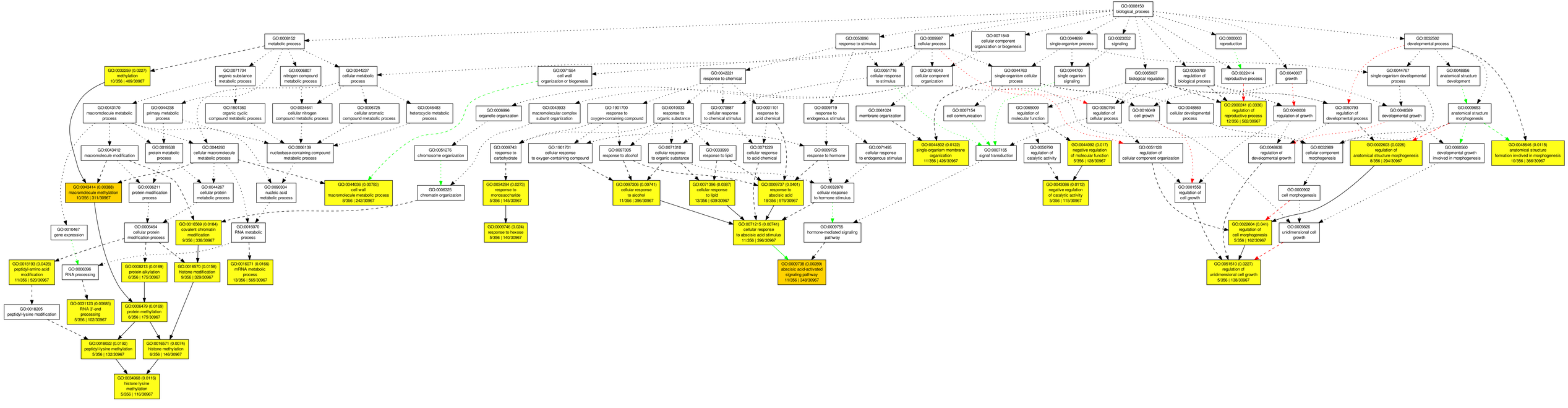
GO:0008150 biological_process
GO:0008152 metabolic process
GO:0050896 response to stimulus
GO:0009987 cellular process
GO:0071840 cellular component organization or biogenesis
GO:0044699 single-organism process
GO:0023052 signaling
GO:0000003 reproduction
GO:0032502 developmental process
GO:0032259 (0.0227) methylation 10/356 | 409/30967
GO:0071704 organic substance metabolic process
GO:0006807 nitrogen compound metabolic process
GO:0044237 cellular metabolic process
GO:0071554 cell wall organization or biogenesis
GO:0042221 response to chemical
GO:0051716 cellular response to stimulus
GO:0016043 cellular component organization
GO:0044763 single-organism cellular process
GO:0044700 single organism signaling
GO:0065007 biological regulation
GO:0050789 regulation of biological process
GO:0022414 reproductive process
GO:0040007 growth
GO:0044767 single-organism developmental process
GO:0048856 anatomical structure development
GO:0043170 macromolecule metabolic process
GO:0044238 primary metabolic process
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:0006996 organelle organization
GO:0043933 macromolecular complex subunit organization
GO:1901700 response to oxygen-containing compound
GO:0010033 response to organic substance
GO:0070887 cellular response to chemical stimulus
GO:0001101 response to acid chemical
GO:0009719 response to endogenous stimulus
GO:0061024 membrane organization
GO:0007154 cell communication
GO:0065009 regulation of molecular function
GO:0050794 regulation of cellular process
GO:0016049 cell growth
GO:0048869 cellular developmental process
GO:2000241 (0.0036) regulation of reproductive process 12/356 | 562/30967
GO:0040008 regulation of growth
GO:0050793 regulation of developmental process
GO:0048589 developmental growth
GO:0009653 anatomical structure morphogenesis
GO:0043412 macromolecule modification
GO:0019538 protein metabolic process
GO:0044260 cellular macromolecule metabolic process
GO:0006139 nucleobase-containing compound metabolic process
GO:0051276 chromosome organization
GO:0009743 response to carbohydrate
GO:1901701 cellular response to oxygen-containing compound
GO:0097305 response to alcohol
GO:0071310 cellular response to organic substance
GO:0033993 response to lipid
GO:0071229 cellular response to acid chemical
GO:0009725 response to hormone
GO:0071495 cellular response to endogenous stimulus
GO:0044802 (0.0122) single-organism membrane organization 11/356 | 426/30967
GO:0007165 signal transduction
GO:0050790 regulation of catalytic activity
GO:0044092 (0.017) negative regulation of molecular function 5/356 | 128/30967
GO:0051128 regulation of cellular component organization
GO:0048638 regulation of developmental growth
GO:0032989 cellular component morphogenesis
GO:0022603 (0.0226) regulation of anatomical structure morphogenesis 8/356 | 294/30967
GO:0060560 developmental growth involved in morphogenesis
GO:0048646 (0.0115) anatomical structure formation involved in morphogenesis 10/356 | 366/30967
GO:0043414 (0.00388) macromolecule methylation 10/356 | 311/30967
GO:0036211 protein modification process
GO:0044267 cellular protein metabolic process
GO:0090304 nucleic acid metabolic process
GO:0044036 (0.00783) cell wall macromolecule metabolic process 9/356 | 242/30967
GO:0006325 chromatin organization
GO:0034284 (0.0273) response to monosaccharide 5/356 | 145/30967
GO:0009746 (0.024) response to hexose 5/356 | 140/30967
GO:0097306 (0.00741) cellular response to alcohol 11/356 | 396/30967
GO:0071396 (0.0387) cellular response to lipid 13/356 | 639/30967
GO:0009737 (0.0401) response to abscisic acid 18/356 | 976/30967
GO:0032870 cellular response to hormone stimulus
GO:0071215 (0.00741) cellular response to abscisic acid stimulus 11/356 | 396/30967
GO:0009755 hormone-mediated signaling pathway
GO:0009738 (0.0089) abscisic acid-activated signaling pathway 11/356 | 348/30967
GO:0043086 (0.0112) negative regulation of catalytic activity 5/356 | 115/30967
GO:0001558 regulation of cell growth
GO:0000902 cell morphogenesis
GO:0022604 (0.041) regulation of cell morphogenesis 5/356 | 162/30967
GO:0009826 unidimensional cell growth
GO:0051510 (0.0227) regulation of unidimensional cell growth 5/356 | 138/30967
GO:0010467 gene expression
GO:0006464 cellular protein modification process
GO:0016569 (0.0184) covalent chromatin modification 9/356 | 338/30967
GO:0016070 RNA metabolic process
GO:0018193 (0.0428) peptidyl-amino acid modification 11/356 | 520/30967
GO:0006396 RNA processing
GO:0008213 (0.0169) protein alkylation 6/356 | 175/30967
GO:0016570 (0.0158) histone modification 9/356 | 329/30967
GO:0016071 (0.0166) mRNA metabolic process 13/356 | 565/30967
GO:0018205 peptidyl-lysine modification
GO:0031123 (0.00685) RNA 3'-end processing 5/356 | 162/30967
GO:0006479 (0.0169) protein methylation 6/356 | 175/30967
GO:0018022 (0.0192) peptidyl-lysine methylation 5/356 | 132/30967
GO:0016571 (0.0074) histone methylation 6/356 | 146/30967
GO:0034968 (0.0116) histone lysine methylation 5/356 | 116/30967