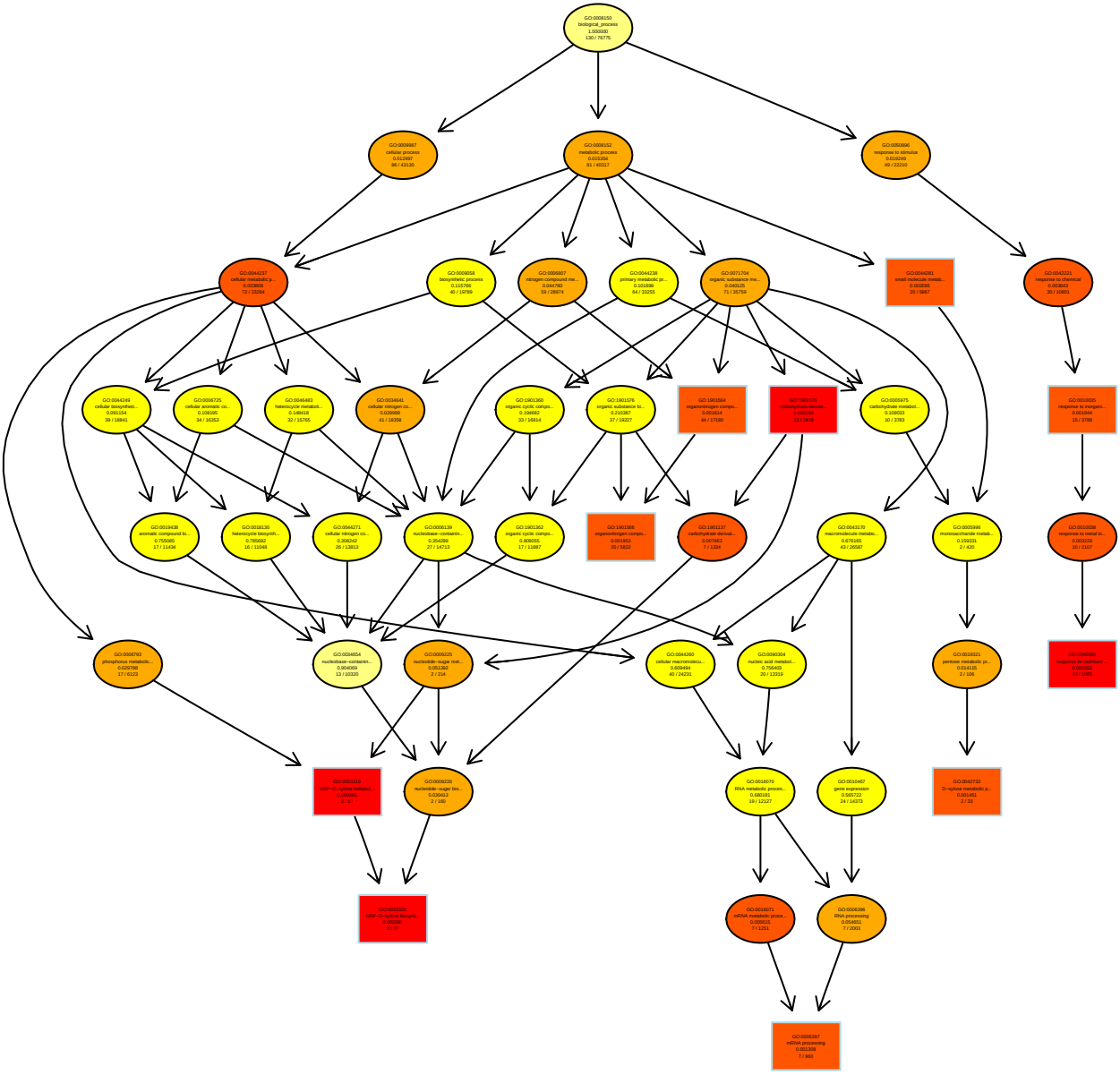
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GO:0008152
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GO:0050896
response to stimulus
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49 / 22210
GO:0044237
cellular metabolic p...
0.903809
72 / 33284
GO:0009058
biosynthetic process
0.112766
45 / 19789
GO:0006807
nitrogen compound me...
