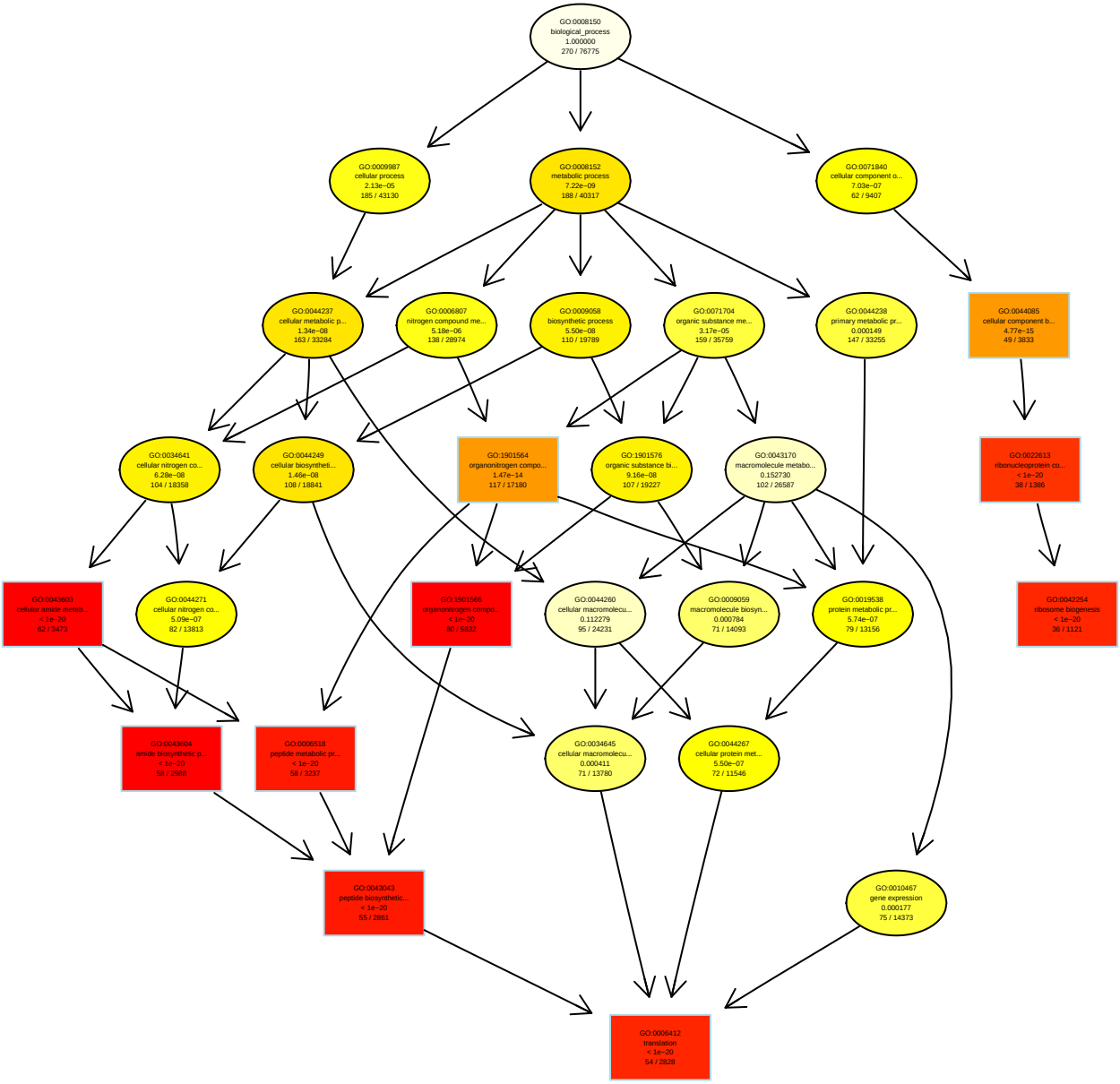
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185 / 43130
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metabolic process
7.22e−09
188 / 40317
GO:0071840
cellular component o...
7.03e−07
62 / 9407
GO:0044237
cellular metabolic p...
1.34e−08
163 / 33284
GO:0006807
nitrogen compound me...
5.18e−06
138 / 28974
GO:0009058
biosynthetic process
5.50e−08
110 / 19789
GO:0071704
organic substance me...
3.17e−05
159 / 35759
GO:0044238
primary metabolic pr...
0.000149
147 / 33255
GO:0044085
cellular component b...
