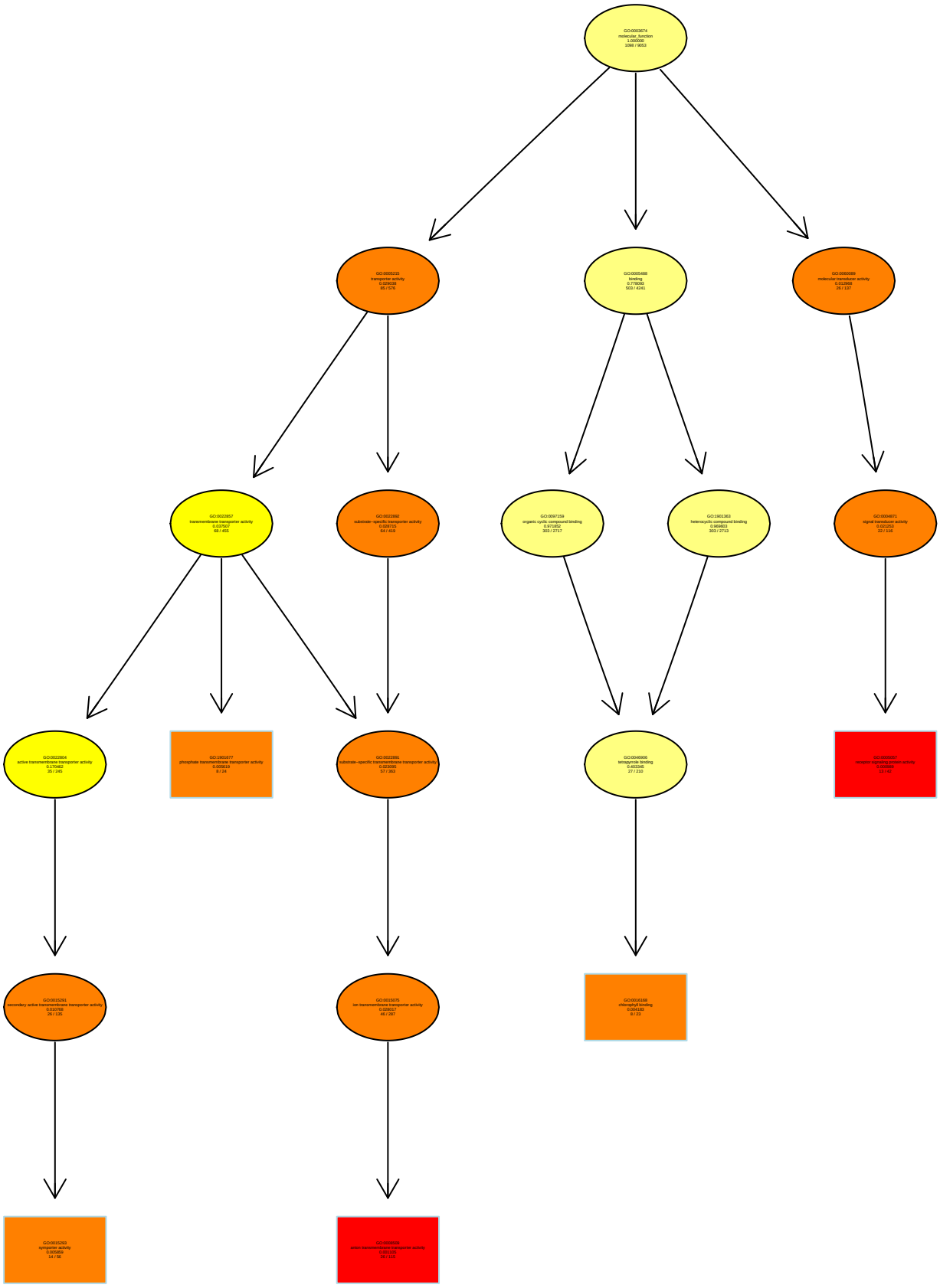
GO:0003674
molecular_function
1.000000
1088 / 9053
GO:0005215
transporter activity
0.026338
65 / 576
GO:0005488
binding
0.779093
503 / 4241
GO:0060089
molecular transducer activity
0.012988
26 / 137
GO:0022857
transmembrane transporter activity
0.037507
58 / 455
GO:0022892
substrate−specific transporter activity
0.028715
54 / 419
GO:0097159
organic cyclic compound binding
0.871052
303 / 2717
GO:1901363
heterocyclic compound binding
0.869803
303 / 2713
GO:0004871
signal transducer activity
0.021253
22 / 116
GO:0022804
active transmembrane transporter activity
0.170442
35 / 245
GO:1901677
phosphate transmembrane transporter activity
0.009519
8 / 24
GO:0022891
substrate−specific transmembrane transporter activity
0.033965
57 / 363
GO:0046906
tetrapyrrole binding
0.402345
27 / 210
GO:0000057
receptor signaling protein activity
0.005869
13 / 42
GO:0015291
secondary active transmembrane transporter activity
0.010759
26 / 135
GO:0015075
ion transmembrane transporter activity
0.028317
46 / 267
GO:0016168
chlorophyll binding
0.004188
8 / 23
GO:0015293
symporter activity
0.005859
14 / 56
GO:0005319
anion transmembrane transporter activity
0.001655
30 / 115