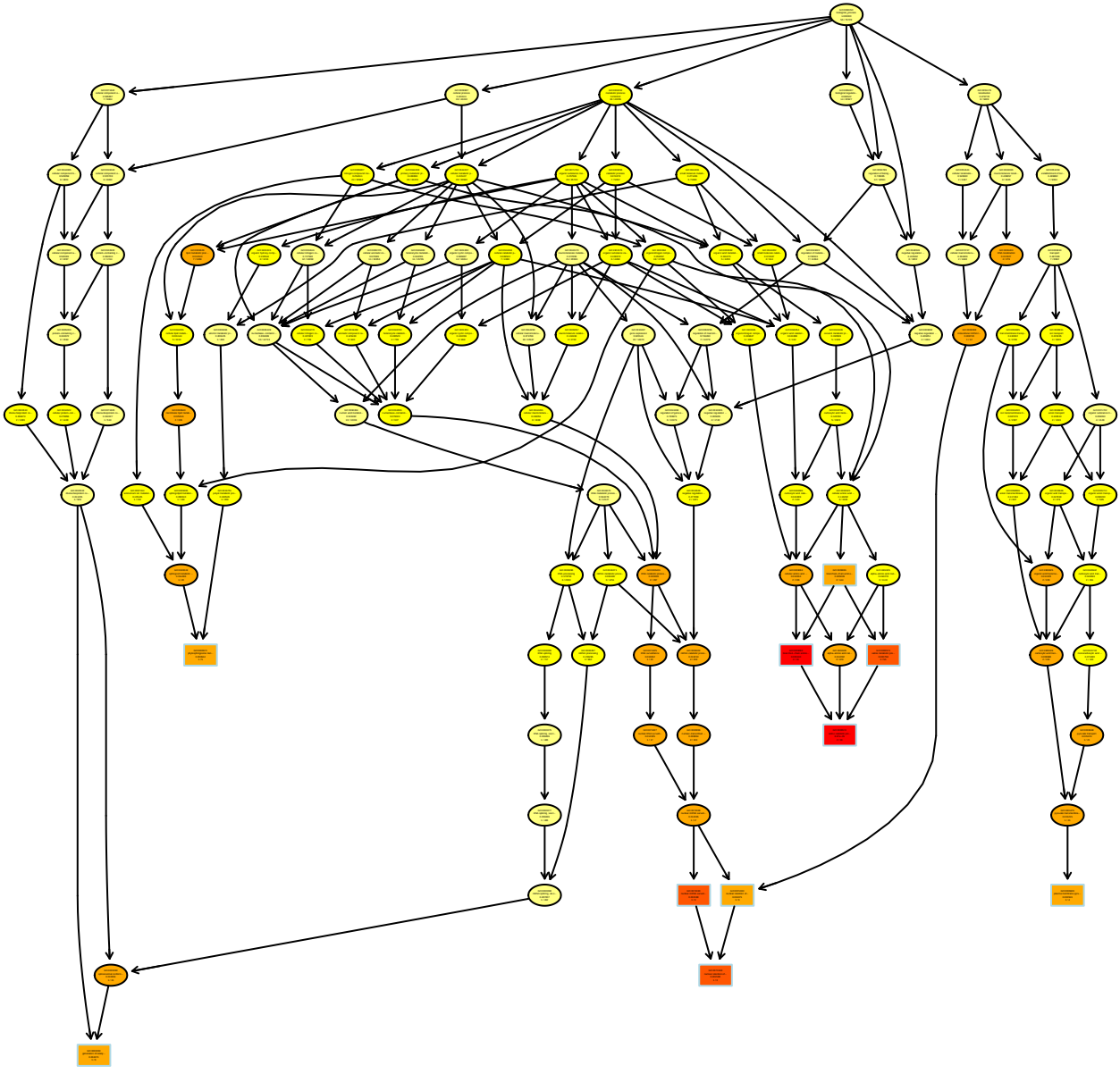

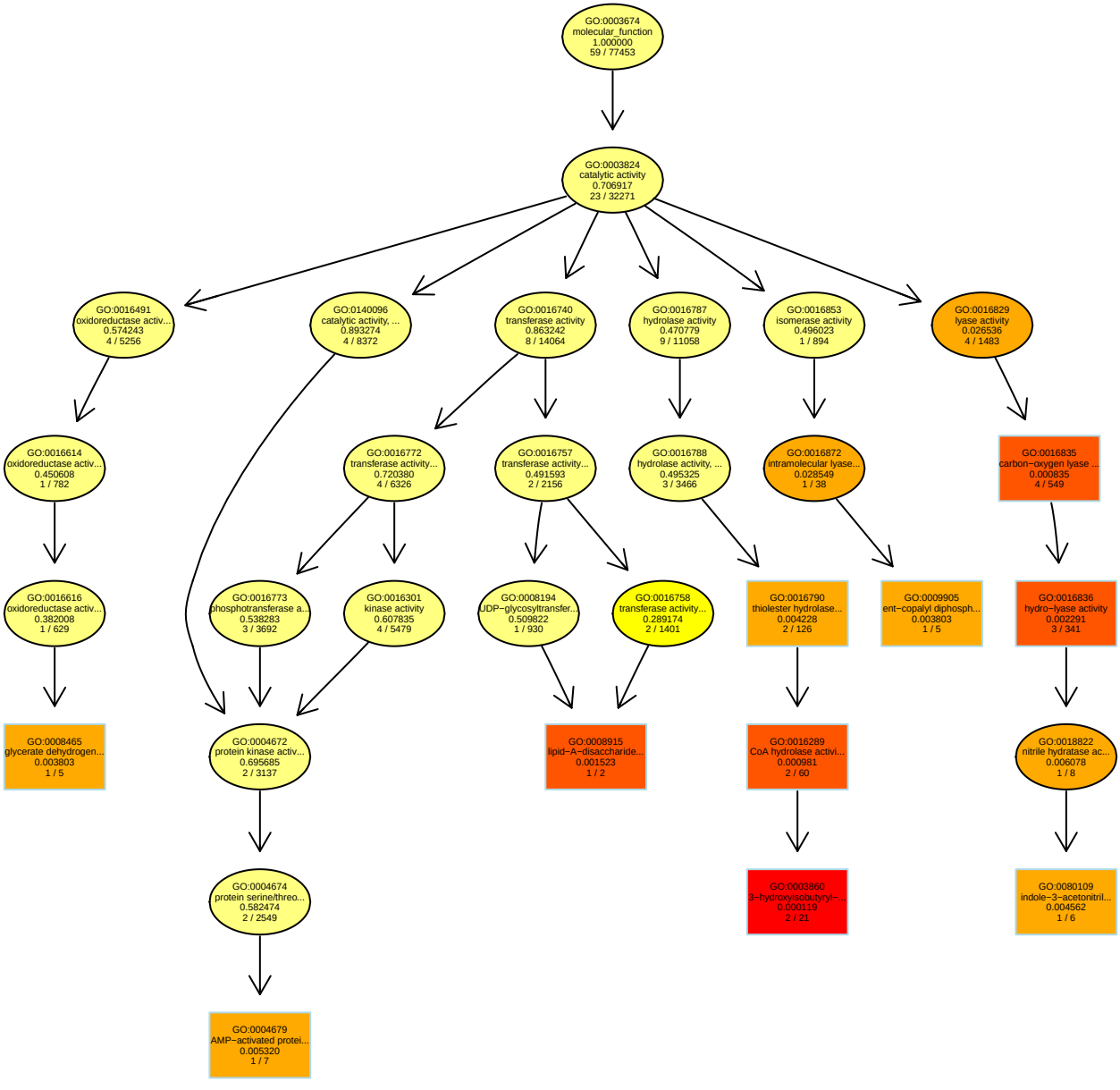
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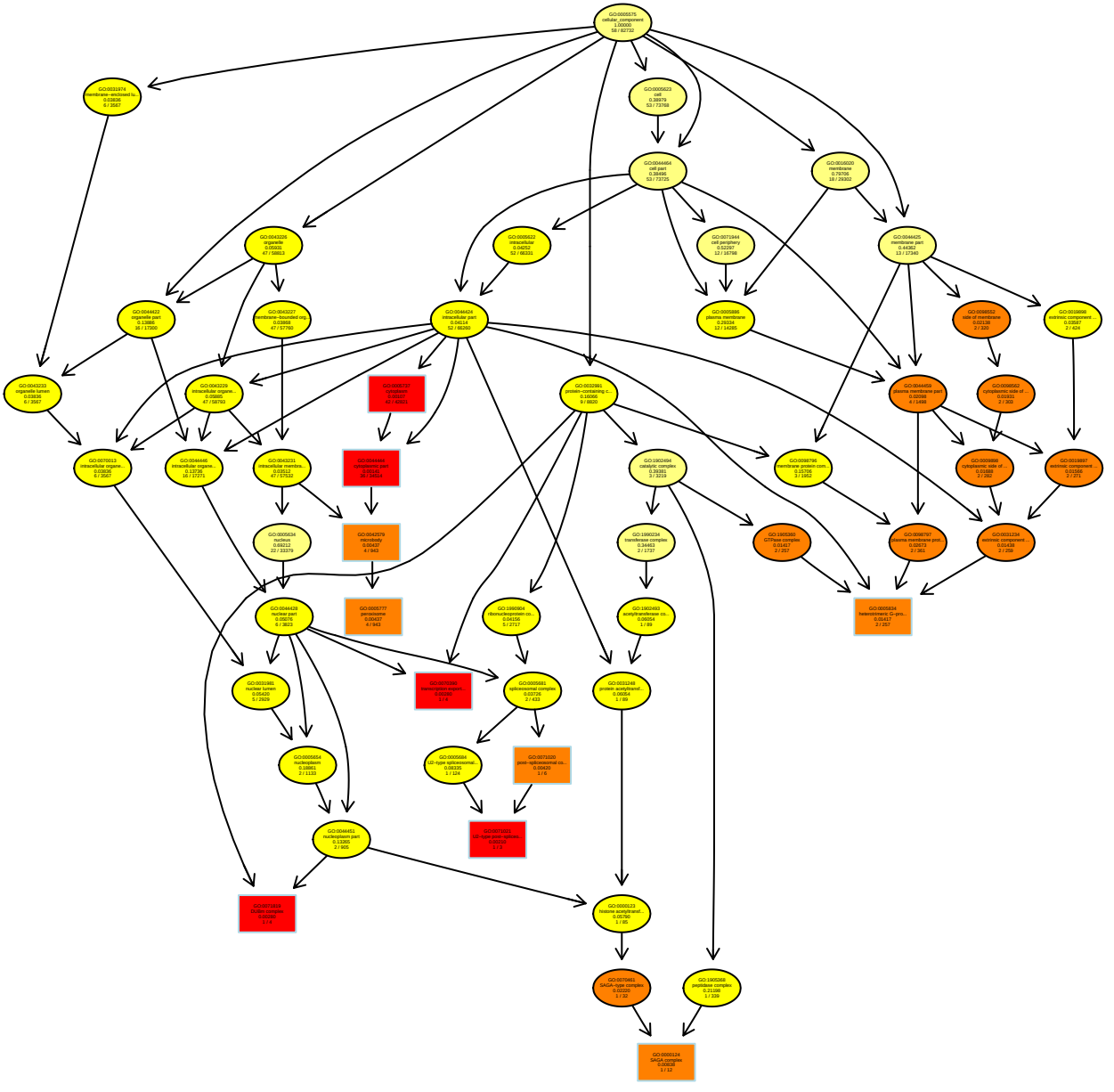
molecular_function
1.000000
59 / 77453
GO:0003824
catalytic activity
0.706917
23 / 32271
GO:0016491
oxidoreductase activ...
0.574243
4 / 5256
GO:0140096
catalytic activity, ...
0.893274
4 / 8372
GO:0016740
transferase activity
0.863242
8 / 14064
GO:0016787
hydrolase activity
0.470779
9 / 11058
GO:0016853
isomerase activity
0.496023
1 / 894
GO:0016829
lyase activity
0.026536
4 / 1483
GO:0016614
oxidoreductase activ...
0.450608
1 / 782
GO:0016772
transferase activity...
0.720380
4 / 6326
GO:0016757
transferase activity...
0.491593
2 / 2156
GO:0016788
hydrolase activity, ...
0.495325
3 / 3466
GO:0016872
intramolecular lyase...
0.028549
1 / 38
GO:0016835
carbon−oxygen lyase ...
0.000835
4 / 549
GO:0016616
oxidoreductase activ...
0.382008
1 / 629
GO:0016773
phosphotransferase a...
0.538283
3 / 3692
GO:0016301
kinase activity
0.607835
4 / 5479
GO:0008194
UDP−glycosyltransfer...
0.509822
1 / 930
GO:0016758
transferase activity...
0.289174
2 / 1401
GO:0016790
thiolester hydrolase...