4.77e−15
49 / 3833
GO:0034641
cellular nitrogen co...
6.28e−08
104 / 18358
GO:0044249
cellular biosynthet...
1.46e−08
108 / 18841
GO:1901564
organonitrogen compo...
1.47e−14
117 / 17180
GO:1901576
organic substance bi...
9.16e−08
107 / 19227
GO:0043170
macromolecule metabo...
0.152730
102 / 26587
GO:0022613
ribonucleoprotein co...
< 1e−20
38 / 1386
GO:0043603
cellular amide metab...
< 1e−20
62 / 3473
GO:0044271
cellular nitrogen co...
5.09e−07
82 / 13813
GO:1901566
organonitrogen compo...
< 1e−20
80 / 5832
GO:0044260
cellular macromolecu...
0.112279
95 / 24231
GO:0009059
macromolecule biosyn...
0.000784
71 / 14093
GO:0019538
protein metabolic pr...
5.74e−07
79 / 13156
GO:0042254
ribosome biogenesis
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36 / 1121
GO:0043604
amide biosynthetic p...
< 1e−20
58 / 2988
GO:0006518
peptide metabolic pr...
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58 / 3237
GO:0034645
cellular macromolecu...
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71 / 13780
GO:0044267
cellular protein met...
5.50e−07
72 / 11546
GO:0043043
peptide biosynthetic...
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55 / 2861
GO:0010467
gene expression
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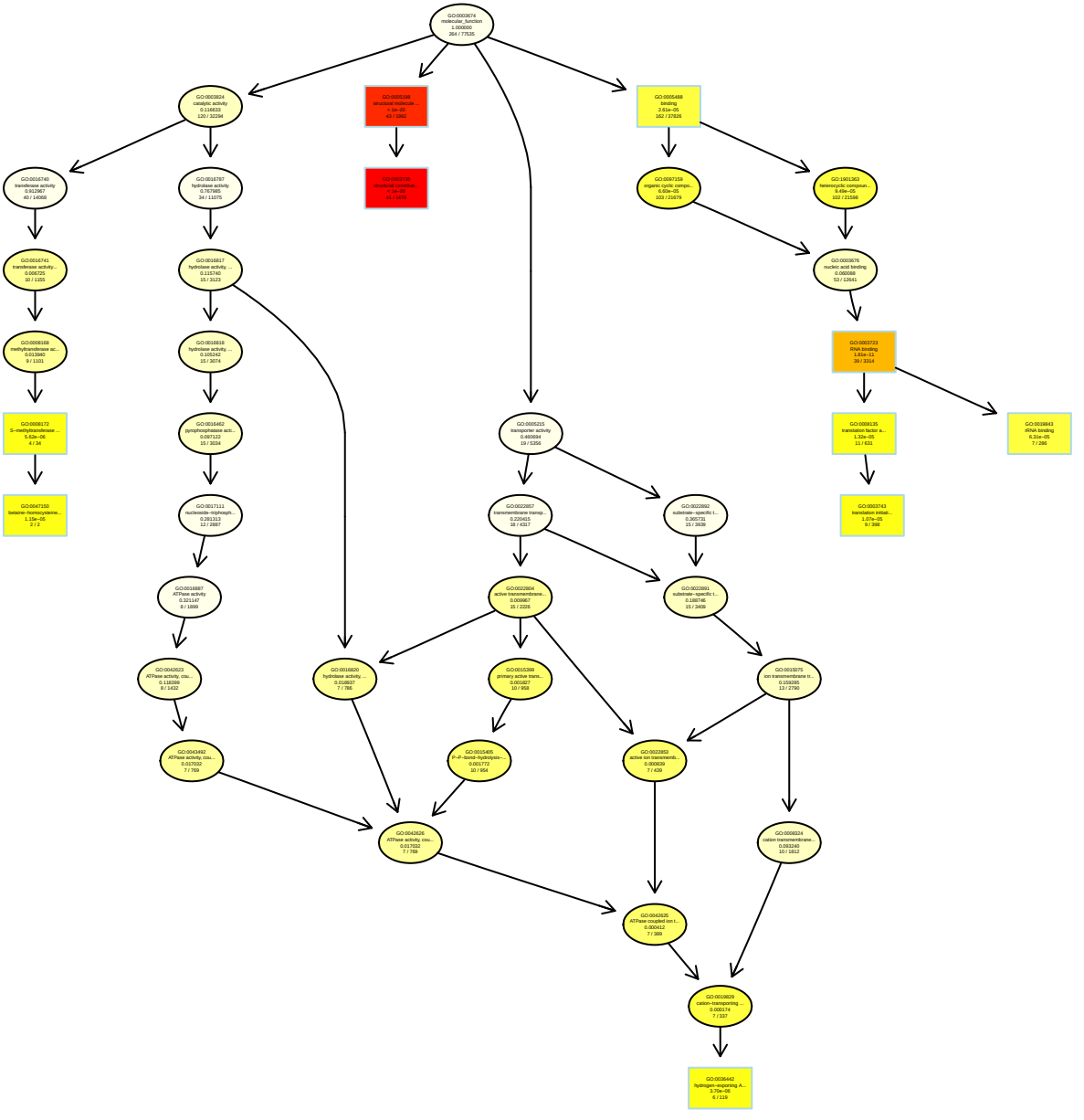
75 / 14373
GO:0006412
translation
< 1e−20
54 / 2828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364 / 77535
GO:0003824
catalytic activity
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120 / 32294
GO:0005198
structural molecule ...
