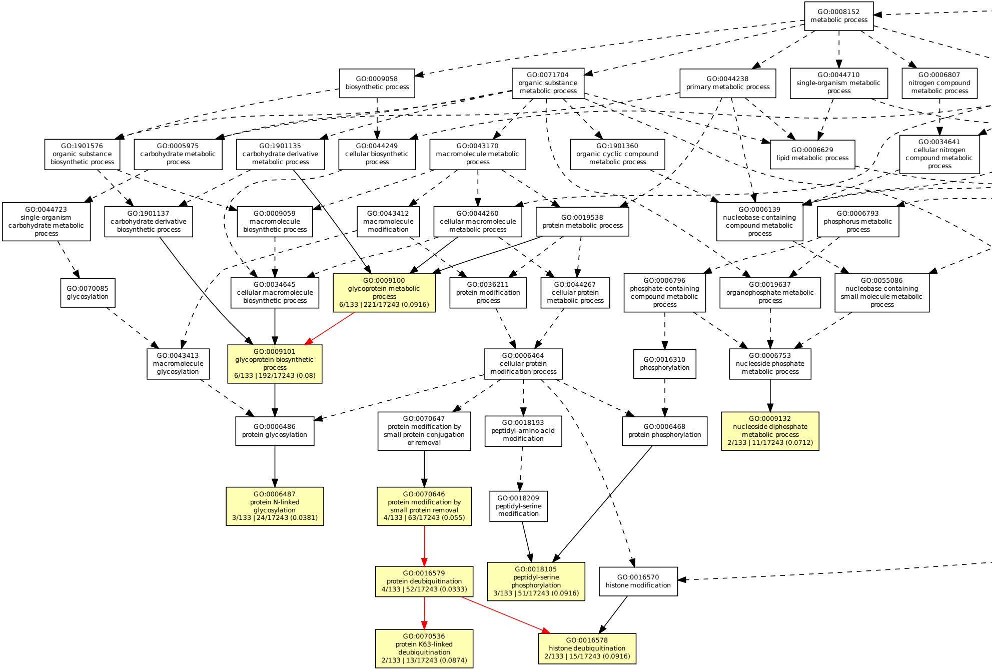
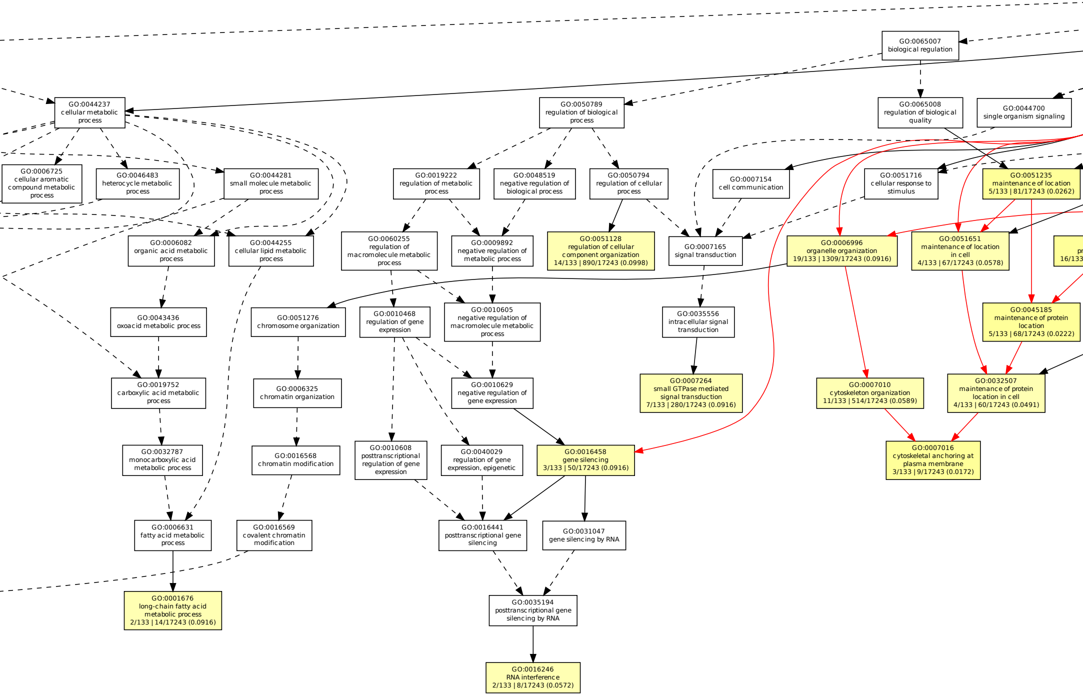
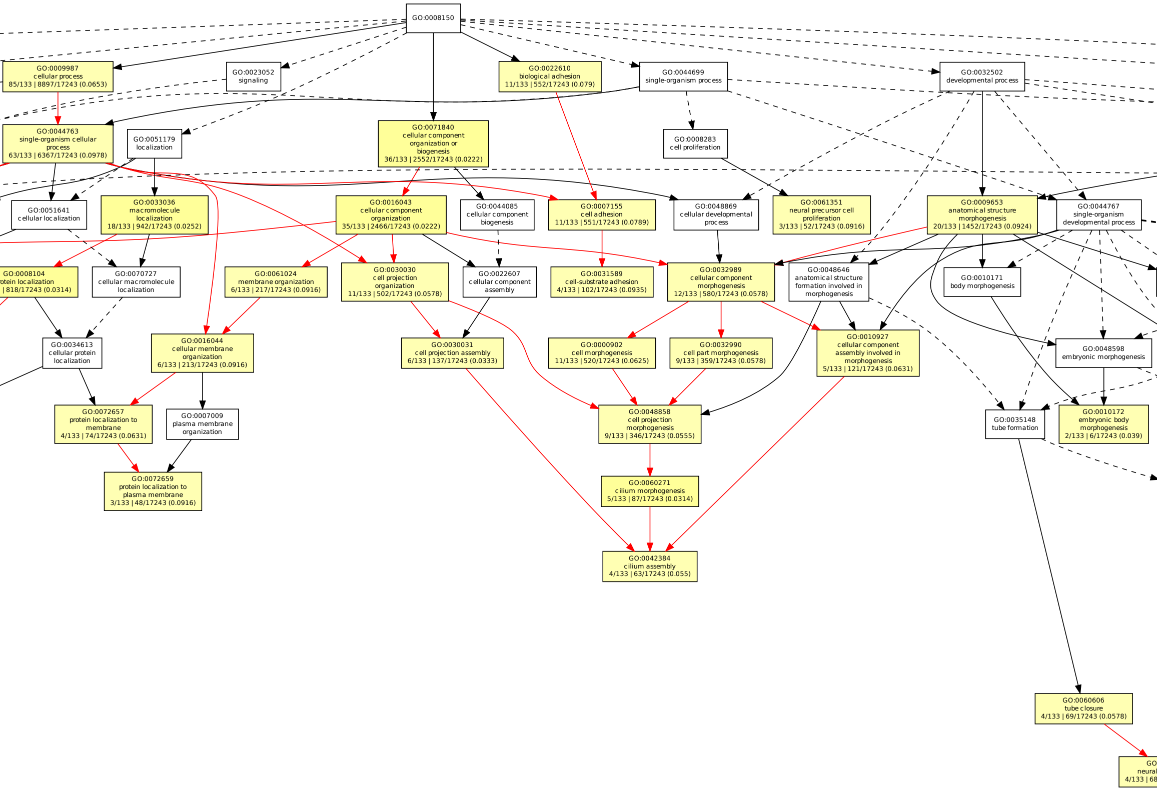
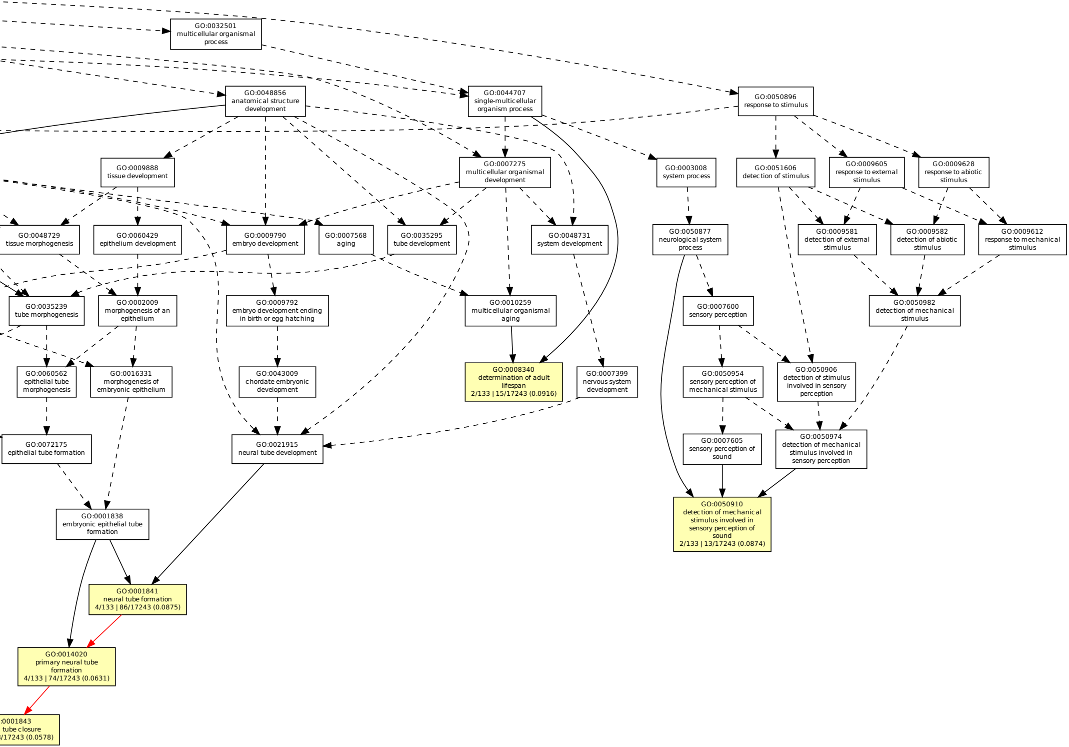

GO:0008152
metabolic process
GO:0009058
biosynthetic process
GO:0071704
organic substance
metabolic process
GO:0044238
primary metabolic process
GO:0044710
single-organism metabolic
process
GO:0006807
nitrogen compound
metabolic process
GO:1901576
organic substance
biosynthetic process
GO:0005975
carbohydrate metabolic
process
GO:1901135
carbohydrate derivative
metabolic process
GO:0044249
cellular biosynthetic
process
GO:0043170
macromolecule metabolic
process
GO:1901360
organic cyclic compound
metabolic process
GO:0006629
lipid metabolic process
GO:0034641
cellular nitrogen
compound metabolic
process
GO:0044723
single-organism
carbohydrate metabolic
process
GO:1901137
carbohydrate derivative
biosynthetic process
GO:0009059
macromolecule
biosynthetic process
GO:0043412
macromolecule
modification
GO:0044260
cellular macromolecule
metabolic process
GO:0019538
protein metabolic process
GO:0006139
nucleobase-containing
compound metabolic
process
GO:0006793
phosphorus metabolic
process
GO:0070085
glycosylation
GO:0034645
cellular macromolecule
biosynthetic process
GO:0009100
glycoprotein metabolic
process
6/133 | 221/17243 (0.0916)
GO:0036211
protein modification
process
GO:0044267
cellular protein
metabolic process
GO:0006796
phosphate-containing
compound metabolic
process
GO:0019637
organophosphate metabolic
process
GO:0055086
nucleobase-containing
small molecule metabolic
process
GO:0043413
macromolecule
glycosylation
GO:0009101
glycoprotein biosynthetic
process
6/133 | 192/17243 (0.08)
GO:0006464
cellular protein
modification process
GO:0016310
phosphorylation
GO:0006753
nucleoside phosphate
metabolic process
GO:0006486
protein glycosylation
