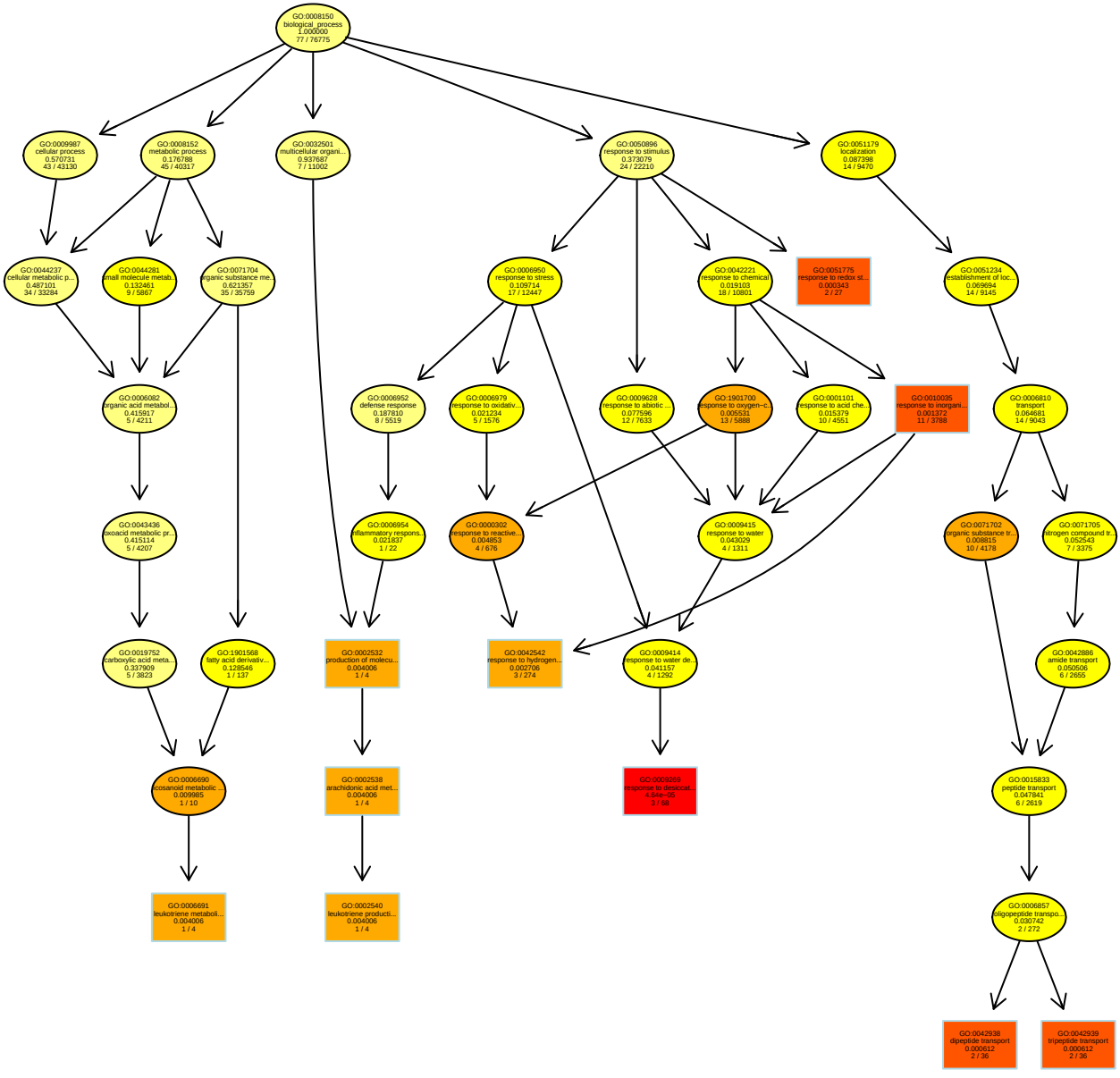
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GO:0009987
cellular process
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43 / 43130
GO:0008152
metabolic process
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45 / 40317
GO:0032501
multicellular organi...
0.937687
7 / 11002
GO:0050896
response to stimulus
0.373079
24 / 22210
GO:0051179
localization
0.087398
14 / 9470
GO:0044237
cellular metabolic p...
0.487101
34 / 33284
GO:0044281
small molecule metab...
0.132461
9 / 5867
GO:0071704
organic substance me...
0.621857
35 / 35759
GO:0006950
response to stress
0.109714
17 / 12447
GO:0042221
response to chemical
0.019103
18 / 10801
GO:0051775
response to redox st...
0.000343
2 / 27
GO:0051234
establishment of loc...
