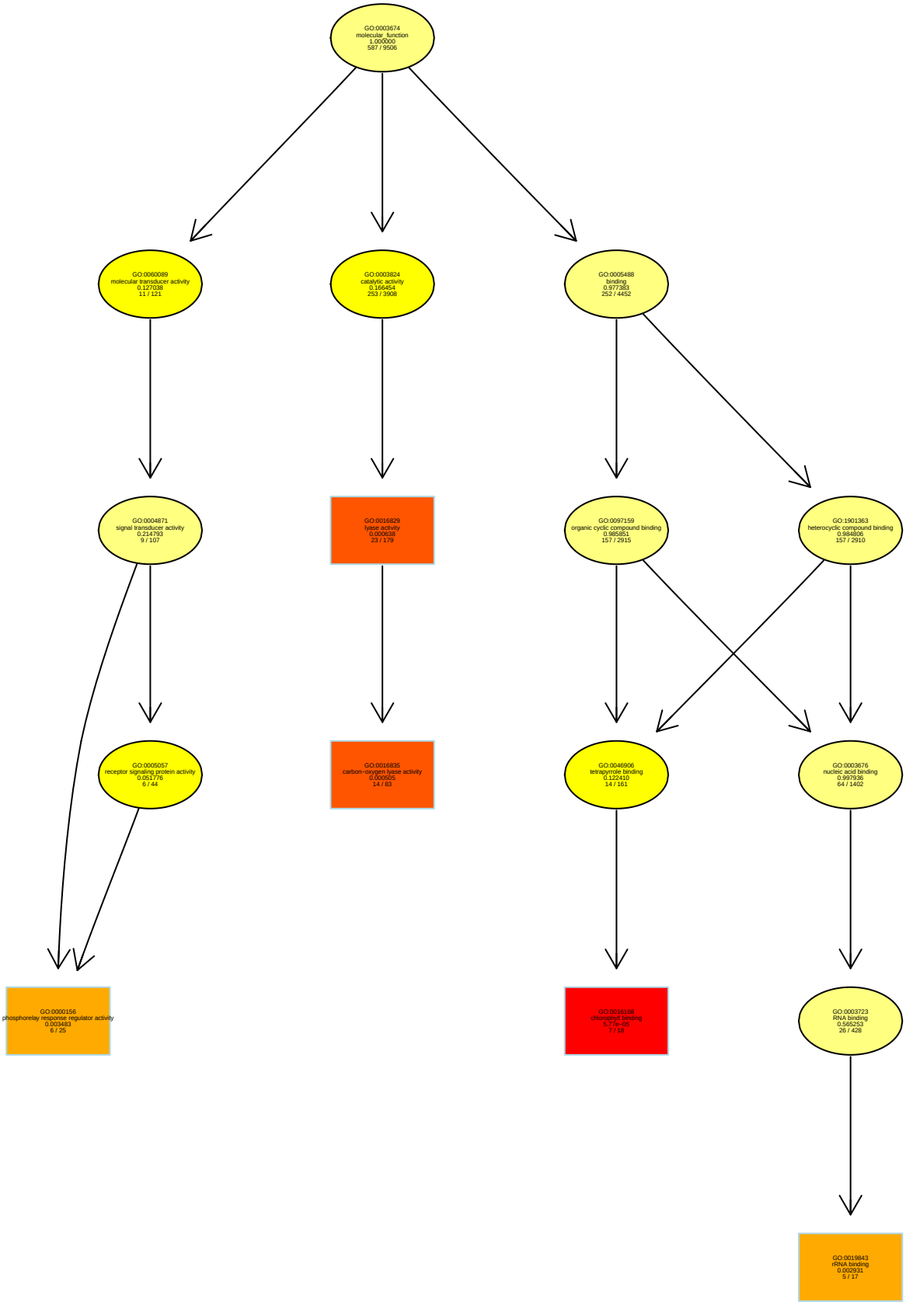
GO:0003674
molecular_function
1.000000
587 / 9506
GO:0060089
molecular transducer activity
0.127038
11 / 121
GO:0003824
catalytic activity
0.166454
253 / 3908
GO:0005488
binding
0.977383
252 / 4452
GO:0004871
signal transducer activity
0.214793
9 / 107
GO:0016829
lyase activity
0.000638
23 / 179
GO:0097159
organic cyclic compound binding
0.985851
157 / 2915
GO:1901363
heterocyclic compound binding
0.984806
157 / 2910
GO:0005057
receptor signaling protein activity
0.051776
6 / 44
GO:0016835
carbon−oxygen lyase activity
0.000505
14 / 83
GO:0046906
tetrapyrrole binding
0.122410
14 / 161
GO:0003676
nucleic acid binding
0.997936
64 / 1402
GO:0000156
phosphorelay response regulator activity
0.003483
6 / 25
GO:0016168
chlorophyll binding
5.77e−05
7 / 18
GO:0003723
RNA binding
0.565253
26 / 428
GO:0019843
rRNA binding
0.002931
5 / 17