0.004228
2 / 126
GO:0009905
ent−copalyl diphosph...
0.003803
1 / 5
GO:0016836
hydro−lyase activity
0.002291
3 / 341
GO:0008465
glycerate dehydrogen...
0.003803
1 / 5
GO:0004672
protein kinase activ...
0.695685
2 / 3137
GO:0008915
lipid−A−disaccharide...
0.001523
1 / 2
GO:0016289
CoA hydrolase activi...
0.000981
2 / 60
GO:0018822
nitrile hydratase ac...
0.006078
1 / 8
GO:0004674
protein serine/threo...
0.582474
2 / 2549
GO:0003860
3−hydroxyisobutyryl−...
0.000119
2 / 21
GO:0080109
indole−3−acetonitril...
0.004562
1 / 6
GO:0004679
AMP−activated protei...
0.005320
1 / 7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58 / 62732
GO:0031974
membrane-enclosed lu...
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1 / 3965
GO:0005623
cell
0.000000
53 / 72768
GO:0044464
cell part
0.00000
53 / 72725
GO:0016020
membrane
0.78706
18 / 29302
GO:0043226
organelle
0.05931
47 / 68813
GO:0005622
intracellular
0.04252
52 / 68321
GO:0071944
cell periphery
0.52297
12 / 16708
GO:0044425
membrane part
0.44362
13 / 17340
GO:0044422
organelle part
0.13986
16 / 17500
GO:0043227
membrane-bounded org...
0.03868
47 / 57765
GO:0044424
intracellular part
0.04114
52 / 68260
GO:0005886
plasma membrane
0.29334
12 / 14235
GO:0098552
side of membrane
0.01138
2 / 320
GO:0019898
extrinsic component ...
0.03597
2 / 424
GO:0043233
organelle lumen
0.03836
6 / 3567
GO:0043229
intracellular organe...
0.05895
47 / 58793
GO:0005737
cytoplasm
0.30107
42 / 42021
GO:0032991
protein-containing c...
0.16066
9 / 8820
GO:0044459
plasma membrane part
0.02998
4 / 1498
GO:0098562
cytoplasmic side of ...
0.01931
2 / 303
GO:0070013
intracellular organe...
0.03836
6 / 3567
GO:0044446
intracellular organe...
0.13736
16 / 17271
GO:0043231
intracellular membra...
0.03512
47 / 57531
GO:0044444
cytoplasmic part
0.26142
36 / 34514
GO:19024694
catalytic complex
0.39381
3 / 3216
GO:0098796
membrane protein com...
0.15706
3 / 1652
GO:0098588
cytoplasmic side of ...
0.01888
2 / 302
GO:0019897
extrinsic component ...
0.01556
2 / 271
GO:0005634
nucleus
0.69212
22 / 13379
GO:0042579
microbody
0.00437
4 / 943
GO:19902234
transferase complex
0.34463
2 / 1737
GO:1905360
GTPase complex
0.01417
2 / 257
GO:0098797
plasma membrane prot...
0.02573
2 / 301
GO:0031234
extrinsic component ...
0.01438
2 / 259
GO:0044428
nuclear part
0.05576
6 / 3623
GO:0005777
peroxisome
0.00437
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GO:1990904
ribonucleoprotein co...
0.04156
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GO:19024493
acetyltransferase co...
0.00054
1 / 89
GO:0005634
heterotrimeric G-pro...
0.01417
2 / 257
GO:0031981
nuclear lumen
0.05420
5 / 2929
GO:0070390
transcription expert...
0.00260
1 / 4
GO:0005681
spliceosomal complex
0.03726
2 / 433
GO:0031248
protein acetyltransf...
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1 / 89
GO:0005654
nucleoplasm
0.18861
2 / 1133
GO:0005684
U2-type spliceosomal ...
0.00335
1 / 128
GO:0071200
post-spliceosomal co...
0.00420
1 / 6
GO:0044452
nucleoplasm part
0.13365
2 / 905
GO:0071021
U2-type post-splicing...
0.00335
1 / 3
GO:0071819
DUBm complex
0.00260
1 / 4
GO:0000523
histone acetyltransf...
0.05760
1 / 85
GO:0070861
MAGA complex
0.01255
1 / 32
GO:1905268
peptidyl-lysine acet...
0.21338
1 / 339
GO:0000124
SAGA complex
0.00208
1 / 12