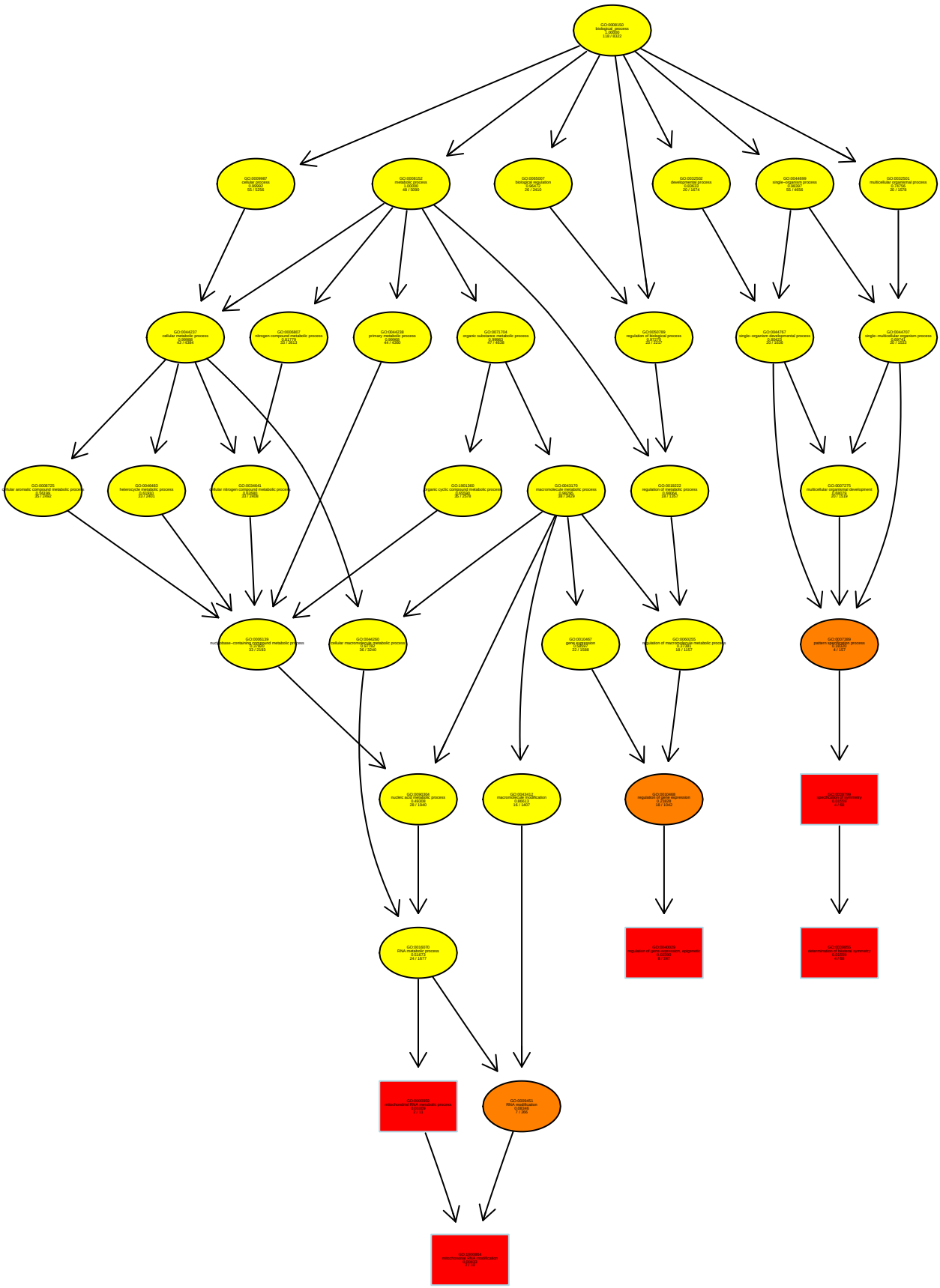
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biological_process
1.00000
119 / 8222
GO:0009987
cellular process
0.99602
55 / 5255
GO:0008152
metabolic process
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49 / 5090
GO:0065007
biological regulation
0.98472
26 / 2610
GO:0032502
developmental process
0.93537
20 / 1674
GO:0044699
single-organism process
0.98287
55 / 4656
GO:0032501
multicellular organismal process
0.78756
30 / 1578
GO:0044237
cellular metabolic process
0.97884
43 / 3884
GO:0006807
nitrogen compound metabolic process
0.97303
35 / 2923
GO:0044238
primary metabolic process
0.97355
44 / 3935
GO:0071704
organic substance metabolic process
0.97355
44 / 3935
GO:0050789
regulation of biological process
0.97223
37 / 2223
GO:0044767
single-organism developmental process
0.97256
30 / 1656
GO:0044707
single-multicellular organism process
0.97223
30 / 1623
GO:0006725
cellular aromatic compound metabolic process
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25 / 1456
GO:0046483
heterocycle metabolic process
0.95201
23 / 1401
GO:0034641
cellular nitrogen compound metabolic process
0.95081
23 / 1408
GO:1901360
organic cyclic compound metabolic process
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25 / 1575
GO:0043170
macromolecule metabolic process
0.94245
28 / 1829
GO:0019222
regulation of metabolic process
0.93824
17 / 1157
GO:0007275
multicellular organismal development
0.94674
20 / 1316
GO:0006139
nucleobase-containing compound metabolic process
0.97922
33 / 2193
GO:0044260
cellular macromolecule metabolic process
0.97532
26 / 1640
GO:0010467
gene expression
0.91910
22 / 1588
GO:0060255
regulation of macromolecule metabolic process
0.91185
18 / 1157
GO:0007389
pattern specification process
0.91910
4 / 157
GO:0090304
nucleic acid metabolic process
0.97532
26 / 1640
GO:0043412
macromolecule modification
0.93910
16 / 1027
GO:0010468
regulation of gene expression
0.91185
16 / 1042
GO:0009799
specification of symmetry
0.91910
4 / 157
GO:0016070
RNA metabolic process
0.51577
24 / 1677
GO:0040029
regulation of gene expression, epigenetic
0.91910
4 / 157
GO:0009855
determination of bilateral symmetry
0.91910
4 / 157
GO:0000963
mitochondrial RNA metabolic process
0.00000
2 / 11
GO:0009451
RNA modification
0.58346
7 / 266
GO:1900864
mitochondrial RNA modification
0.00000
2 / 10