0.044765
59 / 28874
GO:0044238
primary metabolic pr...
0.151599
64 / 33255
GO:0071704
organic substance me...
0.040125
71 / 35759
GO:0044281
small molecule metab...
0.002095
20 / 5867
GO:0042221
response to chemical
0.003543
30 / 10601
GO:0044249
cellular biosynthet...
0.091154
39 / 18841
GO:0006725
cellular aromatic co...
0.188195
34 / 16353
GO:0046483
heterocycle metabol...
0.149418
32 / 15765
GO:0034641
cellular nitrogen co...
0.028988
41 / 18358
GO:1901360
organic cyclic compo...
0.194692
33 / 16814
GO:1901576
organic substance bi...
0.210387
27 / 13227
GO:1901564
organonitrogen compo...
0.001814
44 / 17180
GO:1901135
carbohydrate derivat...
0.000269
13 / 2426
GO:0005975
carbohydrate metabol...
0.109023
10 / 2763
GO:0010035
response to inorganic...
0.001944
31 / 2788
GO:0019438
aromatic compound bi...
0.755085
17 / 11434
GO:0018130
heterocycle biosynth...
0.785692
16 / 10048
GO:0044271
cellular nitrogen co...
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GO:0006139
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GO:0043170
macromolecule metabo...
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43 / 20587
GO:0009996
monosaccharide metab...
0.159331
7 / 420
GO:0010038
response to metal io...
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GO:0006793
phosphorus metabolic...
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17 / 6123
GO:0034654
nucleobase-containin...
0.904069
13 / 10320
GO:0009225
nucleotide-sugar meta...
0.051392
2 / 234
GO:0044260
cellular macromolecu...
0.609494
40 / 24231
GO:0090304
nucleic acid metabol...
0.756403
20 / 13319
GO:0019321
pentose metabolic pr...
0.014135
2 / 106
GO:0046690
response to cadmium ...
0.001052
16 / 1505
GO:0003319
UDP-D-xylose metabol...
0.043391
2 / 17
GO:0009226
nucleotide-sugar bio...
0.020413
2 / 165
GO:0016070
RNA metabolic proces...
0.685181
19 / 12127
GO:0010467
gene expression
0.585723
24 / 14373
GO:0042732
D-xylose metabolic p...
0.001491
