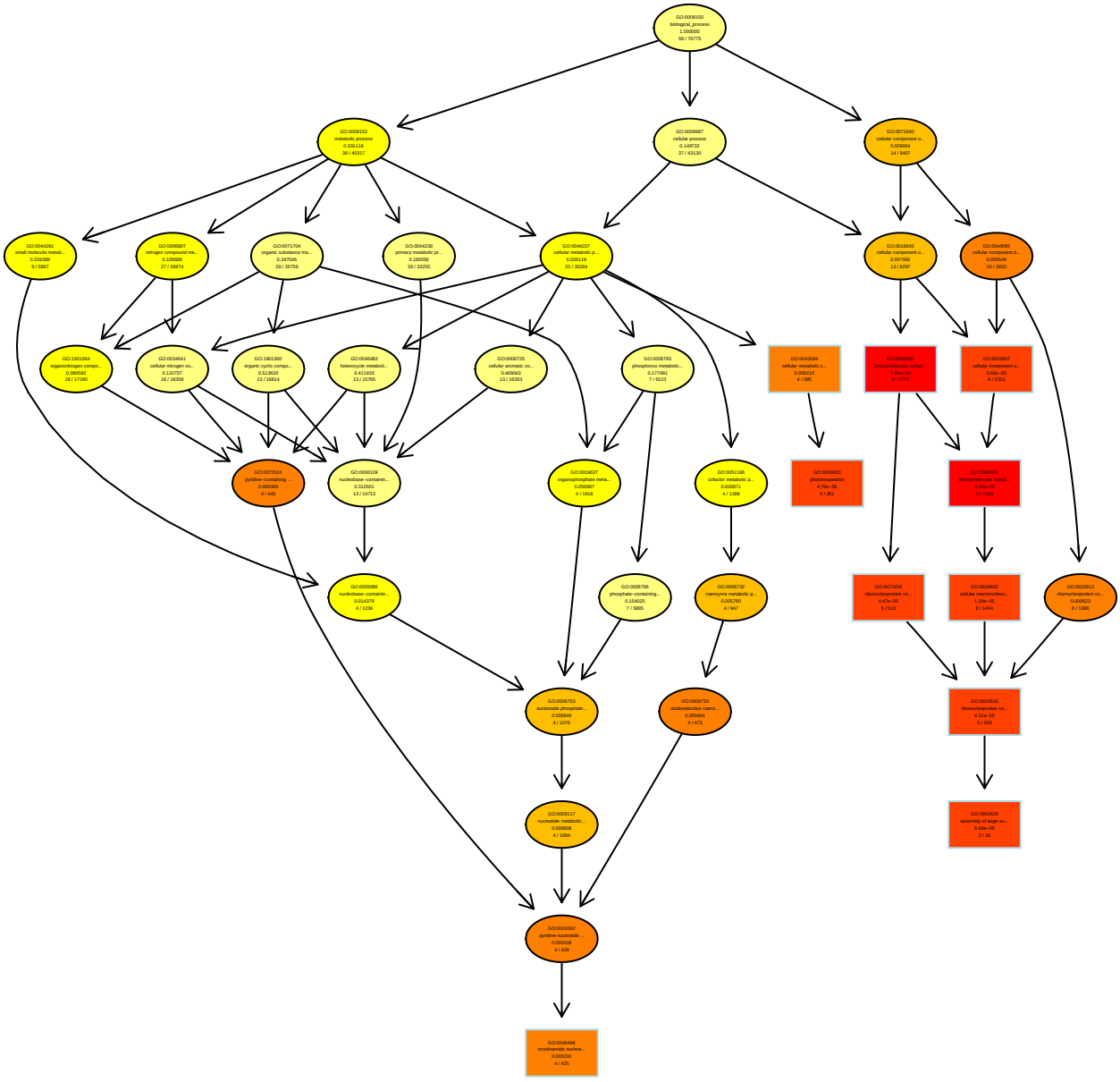
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GO:0009987
cellular process
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37 / 43130
GO:0071840
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14 / 9407
GO:0044281
small molecule metab...
0.031088
9 / 5867
GO:0006807
nitrogen compound me...
