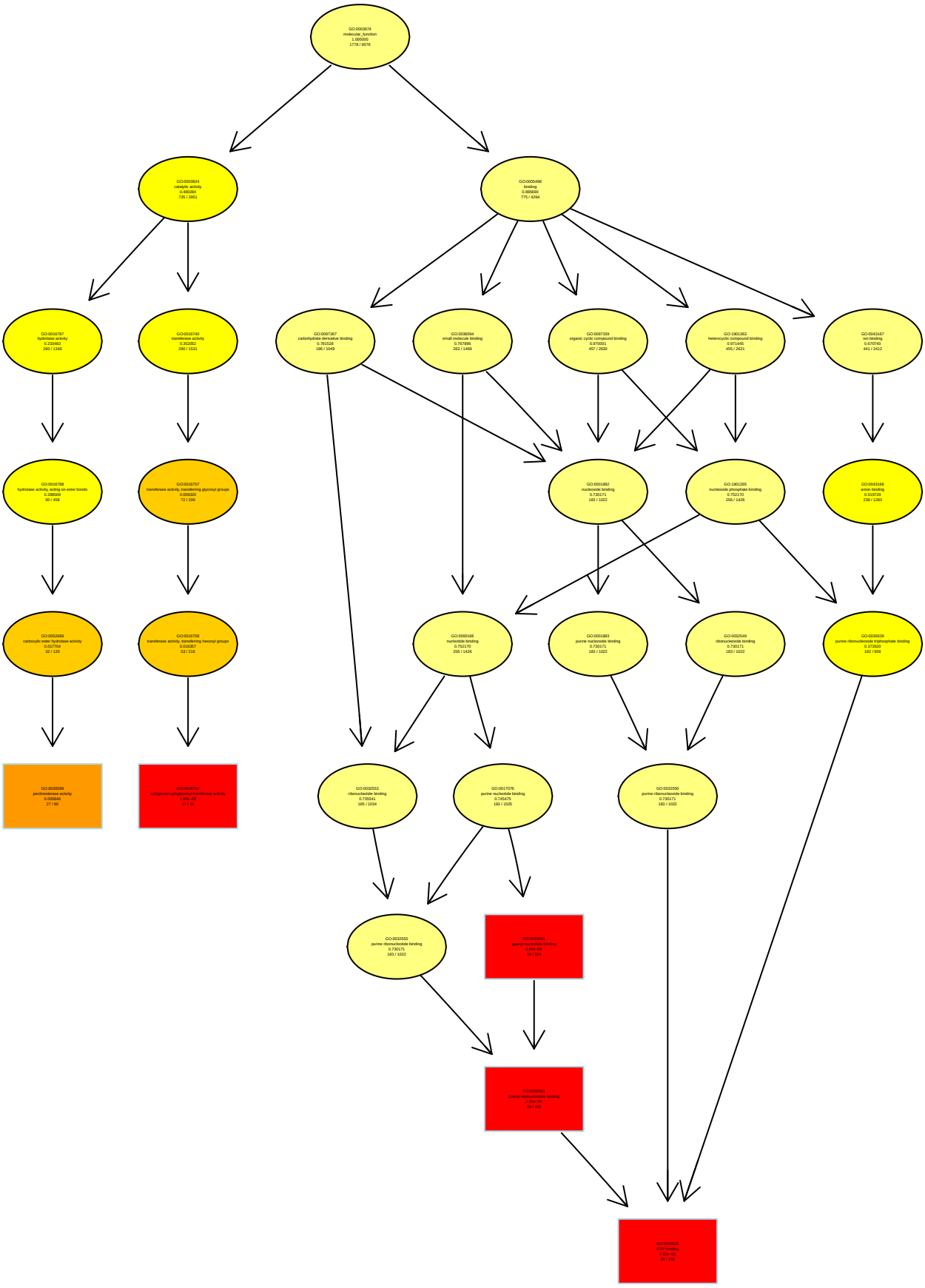
GO:0003674
molecular_function
1.000000
1779 / 9576
GO:0003824
catalytic activity
0.480354
735 / 3951
GO:0005488
binding
0.885590
775 / 4294
GO:0016787
hydrolase activity
0.239463
260 / 1346
GO:0016740
transferase activity
0.352052
290 / 1531
GO:0097367
carbohydrate derivative binding
0.781528
186 / 1049
GO:0036094
small molecule binding
0.767888
263 / 1468
GO:0097159
organic cyclic compound binding
0.970001
457 / 2630
GO:1901363
heterocyclic compound binding
0.971445
455 / 2621
GO:0043167
ion binding
0.670740
461 / 2412
GO:0016788
hydrolase activity, acting on ester bonds
0.288569
90 / 458
GO:0016757
transferase activity, transferring glycosyl groups
0.009320
72 / 299
GO:0001882
nucleoside binding
0.730171
183 / 1022
GO:1901265
nucleoside phosphate binding
0.752170
256 / 1426
GO:0043168
anion binding
0.519729
238 / 1283
GO:0052689
carboxylic ester hydrolase activity
0.027704
32 / 120
GO:0016758
transferase activity, transferring hexosyl groups
0.016287
53 / 216
GO:0000166
nucleotide binding
0.752170
256 / 1426
GO:0001883
purine nucleoside binding
0.730171
183 / 1022
GO:0032549
ribonucleoside binding
0.730171
183 / 1022
GO:0035639
purine ribonucleoside triphosphate binding
0.372820
182 / 958
GO:0030599
pectinesterase activity
0.000846
27 / 80
GO:0016762
xyloglucan:xyloglucosyl transferase activity
1.64e-05
17 / 29
GO:0032553
ribonucleotide binding
0.735541
185 / 1034
GO:0017076
purine nucleotide binding
0.745475
183 / 1025
GO:0032550
purine ribonucleotide binding
0.730171
183 / 1022
GO:0032555
purine ribonucleotide binding
0.730171
183 / 1022
GO:0019001
guanyl nucleotide binding
4.05e-06
29 / 104
GO:0032561
guanyl ribonucleotide binding
2.32e-06
39 / 102
GO:0005525
GTP binding
2.32e-06
39 / 102