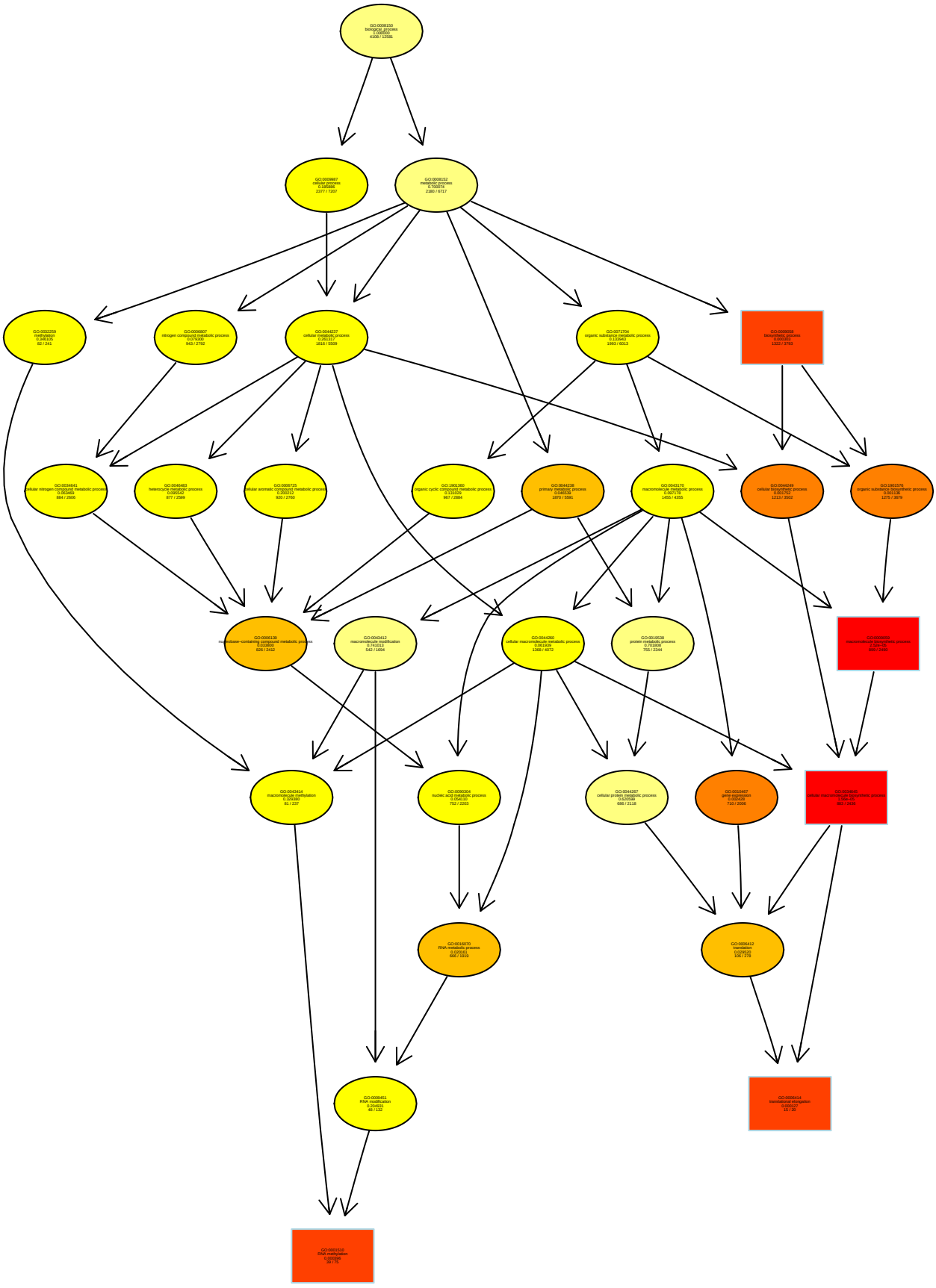
GO:0008150
biological_process
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4108 / 12581
GO:0009987
cellular process
0.185886
2377 / 7207
GO:0008152
metabolic process
0.700074
2180 / 6717
GO:0032259
methylation
0.346105
82 / 241
GO:0006807
nitrogen compound metabolic process
0.079300
943 / 2792
GO:0044237
cellular metabolic process
0.361317
1816 / 5509
GO:0071704
organic substance metabolic process
0.133943
1993 / 6013
GO:0009058
biosynthetic process
0.000333
1322 / 3793
GO:0034641
cellular nitrogen compound metabolic process
0.063469
884 / 2606
GO:0046483
heterocycle metabolic process
0.095542
877 / 2599
GO:0006725
cellular aromatic compound metabolic process
0.200212
920 / 2760
GO:1901360
organic cyclic compound metabolic process
0.131029
967 / 2884
GO:0044238
primary metabolic process
0.046539
1870 / 5591
GO:0043170
macromolecule metabolic process
0.097178
1455 / 4355
GO:0044249
cellular biosynthetic process
0.051792
1213 / 3502
GO:1901576
organic substance biosynthetic process
0.001136
1275 / 3879
GO:0006139
nucleobase-containing compound metabolic process
0.033800
826 / 2412
GO:0043412
macromolecule modification
0.741013
542 / 1694
GO:0044260
cellular macromolecule metabolic process
0.061959
1368 / 4072
GO:0019538
protein metabolic process
0.701808
755 / 2344
GO:0009059
macromolecule biosynthetic process
2.53e-05
869 / 2490
GO:0043414
macromolecule methylation
0.329380
81 / 237
GO:0090304
nucleic acid metabolic process
0.054110
752 / 2203
GO:0044267
cellular protein metabolic process
0.620599
686 / 2118
GO:0010467
gene expression
0.002428
710 / 2006
GO:0034645
cellular macromolecule biosynthetic process
1.56e-05
883 / 2436
GO:0016070
RNA metabolic process
0.020161
666 / 1919
GO:0006412
translation
0.028520
106 / 278
GO:0009451
RNA modification
0.204931
48 / 132
GO:0006414
translational elongation
0.000127
15 / 20
GO:0001510
RNA methylation
0.000396
39 / 75