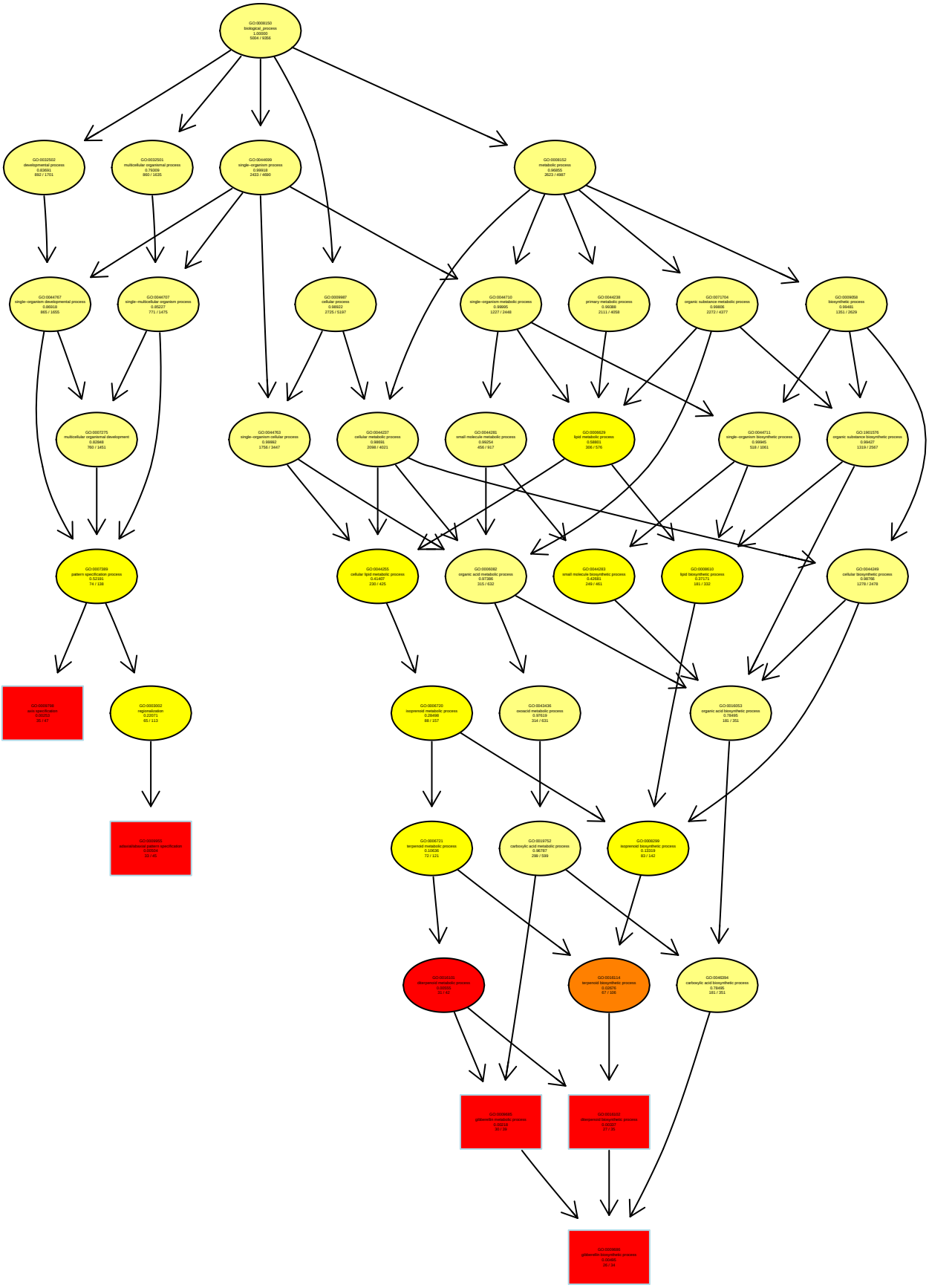
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5304 / 9356
GO:0032502
developmental process
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882 / 1701
GO:0032501
multicellular organismal process
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860 / 1635
GO:0044699
single−organism process
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2433 / 4890
GO:0008152
metabolic process
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2623 / 4987
GO:0044767
single−organism developmental process
0.86918
865 / 1655
GO:0044707
single−multicellular organism process
0.85227
771 / 1475
GO:0009987
cellular process
0.98622
2725 / 5197
GO:0044710
single−organism metabolic process
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1227 / 2448
GO:0044238
primary metabolic process
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2111 / 4058
GO:0071704
organic substance metabolic process
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2272 / 4377
GO:0009058
biosynthetic process
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1351 / 2629
GO:0007275
multicellular organismal development
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760 / 1451
GO:0044763
single−organism cellular process
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1758 / 3447
GO:0044237
cellular metabolic process
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2088 / 4021
GO:0044281
small molecule metabolic process
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456 / 917
GO:0006629
lipid metabolic process
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306 / 576
GO:0044711
single−organism biosynthetic process
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518 / 1061
GO:1901576
organic substance biosynthetic process
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GO:0007389
pattern specification process
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74 / 138
GO:0044255
cellular lipid metabolic process
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230 / 426
GO:0006082
organic acid metabolic process
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315 / 632
GO:0044283
small molecule biosynthetic process
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GO:0008610
lipid biosynthetic process
0.37171
141 / 322
GO:0044249
cellular biosynthetic process
0.98766
1278 / 2478
GO:0009798
axis specification
0.03252
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GO:0003002
regionalization
0.22071
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GO:0006720
isoprenoid metabolic process
0.28456
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GO:0043436
oxoacid metabolic process
0.97619
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GO:0016053
organic acid biosynthetic process
0.78495
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GO:0009951
adaxial/abaxial pattern specification
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GO:0006721
terpenoid metabolic process
0.10636
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GO:0019752
carboxylic acid metabolic process
0.96787
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GO:0008299
isoprenoid biosynthetic process
0.13219
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GO:0016101
diterpenoid metabolic process
0.00555
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GO:0016114
terpenoid biosynthetic process
0.02076
67 / 106
GO:0046394
carboxylic acid biosynthetic process
0.78495
181 / 351
GO:0009685
gibberellin metabolic process
0.00219
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GO:0016102
diterpenoid biosynthetic process
0.00287
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GO:0009686
gibberellin biosynthetic process
0.00485
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