0.106658
27 / 28974
GO:0071704
organic substance me...
0.347046
29 / 35758
GO:0044238
primary metabolic pr...
0.185056
29 / 33255
GO:0044237
cellular metabolic p...
0.026116
33 / 33284
GO:0016043
cellular component o...
0.007592
13 / 8297
GO:0044085
cellular component b...
0.000549
10 / 3833
GO:1901564
organonitrogen compo...
0.080592
18 / 17185
GO:0034641
cellular nitrogen co...
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18 / 18358
GO:1901360
organic cyclic compo...
0.513820
13 / 16814
GO:0046483
heterocycle metaboli...
0.411933
13 / 15765
GO:0006725
cellular aromatic co...
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13 / 16353
GO:0006793
phosphorus metabolic...
0.177481
7 / 6123
GO:0043094
cellular metabolic c...
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4 / 385
GO:0043933
macromolecular compl...
7.04e−06
5 / 1773
GO:0022607
cellular component a...
5.66e−05
9 / 2313
GO:0072524
pyridine−containing ...
0.000369
4 / 445
GO:0006139
nucleobase−containin...
0.312501
13 / 14713
GO:0019637
organophosphate meta...
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6 / 1918
GO:0051186
cofactor metabolic p...
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4 / 1388
GO:0009653
morphogenesis
4.79e−05
4 / 261
GO:0065003
macromolecule compl...
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GO:0055086
nucleobase−containin...
0.014379
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GO:0006796
phosphate−containing...
0.154025
7 / 5885
GO:0006732
coenzyme metabolic p...
0.005760
4 / 947
GO:0071826
ribonucleoprotein co...
4.47e−05
5 / 513
GO:0034622
cellular macromoleca...
1.28e−05
8 / 1444
GO:0022613
ribonucleoprotein co...
0.000023
6 / 1386
GO:0006753
nucleoside phosphate...
0.005844
4 / 1076
GO:0006733
oxidoreduction coenz...
0.005464
4 / 473
GO:0022618
ribonucleoprotein co...
4.31e−05
5 / 509
GO:0009117
nucleotide metabolic...
0.005828
4 / 1064
