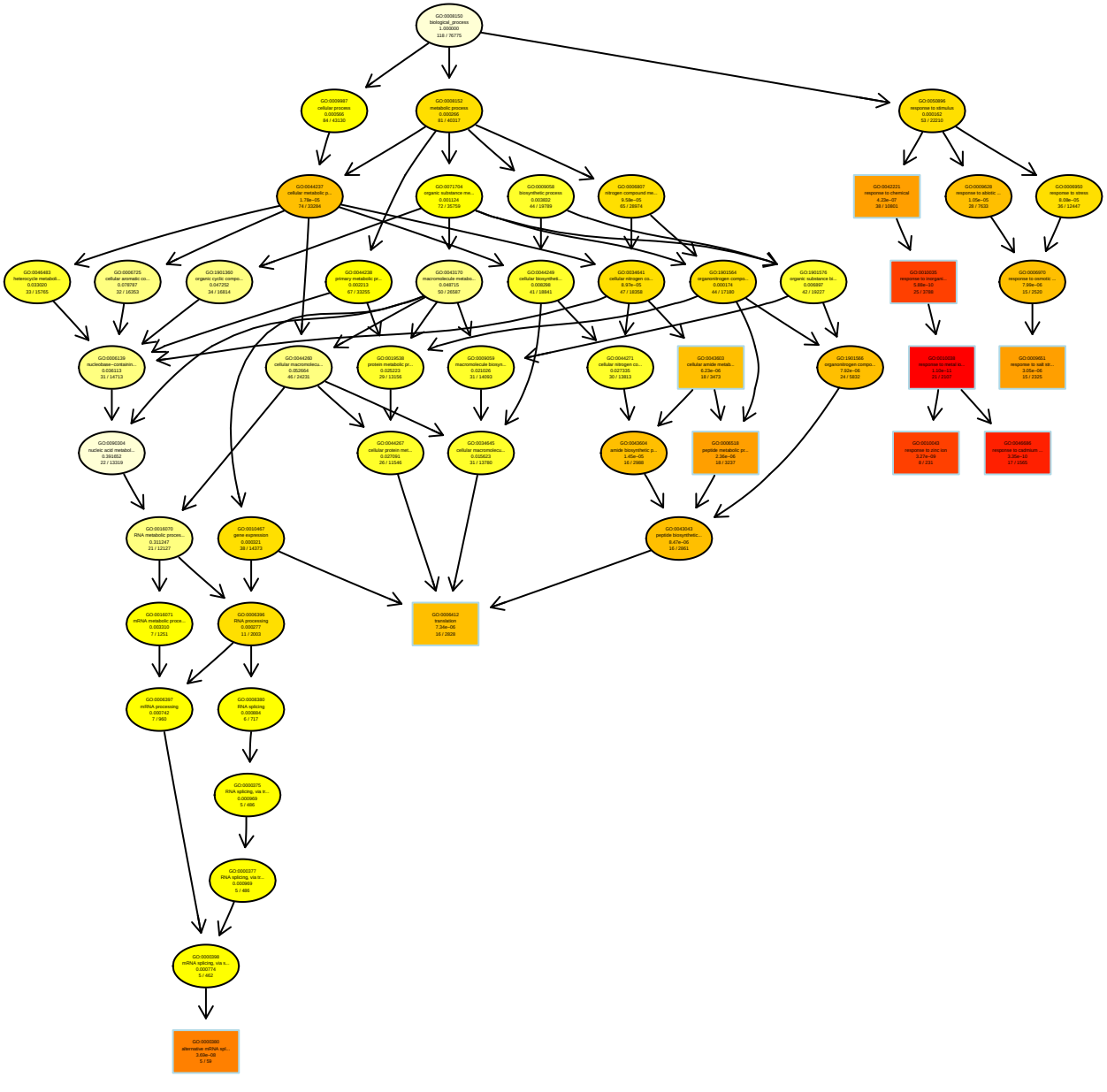
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GO:0009987
cellular process
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GO:0008152
metabolic process
0.000356
81 / 40317
GO:0050896
response to stimulus
0.000152
53 / 22210
GO:0044237
cellular metabolic p...
1.78e−05
74 / 33284
GO:0071704
organic substance me...
0.001124
72 / 35759
GO:0006807
nitrogen compound me...
9.58e−05
65 / 28974
GO:0042221
response to chemical
4.23e−07
38 / 10801
GO:0009628
response to abiotic ...
