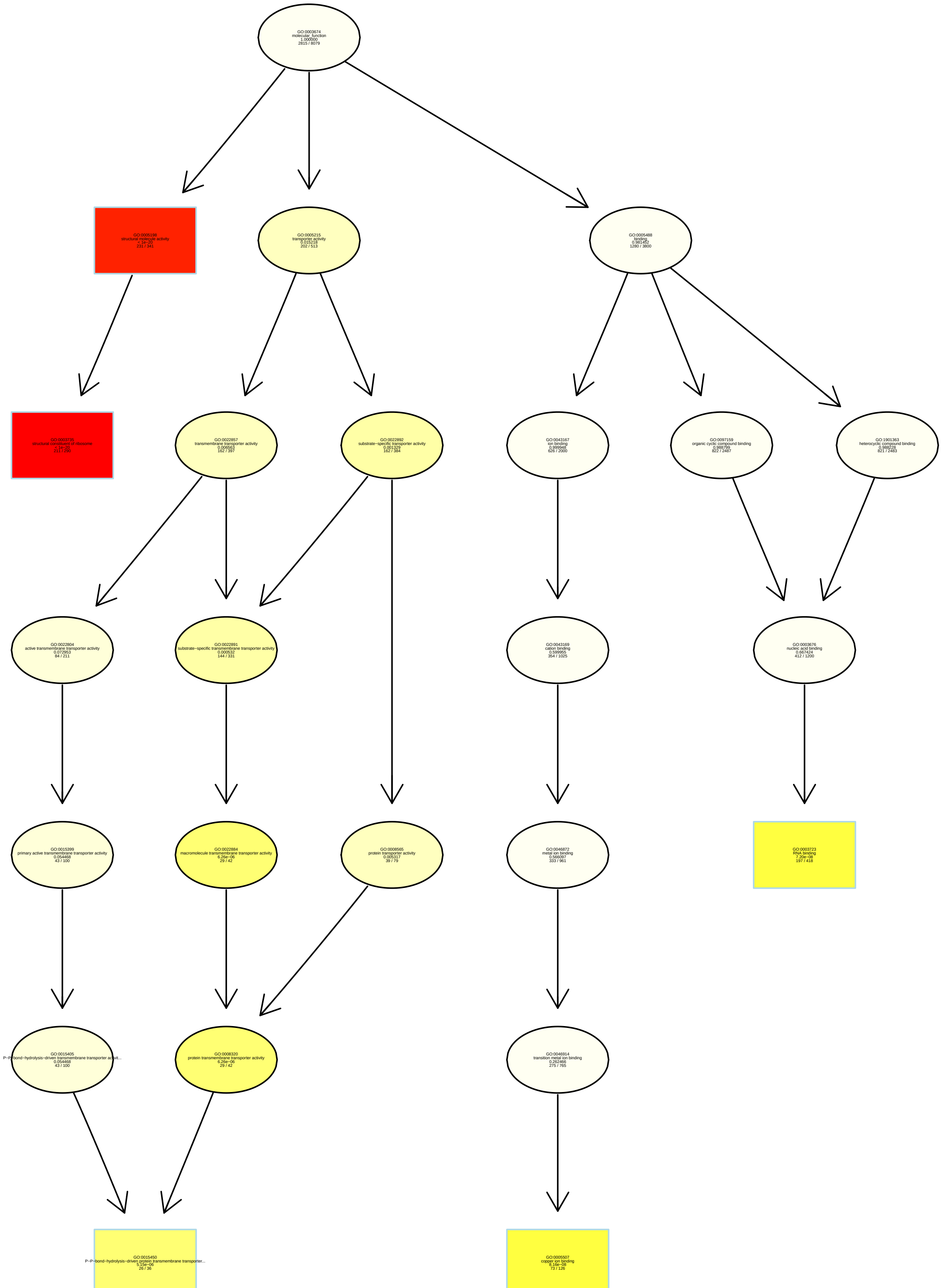

GO:0003674
molecular_function
1.000000
2815 / 8079
GO:0005198
structural molecule activity
< 1e−20
231 / 341
GO:0005215
transporter activity
0.015218
202 / 513
GO:0005488
binding
0.981452
1280 / 3800
GO:0003735
structural constituent of ribosome
< 1e−20
211 / 290
GO:0022857
transmembrane transporter activity
0.006563
162 / 397
GO:0022892
substrate−specific transporter activity
0.001329
162 / 384
GO:0043167
ion binding
0.999948
626 / 2000
GO:0097159
organic cyclic compound binding
0.988799
822 / 2487
GO:1901363
heterocyclic compound binding
0.988228
821 / 2483
GO:0022804
active transmembrane transporter activity
0.072953
84 / 211
GO:0022891
substrate−specific transmembrane transporter activity
0.000532
144 / 331
GO:0043169
cation binding
0.599955
354 / 1025
GO:0003676
nucleic acid binding
0.667424
412 / 1200
GO:0015399
primary active transmembrane transporter activity
0.054468
43 / 100
GO:0022884
macromolecule transmembrane transporter activity
6.26e−06
29 / 42
GO:0008565
protein transporter activity
0.005317
39 / 79
GO:0046872
metal ion binding
0.566097
333 / 961
GO:0003723
RNA binding
7.20e−08
197 / 418
GO:0015405
P−P−bond−hydrolysis−driven transmembrane transporter activit...
0.054468
43 / 100
GO:0008320
protein transmembrane transporter activity
6.26e−06
29 / 42
GO:0046914
transition metal ion binding
0.262466
275 / 765
GO:0015450
P−P−bond−hydrolysis−driven protein transmembrane transporter...
5.15e−06
26 / 36
GO:0005507
copper ion binding
8.16e−08
73 / 126