GO:1902626
assembly of large su...
6.68e−05
2 / 16
GO:0019362
pyridine nucleotide ...
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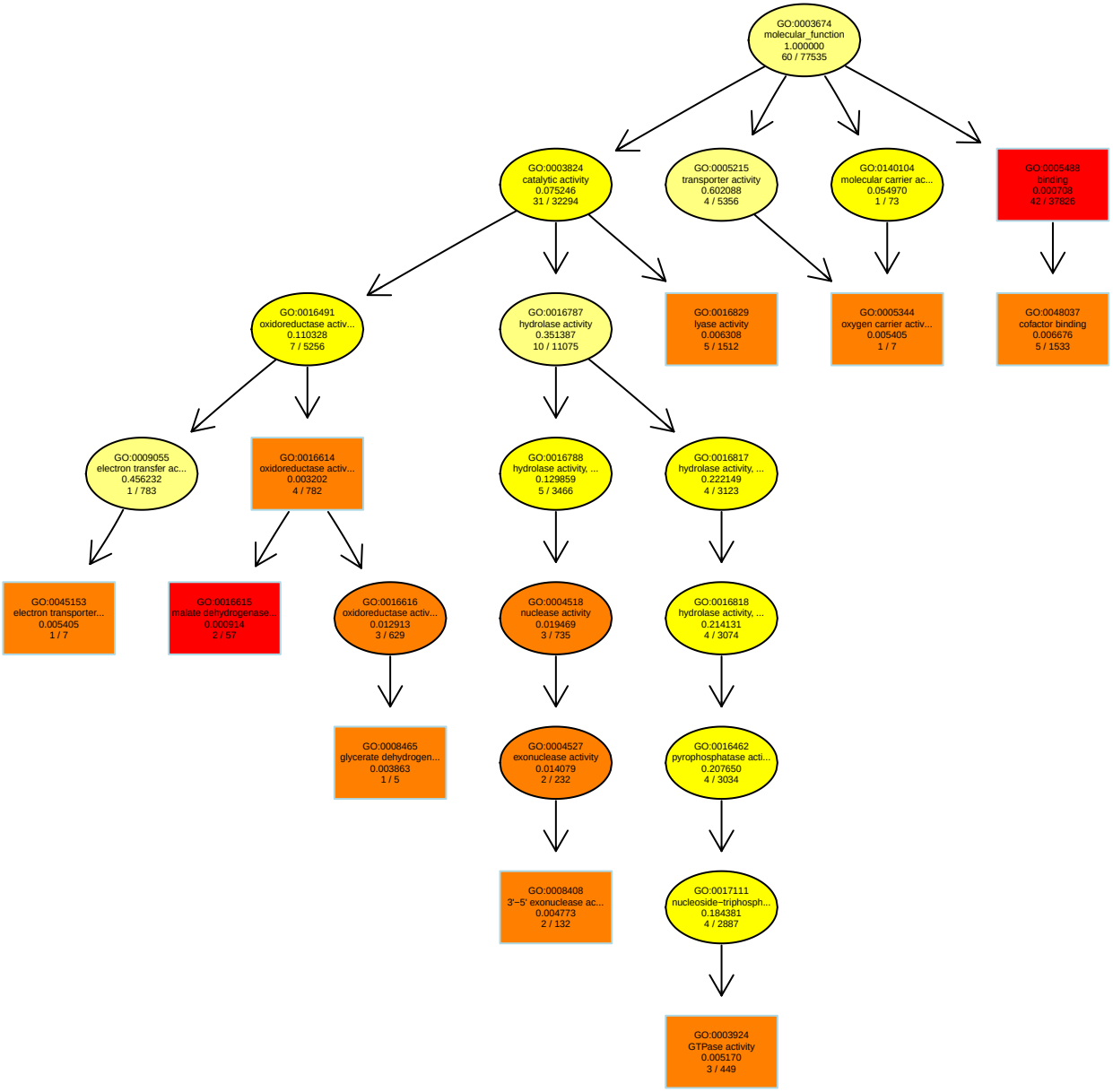
4 / 428
GO:0046496
nicotinamide nucleot...
0.000310
4 / 425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transporter activity
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4 / 5356
GO:0140104
molecular carrier ac...
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1 / 73
GO:0005488
binding
0.000708
42 / 37826
GO:0016491
oxidoreductase activ...
0.110328
7 / 5256
GO:0016787
hydrolase activity
0.351387
10 / 11075
GO:0016829
lyase activity
0.006308
5 / 1512
GO:0005344
oxygen carrier activ...
0.005405
1 / 7
GO:0048037
cofactor binding
0.006676
5 / 1533
GO:0009055
electron transfer ac...
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1 / 783
GO:0016614
oxidoreductase activ...
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GO:0016788
hydrolase activity, ...
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5 / 3466
GO:0016817
hydrolase activity, ...
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electron transporter...
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1 / 7
GO:0016615
malate dehydrogenase...
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2 / 57
GO:0016616
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3 / 629
GO:0004518
nuclease activity
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3 / 735
GO:0016818
hydrolase activity, ...
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4 / 3074
GO:0008465
glycerate dehydrogen...
0.003863
1 / 5
GO:0004527
exonuclease activity
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2 / 232
GO:0016462
pyrophosphatase acti...
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GO:0008408
3'−5' exonuclease ac...
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GO:0017111
nucleoside−triphosph...
