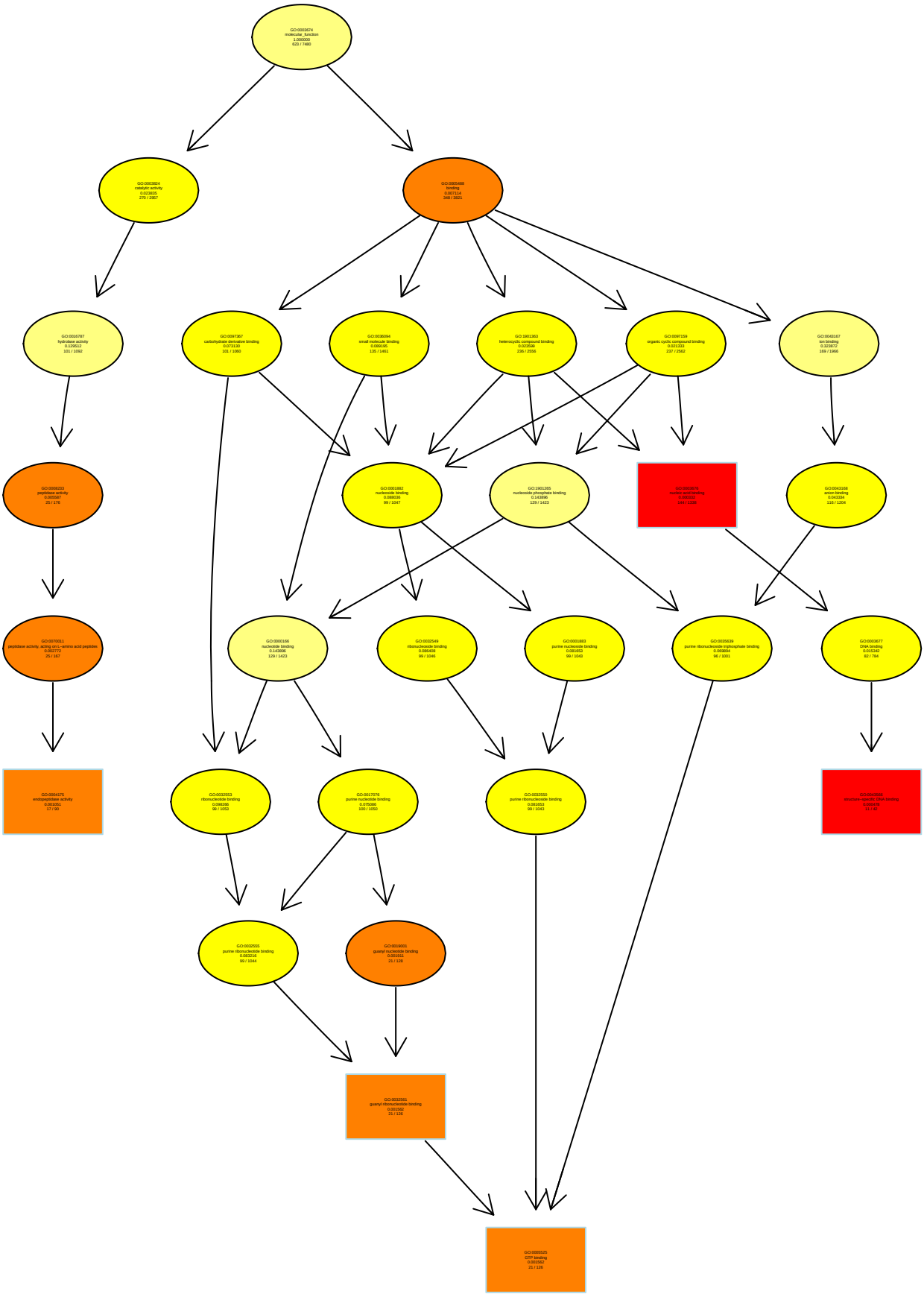
GO:0003674
molecular_function
1.000000
623 / 7480
GO:0003824
catalytic activity
0.433835
270 / 2957
GO:0005488
binding
0.607114
348 / 3821
GO:0016787
hydrolase activity
0.129512
101 / 1092
GO:0097367
carbohydrate derivative binding
0.073130
101 / 1060
GO:0036094
small molecule binding
0.088195
135 / 1461
GO:1901363
heterocyclic compound binding
0.022556
236 / 2556
GO:0097159
organic cyclic compound binding
0.021333
237 / 2562
GO:0043167
ion binding
0.323872
169 / 1966
GO:0008233
peptidase activity
0.005287
25 / 176
GO:0001882
nucleoside binding
0.088036
96 / 1047
GO:1901265
nucleoside phosphate binding
0.143896
129 / 1423
GO:0003676
nucleic acid binding
0.093332
144 / 1538
GO:0043168
anion binding
0.043234
116 / 1204
GO:0070011
peptidase activity, acting on L-amino acid peptides
0.002772
25 / 167
GO:0000166
nucleotide binding
0.143896
129 / 1423
GO:0032549
ribonucleoside binding
0.086408
99 / 1046
GO:0001883
purine nucleoside binding
0.081603
99 / 1043
GO:0035639
purine ribonucleoside triphosphate binding
0.068804
96 / 1001
GO:0003677
DNA binding
0.015342
82 / 784
GO:0004175
endopeptidase activity
0.001001
17 / 90
GO:0032553
ribonucleotide binding
0.088036
96 / 1053
GO:0017076
purine nucleotide binding
0.070086
100 / 1058
GO:0032550
purine ribonucleoside binding
0.081603
99 / 1043
GO:0043565
sequence-specific DNA binding
0.000476
11 / 23
GO:0032555
purine ribonucleotide binding
0.089216
99 / 1044
GO:0019001
guanyl nucleotide binding
0.001931
21 / 128
GO:0032561
guanyl ribonucleotide binding
0.001562
21 / 126
GO:0005525
GTP binding
0.001562
21 / 126