0.069694
14 / 9145
GO:0006082
organic acid metabol...
0.415917
5 / 4211
GO:0006952
defense response
0.187810
8 / 5519
GO:0006979
response to oxidativ...
0.021234
5 / 1576
GO:0009628
response to abiotic ...
0.077596
12 / 7633
GO:1901700
response to oxygen-c...
0.005531
13 / 5888
GO:0001101
response to acid che...
0.015379
10 / 4551
GO:0010035
response to inorgani...
0.001372
11 / 3788
GO:0006810
transport
0.064681
14 / 9043
GO:0043436
oxoacid metabolic pr...
0.415114
5 / 4207
GO:0006954
inflammatory respons...
0.021837
1 / 22
GO:0000302
response to reactive...
0.004853
4 / 676
GO:0009415
response to water
0.043029
4 / 1311
GO:0071702
organic substance tr...
0.008815
10 / 4178
GO:0071705
nitrogen compound tr...
0.052543
7 / 3375
GO:0019752
carboxylic acid meta...
0.337909
5 / 3823
GO:1901568
fatty acid derivativ...
0.128546
1 / 137
GO:0002532
production of molecu...
0.004006
1 / 4
GO:0042542
response to hydrogen...
0.002706
3 / 274
GO:0009414
response to water de...
0.041157
4 / 1292
GO:0042886
amide transport
0.050506
6 / 2655
GO:0006690
icosanoid metabolic ...
0.009985
1 / 10
GO:0002538
arachidonic acid met...
0.004006
1 / 4
GO:0009269
response to desiccat...
4.64e-05
3 / 68
GO:0015833
peptide transport
0.047841
6 / 2619
GO:0006691
leukotriene metaboli...
0.004006
1 / 4
GO:0002540
leukotriene producti...
0.004006
1 / 4
GO:0006857
oligopeptide transpo...
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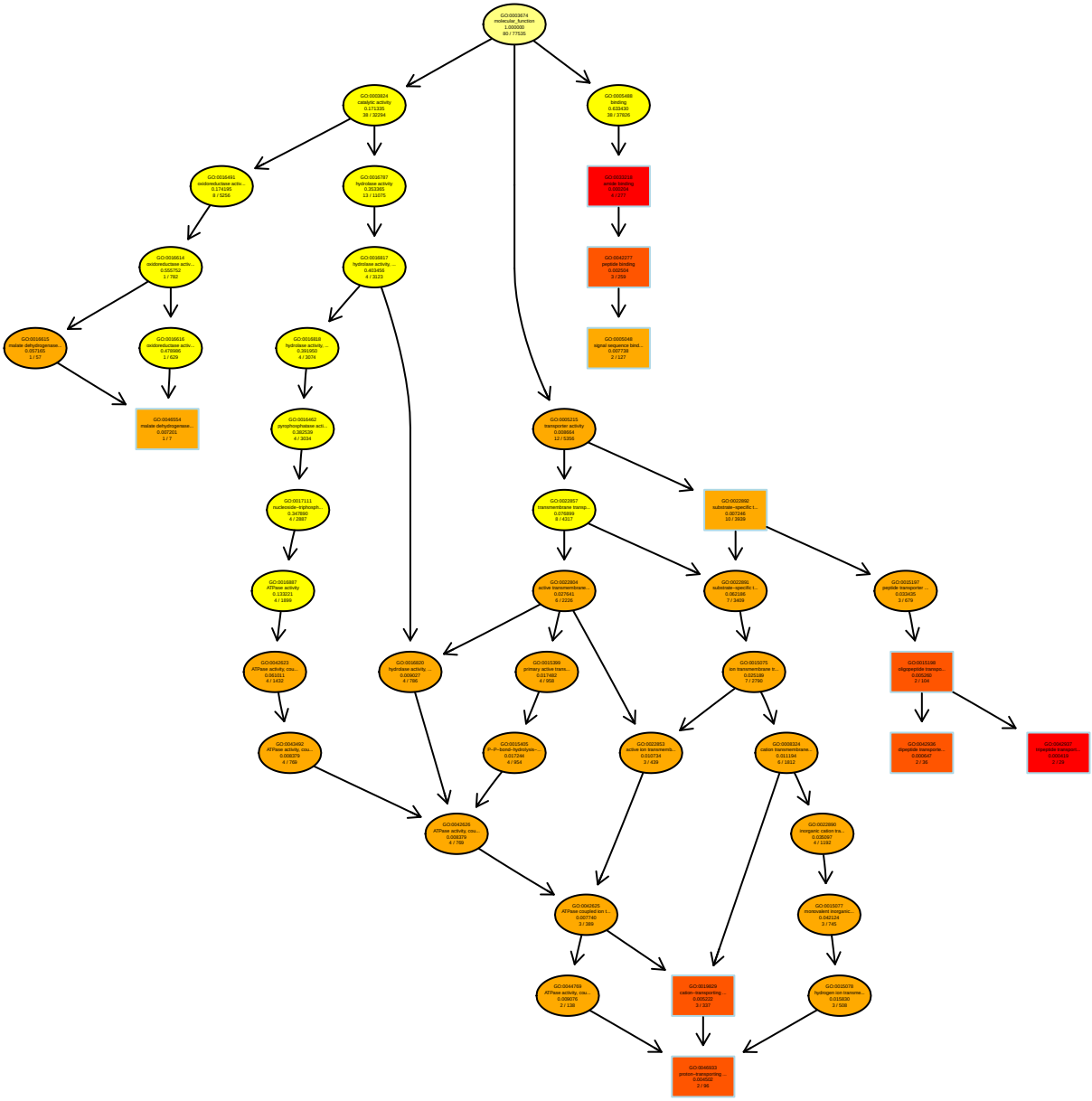
GO:0042938
dipeptide transport
0.000612
2 / 36
GO:0042939
tripeptide transport
0.000612
2 / 36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GO:0003824
catalytic activity
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38 / 32264
GO:0005488
binding
0.833430
38 / 27535
GO:0016491
oxidoreductase activ...
0.057266