x 1e-20
43 / 5002
GO:0005488
binding
2.61e-05
162 / 27826
GO:0016740
transferase activity
0.012938
40 / 14068
GO:0016787
hydrolase activity
0.087982
34 / 11075
GO:0003735
structural constituent...
0.000128
41 / 5476
GO:0097159
organic cyclic compo...
0.000426
103 / 21675
GO:1901363
heterocyclic compoun...
0.000426
102 / 21548
GO:0016741
transferase activity,...
0.006725
10 / 1355
GO:0016817
hydrolase activity, ...
0.115740
15 / 3123
GO:0003676
nucleic acid binding
0.060088
53 / 12641
GO:0008168
methyltransferase ac...
0.013940
9 / 1101
GO:0016818
hydrolase activity, ...
0.105242
15 / 3074
GO:0003723
RNA binding
1.81e-11
39 / 3314
GO:0008172
S-methyltransferase ...
5.63e-06
4 / 34
GO:0016462
pyrophosphatase acti...
0.087122
15 / 3016
GO:0005215
transporter activity
0.460934
19 / 5358
GO:0008135
translation factor a...
1.13e-05
11 / 631
GO:0019843
rRNA binding
6.31e-08
7 / 286
GO:0047150
betaine-homocysteine...
1.15e-05
2 / 2
GO:0017111
nucleoside-triphosph...
0.281313
12 / 2887
GO:0022857
transmembrane transp...
0.226415
18 / 4317
GO:0022892
substrate-specific t...
0.365731
15 / 3639
GO:0003743
translation initiat...
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9 / 398
GO:0016887
ATPase activity
0.321147
8 / 1899
GO:0022804
active transmembrane ...
0.009967
25 / 2326
GO:0022891
substrate-specific t...
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GO:0042623
ATPase activity, cou...
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GO:0016820
hydrolase activity, ...
0.016637
7 / 786
GO:0015399
primary active trans...
0.001827
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GO:0015075
ion transmembrane tr...
0.159285
13 / 2790
GO:0043492
ATPase activity, cou...
0.017032
7 / 769
GO:0015405
P-P-bond-hydrolysis-...
