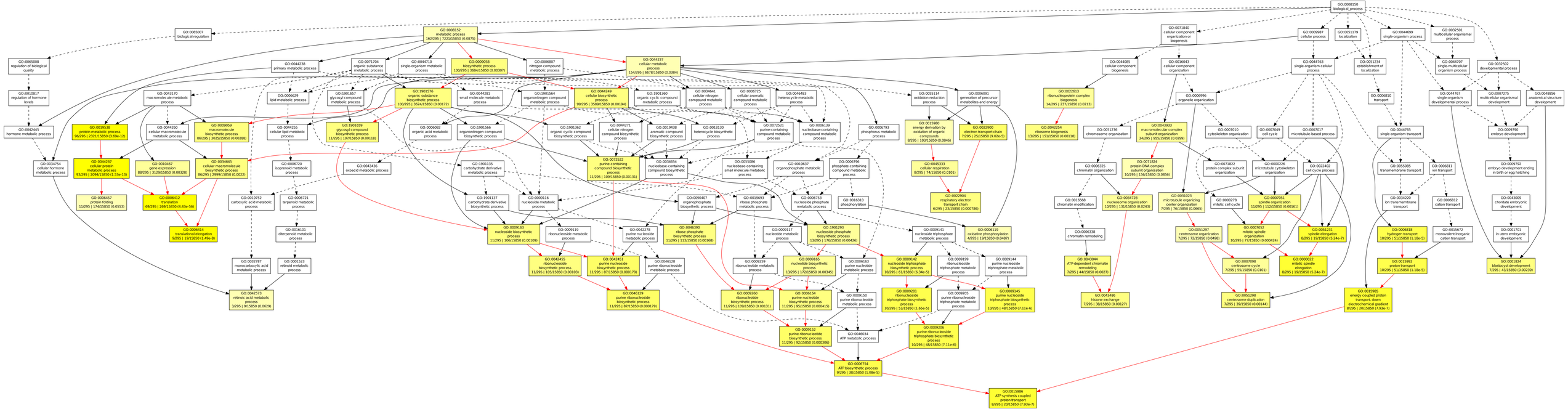
GO:0008150
biological_process
GO:0065007
biological regulation
GO:0008152
metabolic process
162/295 | 7221/15850 (0.0875)
GO:0071840
cellular component organization or biogenesis
GO:0009987
cellular process
GO:0051179
localization
GO:0044699
single-organism process
GO:0032501
multicellular organismal process
GO:0065008
regulation of biological quality
GO:0044238
primary metabolic process
GO:0071704
organic substance metabolic process
GO:0044710
single-organism metabolic process
GO:0009058
biosynthetic process
100/295 | 3684/15850 (0.00307)
GO:0006807
nitrogen compound metabolic process
GO:0044237
cellular metabolic process
154/295 | 6678/15850 (0.0384)
GO:0044085
cellular component biogenesis
GO:0016043
cellular component organization
GO:0044763
single-organism cellular process
GO:0051234
establishment of localization
GO:0044707
single-multicellular organism process
GO:0032502
developmental process
GO:0010817
regulation of hormone levels
GO:0043170
macromolecule metabolic process
GO:0006629
lipid metabolic process
GO:1901657
glycosyl compound metabolic process
GO:1901576
organic substance biosynthetic process
100/295 | 3624/15850 (0.00172)
GO:0044281
small molecule metabolic process
GO:1901564
organonitrogen compound metabolic process
GO:0044249
cellular biosynthetic process
99/295 | 3589/15850 (0.00194)
GO:1901360
organic cyclic compound metabolic process
GO:0034641
cellular nitrogen compound metabolic process
GO:0006725
cellular aromatic compound metabolic process
GO:0046483
heterocycle metabolic process
GO:0055114
oxidation-reduction process
GO:0006091
generation of precursor metabolites and energy
GO:0022613
ribonucleoprotein complex biogenesis
14/295 | 237/15850 (0.0213)
GO:0006996
organelle organization
GO:0006810
transport
GO:0044767
single-organism developmental process
GO:0007275
multicellular organismal development
GO:0048856
anatomical structure development
GO:0042445
hormone metabolic process
GO:0019538
protein metabolic process
96/295 | 2321/15850 (3.69e-12)
GO:0009059
macromolecule biosynthetic process
86/295 | 3025/15850 (0.00288)
GO:0044260
cellular macromolecule metabolic process
GO:0044255
cellular lipid metabolic process
GO:1901659
glycosyl compound biosynthetic process
11/295 | 107/15850 (0.00118)
GO:0006082
organic acid metabolic process
GO:1901566
organonitrogen compound biosynthetic process
GO:1901362
organic cyclic compound biosynthetic process
GO:0044271
cellular nitrogen compound biosynthetic process
GO:0019438
aromatic compound biosynthetic process
GO:0018130
heterocycle biosynthetic process
GO:0072521
purine-containing compound metabolic process
GO:0006139
nucleobase-containing compound metabolic process
GO:0006793
phosphorus metabolic process
GO:0015980
energy derivation by oxidation of organic compounds
8/295 | 103/15850 (0.0846)
GO:0022900
electron transport chain
