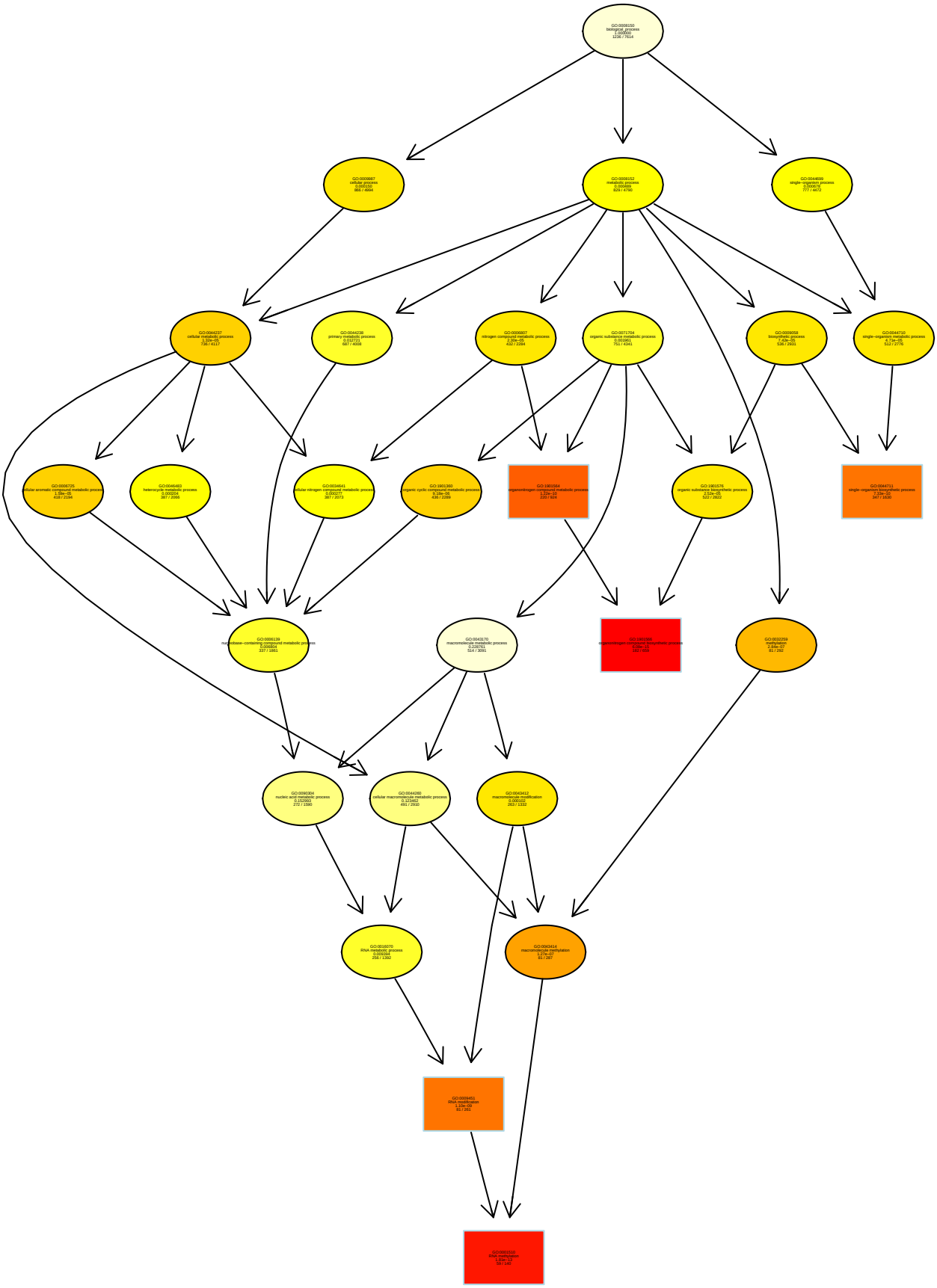
GO:0008150
biological_process
1.000000
1236 / 7614
GO:0009987
cellular process
0.550150
866 / 4994
GO:0008152
metabolic process
0.000089
829 / 4790
GO:0044699
single−organism process
0.700076
777 / 4472
GO:0044237
cellular metabolic process
0.000000
730 / 4123
GO:0044238
primary metabolic process
0.000000
687 / 4028
GO:0006807
nitrogen compound metabolic process
0.000000
622 / 2954
GO:0071704
organic substance metabolic process
0.000000
751 / 4341
GO:0009058
biosynthetic process
1.11e−16
538 / 2601
GO:0044710
single−organism metabolic process
0.000000
517 / 2776
GO:0006725
cellular aromatic compound metabolic process
1.19e−05
419 / 2194
GO:0046483
heterocycle metabolic process
0.000204
397 / 2066
GO:0034641
cellular nitrogen compound metabolic process
0.000277
387 / 2073
GO:1901360
organic cyclic compound metabolic process
9.18e−05
436 / 2269
GO:1901564
organonitrogen compound metabolic process
1.23e−10
265 / 924
GO:1901576
organic substance biosynthetic process
2.52e−05
522 / 2625
GO:0044711
single−organism biosynthetic process
7.19e−16
247 / 1430
GO:0006139
nucleobase−containing compound metabolic process
0.000004
337 / 1861
GO:0043170
macromolecule metabolic process
0.230761
514 / 3091
GO:1901565
organonitrogen compound biosynthetic process
2.09e−12
182 / 659
GO:0032259
methylation
2.84e−27
81 / 292
GO:0090304
nucleic acid metabolic process
0.000000
272 / 1586
GO:0044260
cellular macromolecule metabolic process
0.000000
681 / 2955
GO:0043412
macromolecule modification
0.000000
263 / 1122
GO:0016070
RNA metabolic process
0.000384
256 / 1392
GO:0043414
macromolecule methylation
1.27e−07
61 / 267
GO:0009451
RNA modification
1.19e−09
51 / 261
GO:0001510
RNA methylation
1.01e−12
50 / 187