8 / 5258
GO:0016787
hydrolase activity
0.000204
13 / 11075
GO:0033218
amide binding
0.000204
4 / 177
GO:0016614
oxidoreductase activ...
0.055752
1 / 782
GO:0016817
hydrolase activity, ...
0.403456
4 / 3122
GO:0042277
peptide binding
0.000204
3 / 259
GO:0016615
malate dehydrogenase
0.057165
1 / 57
GO:0016616
oxidoreductase activ...
0.479686
1 / 629
GO:0016818
hydrolase activity, ...
0.391950
4 / 3076
GO:0005048
signal sequence bind...
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2 / 127
GO:0046554
malate dehydrogenase...
0.007201
1 / 7
GO:0016462
pyrophosphatase acti...
0.382539
4 / 3034
GO:0005215
transporter activity
0.558664
12 / 5355
GO:0017111
nucleoside-triphosph...
0.347860
4 / 2887
GO:0022857
transmembrane transp...
0.576699
8 / 4317
GO:0022892
substrate-specific t...
0.667246
10 / 3639
GO:0016887
ATPase activity
0.133211
4 / 1899
GO:0022804
active transmembrane...
0.027641
6 / 2235
GO:0022891
substrate-specific t...
0.062166
7 / 3439
GO:0015197
peptide transporter ...
0.034435
3 / 679
GO:0042623
ATPase activity, cou...
0.061011
4 / 1432
GO:0016820
hydrolase activity, ...
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4 / 786
GO:0015399
primary active trans...
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4 / 958
GO:0015075
ion transmembrane tr...
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GO:0015198
oligopeptide transpo...
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2 / 104
GO:0043492
ATPase activity, cou...
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4 / 759
GO:0015405
P-P-bond-hydrolysis-...
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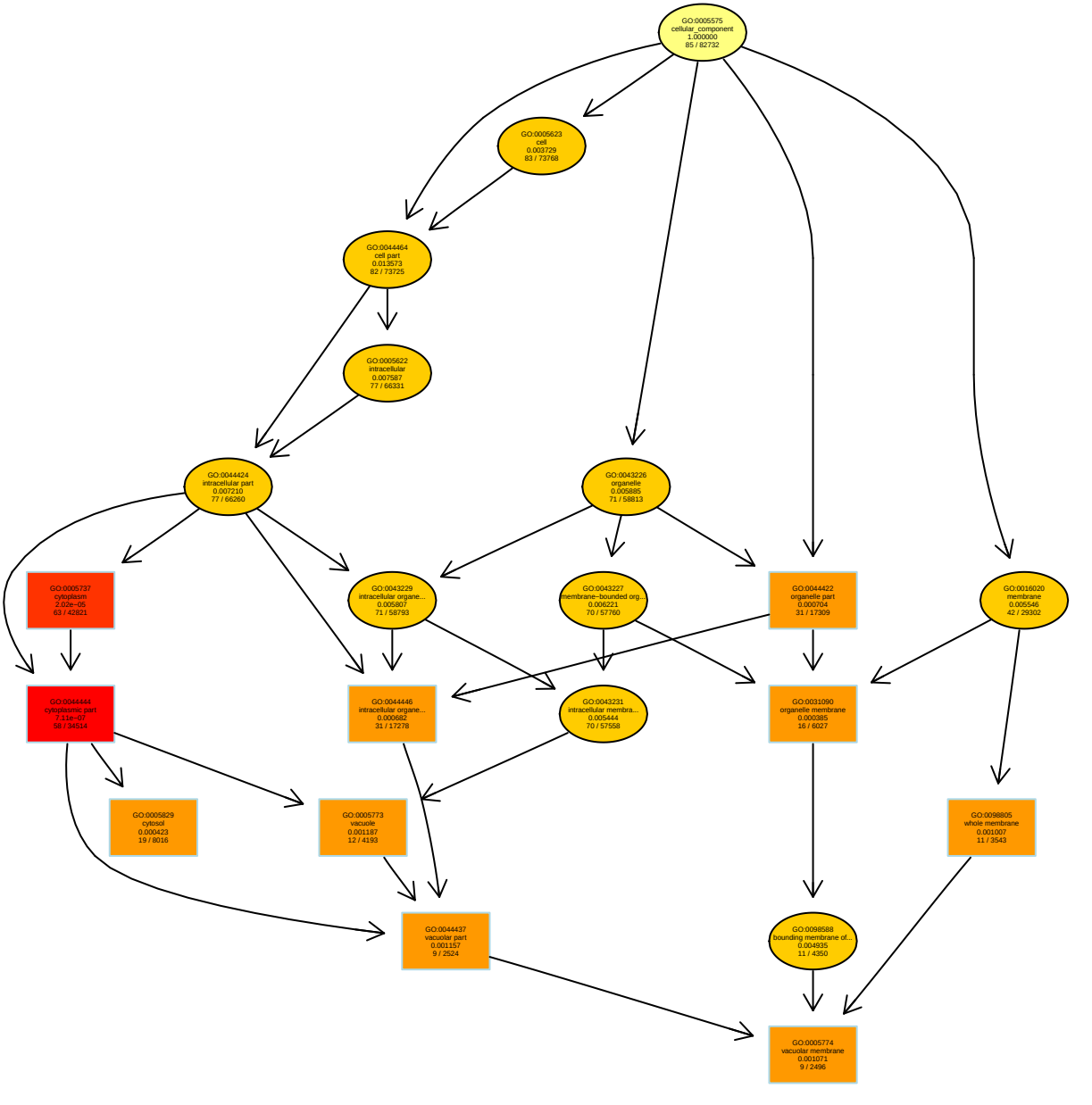
4 / 954
GO:0022853
active ion transmemb...
0.010734
3 / 439
GO:0008324
cation transmembrane...
0.011194
6 / 1812
GO:0042936
