```
R-MMU-9711123: Cellular response to chemical stress
                                                       R-MMU-6798695: Neutrophil degranulation
                                                       ko04142: Lysosome
                                                       ko00190: Oxidative phosphorylation
                                                       GO:0007005: mitochondrion organization
                                                       R-MMU-917937: Iron uptake and transport
                                                       GO:0006457: protein folding
                                                       GO:0045454: cell redox homeostasis
                                                       R-MMU-72766: Translation
                                                       GO:1902600: proton transmembrane transport
                                                       GO:0001817: regulation of cytokine production
                                                       GO:0044403: biological process involved in symbiotic interaction
                                                       GO:0019884: antigen processing and presentation of exogenous antigen
                                                       ko04141: Protein processing in endoplasmic reticulum
                                                       GO:0006839: mitochondrial transport
                                                       GO:0034341: response to interferon-gamma
                                                       R-MMU-5653656: Vesicle-mediated transport
                                                       GO:0048102: autophagic cell death
                                                       GO:0072593: reactive oxygen species metabolic process
                                                       GO:0045851: pH reduction
                                                       WP2271: Macrophage markers
                                                       GO:2001242: regulation of intrinsic apoptotic signaling pathway
                                                       R-MMU-2132295: MHC class II antigen presentation
                                                       R-MMU-432720: Lysosome Vesicle Biogenesis
                                                       mmu05152: Tuberculosis
                                                       GO:0042440: pigment metabolic process
                                                       GO:0002274: myeloid leukocyte activation
                                                       mmu05169: Epstein-Barr virus infection
                                                       GO:0036296: response to increased oxygen levels
                                                       R-MMU-5218859: Regulated Necrosis
                                                       GO:0030162: regulation of proteolysis
                                                       GO:0042590: antigen processing and presentation of exogenous peptide antigen via MHC class I
                                                       R-MMU-77286: mitochondrial fatty acid beta-oxidation of saturated fatty acids
                                                       GO:0008637: apoptotic mitochondrial changes
                                                       WP3626: Microglia Pathogen Phagocytosis Pathway
                                                       GO:0034599: cellular response to oxidative stress
                                                       GO:0002697: regulation of immune effector process
                                                       R-MMU-168898: Toll-like Receptor Cascades
                                                       GO:0043254: regulation of protein-containing complex assembly
                                                       R-MMU-8876725: Protein methylation
                                                       R-MMU-1632852: Macroautophagy
                                                       GO:0009615: response to virus
                                                       GO:0051054: positive regulation of DNA metabolic process
                                                       GO:0045730: respiratory burst
                                                       GO:0030595: leukocyte chemotaxis
                                                       GO:2000106: regulation of leukocyte apoptotic process
                                                       GO:0050764: regulation of phagocytosis
                                                       GO:0071402: cellular response to lipoprotein particle stimulus
                                                       R-MMU-2029480: Fcgamma receptor (FCGR) dependent phagocytosis
                                                       M00002: Glycolysis, core module involving three-carbon compounds
                                                       GO:0002683: negative regulation of immune system process
                                                       ko04216: Ferroptosis
                                                       GO:0006873: cellular ion homeostasis
                                                       GO:0051235: maintenance of location
                                                       ko04210: Apoptosis
                                                       mmu04621: NOD-like receptor signaling pathway
                                                       GO:0035967: cellular response to topologically incorrect protein
                                                       R-MMU-72649: Translation initiation complex formation
                                                       R-MMU-5628897: TP53 Regulates Metabolic Genes
                                                       GO:0090151: establishment of protein localization to mitochondrial membrane
                                                       GO:0048193: Golgi vesicle transport
                                                       GO:0042110: T cell activation
                                                       GO:0098760: response to interleukin-7
                                                       CORUM:413: (ER)-localized multiprotein complex, Ig heavy chains associated
                                                       CORUM:495: Cytochrome bc1-complex, mitochondrial
                                                       GO:0000338: protein deneddylation
                                                       GO:0007596: blood coagulation
                                                       GO:1903827: regulation of cellular protein localization
                                                       GO:0051090: regulation of DNA-binding transcription factor activity
                                                       GO:0006575: cellular modified amino acid metabolic process
                                                       R-MMU-109582: Hemostasis
                                                       WP87: Nucleotide Metabolism
                                                       GO:0043535: regulation of blood vessel endothelial cell migration
                                                       GO:0002224: toll-like receptor signaling pathway
                                                       GO:0045899: positive regulation of RNA polymerase II transcription preinitiation complex assembly
                                                       GO:0045921: positive regulation of exocytosis
                                                       mmu05133: Pertussis
                                                       GO:0071219: cellular response to molecule of bacterial origin
                                                       R-MMU-556833: Metabolism of lipids
                                                       GO:0019262: N-acetylneuraminate catabolic process
                                                       R-MMU-156590: Glutathione conjugation
                                                       GO:0051604: protein maturation
                                                       GO:0033032: regulation of myeloid cell apoptotic process
                                                       GO:0051271: negative regulation of cellular component movement
                                                       R-MMU-174824: Plasma lipoprotein assembly, remodeling, and clearance
                                                       GO:0042559: pteridine-containing compound biosynthetic process
                                                       GO:0042180: cellular ketone metabolic process
                                                       GO:0045577: regulation of B cell differentiation
                                                       GO:0072657: protein localization to membrane
                                                       M00072: N-glycosylation by oligosaccharyltransferase
                                                       GO:1903334: positive regulation of protein folding
                                                       GO:0002317: plasma cell differentiation
                                                       mmu05134: Legionellosis
                                                       GO:0070936: protein K48-linked ubiquitination
                                                       GO:0008630: intrinsic apoptotic signaling pathway in response to DNA damage
                                                       ko04140: Autophagy - animal
                                                       GO:0018119: peptidyl-cysteine S-nitrosylation
                                                       GO:0006914: autophagy
                                                       GO:0051817: modulation of process of other organism involved in symbiotic interaction
                                                       ko00740: Riboflavin metabolism
5
        10
                  15
                                     25
                                              30
                           20
                  -log10(P)
```