GO:0070647
protein modification by
small protein conjugation
or removal
GO:0018193
peptidyl-amino acid
modification
GO:0006468
protein phosphorylation
GO:0009132
nucleoside diphosphate
metabolic process
2/133 | 11/17243 (0.0712)
GO:0006487
protein N-linked
glycosylation
3/133 | 24/17243 (0.0381)
GO:0070646
protein modification by
small protein removal
4/133 | 63/17243 (0.055)
GO:0018209
peptidyl-serine
modification
GO:0016579
protein deubiquitination
4/133 | 52/17243 (0.0333)
GO:0018105
peptidyl-serine
phosphorylation
3/133 | 51/17243 (0.0916)
GO:0016570
histone modification
GO:0070536
protein K63-linked
deubiquitination
2/133 | 13/17243 (0.0874)
GO:0016578
histone deubiquitination
2/133 | 15/17243 (0.0916)


GO:0065007
biological regulation
GO:0044237
cellular metabolic process
GO:0050789
regulation of biological process
GO:0065008
regulation of biological quality
GO:0044700
single organism signaling
GO:0006725
cellular aromatic compound metabolic process
GO:0046483
heterocycle metabolic process
GO:0044281
small molecule metabolic process
GO:0019222
regulation of metabolic process
GO:0048519
negative regulation of biological process
GO:0050794
regulation of cellular process
GO:0007154
cell communication
GO:0051716
cellular response to stimulus
GO:0051235
maintenance of location
5/133 | 81/17243 (0.0262)
GO:0006082
organic acid metabolic process
GO:0044255
cellular lipid metabolic process
GO:0060255
regulation of macromolecule metabolic process
GO:0009892
negative regulation of metabolic process
GO:0051128
regulation of cellular component organization
14/133 | 890/17243 (0.0998)
GO:0007165
signal transduction
GO:0006996
organelle organization
19/133 | 1309/17243 (0.0916)
GO:0051651
maintenance of location in cell
4/133 | 67/17243 (0.0578)
pr
16/133
GO:0043436
oxoacid metabolic process
GO:0051276