2 / 33
GO:0003320
UDP-D-xylose biosynt...
0.000391
2 / 17
GO:0016071
mRNA metabolic proce...
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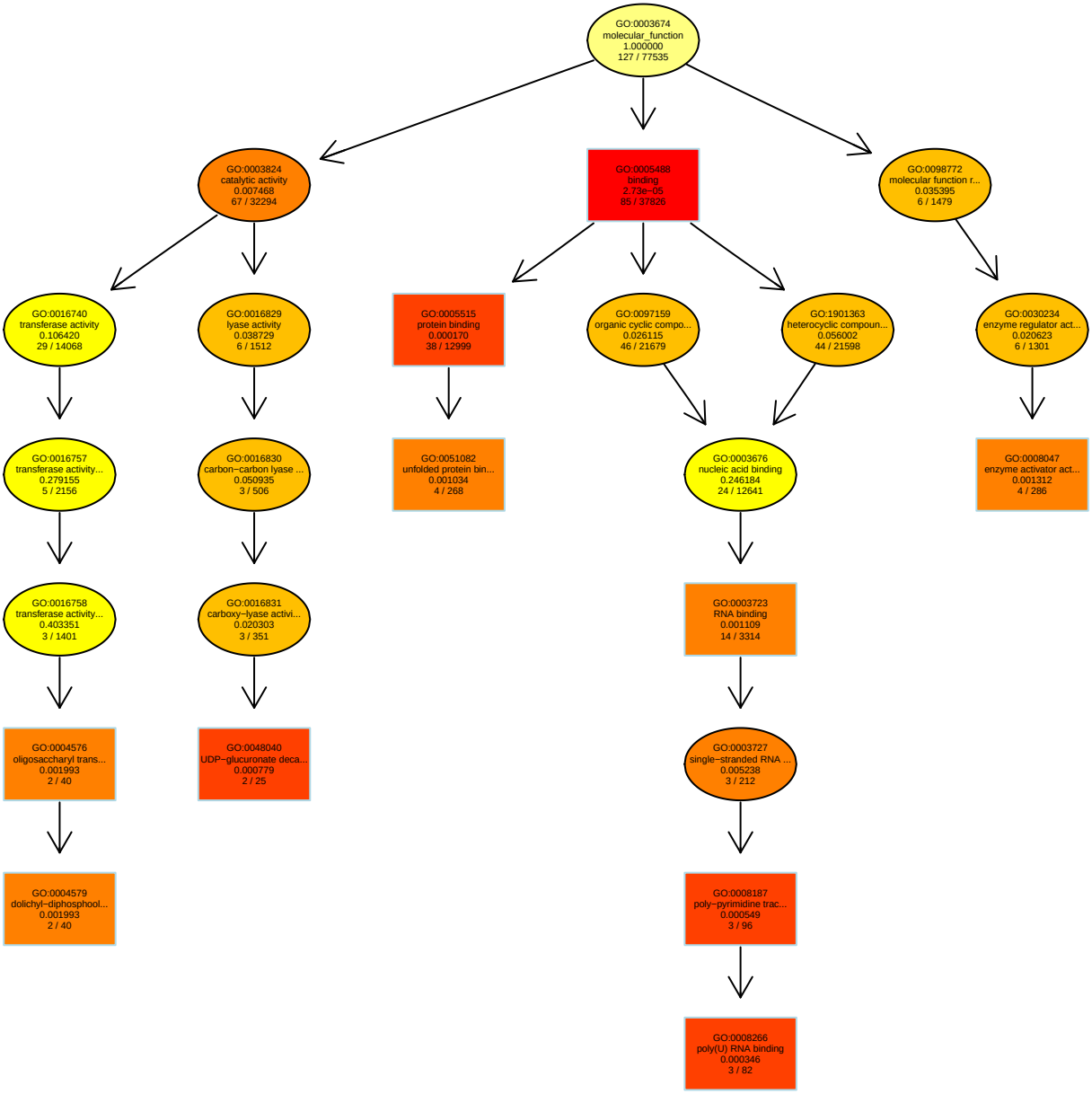
7 / 1251
GO:0006396
RNA processing
0.654651
7 / 2003
GO:0006397
mRNA processing
0.551108
7 / 960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67 / 32294
GO:0005488
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85 / 37826
GO:0098772
molecular function r...
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6 / 1479
GO:0016740
transferase activity
0.106420
29 / 14068
GO:0016829
lyase activity
0.038729
6 / 1512
GO:0005515
protein binding
0.000170
38 / 12999
GO:0097159
organic cyclic compo...
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46 / 21679
GO:1901363
heterocyclic compoun...
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44 / 21598
GO:0030234
enzyme regulator act...
0.020623
6 / 1301
GO:0016757
transferase activity...
0.279155
5 / 2156
GO:0016830
carbon−carbon lyase ...
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3 / 506
GO:0051082
unfolded protein bin...
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4 / 268
GO:0003676
nucleic acid binding
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24 / 12641
GO:0009047
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0.001312
4 / 286
GO:0016758
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GO:0016831
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3 / 351
GO:0003723
RNA binding
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14 / 3314
GO:0004576
oligosaccharyl trans...
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GO:0048040
UDP−glucuronate deca...
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GO:0003727
single−stranded RNA ...
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3 / 212
GO:0004579
dolichyl−diphosphool...
0.001993
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GO:0008187
poly−pyrimidine trac...