1.05e−05
28 / 7632
GO:0006950
response to stress
8.08e−05
36 / 12447
GO:0044483
heterocyclic metabol...
0.033005
33 / 15765
GO:0006725
cellular aromatic co...
0.078797
22 / 16253
GO:1901360
organic cyclic compo...
0.047252
34 / 16814
GO:0044238
primary metabolic pr...
0.002313
47 / 33255
GO:0043170
macromolecule metabo...
0.048715
50 / 26587
GO:0044249
cellular biosynthet...
0.006398
41 / 18841
GO:0034641
cellular nitrogen co...
8.97e−05
47 / 18358
GO:1901564
organonitrogen compo...
0.000174
46 / 17180
GO:1901576
organic substance bi...
0.006897
42 / 19227
GO:0010035
response to inorganic...
6.88e−10
25 / 3788
GO:0006970
response to osmotic ...
7.99e−06
15 / 2539
GO:0006139
nucleobase−containin...
0.036113
31 / 14713
GO:0044260
cellular macromolecu...
0.052664
46 / 24231
GO:0019538
protein metabolic pr...
0.035223
29 / 13156
GO:0009059
macromolecule biosyn...
0.021026
31 / 14063
GO:0044271
cellular nitrogen co...
0.027335
30 / 13813
GO:0043603
cellular amide metab...
6.23e−06
18 / 3473
GO:1901566
organonitrogen compo...
7.92e−06
24 / 5832
GO:0010038
response to metal io...
1.10e−11
21 / 1027
GO:0009651
response to salt str...
3.05e−06
15 / 2325
GO:0090304
nucleic acid metabol...
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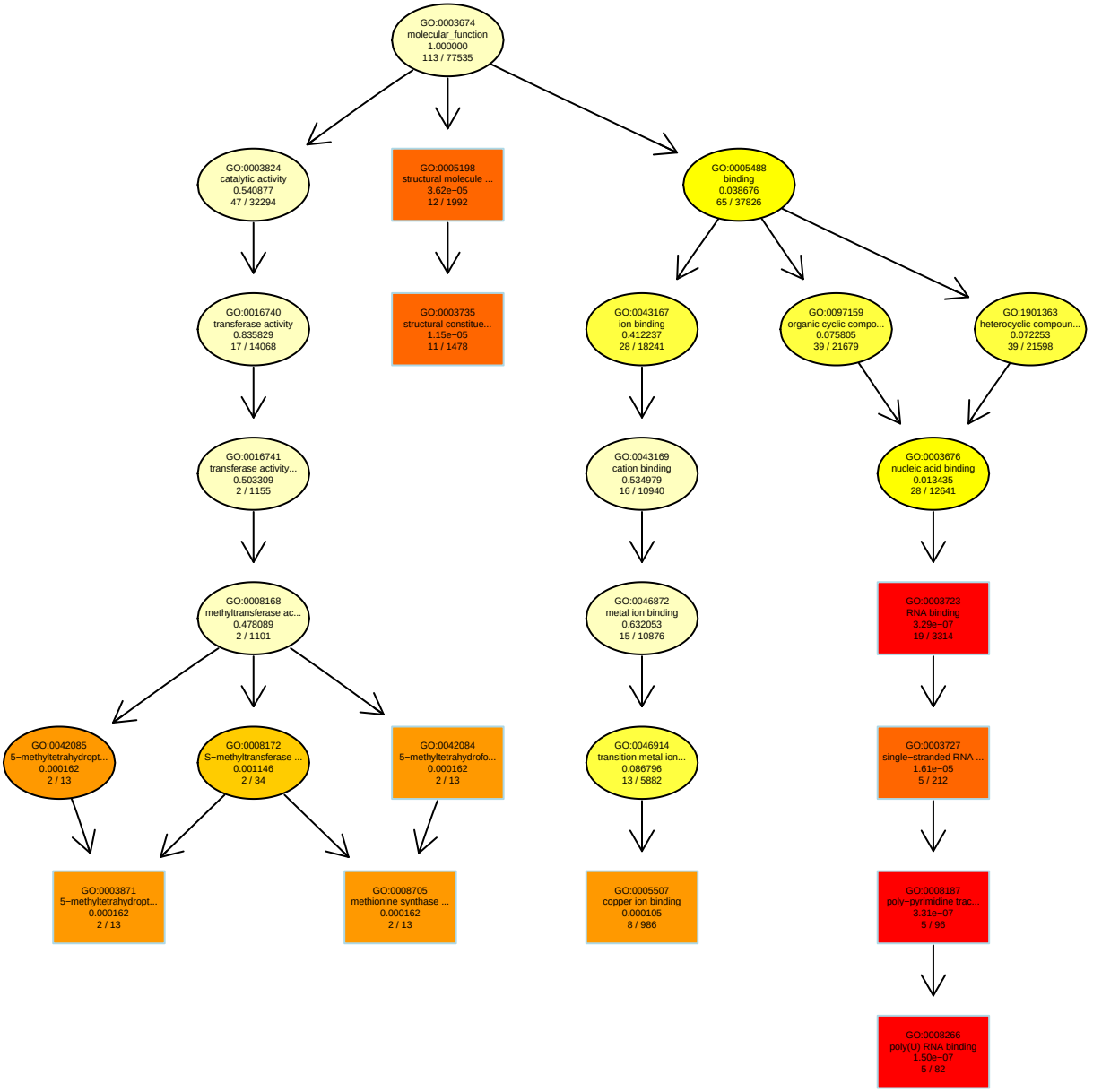
22 / 13310
GO:0044267
cellular protein met...
