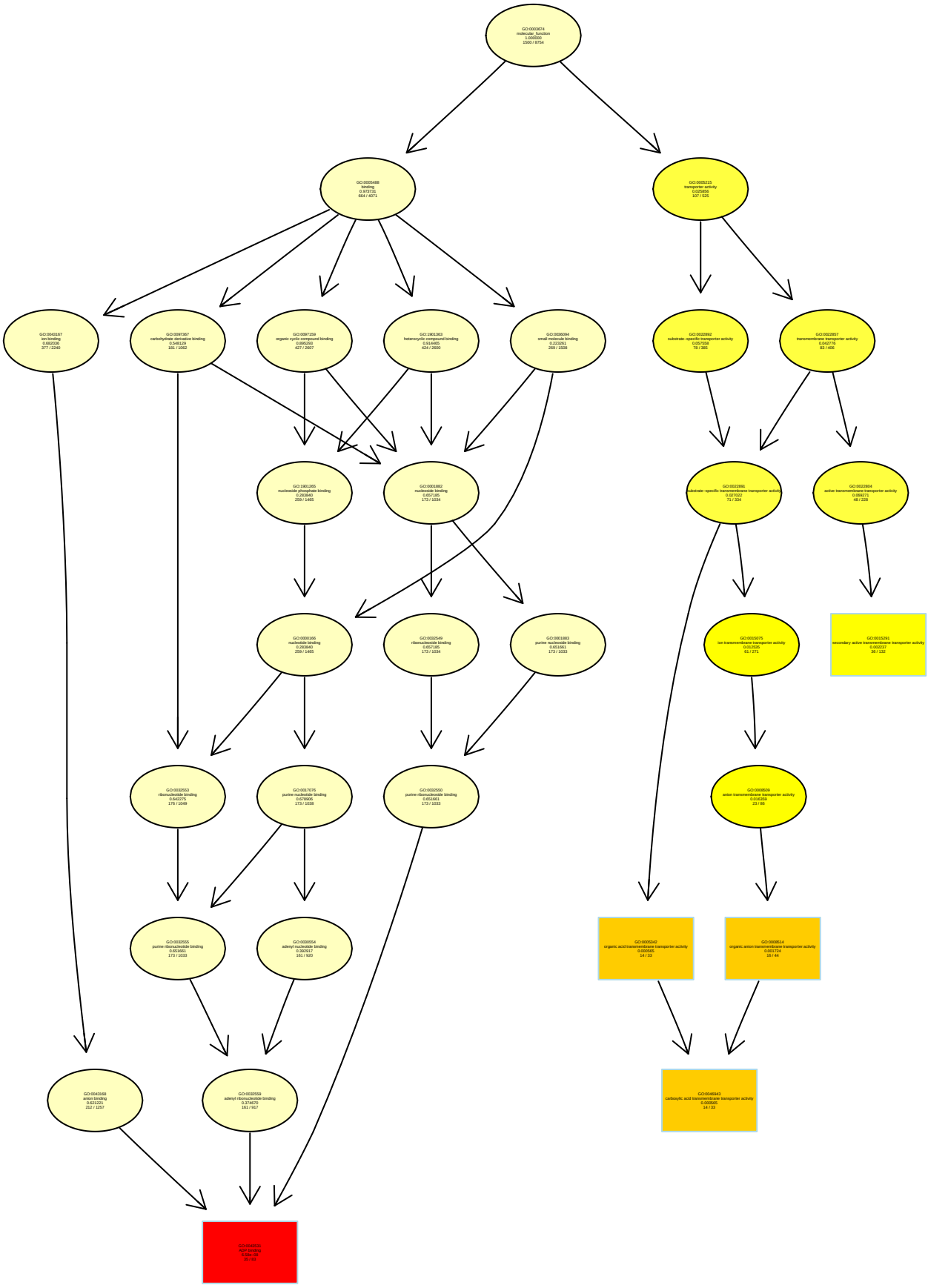
GO:0003674
molecular_function
1.000000
1000 / 8754
GO:0005488
binding
0.973731
664 / 4071
GO:0005215
transporter activity
0.026856
107 / 525
GO:0043167
ion binding
0.660206
377 / 2240
GO:0097367
carbohydrate derivative binding
0.546120
181 / 1082
GO:0097159
organic cyclic compound binding
0.695293
427 / 2607
GO:1901363
heterocyclic compound binding
0.914465
424 / 2600
GO:0036094
small molecule binding
0.223051
269 / 1508
GO:0022892
substrate-specific transporter activity
0.057558
78 / 385
GO:0022857
transmembrane transporter activity
0.042776
83 / 406
GO:1901265
nucleoside phosphate binding
0.283640
259 / 1465
GO:0001882
nucleoside binding
0.657185
173 / 1034
GO:0022891
substrate-specific transmembrane transporter activity
0.027022
71 / 336
GO:0022804
active transmembrane transporter activity
0.069271
48 / 229
GO:0000166
nucleotide binding
0.283640
259 / 1465
GO:0032549
ribonucleoside binding
0.657185
173 / 1034
GO:0001883
purine nucleoside binding
0.651661
173 / 1033
GO:0015075
ion transmembrane transporter activity
0.012926
61 / 271
GO:0015291
secondary active transmembrane transporter activity
0.002227
36 / 132
GO:0032553
ribonucleotide binding
0.642275
176 / 1049
GO:0017076
purine nucleotide binding
0.679906
173 / 1038
GO:0032550
purine ribonucleoside binding
0.651661
173 / 1033
GO:0008509
anion transmembrane transporter activity
0.014534
23 / 86
GO:0032555
purine ribonucleotide binding
0.651661
173 / 1033
GO:0030554
adenyl nucleotide binding
0.392917
161 / 920
GO:0005342
organic acid transmembrane transporter activity
0.008585
14 / 33
GO:0008514
organic anion transmembrane transporter activity
0.013724
16 / 44
GO:0043168
anion binding
0.621221
212 / 1257
GO:0032559
adenyl ribonucleotide binding
0.374670
161 / 917
GO:0046943
carboxylic acid transmembrane transporter activity
0.008585
14 / 33
GO:0005524
ATP binding
0.538e-09
20 / 33