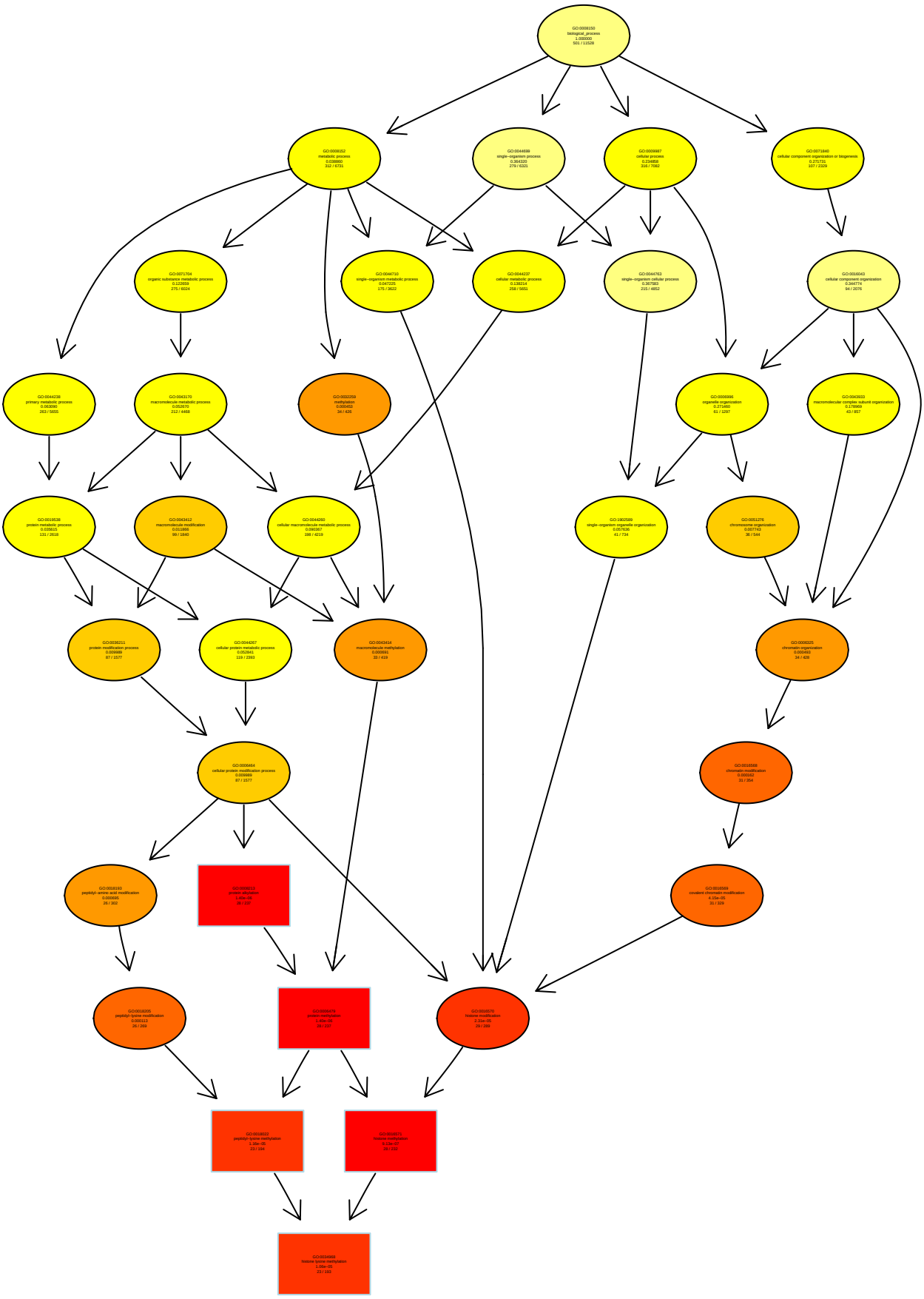
GO:0008150
biological_process
1.000000
501 / 11528
GO:0008152
metabolic process
0.038860
312 / 6731
GO:0044699
single-organism process
0.364320
279 / 6321
GO:0009987
cellular process
0.234858
316 / 7082
GO:0071840
cellular component organization or biogenesis
0.271731
107 / 2329
GO:0071704
organic substance metabolic process
0.122659
275 / 6024
GO:0044710
single-organism metabolic process
0.047225
175 / 3622
GO:0044237
cellular metabolic process
0.138214
258 / 5651
GO:0044763
single-organism cellular process
0.367583
215 / 4852
GO:0016043
cellular component organization
0.344774
94 / 2076
GO:0044238
primary metabolic process
0.063090
263 / 5655
GO:0043170
macromolecule metabolic process
0.053570
212 / 4468
GO:0032259
methylation
0.000453
34 / 426
GO:0006996
organelle organization
0.271460
61 / 1297
GO:0043933
macromolecular complex subunit organization
0.178969
43 / 857
GO:0019538
protein metabolic process
0.035615
131 / 2618
GO:0043412
macromolecule modification
0.011895
99 / 1840
GO:0044260
cellular macromolecule metabolic process
0.090367
198 / 4219
GO:1902589
single-organism organelle organization
0.057636
41 / 734
GO:0051276
chromosome organization
0.007743
36 / 544
GO:0036211
protein modification process
0.009989
87 / 1577
GO:0044267
cellular protein metabolic process
0.052841
119 / 2393
GO:0043414
macromolecule methylation
0.000691
33 / 419
GO:0006325
chromatin organization
0.000493
34 / 428
GO:0006464
cellular protein modification process
0.009989
87 / 1577
GO:0016568
chromatin modification
0.000162
31 / 354
GO:0018193
peptidyl-amino acid modification
0.000695
26 / 302
GO:0006313
protein alkylation
1.40e-06
28 / 237
GO:0016569
covalent chromatin modification
4.15e-05
31 / 329
GO:0018205
peptidyl-lysine modification
0.000113
26 / 269
GO:0006479
protein methylation
1.40e-06
28 / 237
GO:0016570
histone modification
2.31e-05
29 / 289
GO:0018022
peptidyl-lysine methylation
1.16e-05
23 / 194
GO:0016571
histone methylation
9.13e-07
28 / 232
GO:0034968
histone lysine methylation
1.06e-05
23 / 193