GO:0008150 biological_process
GO:0050896 response to stimulus
GO:0044699 single-organism process
GO:0009987 cellular process
GO:0023052 signaling
GO:0065007 biological regulation
GO:0022610 biological adhesion
GO:0051179 localization
GO:0002376 immune system process
GO:0008152 metabolic process
GO:0008283 cell proliferation
GO:0009628 response to abiotic stimulus
GO:0006950 response to stress
GO:0009758 carbohydrate utilization
GO:0042221 response to chemical
GO:0051716 cellular response to stimulus
GO:0009605 response to external stimulus
GO:0007154 cell communication
GO:0044763 single-organism cellular process
GO:0044700 single organism signaling
GO:0050789 regulation of biological process
GO:0007155 cell adhesion
GO:1902578 single-organism localization
GO:0051234 establishment of localization
GO:0033036 macromolecule localization
GO:0044237 cellular metabolic process
GO:0044238 primary metabolic process
GO:0071704 organic substance metabolic process
GO:0009058 biosynthetic process
GO:0006807 nitrogen compound metabolic process
GO:0050673 epithelial cell proliferation
GO:0009266 response to temperature stimulus
GO:0033554 cellular response to stress
GO:0010033 response to organic substance
GO:0048583 regulation of response to stimulus
GO:0009991 response to extracellular stimulus
GO:0071496 cellular response to external stimulus
GO:0002682 regulation of immune system process
GO:0001775 cell activation
GO:0050794 regulation of cellular process
GO:0098602 single organism cell adhesion
GO:0098609 cell-cell adhesion
GO:0030155 regulation of cell adhesion
GO:0048518 positive regulation of biological process
GO:0006810 transport
GO:0032879 regulation of localization
GO:0008104 protein localization
GO:0006793 phosphorus metabolic process
GO:0006796 phosphate-containing compound metabolic process
GO:0043170 macromolecule metabolic process
GO:0044249 cellular biosynthetic process
GO:0019222 regulation of metabolic process
GO:0006725 cellular aromatic compound metabolic process
GO:0046483 heterocycle metabolic process
GO:1901576 organic substance biosynthetic process
GO:1901360 organic cyclic compound metabolic process
GO:0034641 cellular nitrogen compound metabolic process
GO:0001935 endothelial cell proliferation
GO:0042127 regulation of cell proliferation
GO:0080134 regulation of stress response
GO:0031667 response to nutrient levels
GO:0071468 cellular response to acidic pH
GO:0045321 leukocyte activation
GO:0002684 regulation of immune system process
GO:0050865 regulation of cell activation
GO:0007165 signal transduction
GO:0016337 single organismal cell-cell adhesion
GO:0048522 positive regulation of cellular process
GO:0045785 positive regulation of cell adhesion
GO:0015031 protein transport
GO:0070201 regulation of establishment of protein localization
GO:0032880 regulation of protein localization
GO:0044765 single-organism transport
GO:0051049 regulation of transport
GO:0017702 organic hydroxy compound transport
GO:0045184 establishment of protein localization
GO:0019538 protein metabolic process
GO:0044412 macromolecule modification
GO:0031327 negative regulation of cellular biosynthetic process
GO:0044260 cellular macromolecule metabolic process
GO:0010467 gene expression
GO:0060255 regulation of macromolecule metabolic process
GO:0080090 regulation of primary metabolic process
GO:0009889 regulation of biosynthetic process
GO:0009893 positive regulation of metabolic process
GO:0018130 heterocycle biosynthetic process
GO:0019438 aromatic compound biosynthetic process
GO:0009059 macromolecule biosynthetic process
GO:0044271 cellular nitrogen compound biosynthetic process
GO:1901362 organic cyclic compound biosynthetic process
GO:0006139 nucleobase-containing compound metabolic process
GO:0051171 regulation of nitrogen compound metabolic process
GO:0065008 regulation of biological quality
GO:0006355 regulation of transcription, DNA-templated
GO:0006351 transcription, DNA-templated
GO:0045893 positive regulation of transcription, DNA-templated
GO:0045944 positive regulation of transcription from RNA polymerase II promoter
GO:0006359 regulation of transcription from RNA polymerase III promoter
GO:0006383 transcription from RNA polymerase III promoter
GO:0001934 positive regulation of protein phosphorylation
GO:0019220 regulation of phosphate metabolic process
GO:0051174 regulation of phosphorus metabolic process
GO:0016310 phosphorylation
GO:0019538 protein metabolic process
GO:0044267 cellular protein metabolic process
GO:0051246 protein metabolic process
GO:0036211 protein modification process
GO:0031325 positive regulation of cellular metabolic process
GO:0034645 cellular macromolecule biosynthetic process
GO:0010468 regulation of gene expression
GO:0031329 regulation of cellular biosynthetic process
GO:0010604 positive regulation of macromolecule metabolic process