chromosome organization
GO:0010468
regulation of gene expression
GO:0010605
negative regulation of macromolecule metabolic process
GO:0035556
intracellular signal transduction
GO:0045185
maintenance of protein location
5/133 | 68/17243 (0.0222)
GO:0019752
carboxylic acid metabolic process
GO:0006325
chromatin organization
GO:0010629
negative regulation of gene expression
GO:0007264
small GTPase mediated signal transduction
7/133 | 280/17243 (0.0916)
GO:0007010
cytoskeleton organization
11/133 | 514/17243 (0.0589)
GO:0032507
maintenance of protein location in cell
4/133 | 60/17243 (0.0491)
GO:0032787
monocarboxylic acid metabolic process
GO:0016568
chromatin modification
GO:0010608
posttranscriptional regulation of gene expression
GO:0040029
regulation of gene expression, epigenetic
GO:0016458
gene silencing
3/133 | 50/17243 (0.0916)
GO:0007016
cytoskeletal anchoring at plasma membrane
3/133 | 9/17243 (0.0172)
GO:0006631
fatty acid metabolic process
GO:0016569
covalent chromatin modification
GO:0016441
posttranscriptional gene silencing
GO:0031047
gene silencing by RNA
GO:0001676
long-chain fatty acid metabolic process
2/133 | 14/17243 (0.0916)
GO:0035194
posttranscriptional gene silencing by RNA
GO:0016246
RNA interference
2/133 | 8/17243 (0.0572)


GO:0008150
GO:0009987
cellular process
85/133 | 8897/17243 (0.0653)
GO:0023052
signaling
GO:0022610
biological adhesion
11/133 | 552/17243 (0.079)
GO:0044699
single-organism process
GO:0032502
developmental process
GO:0044763
single-organism cellular
process
63/133 | 6367/17243 (0.0978)
GO:0051179
localization
GO:0071840
cellular component
organization or
biogenesis
36/133 | 2552/17243 (0.0222)
GO:0008283
cell proliferation
GO:0051641
cellular localization
GO:0033036
macromolecule
localization
18/133 | 942/17243 (0.0252)
GO:0016043
cellular component
