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WP163: Cytoplasmic Ribosomal Proteins
                                              GO:0043043: peptide biosynthetic process
                                              mmu00190: Oxidative phosphorylation
                                              WP103: Cholesterol Biosynthesis
                                              GO:0031329: regulation of cellular catabolic process
                                              ko04141: Protein processing in endoplasmic reticulum
                                              WP1248: Oxidative phosphorylation
                                              GO:0009895: negative regulation of catabolic process
                                              mmu05206: MicroRNAs in cancer
                                              GO:2000377: regulation of reactive oxygen species metabolic process
                                              GO:0043281: regulation of cysteine-type endopeptidase activity involved in apoptotic process
                                              GO:0097193: intrinsic apoptotic signaling pathway
                                              WP310: mRNA processing
                                              GO:0043632: modification-dependent macromolecule catabolic process
                                              GO:0032922: circadian regulation of gene expression
                                              GO:0062197: cellular response to chemical stress
                                              GO:0036003: positive regulation of transcription from RNA polymerase II promoter in response to stress
                                              GO:0051345: positive regulation of hydrolase activity
                                              WP1763: PluriNetWork: mechanisms associated with pluripotency
                                              GO:0006984: ER-nucleus signaling pathway
                                              GO:0033002: muscle cell proliferation
                                              ko04657: IL-17 signaling pathway
                                              GO:0043408: regulation of MAPK cascade
                                              R-MMU-373755: Semaphorin interactions
                                              GO:0007159: leukocyte cell-cell adhesion
                                              GO:0001503: ossification
                                              WP307: Translation Factors
                                              GO:0006413: translational initiation
                                              GO:0048589: developmental growth
                                              GO:0001817: regulation of cytokine production
                                              GO:0051098: regulation of binding
                                              GO:0071214: cellular response to abiotic stimulus
                                              GO:0031400: negative regulation of protein modification process
                                              GO:1901653: cellular response to peptide
                                              GO:0048771: tissue remodeling
                                              GO:0048732: gland development
                                              GO:1902175: regulation of oxidative stress-induced intrinsic apoptotic signaling pathway
                                              GO:0097242: amyloid-beta clearance
                                              GO:2000278: regulation of DNA biosynthetic process
                                              GO:0001837: epithelial to mesenchymal transition
                                              WP447: Adipogenesis genes
                                              GO:1903844: regulation of cellular response to transforming growth factor beta stimulus
                                              GO:1902532: negative regulation of intracellular signal transduction
                                              GO:0022618: ribonucleoprotein complex assembly
                                              GO:0002521: leukocyte differentiation
                                              GO:0048569: post-embryonic animal organ development
                                              GO:0010922: positive regulation of phosphatase activity
                                              GO:0060212: negative regulation of nuclear-transcribed mRNA poly(A) tail shortening
                                              mmu05205: Proteoglycans in cancer
                                              GO:0071496: cellular response to external stimulus
                                              GO:0022604: regulation of cell morphogenesis
                                              mmu04151: PI3K-Akt signaling pathway
                                              GO:0001667: ameboidal-type cell migration
                                              GO:0006954: inflammatory response
                                              WP373: IL-3 Signaling Pathway
                                              GO:0031204: posttranslational protein targeting to membrane, translocation
                                              GO:0010656: negative regulation of muscle cell apoptotic process
                                              GO:0007566: embryo implantation
                                              GO:1903828: negative regulation of cellular protein localization
                                              R-MMU-166520: Signaling by NTRKs
                                              ko04550: Signaling pathways regulating pluripotency of stem cells
                                              R-MMU-1280215: Cytokine Signaling in Immune system
                                              CORUM:414: (ER)-localized multiprotein complex, in absence of Ig heavy chains
                                              R-MMU-5218859: Regulated Necrosis
                                              CORUM:25: 9S-cytosolic aryl hydrocarbon (Ah) receptor non-ligand activated complex
                                              GO:0045338: farnesyl diphosphate metabolic process
                                              GO:0010748: negative regulation of long-chain fatty acid import across plasma membrane
                                              GO:0032103: positive regulation of response to external stimulus
                                              GO:0043491: protein kinase B signaling
                                              GO:0071347: cellular response to interleukin-1
                                              GO:0032651: regulation of interleukin-1 beta production
                                              CORUM:382: Respiratory chain complex I, mitochondrial
                                              GO:0071634: regulation of transforming growth factor beta production
                                              GO:0045862: positive regulation of proteolysis
                                              GO:1904994: regulation of leukocyte adhesion to vascular endothelial cell
                                              GO:0033119: negative regulation of RNA splicing
                                              GO:0002573: myeloid leukocyte differentiation
                                              GO:0002262: myeloid cell homeostasis
                                               R-MMU-3700989: Transcriptional Regulation by TP53
                                              mmu05164: Influenza A
                                              GO:0045834: positive regulation of lipid metabolic process
                                              WP2084: SREBF and miR33 in cholesterol and lipid homeostasis
                                              GO:0075522: IRES-dependent viral translational initiation
                                              GO:0051090: regulation of DNA-binding transcription factor activity
                                              R-MMU-8953897: Cellular responses to external stimuli
                                              GO:0001922: B-1 B cell homeostasis
                                              GO:0030111: regulation of Wnt signaling pathway
                                              GO:2000378: negative regulation of reactive oxygen species metabolic process
                                              GO:0001704: formation of primary germ layer
                                              GO:0003300: cardiac muscle hypertrophy
                                              GO:0032527: protein exit from endoplasmic reticulum
                                              GO:0045599: negative regulation of fat cell differentiation
                                              GO:0030335: positive regulation of cell migration
                                              CORUM:538: Cytochrome c oxidase, mitochondrial
                                              GO:0007568: aging
                                              GO:0046777: protein autophosphorylation
                                              mmu05202: Transcriptional misregulation in cancer
                                              GO:0061458: reproductive system development
                                              GO:0006325: chromatin organization
                                              R-MMU-5689880: Ub-specific processing proteases
         15
                   20
                                      30
10
                             25
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