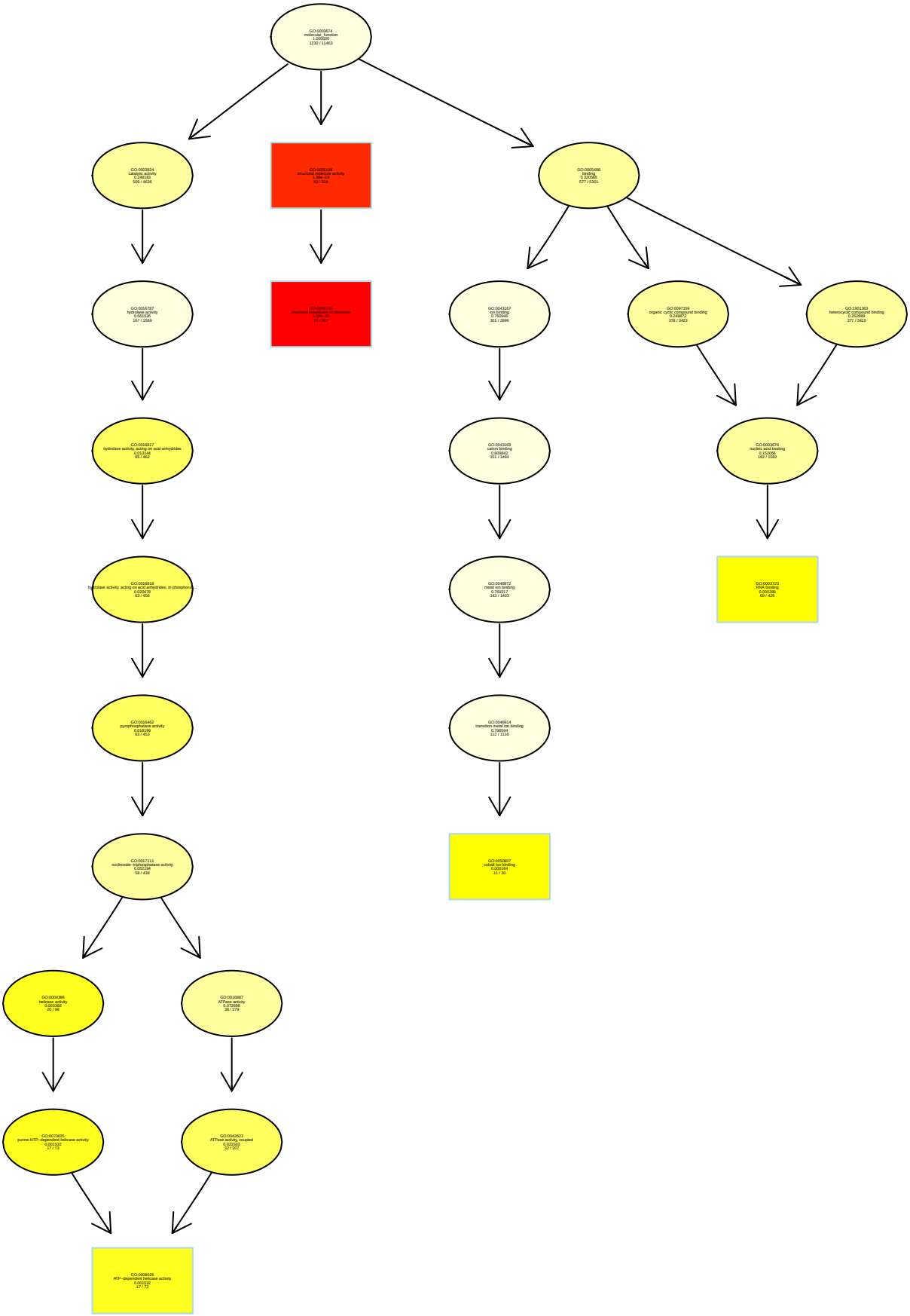
GO:0003674
molecular_function
1.000000
1230 / 11463
GO:0003824
catalytic activity
0.396325
509 / 4636
GO:0005198
structural molecule activity
0.256410
83 / 324
GO:0005488
binding
0.335554
577 / 5301
GO:0016787
hydrolase activity
0.561536
167 / 1569
GO:0003735
structural constituent of ribosome
1.39e-16
70 / 257
GO:0043167
ion binding
0.765946
301 / 2846
GO:0097159
organic cyclic compound binding
0.249872
378 / 3423
GO:1901363
heterocyclic compound binding
0.252689
377 / 3415
GO:0016817
hydrolase activity, acting on acid anhydrides
0.013144
65 / 462
GO:0043169
cation binding
0.809842
251 / 1494
GO:0003676
nucleic acid binding
0.152066
182 / 2162
GO:0016818
hydrolase activity, acting on acid anhydrides, in phosphorus-c
0.020678
61 / 456
GO:0046872
metal ion binding
0.760317
243 / 1463
GO:0003723
RNA binding
0.000299
69 / 426
GO:0016462
pyrophosphatase activity
0.018199
61 / 453
GO:0046914
transition metal ion binding
0.798594
112 / 1115
GO:0017111
nucleoside-triphosphatase activity
0.013486
58 / 438
GO:0050897
cobalt ion binding
0.033333
11 / 30
GO:0004386
helicase activity
0.010152
20 / 98
GO:0016887
ATPase activity
0.055132
38 / 279
GO:0070035
purine NTP-dependent helicase activity
0.010152
17 / 72
GO:0042623
ATPase activity, coupled
0.010152
32 / 207
GO:0008026
ATP-dependent helicase activity
0.010152
17 / 72