organization
35/133 | 2466/17243 (0.0222)
GO:0044085
cellular component
biogenesis
GO:0007155
cell adhesion
11/133 | 551/17243 (0.0789)
GO:0048869
cellular developmental
process
GO:0061351
neural precursor cell
proliferation
3/133 | 52/17243 (0.0916)
GO:0009653
anatomical structure
morphogenesis
20/133 | 1452/17243 (0.0924)
GO:0044767
single-organism
developmental process
GO:0008104
otein localization
| 818/17243 (0.0314)
GO:0070727
cellular macromolecule
localization
GO:0061024
membrane organization
6/133 | 217/17243 (0.0916)
GO:0030030
cell projection
organization
11/133 | 502/17243 (0.0578)
GO:0022607
cellular component
assembly
GO:0031589
cell-substrate adhesion
4/133 | 102/17243 (0.0935)
GO:0032989
cellular component
morphogenesis
12/133 | 580/17243 (0.0578)
GO:0048646
anatomical structure
formation involved in
morphogenesis
GO:0010171
body morphogenesis
GO:0034613
cellular protein
localization
GO:0016044
cellular membrane
organization
6/133 | 213/17243 (0.0916)
GO:0030031
cell projection assembly
6/133 | 137/17243 (0.0333)
GO:0000902
cell morphogenesis
11/133 | 520/17243 (0.0625)
GO:0032990
cell part morphogenesis
9/133 | 359/17243 (0.0578)
GO:0010927
cellular component
assembly involved in
morphogenesis
5/133 | 121/17243 (0.0631)
GO:0048598
embryonic morphogenesis
GO:0072657
protein localization to
membrane
4/133 | 74/17243 (0.0631)
GO:0007009
plasma membrane
organization
GO:0048858
cell projection
morphogenesis
9/133 | 346/17243 (0.0555)
GO:0035148
tube formation
GO:0010172
embryonic body
morphogenesis
2/133 | 6/17243 (0.039)
GO:0072659
protein localization to
plasma membrane
3/133 | 48/17243 (0.0916)
GO:0060271
cilium morphogenesis
5/133 | 87/17243 (0.0314)
GO:0042384
cilium assembly
4/133 | 63/17243 (0.055)
GO:0060606
tube closure
4/133 | 69/17243 (0.0578)
GO
neural
4/133 | 68


GO:0032501
