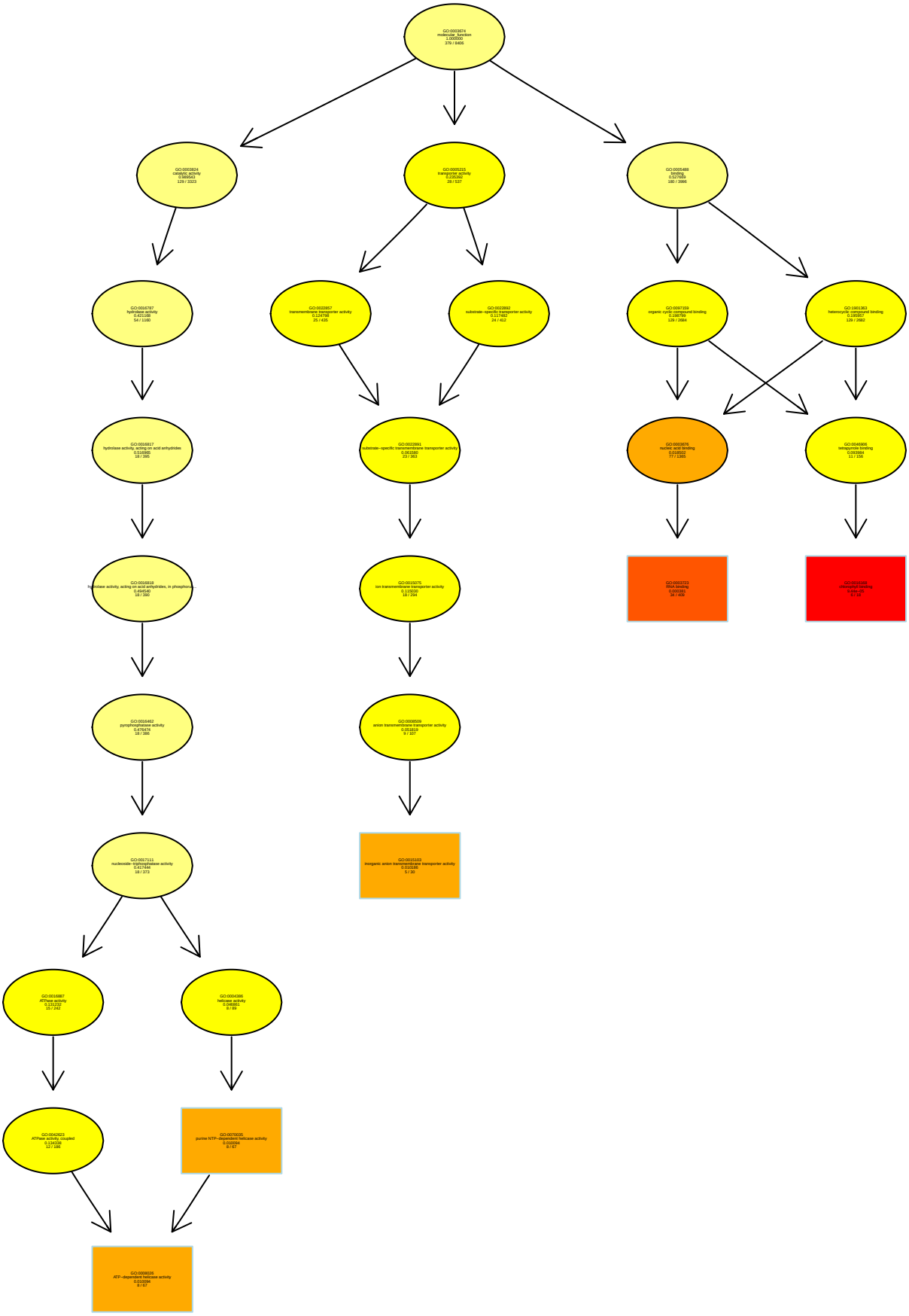
GO:0003824
catalytic activity
1.000000
379 / 8406
GO:0003824
catalytic activity
0.000000
129 / 3323
GO:0005215
transporter activity
0.000000
28 / 537
GO:0005488
binding
0.000000
180 / 3966
GO:0016787
hydrolase activity
0.421198
54 / 1180
GO:0022857
transmembrane transporter activity
0.124798
25 / 435
GO:0022892
substrate-specific transporter activity
0.117482
24 / 412
GO:0097159
organic cyclic compound binding
0.198790
129 / 2684
GO:1901363
heterocyclic compound binding
0.195957
129 / 2682
GO:0016817
hydrolase activity, acting on acid anhydrides
0.516965
18 / 365
GO:0022891
substrate-specific transmembrane transporter activity
0.061580
23 / 363
GO:0003676
nucleic acid binding
0.018932
77 / 1365
GO:0046906
tetrapyrrole binding
0.093984
11 / 155
GO:0016818
hydrolase activity, acting on acid anhydrides, in phosphorus-
0.494540
18 / 360
GO:0015075
ion transmembrane transporter activity
0.119330
18 / 264
GO:0003723
RNA binding
0.000381
38 / 409
GO:0016163
nitrogenase activity
9.44e-05
6 / 19
GO:0016462
pyrophosphatase activity
0.476674
18 / 365
GO:0008509
anion transmembrane transporter activity
0.051819
9 / 107
GO:0017111
nucleoside-triphosphatase activity
0.421784
18 / 373
GO:0015103
inorganic anion transmembrane transporter activity
0.012486
5 / 30
GO:0016887
ATPase activity
0.134339
12 / 242
GO:0004386
helicase activity
0.010294
8 / 69
GO:0042623
ATPase activity, coupled
0.134339
12 / 186
GO:0070035
purine NTP-dependent helicase activity
0.010294
8 / 67
GO:0008026
ATP-dependent helicase activity
0.010294
8 / 67