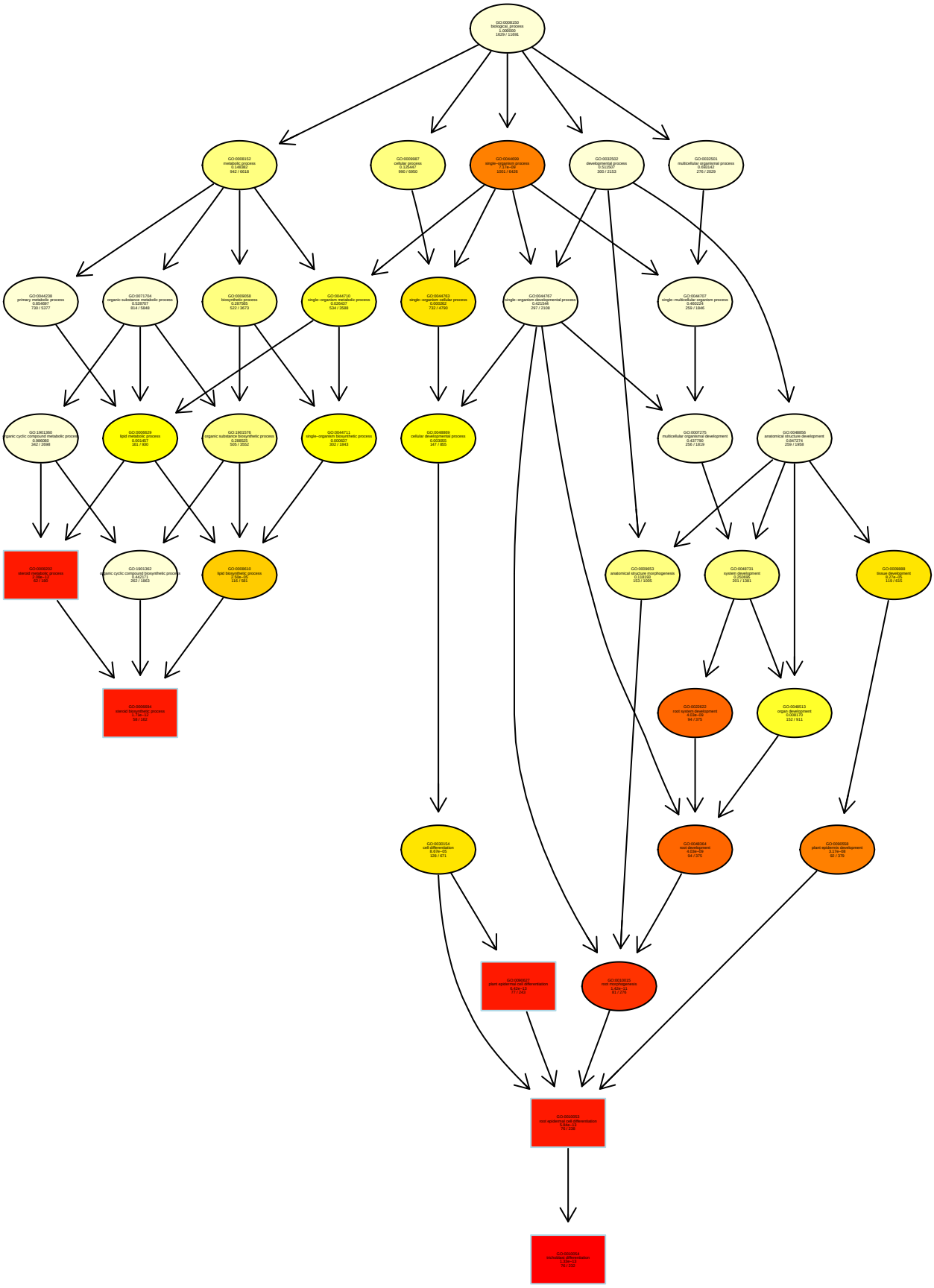
GO:0008150
biological_process
1.000000
1629 / 11691
GO:0008152
metabolic process
0.146362
942 / 6618
GO:0009987
cellular process
0.185447
990 / 6950
GO:0044699
single-organism process
2.17e-05
1031 / 6436
GO:0032502
developmental process
0.311052
300 / 2153
GO:0032501
multicellular organismal process
0.081142
276 / 2029
GO:0044238
primary metabolic process
0.056802
730 / 5377
GO:0071704
organic substance metabolic process
0.126707
814 / 5848
GO:0009058
biosynthetic process
0.080585
522 / 3673
GO:0044710
single-organism biosynthetic process
0.006407
534 / 3588
GO:0044763
single-organism cellular process
0.000000
732 / 4790
GO:0044767
single-organism developmental process
0.421548
297 / 2108
GO:0044707
single-multicellular organism process
0.466228
259 / 1846
GO:1901360
organic cyclic compound metabolic process
0.980060
342 / 2698
GO:0006629
lipid metabolic process
0.001457
161 / 760
GO:1901576
organic substance biosynthetic process
0.268825
555 / 3552
GO:0044711
single-organism biosynthetic process
0.000637
302 / 1843
GO:0048869
cellular developmental process
0.003855
147 / 852
GO:0007275
multicellular organismal development
0.437790
258 / 1819
GO:0048856
anatomical structure development
0.847274
259 / 1858
GO:0009202
steroid metabolic process
2.09e-12
62 / 180
GO:1901362
organic cyclic compound biosynthetic process
0.442171
262 / 1863
GO:0008610
lipid biosynthetic process
2.53e-05
116 / 591
GO:0009653
anatomical structure morphogenesis
0.118193
153 / 1035
GO:0048731
system development
0.250695
201 / 1381
GO:0009888
tissue development
0.27e-05
119 / 815
GO:0006694
steroid biosynthetic process
1.00e-12
59 / 162
GO:0022622
root system development
5.61e-09
64 / 275
GO:0048513
organ development
0.006176
152 / 911
GO:0030154
cell differentiation
0.000000
126 / 671
GO:0048364
root development
5.61e-09
64 / 275
GO:0090558
plant epidermis development
3.27e-09
62 / 275
GO:0009827
plant epidermal cell differentiation
6.43e-13
77 / 243
GO:0010015
root morphogenesis
1.47e-11
51 / 175
GO:0010053
root epidermal cell differentiation
5.84e-13
76 / 238
GO:0010054
trichoblast differentiation
1.13e-13
76 / 232