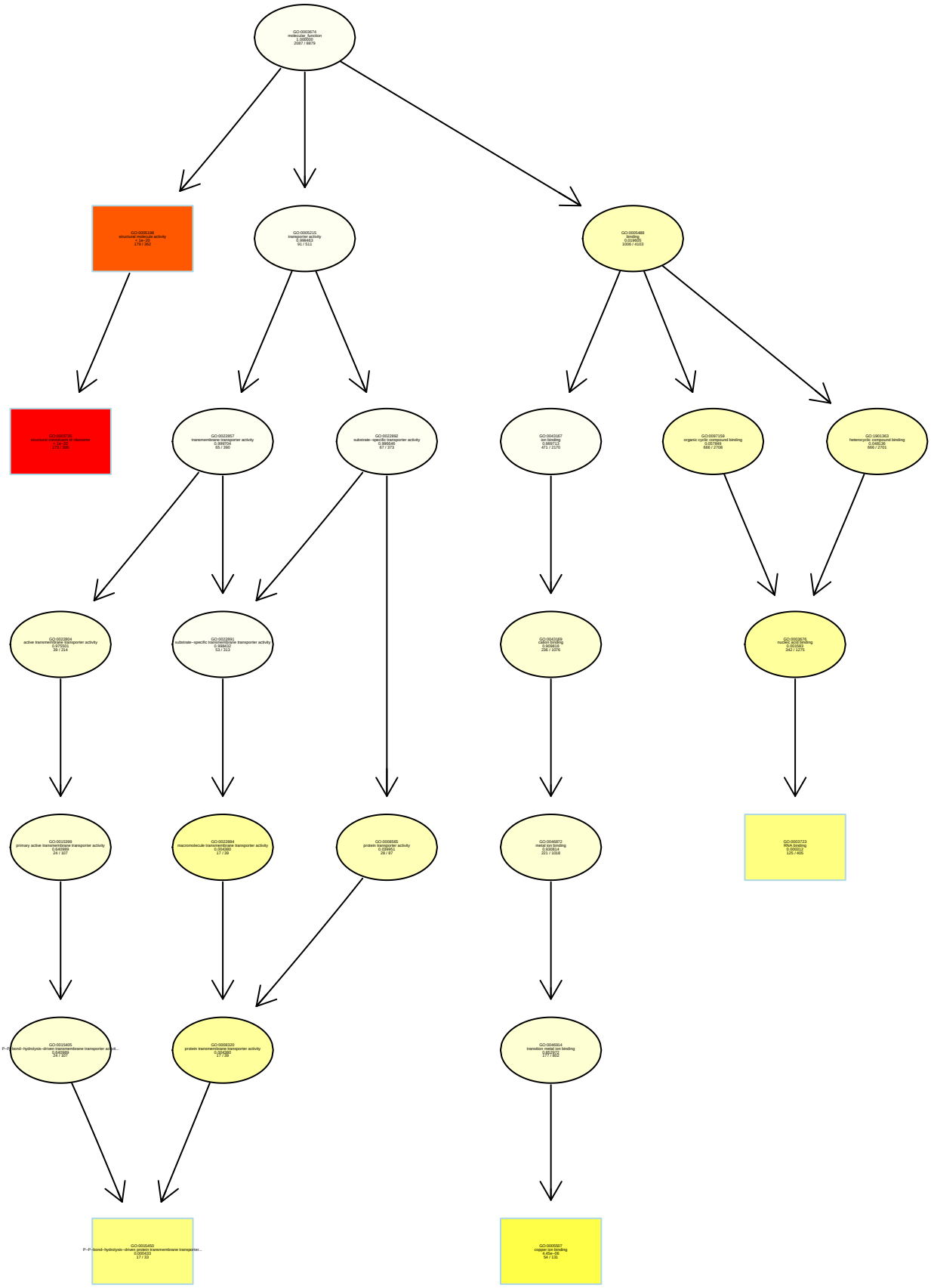
GO:0003674
molecular_function
1.000000
3387 / 8879
GO:0005198
structural molecule activity
0.000000
178 / 262
GO:0005215
transporter activity
0.000000
91 / 511
GO:0005488
binding
0.000000
1006 / 4103
GO:0003735
structural constituent of ribosome
1.1e-20
170 / 253
GO:0022857
transmembrane transporter activity
0.999500
65 / 390
GO:0022892
substrate-specific transporter activity
0.999549
57 / 373
GO:0043167
ion binding
0.000000
471 / 2170
GO:0097159
organic cyclic compound binding
0.057369
666 / 2708
GO:1901363
heterocyclic compound binding
0.048136
666 / 2701
GO:0022804
active transmembrane transporter activity
0.975553
39 / 214
GO:0022891
substrate-specific transmembrane transporter activity
0.998022
53 / 313
GO:0046872
metal ion binding
0.000819
236 / 1076
GO:0003676
nucleic acid binding
0.000000
342 / 1275
GO:0015399
primary active transmembrane transporter activity
0.940889
24 / 107
GO:0022884
macromolecule transmembrane transporter activity
0.994980
17 / 39
GO:0008565
protein transporter activity
0.999653
28 / 87
GO:0046872
metal ion binding
0.000814
221 / 1018
GO:0003723
RNA binding
0.000017
125 / 405
GO:0015405
P-P-bond-hydrolysis-driven transmembrane transporter activ...
0.940889
24 / 107
GO:0008320
protein transmembrane transporter activity
0.994980
17 / 39
GO:0046914
transition metal ion binding
0.897972
177 / 802
GO:0015450
P-P-bond-hydrolysis-driven protein transmembrane transporter...
0.000423
17 / 33
GO:0005507
copper ion binding
4.45e-06
54 / 131