0.001772
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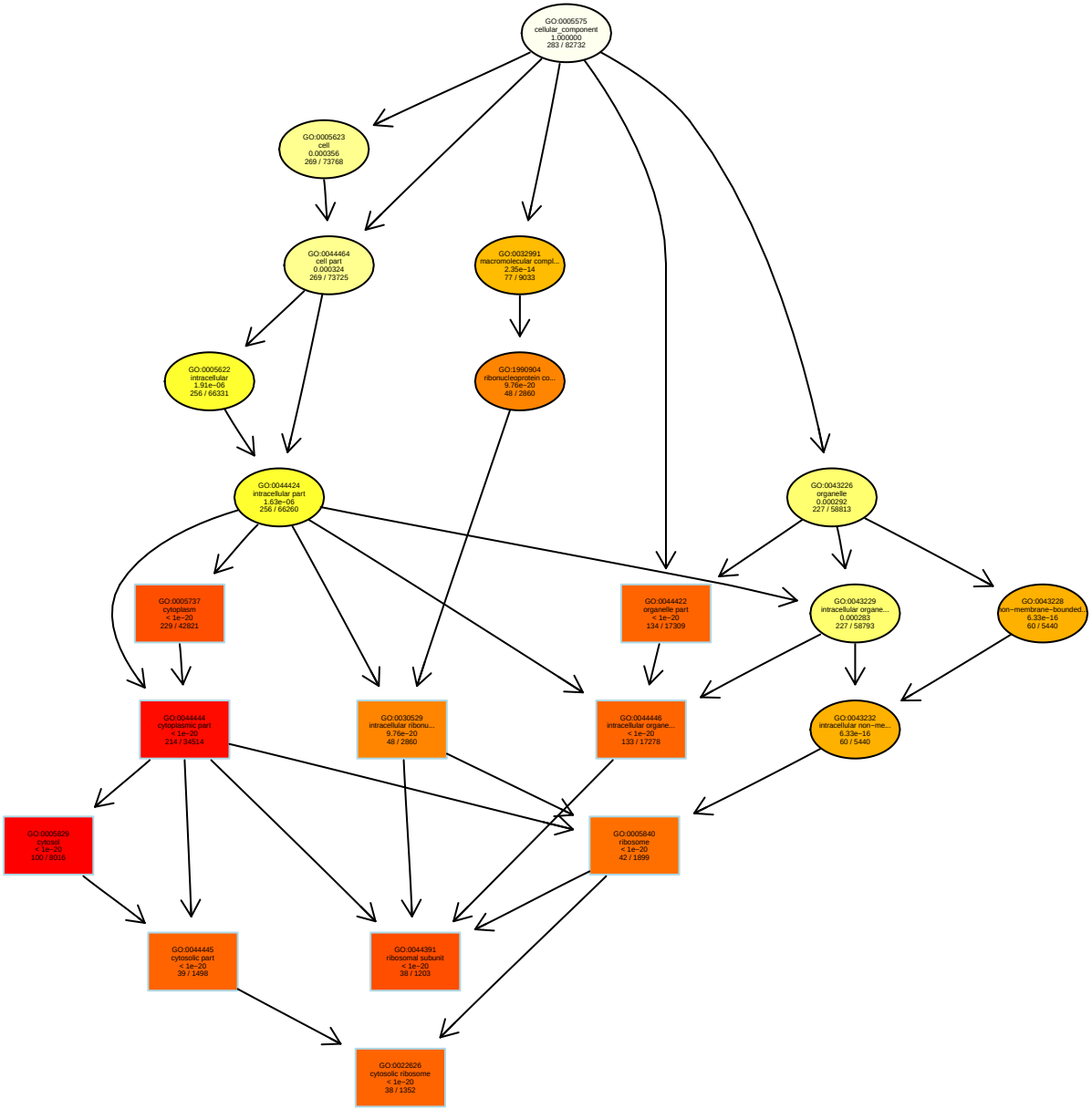
GO:0022853
active ion transmemb...
0.000839
7 / 439
GO:0042626
ATPase activity, cou...
0.017032
7 / 769
GO:0008324
cation transmembrane...
0.092240
10 / 1812
GO:0042625
ATPase coupled ion t...
0.000412
7 / 389
GO:0019829
cation-transporting ...
0.00374
7 / 337
GO:0036442
hydrogen-exporting A...
3.70e-08
6 / 119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GO:0005623
cell
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269 / 73768
GO:0044464
cell part
0.000324
269 / 73725
GO:0032991
macromolecular compl...
2.35e−14
77 / 9033
GO:0005622
intracellular
1.91e−06
256 / 66331
GO:1990904
ribonucleoprotein co...
9.76e−20
48 / 2860
GO:0044424
intracellular part
1.63e−06
256 / 66260
GO:0043226
organelle
0.000292
227 / 58813
GO:0005737
cytoplasm
< 1e−20
229 / 42821
GO:0044422
organelle part
< 1e−20
134 / 17309
GO:0043229
intracellular organe...
0.000293
227 / 58793
GO:0043228
non−membrane−bounded...
6.33e−16
60 / 5440
GO:0044444
cytoplasmic part
< 1e−20
214 / 34514
GO:0030529
intracellular ribonu...
9.76e−20
48 / 2860
GO:0044446
intracellular organe...
< 1e−20
133 / 17278
GO:0043232
intracellular non−me...
6.33e−16
60 / 5440
GO:0005829
cytosol
< 1e−20
100 / 8016
GO:0005840
ribosome
< 1e−20
42 / 1899
GO:0044445
cytosolic part
< 1e−20
39 / 1498
GO:0044391
ribosomal subunit
< 1e−20
38 / 1203
GO:0022626
cytosolic ribosome
< 1e−20
38 / 1352