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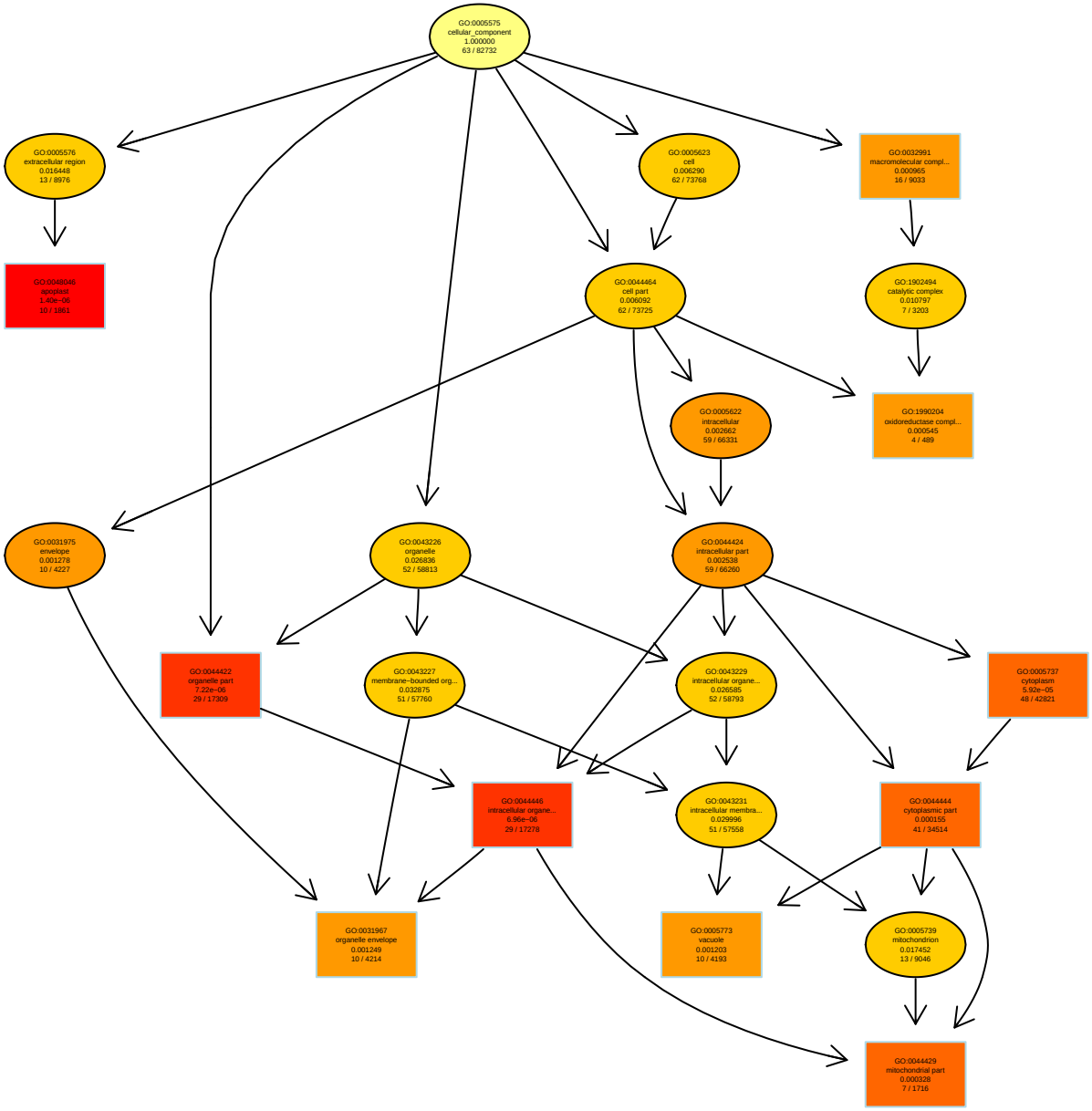
4 / 2887
GO:0003924
GTPase activity
0.005170
3 / 449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GO:0032991
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GO:0048046
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GO:0044464
cell part
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62 / 73725
GO:1902494
catalytic complex
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GO:0005622
intracellular
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59 / 66331
GO:1990204
oxidoreductase compl...
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4 / 489
GO:0031975
envelope
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GO:0043226
organelle
0.026836
52 / 58813
GO:0044424
intracellular part
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59 / 66260
GO:0044422
organelle part
7.22e−06
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GO:0043227
membrane−bounded org...
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GO:0043229
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52 / 58793
GO:0005737
cytoplasm
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48 / 42821
GO:0044446
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6.96e−06
29 / 17278
GO:0043231
intracellular membra...
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51 / 57558
GO:0044444
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41 / 34514
GO:0031967
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GO:0005773
vacuole
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10 / 4193
GO:0005739
mitochondrion
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13 / 9046
GO:0044429
mitochondrial part
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7 / 1716