0.027691
26 / 11546
GO:0034645
cellular macromoleca...
0.015623
31 / 12780
GO:0043604
amide biosynthetic p...
1.45e−05
16 / 2968
GO:0006518
peptide metabolic pr...
2.36e−06
18 / 3237
GO:0010043
response to zinc ion
3.27e−09
8 / 231
GO:0046686
response to cadmium ...
3.05e−10
17 / 1565
GO:0016070
RNA metabolic proce...
0.311247
21 / 12127
GO:0010467
gene expression
0.000321
20 / 14373
GO:0043043
peptide biosynthetic...
8.47e−06
16 / 2861
GO:0006412
translation
7.34e−06
16 / 2826
GO:0016071
mRNA metabolic proce...
0.003310
7 / 1251
GO:0006396
RNA processing
0.000277
11 / 2003
GO:0006397
mRNA processing
0.000742
7 / 980
GO:0008380
RNA splicing
0.000394
6 / 717
GO:0000375
RNA splicing, via tr...
0.000969
5 / 486
GO:0000377
RNA splicing, via tr...
0.000969
5 / 486
GO:0000398
mRNA splicing, via s...
0.000774
5 / 462
GO:0000380
alternative mRNA spl...
3.08e−08
5 / 59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GO:0003824
catalytic activity
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47 / 32294
GO:0005198
structural molecule ...
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12 / 1992
GO:0005488
binding
0.038676
65 / 37826
GO:0016740
transferase activity
0.835829
17 / 14068
GO:0003735
structural constitue...
1.15e−05
11 / 1478
GO:0043167
ion binding
0.412237
28 / 18241
GO:0097159
organic cyclic compo...
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39 / 21679
GO:1901363
heterocyclic compoun...
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39 / 21598
GO:0016741
transferase activity...
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2 / 1155
GO:0043169
cation binding
0.534979
16 / 10940
GO:0003676
nucleic acid binding
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28 / 12641
GO:0008168
methyltransferase ac...
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2 / 1101
GO:0046872
metal ion binding
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15 / 10876
GO:0003723
RNA binding
3.29e−07
19 / 3314
GO:0042085
5−methyltetrahydropt...
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2 / 13
GO:0008172
S−methyltransferase ...
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2 / 34
GO:0042084
5−methyltetrahydrofo...
0.000162
2 / 13
GO:0046914
transition metal ion...