dipeptide transpore...
0.000647
2 / 38
GO:0042937
tripeptide transport...
0.000419
2 / 29
GO:0042626
ATPase activity, cou...
0.000379
4 / 759
GO:0022890
inorganic cation tra...
0.036267
4 / 1292
GO:0042625
ATPase coupled ion t...
0.007740
3 / 389
GO:0015077
monovalent inorganic...
0.042134
3 / 745
GO:0044769
ATPase activity, cou...
0.009076
3 / 336
GO:0018829
cation-transporting ...
0.005222
3 / 327
GO:0015078
hydrogen ion transme...
0.015830
3 / 508
GO:0046933
proton-transporting ...
0.004502
2 / 96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GO:0005623
cell
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83 / 73768
GO:0044464
cell part
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82 / 73725
GO:0005622
intracellular
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77 / 66331
GO:0044424
intracellular part
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77 / 66260
GO:0043226
organelle
0.005885
71 / 58813
GO:0005737
cytoplasm
2.02e−05
63 / 42821
GO:0043229
intracellular organe...
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71 / 58793
GO:0043227
membrane−bounded org...
0.006221
70 / 57760
GO:0044422
organelle part
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31 / 17309
GO:0016020
membrane
0.005546
42 / 29302
GO:0044444
cytoplasmic part
7.11e−07
58 / 34514
GO:0044446
intracellular organe...
0.000682
31 / 17278
GO:0043231
intracellular membra...
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70 / 57558
GO:0031090
organelle membrane
0.000385
16 / 6027
GO:0005829
cytosol
0.000423
19 / 8016
GO:0005773
vacuole
0.001187
12 / 4193
GO:0098805
whole membrane
0.001007
11 / 3543
GO:0044437
vacuolar part
0.001157
9 / 2524
GO:0098588
bounding membrane of...
0.004935
11 / 4350
GO:0005774
vacuolar membrane
0.001071
9 / 2496