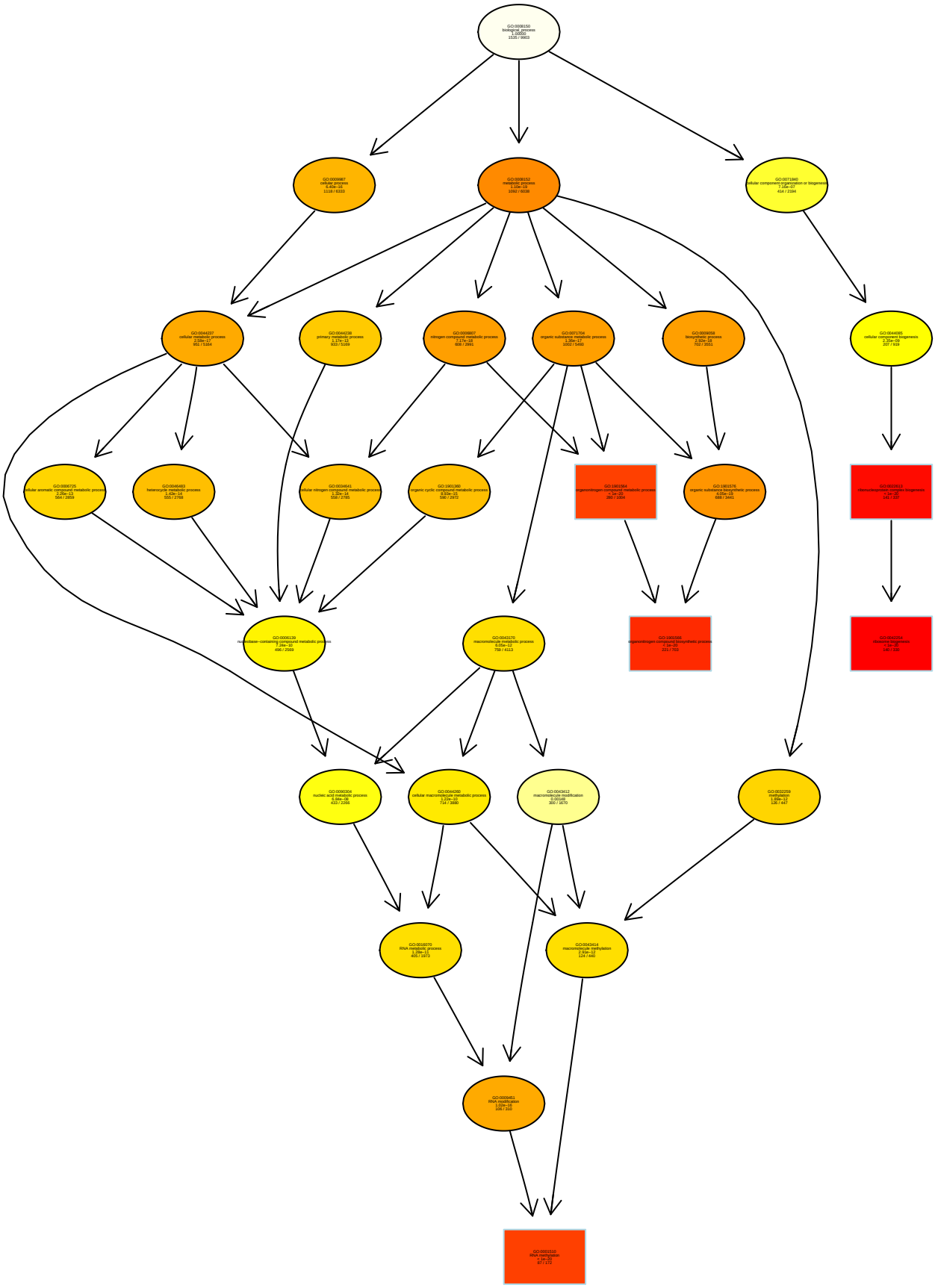
GO:0008150
biological_process
1.00000
1135 / 9903
GO:0009987
cellular process
6.48e−18
1118 / 6333
GO:0008152
metabolic process
1.10e−19
1092 / 6038
GO:0071840
cellular component organization or biogenesis
7.16e−27
414 / 2194
GO:0044237
cellular metabolic process
1.00e−19
821 / 5164
GO:0044238
primary metabolic process
1.00e−18
810 / 5109
GO:0006807
nitrogen compound metabolic process
1.00e−18
659 / 2991
GO:0071704
organic substance metabolic process
1.00e−18
1002 / 5491
GO:0009058
biosynthetic process
1.00e−18
707 / 3551
GO:0044085
cellular component biogenesis
2.27e−23
327 / 918
GO:0006725
cellular aromatic compound metabolic process
7.29e−13
564 / 2859
GO:0046483
heterocycle metabolic process
1.43e−14
555 / 2768
GO:0034641
cellular nitrogen compound metabolic process
1.20e−14
558 / 2785
GO:1901360
organic cyclic compound metabolic process
8.93e−15
566 / 2972
GO:1901564
organonitrogen compound metabolic process
< 1e−30
280 / 1004
GO:1901576
organic substance biosynthetic process
4.05e−19
688 / 3441
GO:0022613
ribonucleoprotein complex biogenesis
< 1e−20
141 / 337
GO:0006139
nucleobase−containing compound metabolic process
7.24e−10
496 / 2569
GO:0043170
macromolecule metabolic process
6.05e−12
759 / 4113
GO:1901566
organonitrogen compound biosynthetic process
< 1e−30
221 / 703
GO:0042254
ribosome biogenesis
< 1e−20
140 / 336
GO:0090304
nucleic acid metabolic process
2.84e−10
433 / 2266
GO:0044260
cellular macromolecule metabolic process
1.00e−12
717 / 3880
GO:0043412
macromolecule modification
0.00148
300 / 1670
GO:0033259
rna biosynthesis
1.69e−12
126 / 447
GO:0016070
RNA metabolic process
1.29e−10
405 / 1973
GO:0043414
macromolecule methylation
1.29e−12
124 / 440
GO:0009451
RNA modification
1.02e−12
106 / 319
GO:0001510
RNA methylation
1.02e−12
67 / 172