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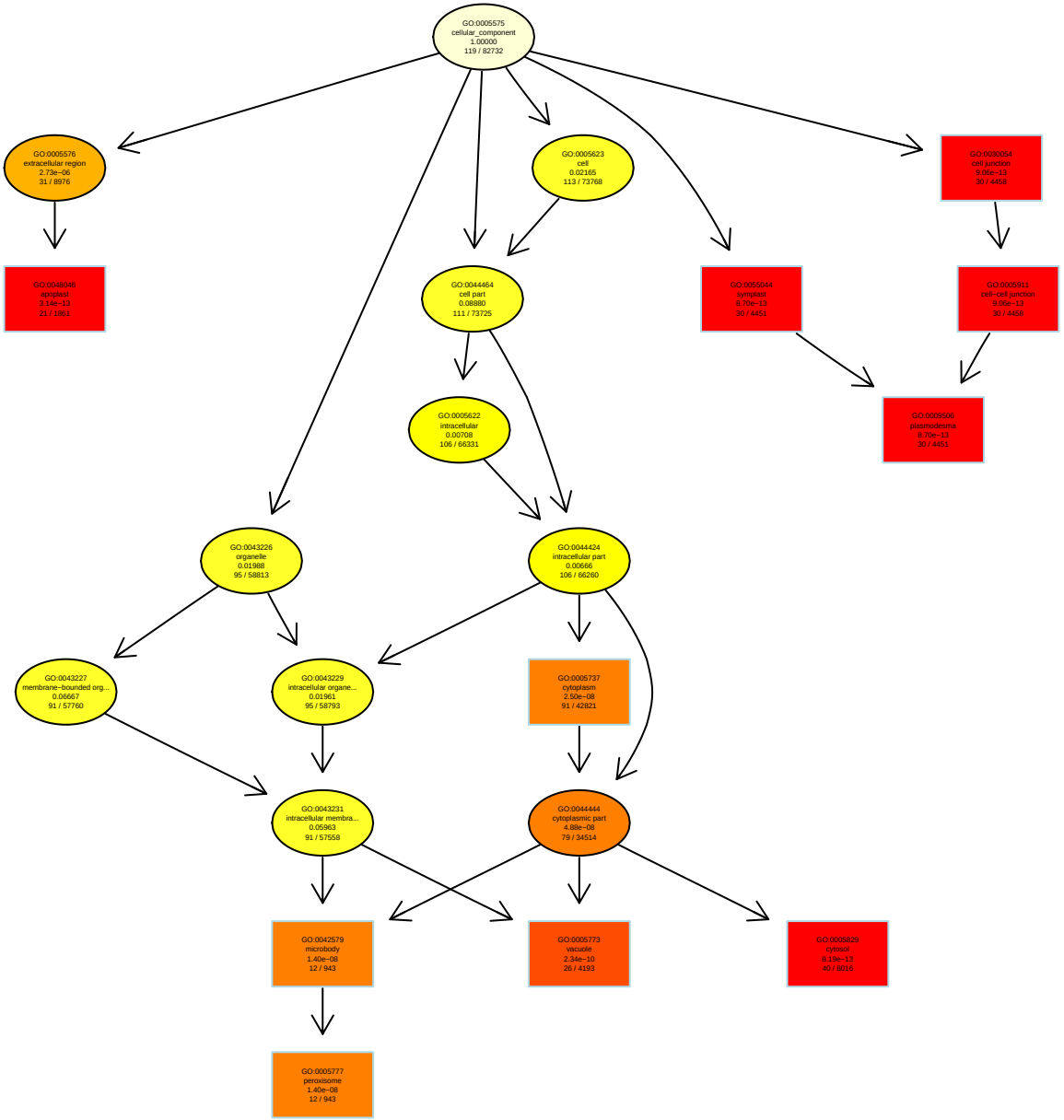
13 / 5882
GO:0003727
single−stranded RNA ...
1.61e−05
5 / 212
GO:0003871
5−methyltetrahydropt...
0.000162
2 / 13
GO:0008705
methionine synthase ...
0.000162
2 / 13
GO:0005507
copper ion binding
0.000105
8 / 986
GO:0008187
poly−pyrimidine trac...
3.31e−07
5 / 96
GO:0008266
poly(U) RNA binding
1.50e−07
5 / 82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GO:0005576
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31 / 8976
GO:0005623
cell
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113 / 73768
GO:0030054
cell junction
9.06e−13
30 / 4458
GO:0048046
apoplast
3.14e−13
21 / 1861
GO:0044464
cell part
0.08880
111 / 73725
GO:0055044
symplast
8.70e−13
30 / 4451
GO:0005911
cell−cell junction
9.06e−13
30 / 4458
GO:0005622
intracellular
0.00708
106 / 66331
GO:0009506
plasmodesma
8.70e−13
30 / 4451
GO:0043226
organelle
0.01988
95 / 58813
GO:0044424
intracellular part
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106 / 66260
GO:0043227
membrane−bounded org...
0.06667
91 / 57760
GO:0043229
intracellular organe...
0.01961
95 / 58793
GO:0005737
cytoplasm
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91 / 42821
GO:0043231
intracellular membra...
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91 / 57558
GO:0044444
cytoplasmic part
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79 / 34514
GO:0042579
microbody
1.40e−08
12 / 943
GO:0005773
vacuole
2.34e−10
26 / 4193
GO:0005829
cytosol
8.19e−13
40 / 8016
GO:0005777
peroxisome
1.40e−08
12 / 943