198
structural molecule activity
0.000000
163 / 652
GO:0005215
transporter activity
0.000000
58 / 642
GO:0005488
binding
0.000000
846 / 5597
GO:0016787
hydrolase activity
0.351651
276 / 1673
GO:0003735
structural constituent of ribosome
0.000000
57 / 316
GO:0022857
transmembrane transporter activity
0.953868
64 / 490
GO:0022892
substrate-specific transporter activity
0.945032
50 / 469
GO:0097159
organic cyclic compound binding
0.000189
637 / 3658
GO:1901363
heterocyclic compound binding
0.000151
638 / 3652
GO:0008233
peptidase activity
0.353989
51 / 313
GO:0022804
active transmembrane transporter activity
0.450070
43 / 272
GO:0022891
substrate-specific transmembrane transporter activity
0.567001
50 / 299
GO:0003676
nucleic acid binding
0.000163
341 / 1845
GO:0070011
peptidase activity, acting on L-amino acid peptides
0.341454
48 / 299
GO:0015399
primary active transmembrane transporter activity
0.075661
27 / 131
GO:0022884
macromolecule transmembrane transporter activity
0.000823
16 / 44
GO:0008565
protein transporter activity
0.038108
21 / 91
GO:0003723
RNA binding
0.000000
128 / 515
GO:0004175
endopeptidase activity
0.264482
37 / 165
GO:0008237
metallopeptidase activity
0.309184
19 / 62
P-P-bond-hydrolysis-driven transmembrane transporter acti...
GO:0015405
0.075661
27 / 131
GO:0008320
protein transmembrane transporter activity
0.000213
16 / 44
GO:0004222
metalloendopeptidase activity
0.500142
14 / 32
P-P-bond-hydrolysis-driven protein transmembrane transporter...
GO:0015450
0.500122
16 / 28