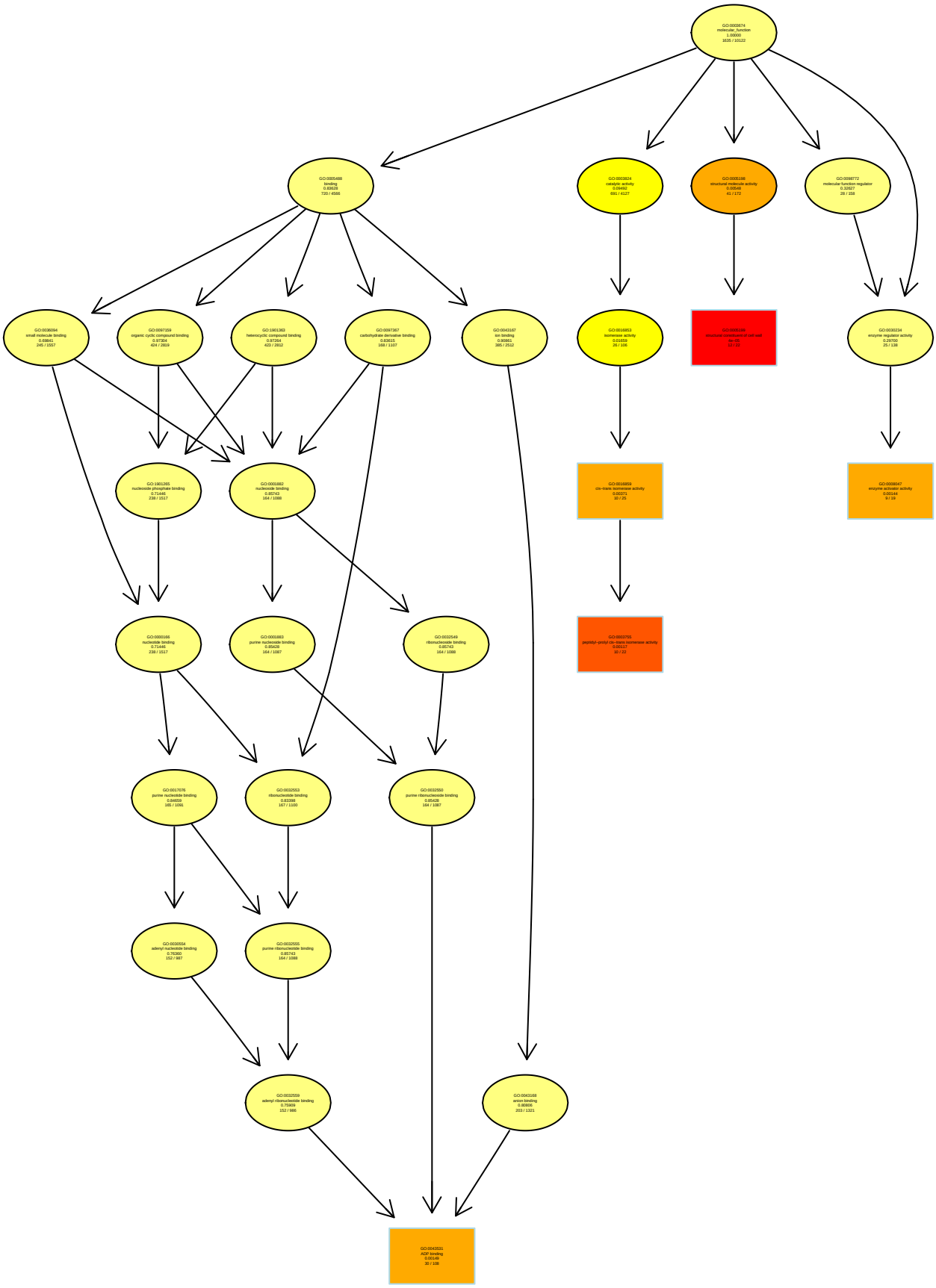
GO:0003674
molecular_function
1.00000
1635 / 10122
GO:0005488
binding
0.93628
720 / 4566
GO:0003824
catalytic activity
0.69492
691 / 4127
GO:0005198
structural molecule activity
0.05048
41 / 172
GO:0098772
molecular function regulator
0.33027
28 / 159
GO:0036094
small molecule binding
0.69841
245 / 1557
GO:0097159
organic cyclic compound binding
0.67284
424 / 2819
GO:1901363
heterocyclic compound binding
0.67264
423 / 2812
GO:0097367
carbohydrate derivative binding
0.63615
168 / 1107
GO:0043167
ion binding
0.90861
385 / 2512
GO:0016853
isomerase activity
0.01656
16 / 106
GO:0005199
structural constituent of cell wall
0.<0.01
0 / 21
GO:0030234
enzyme regulator activity
0.29755
25 / 138
GO:1901265
nucleoside phosphate binding
0.71446
238 / 1517
GO:0001882
nucleoside binding
0.85743
164 / 1088
GO:0016859
cis–trans isomerase activity
0.00371
02 / 25
GO:0008047
enzyme activator activity
0.00144
0 / 19
GO:0000166
nucleotide binding
0.71446
238 / 1517
GO:0001883
purine nucleoside binding
0.85428
164 / 1087
GO:0032549
ribonucleoside binding
0.85743
164 / 1088
GO:0003755
peptidyl–prolyl cis–trans isomerase activity
0.00117
10 / 22
GO:0017076
purine nucleotide binding
0.84659
161 / 1091
GO:0032553
ribonucleotide binding
0.83398
167 / 1100
GO:0032550
purine ribonucleoside binding
0.85428
164 / 1087
GO:0030554
adenyl nucleotide binding
0.76360
152 / 987
GO:0032555
purine ribonucleotide binding
0.85743
164 / 1088
GO:0032559
adenyl ribonucleotide binding
0.75909
152 / 986
GO:0043168
anion binding
0.80606
203 / 1321
GO:0043531
ADP binding
0.00149
02 / 108