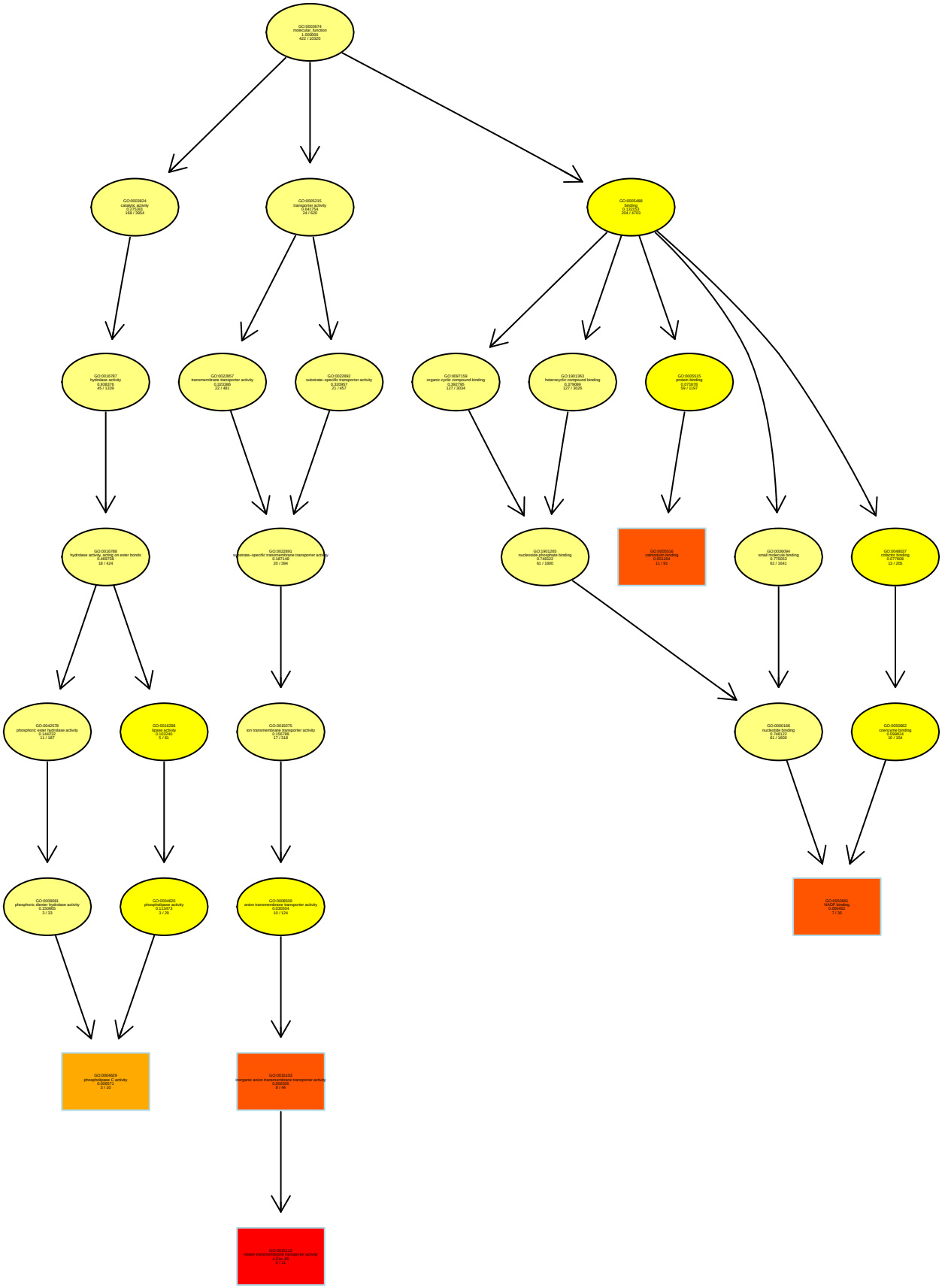
GO:0003674
molecular_function
1.000000
422 / 10320
GO:0003824
catalytic activity
0.275300
168 / 3954
GO:0005215
transporter activity
0.641754
24 / 620
GO:0005488
binding
0.132153
204 / 4703
GO:0016787
hydrolase activity
0.036376
45 / 1339
GO:0022857
transmembrane transporter activity
0.422586
22 / 481
GO:0022892
substrate-specific transporter activity
0.329903
21 / 457
GO:0097159
organic cyclic compound binding
0.282795
127 / 3034
GO:1901363
heterocyclic compound binding
0.279089
127 / 3026
GO:0036516
protein tetramerization
0.073278
59 / 1197
GO:0016788
hydrolase activity, acting on ester bonds
0.042232
18 / 424
GO:0022891
substrate-specific transmembrane transporter activity
0.329903
20 / 394
GO:1901265
nucleoside phosphate binding
0.282795
61 / 1600
GO:0008516
calmodulin binding
0.000000
11 / 91
GO:0036094
small molecule binding
0.282795
62 / 1641
GO:0048037
cofactor binding
0.042232
13 / 295
GO:0042578
phosphoric ester hydrolase activity
0.144232
11 / 187
GO:0016298
lipase activity
0.193240
9 / 61
GO:0015075
ion transmembrane transporter activity
0.156788
17 / 318
GO:0000166
nucleotide binding
0.748122
61 / 1600
GO:0050662
coenzyme binding
0.099624
10 / 154
GO:0008081
phosphoric diester hydrolase activity
0.150850
3 / 33
GO:0004620
phospholipase activity
0.113473
3 / 29
GO:0008509
anion transmembrane transporter activity
0.035054
10 / 124
GO:0050661
NADP binding
0.065452
7 / 36
GO:0004629
phospholipase C activity
0.006571
3 / 10
GO:0015103
inorganic anion transmembrane transporter activity
0.000355
8 / 66
GO:0015112
nitrate transmembrane transporter activity
4.15e-05
5 / 11