7/295 | 25/15850 (9.02e-5)
GO:0042254
ribosome biogenesis
13/295 | 151/15850 (0.00118)
GO:0051276
chromosome organization
GO:0043933
macromolecular complex subunit organization
34/295 | 955/15850 (0.0299)
GO:0007010
cytoskeleton organization
GO:0007049
cell cycle
GO:0007017
microtubule-based process
GO:0006810
transport
GO:0044765
single-organism transport
GO:0007030
Golgi organization
GO:0019915
lipid storage
GO:0044262
cellular carbohydrate metabolic process
GO:0034645
cellular macromolecule biosynthetic process
86/295 | 2999/15850 (0.0022)
GO:0034654
nucleobase-containing compound biosynthetic process
GO:0055086
nucleobase-containing small molecule metabolic process
GO:0019637
organophosphate metabolic process
GO:0006796
phosphate-containing compound metabolic process
GO:0006753
nucleoside phosphate metabolic process
GO:0016310
phosphorylation
GO:0022904
respiratory electron transport chain
6/295 | 23/15850 (0.000786)
GO:0045333
cellular respiration
8/295 | 74/15850 (0.0101)
GO:0051641
cellular localization
GO:0006886
intracellular protein transport
GO:0006605
protein targeting
GO:0015031
protein transport
GO:0009790
embryo development
GO:0044267
cellular protein metabolic process
93/295 | 2094/15850 (1.53e-13)
GO:0010467
gene expression
88/295 | 3129/15850 (0.0032)
GO:0006457
protein folding
11/295 | 174/15850 (0.0553)
GO:0006412
translation
69/295 | 269/15850 (4.43e-56)
GO:0006414
translational elongation
9/295 | 19/15850 (1.49e-8)
GO:0006720
isoprenoid metabolic process
GO:0043436
oxoacid metabolic process
GO:0019752
carboxylic acid metabolic process
GO:0006721
terpenoid metabolic process
GO:0016101
diterpenoid metabolic process
GO:0032787
monocarboxylic acid metabolic process
GO:0001523
retinoid metabolic process
GO:0042573
retinoic acid metabolic process
3/295 | 9/15850 (0.0629)
GO:1901135
carbohydrate derivative metabolic process
GO:1901137
carbohydrate derivative biosynthetic process
GO:0009116
nucleoside metabolic process
GO:0009693
ribose phosphate metabolic process
GO:0009117
nucleotide metabolic process
GO:1901293
nucleoside phosphate biosynthetic process
13/295 | 176/15850 (0.00426)
GO:0009141
nucleoside triphosphate metabolic process
GO:0006119
oxidative phosphorylation
4/295 | 19/15850 (0.0487)
GO:0009618
nucleoside biosynthetic process
11/295 | 106/15850 (0.00109)
GO:0009119
ribonucleoside metabolic process
GO:0042278
purine nucleoside metabolic process
GO:0046390
ribose phosphate biosynthetic process
11/295 | 113/15850 (0.00168)
GO:0009142
nucleoside triphosphate biosynthetic process
10/295 | 61/15850 (6.34e-5)
GO:0009199
ribonucleoside triphosphate metabolic process
GO:0009144
purine nucleoside triphosphate metabolic process
GO:0009201
ribonucleoside triphosphate biosynthetic process
10/295 | 53/15850 (1.65e-5)
GO:0009205
purine ribonucleoside triphosphate metabolic process
GO:0009145