GO:0010556 regulation of macromolecule biosynthetic process
GO:0090304 nucleic acid metabolic process
GO:0019219 regulation of nucleobase-containing compound metabolic process
GO:0009891 positive regulation of biosynthetic process
GO:0051173 positive regulation of nitrogen compound metabolic process
GO:0051172 negative regulation of nitrogen compound metabolic process
GO:0049935 positive regulation of nucleobase-containing compound metabolic process
GO:0010557 positive regulation of macromolecule biosynthetic process
GO:0016070 RNA metabolic process
GO:0032774 RNA biosynthetic process
GO:0051252 regulation of RNA metabolic process
GO:2001141 regulation of RNA biosynthetic process
GO:0051254 positive regulation of RNA metabolic process
GO:1902680 positive regulation of RNA biosynthetic process
GO:1903508 positive regulation of nucleic acid-templated transcription
GO:1903506 regulation of nucleic acid-templated transcription
GO:0097659 nucleic acid-templated transcription
GO:0006412 translation
GO:0006414 translation elongation
GO:0006415 translation termination
GO:0006417 regulation of translation
GO:0006418 tRNA aminoacylation for protein translation
GO:0006419 alanyl-tRNA aminoacylation
GO:0006420 arginyl-tRNA aminoacylation
GO:0006421 asparaginyl-tRNA aminoacylation
GO:0006422 aspartyl-tRNA aminoacylation
GO:0006423 cysteinyl-tRNA aminoacylation
GO:0006424 glutamyl-tRNA aminoacylation
GO:0006425 glutaminyl-tRNA aminoacylation
GO:0006426 glycyl-tRNA aminoacylation
GO:0006427 histidyl-tRNA aminoacylation
GO:0006428 isoleucyl-tRNA aminoacylation
GO:0006429 leucyl-tRNA aminoacylation
GO:0006430 lysyl-tRNA aminoacylation
GO:0006431 methionyl-tRNA aminoacylation
GO:0006432 phenylalanyl-tRNA aminoacylation
GO:0006433 prolyl-tRNA aminoacylation
GO:0006434 seryl-tRNA aminoacylation
GO:0006435 threonyl-tRNA aminoacylation
GO:0006436 tryptophanyl-tRNA aminoacylation
GO:0006437 tyrosyl-tRNA aminoacylation
GO:0006438 valyl-tRNA aminoacylation
GO:0006439 ATP-independent chaperonin-containing T-complex
GO:0006440 protein folding
GO:0006441 regulation of protein folding
GO:0006442 regulation of protein folding
GO:0006443 positive regulation of protein folding
GO:0006444 regulation of protein import into nucleus
GO:0006445 regulation of translation
GO:0006446 regulation of translation
GO:0006447 regulation of translation
GO:0006448 regulation of translation
GO:0006449 regulation of translation
GO:0006450 regulation of translation
GO:0006451 translational frameshifting
GO:0006452 translational frameshifting
GO:0006453 regulation of protein secretion
GO:0006454 regulation of protein secretion
GO:0006455 regulation of protein secretion
GO:0006456 regulation of protein secretion
GO:0006457 protein folding
GO:0006458 protein folding
GO:0006459 protein folding
GO:0006460 protein folding
GO:0006461 protein complex assembly
GO:0006462 protein complex assembly
GO:0006463 protein complex assembly
GO:0006464 protein modification process
GO:0006465 signal peptide processing
GO:0006466 signal peptide processing
GO:0006467 signal peptide processing
GO:0006468 protein phosphorylation
GO:0006469 negative regulation of protein phosphorylation
GO:0006470 protein dephosphorylation
GO:0006471 protein ADP-ribosylation
GO:0006472 protein ADP-ribosylation
GO:0006473 protein ADP-ribosylation
GO:0006474 protein ADP-ribosylation
GO:0006475 internal protein amino acid acetylation
GO:0006476 protein deacetylation
GO:0006477 protein sulfation
GO:0006478 protein sulfation
GO:0006479 protein methylation
GO:0006480 protein methylation
GO:0006481 protein methylation
GO:0006482 protein methylation
GO:0006483 protein methylation
GO:0006484 protein methylation
GO:0006485 protein methylation
GO:0006486 protein glycosylation
GO:0006487 protein glycosylation
GO:0006488 protein glycosylation
GO:0006489 protein glycosylation
GO:0006490 protein glycosylation
GO:0006491 protein glycosylation
GO:0006492 protein glycosylation
GO:0006493 protein glycosylation
GO:0006494 protein glycosylation
GO:0006495 protein glycosylation
GO:0006496 protein glycosylation
GO:0006497 protein lipidation
GO:0006498 protein lipidation
GO:0006499 protein lipidation
GO:0006500 protein lipidation
GO:0006501 protein lipidation
GO:0006502 protein lipidation
GO:0006503 protein lipidation
GO:0006504 protein lipidation
GO:0006505 protein lipidation
GO:0006506 protein lipidation
GO:0006507 protein lipidation
GO:0006508 proteolysis
GO:0006509 proteolysis
GO:0006510 proteolysis
GO:0006511 proteolysis
GO:0006512 proteolysis
GO:0006513 proteolysis
GO:0006514 proteolysis
GO:0006515 proteolysis
GO:0006516 proteolysis
GO:0006517 proteolysis
GO:0006518 proteolysis
GO:0006519 proteolysis
GO:0006520 proteolysis
GO:0006521 proteolysis
GO:0006522 proteolysis
GO:0006523 proteolysis
GO:0006524 proteolysis
GO:0006525 proteolysis
GO:0006526 proteolysis
GO:0006527 proteolysis