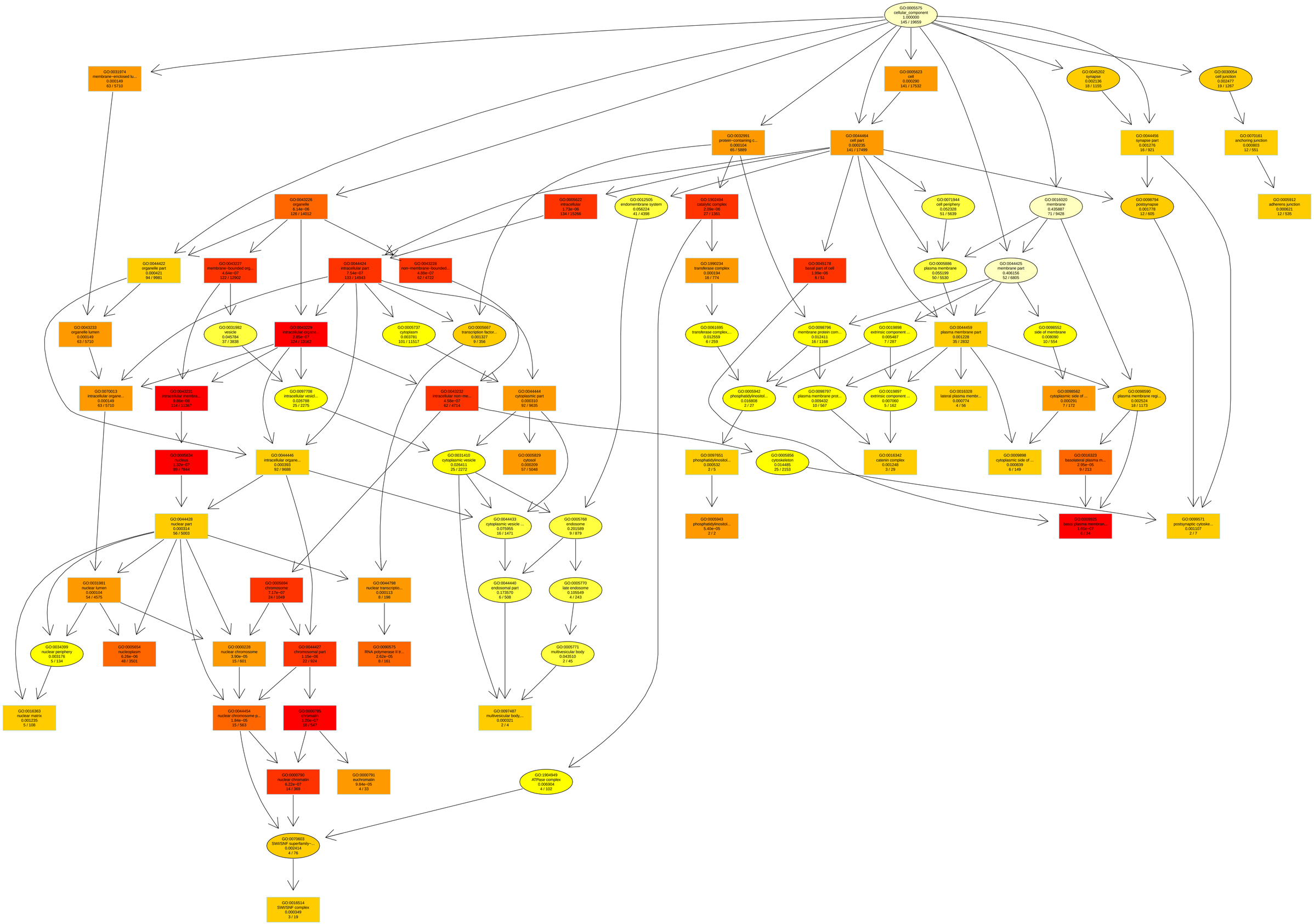
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145 / 19659
GO:0031974
membrane−enclosed lu...
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63 / 5710
GO:0005623
cell
0.000290
141 / 17532
GO:0045202
synapse
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18 / 1155
GO:0030054
cell junction
0.002477
19 / 1267
GO:0032991
protein−containing c...
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65 / 5889
GO:0044464
cell part
0.000235
141 / 17499
GO:0044456
synapse part
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16 / 921
GO:0070161
anchoring junction
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12 / 551
GO:0043226
organelle
6.14e−06
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GO:0005622
intracellular
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134 / 15266
GO:0012505
endomembrane system
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GO:1902494
catalytic complex
2.03e−06
27 / 1361
GO:0071944
cell periphery
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51 / 5639
GO:0016020
membrane
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71 / 9428
GO:0098794
postsynapse
0.001778
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GO:0005912
adherens junction
0.000621
12 / 535
GO:0044422
organelle part
0.000421
94 / 9981
GO:0043227
membrane−bounded org...
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122 / 12902
GO:0044424
intracellular part
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133 / 14943
GO:0043228
non−membrane−bounded...
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GO:1990234
transferase complex
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GO:0045178
basal part of cell
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GO:0005886
plasma membrane
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GO:0044425
membrane part
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52 / 6805
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63 / 5710
GO:0031982
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GO:0043229
intracellular organe...
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GO:0005667
transcription factor...
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GO:0061695
transferase complex,...
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GO:0098796
membrane protein com...
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GO:0019898
extrinsic component ...
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7 / 287
GO:0044459
plasma membrane part
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35 / 2832
GO:0098552
side of membrane
0.008090
10 / 554
GO:0070013
intracellular organe...
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63 / 5710
GO:0043231
intracellular membra...
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114 / 11367
GO:0097708
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GO:0043232
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GO:0044444
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GO:0005942
phosphatidylinositol...
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GO:0098797
plasma membrane prot...
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GO:0019897
extrinsic component ...
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5 / 162
GO:0016328
lateral plasma membr...
0.000774
4 / 56
GO:0098562
cytoplasmic side of ...
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GO:0098590
plasma membrane regi...
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nucleus
1.32e−07
89 / 7844
GO:0044446
intracellular organe...
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92 / 9688
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cytoplasmic vesicle
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25 / 2272
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57 / 5048
GO:0097651
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GO:0005856
cytoskeleton
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25 / 2153
GO:0016342
catenin complex
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GO:0009898
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GO:0016323
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GO:0099571
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GO:0031981
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GO:0005694
chromosome
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GO:0044798
nuclear transcriptio...
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GO:0044440
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GO:0005770
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0.105549
4 / 243
GO:0034399
nuclear periphery
0.003176
5 / 134
GO:0005654
nucleoplasm
6.26e−06
48 / 3501
GO:0000228
nuclear chromosome
3.90e−05
15 / 601
GO:0044427
chromosomal part
1.15e−06
22 / 924
GO:0090575
RNA polymerase II tr...
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8 / 161
GO:0005771
multivesicular body
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2 / 45
GO:0016363
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GO:0044454
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15 / 563
GO:0000785
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1.20e−07
18 / 547
GO:0097487
multivesicular body,...
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GO:0000790
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6.22e−07
14 / 369
GO:0000791
euchromatin
9.84e−05
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GO:1904949
ATPase complex
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GO:0070603
SWI/SNF superfamily−...
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GO:0016514
SWI/SNF complex
0.000349
3 / 19