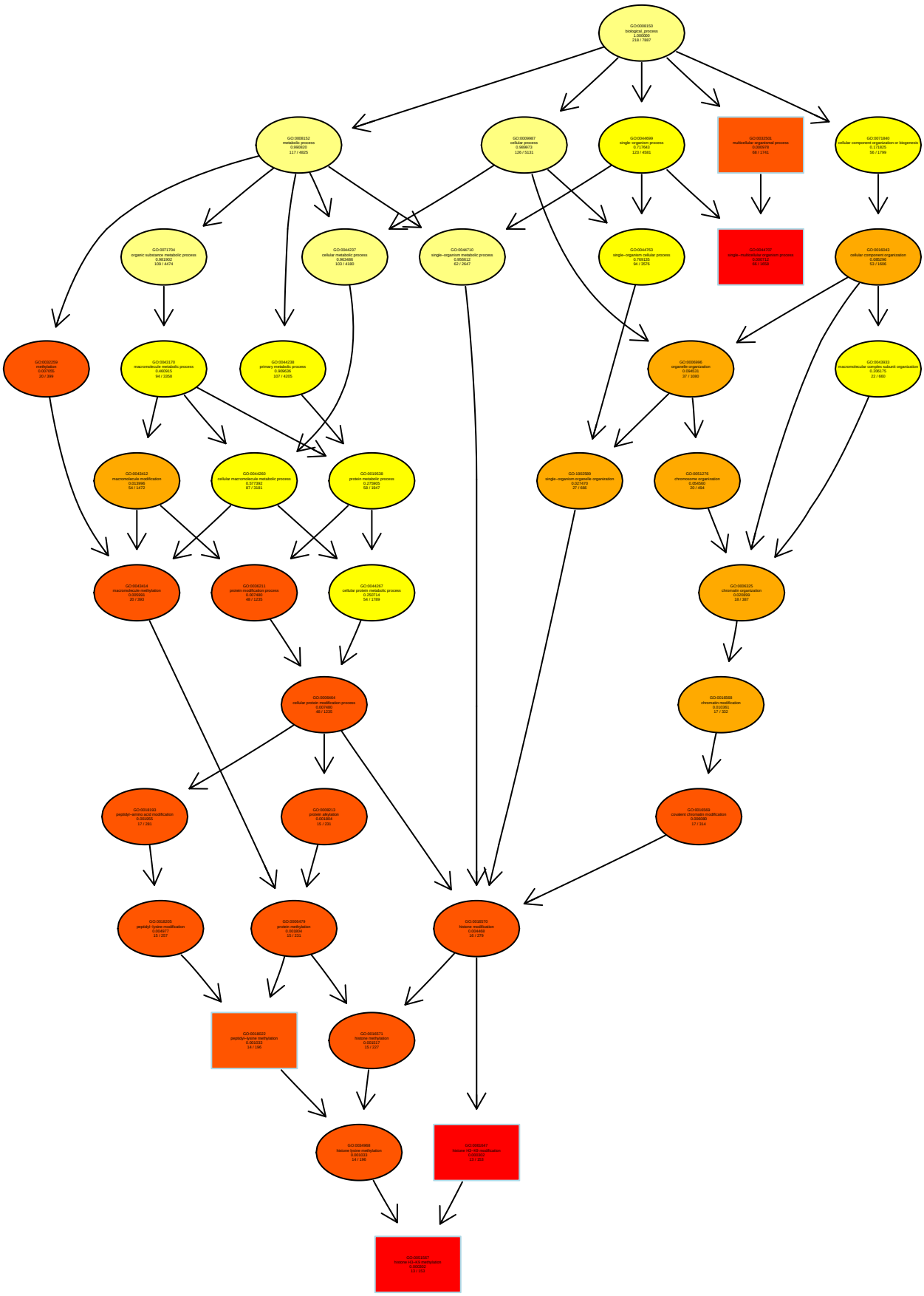
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biological_process
1.000000
218 / 7887
GO:0008152
metabolic process
0.990820
117 / 4825
GO:0009987
cellular process
0.989973
126 / 5131
GO:0044699
single-organism process
0.717643
123 / 4581
GO:0032501
multicellular organismal process
0.000978
68 / 1741
GO:0071840
cellular component organization or biogenesis
0.171825
56 / 1799
GO:0071704
organic substance metabolic process
0.991902
109 / 4474
GO:0044237
cellular metabolic process
0.963495
103 / 4180
GO:0044710
single-organism metabolic process
0.955612
62 / 2647
GO:0044763
single-organism cellular process
0.799135
94 / 3576
GO:0044707
single-multicellular organism process
0.000712
68 / 1658
GO:0016043
cellular component organization
0.085296
53 / 1606
GO:0032259
methylation
0.007056
20 / 399
GO:0043170
macromolecule metabolic process
0.460915
94 / 3358
GO:0044238
primary metabolic process
0.909636
107 / 4205
GO:0006996
organelle organization
0.084531
37 / 1080
GO:0043933
macromolecular complex subunit organization
0.206175
22 / 660
GO:0043412
macromolecule modification
0.013996
54 / 1472
GO:0044260
cellular macromolecule metabolic process
0.577392
87 / 3181
GO:0019538
protein metabolic process
0.275905
58 / 1947
GO:1902589
single-organism organelle organization
0.027470
27 / 666
GO:0051276
chromosome organization
0.054560
20 / 494
GO:0043414
macromolecule methylation
0.005991
20 / 393
GO:0036211
protein modification process
0.007480
48 / 1235
GO:0044267
cellular protein metabolic process
0.250714
54 / 1789
GO:0006325
chromatin organization
0.020899
18 / 387
GO:0006464
cellular protein modification process
0.007480
48 / 1235
GO:0016568
chromatin modification
0.010361
17 / 332
GO:0018193
peptidyl-amino acid modification
0.001955
17 / 281
GO:0008213
protein alkylation
0.001804
15 / 231
GO:0016569
covalent chromatin modification
0.006080
17 / 314
GO:0018205
peptidyl-lysine modification
0.004977
15 / 257
GO:0006479
protein methylation
0.001804
15 / 231
GO:0016570
histone modification
0.004468
14 / 279
GO:0018022
peptidyl-lysine methylation
0.001033
14 / 196
GO:0016571
histone methylation
0.001517
15 / 227
GO:0034968
histone lysine methylation
0.001033
14 / 196
GO:0061647
histone H3-K9 modification
0.000302
13 / 153
GO:0051567
histone H3-K9 methylation
0.000302
13 / 153