multicellular organismal process
GO:0048856
anatomical structure development
GO:0044707
single-multicellular organism process
GO:0050896
response to stimulus
GO:0009888
tissue development
GO:0007275
multicellular organismal development
GO:0003008
system process
GO:0051606
detection of stimulus
GO:0009605
response to external stimulus
GO:0009628
response to abiotic stimulus
GO:0048729
tissue morphogenesis
GO:0060429
epithelium development
GO:0009790
embryo development
GO:0007568
aging
GO:0035295
tube development
GO:0048731
system development
GO:0050877
neurological system process
GO:0009581
detection of external stimulus
GO:0009582
detection of abiotic stimulus
GO:0009612
response to mechanical stimulus
GO:0035239
tube morphogenesis
GO:0002009
morphogenesis of an epithelium
GO:0009792
embryo development ending in birth or egg hatching
GO:0010259
multicellular organismal aging
GO:0007600
sensory perception
GO:0050982
detection of mechanical stimulus
GO:0060562
epithelial tube morphogenesis
GO:0016331
morphogenesis of embryonic epithelium
GO:0043009
chordate embryonic development
GO:0008340
determination of adult lifespan
2/133 | 15/17243 (0.0916)
GO:0007399
nervous system development
GO:0050954
sensory perception of mechanical stimulus
GO:0050906
detection of stimulus involved in sensory perception
GO:0072175
epithelial tube formation
GO:0021915
neural tube development
GO:0007605
sensory perception of sound
GO:0050974
detection of mechanical stimulus involved in sensory perception
GO:0050910
detection of mechanical stimulus involved in sensory perception of sound
2/133 | 13/17243 (0.0874)
GO:0001838
embryonic epithelial tube formation
GO:0001841
neural tube formation
4/133 | 86/17243 (0.0875)
GO:0014020
primary neural tube formation
4/133 | 74/17243 (0.0631)
:0001843
tube closure
/17243 (0.0578)