purine nucleoside triphosphate biosynthetic process
10/295 | 48/15850 (7.11e-6)
GO:0009206
purine ribonucleoside triphosphate biosynthetic process
10/295 | 48/15850 (7.11e-6)
GO:0006754
ATP biosynthetic process
9/295 | 38/15850 (1.08e-5)
GO:0015986
ATP synthesis coupled proton transport
8/295 | 20/15850 (7.93e-7)
GO:0042455
ribonucleoside biosynthetic process
11/295 | 105/15850 (0.00103)
GO:0042451
purine nucleoside biosynthetic process
11/295 | 87/15850 (0.000179)
GO:0046128
purine ribonucleoside metabolic process
GO:0046129
purine ribonucleoside biosynthetic process
11/295 | 87/15850 (0.000179)
GO:0009163
nucleoside biosynthetic process
11/295 | 108/15850 (0.00131)
GO:0009165
nucleotide biosynthetic process
13/295 | 172/15850 (0.00345)
GO:0009163
purine nucleotide metabolic process
GO:0009260
ribonucleotide biosynthetic process
11/295 | 108/15850 (0.00131)
GO:0006164
purine nucleotide biosynthetic process
11/295 | 95/15850 (0.000415)
GO:0009150
purine ribonucleotide metabolic process
GO:0009152
purine ribonucleotide biosynthetic process
11/295 | 92/15850 (0.000306)
GO:0046034
ATP metabolic process
GO:0009239
ribonucleotide metabolic process
GO:0009260
ribonucleotide biosynthetic process
11/295 | 108/15850 (0.00131)
GO:0016568
chromatin modification
GO:0006325
chromatin organization
GO:0071824
protein-DNA complex subunit organization
10/295 | 156/15850 (0.0856)
GO:0071822
protein complex subunit organization
GO:0000226
microtubule cytoskeleton organization
GO:0022402
cell cycle process
GO:0034728
nucleosome organization
10/295 | 131/15850 (0.0243)
GO:0006338
chromatin remodeling
GO:0043044
ATP-dependent chromatin remodeling
7/295 | 44/15850 (0.0027)
GO:0043486
histone exchange
7/295 | 38/15850 (0.00127)
GO:0031023
microtubule organizing center organization
7/295 | 76/15850 (0.0665)
GO:0000278
mitotic cell cycle
GO:0007051
spindle organization
11/295 | 112/15850 (0.00161)
GO:0051297
centrosome organization
7/295 | 72/15850 (0.0498)
GO:0007052
mitotic spindle organization
10/295 | 77/15850 (0.000424)
GO:0051231
spindle elongation
8/295 | 19/15850 (5.24e-7)
GO:0007098
centrosome cycle
7/295 | 55/15850 (0.0101)
GO:0000022
mitotic spindle elongation
8/295 | 19/15850 (5.24e-7)
GO:0051298
centrosome duplication
7/295 | 38/15850 (0.00144)
GO:0055085
transmembrane transport
GO:0006811
ion transport
GO:0034220
ion transmembrane transport
GO:0006812
cation transport
GO:0006818
hydrogen transport
10/295 | 51/15850 (1.18e-5)
GO:0015672
monovalent inorganic cation transport
GO:0015992
proton transport
10/295 | 51/15850 (1.18e-5)
GO:0015985
energy coupled proton transport, down electrochemical gradient
8/295 | 20/15850 (7.93e-7)
GO:0009790
embryo development
GO:0009792
embryo development ending in birth or egg hatching
GO:0043009
chordate embryonic development
GO:0001701
in utero embryonic development
GO:0001824
blastocyst development
7/295 | 43/15850 (0.00239)