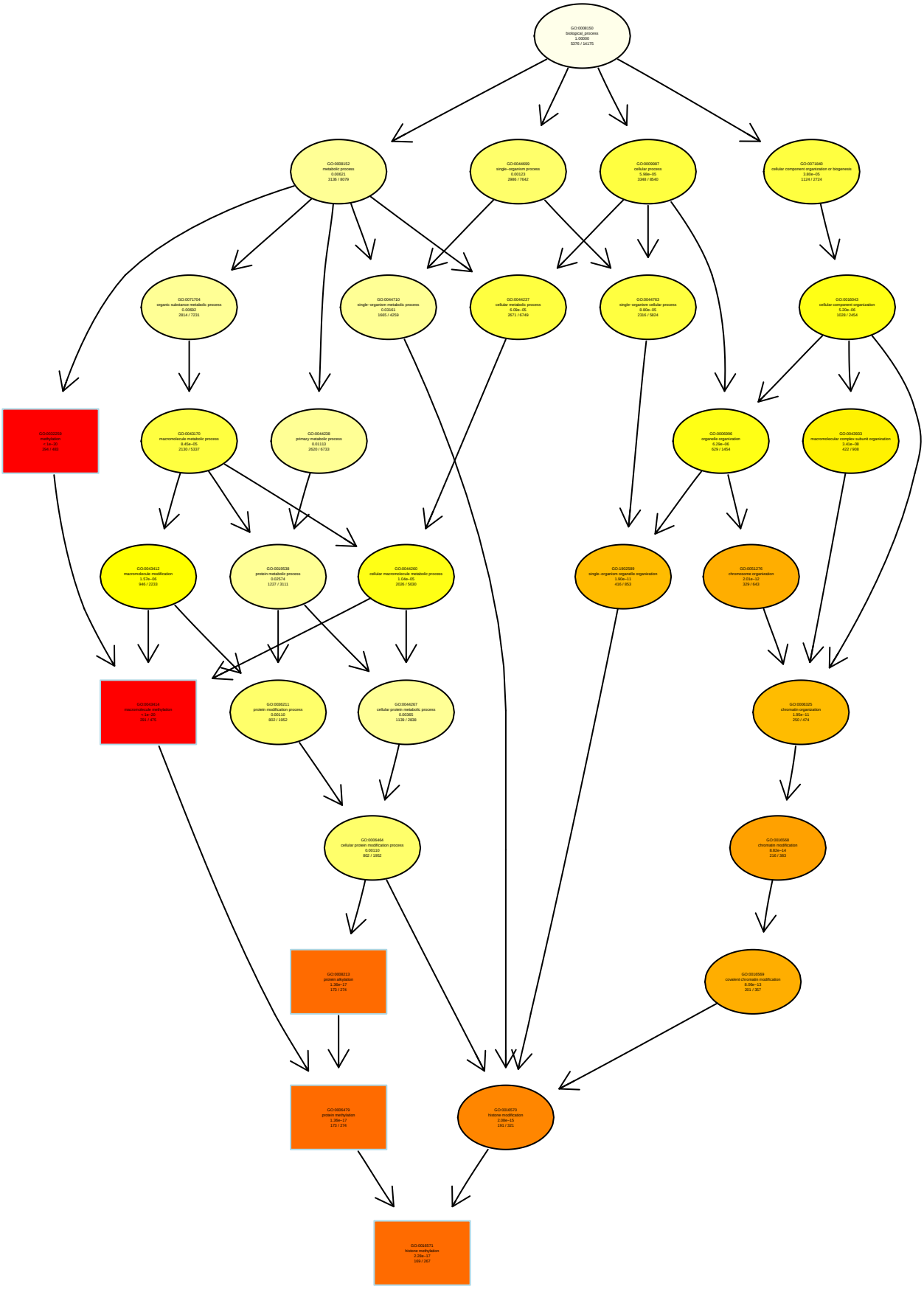
GO:0008150
biological_process
1.00000
5376 / 14175
GO:0008152
metabolic process
0.00821
3136 / 8079
GO:0044699
single−organism process
0.00123
2986 / 7642
GO:0009987
cellular process
5.98e−05
3248 / 8540
GO:0071840
cellular component organization or biogenesis
1.80e−05
1124 / 2724
GO:0071704
organic substance metabolic process
0.00892
2814 / 7231
GO:0044710
single−organism metabolic process
0.03161
1665 / 4259
GO:0044237
cellular metabolic process
6.09e−05
2671 / 6749
GO:0044763
single−organism cellular process
8.80e−05
2316 / 5824
GO:0016043
cellular component organization
5.20e−05
1028 / 2454
GO:0032259
methylation
< 1e−20
294 / 489
GO:0043170
macromolecule metabolic process
8.45e−05
2130 / 5337
GO:0044238
primary metabolic process
0.01113
2620 / 6733
GO:0006996
organelle organization
6.29e−06
629 / 1454
GO:0043933
macromolecular complex subunit organization
3.41e−08
422 / 908
GO:0043412
macromolecule modification
1.37e−06
946 / 2233
GO:0019538
protein metabolic process
0.02574
1227 / 3111
GO:0044260
cellular macromolecule metabolic process
1.04e−05
2026 / 5030
GO:1902589
single−organism organelle organization
1.90e−11
416 / 853
GO:0051276
chromosome organization
2.01e−12
329 / 643
GO:0043414
macromolecule methylation
< 1e−20
254 / 475
GO:0036211
protein modification process
0.00110
802 / 1952
GO:0044267
cellular protein metabolic process
0.00365
1139 / 2838
GO:0006325
chromatin organization
1.95e−11
250 / 474
GO:0006464
cellular protein modification process
0.00110
802 / 1952
GO:0016568
chromatin modification
8.62e−14
216 / 383
GO:0008213
protein alkylation
1.36e−17
173 / 274
GO:0016569
covalent chromatin modification
8.09e−13
201 / 357
GO:0006479
protein methylation
1.36e−17
173 / 274
GO:0016570
histone modification
2.08e−15
191 / 321
GO:0016571
histone methylation
2.28e−17
169 / 267