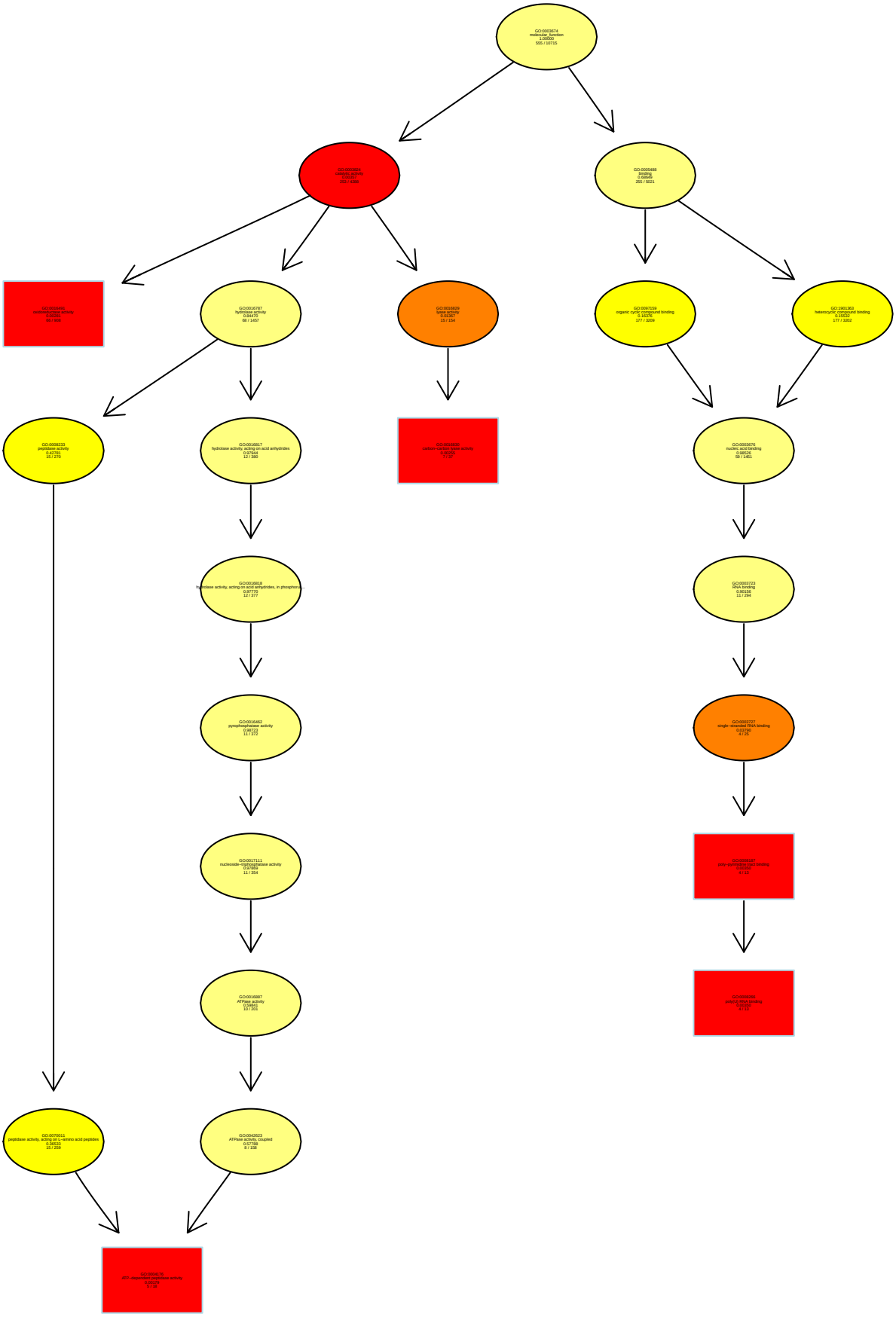
GO:0003674
molecular_function
1.00000
555 / 10715
GO:0003824
catalytic activity
0.99000
253 / 4268
GO:0005488
binding
0.68649
255 / 5021
GO:0016491
oxidoreductase activity
0.00181
56 / 906
GO:0016787
hydrolase activity
0.84470
88 / 1457
GO:0016829
lyase activity
0.61367
15 / 154
GO:0097159
organic cyclic compound binding
0.16376
177 / 3209
GO:1901363
heterocyclic compound binding
0.16532
177 / 3202
GO:0008233
peptidase activity
0.42781
10 / 270
GO:0016817
hydrolase activity, acting on acid anhydrides
0.97944
12 / 380
GO:0016830
carbon-carbon lyase activity
0.00250
7 / 27
GO:0003676
nucleic acid binding
0.98526
59 / 1451
GO:0016818
hydrolase activity, acting on acid anhydrides, in phosphorus-...
0.97770
12 / 377
GO:0003723
RNA binding
0.96156
11 / 294
GO:0016462
pyrophosphatase activity
0.96723
11 / 372
GO:0003727
single-stranded RNA binding
0.03790
4 / 15
GO:0017111
nucleoside-triphosphatase activity
0.96553
11 / 364
GO:0008187
poly-pyrimidine tract binding
0.00330
4 / 12
GO:0016887
ATPase activity
0.96553
12 / 201
GO:0008266
poly(U) RNA binding
0.00330
4 / 12
GO:0070011
peptidase activity, acting on L-amino acid peptides
0.38533
13 / 258
GO:0042623
ATPase activity, coupled
0.57788
6 / 158
GO:0004176
ATP-dependent peptidase activity
0.00279
5 / 18