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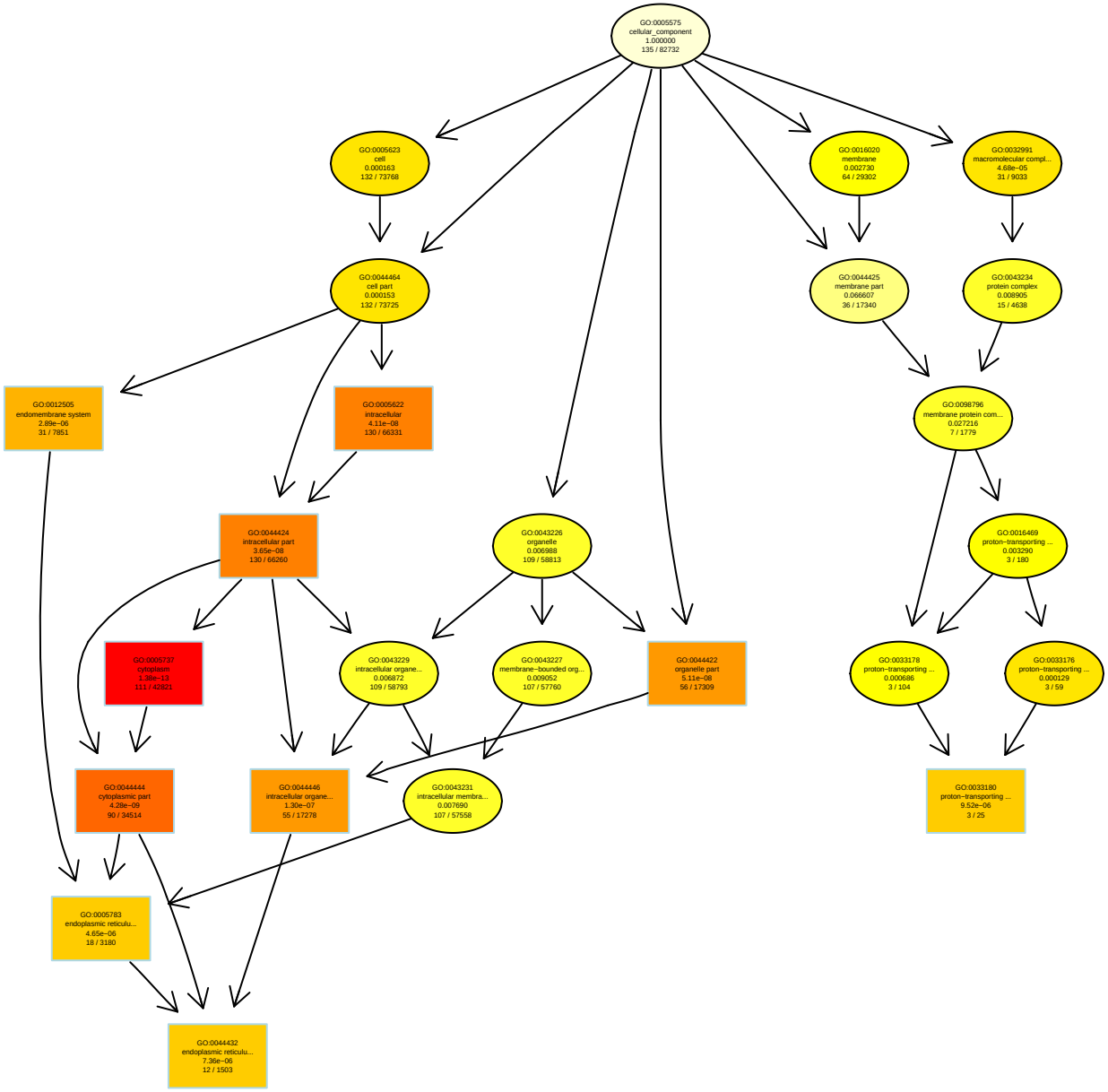
3 / 96
GO:0008266
poly(U) RNA binding
0.000346
3 / 82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GO:0016020
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64 / 29302
GO:0032991
macromolecular compl...
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31 / 9033
GO:0044464
cell part
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132 / 73725
GO:0044425
membrane part
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36 / 17340
GO:0043234
protein complex
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15 / 4638
GO:0012505
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GO:0005622
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GO:0098796
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7 / 1779
GO:0044424
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130 / 66260
GO:0043226
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GO:0016469
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3 / 180
GO:0005737
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109 / 58793
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GO:0033178
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GO:0033176
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3 / 59
GO:0044444
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GO:0044446
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GO:0043231
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GO:0033180
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GO:0005783
endoplasmic reticulu...
4.65e−06
18 / 3180
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endoplasmic reticulu...
7.36e−06
12 / 1503