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hsa04141: Protein processing in endoplasmic reticulum
                                          R-HSA-5653656: Vesicle-mediated transport
                                          R-HSA-1474244: Extracellular matrix organization
                                          R-HSA-69278: Cell Cycle, Mitotic
                                          GO:1901137: carbohydrate derivative biosynthetic process
                                           GO:0009123: nucleoside monophosphate metabolic process
                                          GO:0002263: cell activation involved in immune response
                                          GO:0046700: heterocycle catabolic process
                                          R-HSA-72766: Translation
                                          R-HSA-109582: Hemostasis
                                          R-HSA-381119: Unfolded Protein Response (UPR)
                                          GO:0007265: Ras protein signal transduction
                                          GO:0060627: regulation of vesicle-mediated transport
                                          R-HSA-15869: Metabolism of nucleotides
                                          GO:0072657: protein localization to membrane
                                          GO:0035690: cellular response to drug
                                          GO:0030433: ubiquitin-dependent ERAD pathway
                                          hsa05012: Parkinson's disease
                                          GO:0045056: transcytosis
GO:0019058: viral life cycle
                                          GO:0071417: cellular response to organonitrogen compound
                                          GO:0050878: regulation of body fluid levels
                                          R-HSA-9675108: Nervous system development
                                          R-HSA-2262752: Cellular responses to stress
                                          hsa04611: Platelet activation
                                          hsa04110: Cell cycle
                                          GO:0007568: aging
                                          hsa04914: Progesterone-mediated oocyte maturation
                                          GO:0032386: regulation of intracellular transport
                                          R-HSA-69190: DNA strand elongation
                                          GO:0032799: low-density lipoprotein receptor particle metabolic process
                                          R-HSA-1566948: Elastic fibre formation
                                          R-HSA-382551: Transport of small molecules
                                          GO:0043687: post-translational protein modification
                                          GO:1903510: mucopolysaccharide metabolic process
                                          GO:0044772: mitotic cell cycle phase transition
                                          hsa04142: Lysosome
                                          GO:0051129: negative regulation of cellular component organization
                                          hsa00240: Pyrimidine metabolism
                                           M237: PID VEGFR1 2 PATHWAY
                                          R-HSA-73817: Purine ribonucleoside monophosphate biosynthesis
                                          GO:0031667: response to nutrient levels
                                          hsa04962: Vasopressin-regulated water reabsorption
                                          GO:0046040: IMP metabolic process
                                          GO:1904666: regulation of ubiquitin protein ligase activity
                                          M3468: NABA EČM REGULATORS
                                          hsa04933: AGE-RAGE signaling pathway in diabetic complications
                                          GO:0016482: cytosolic transport
                                          GO:0009895: negative regulation of catabolic process
                                          GO:0048284: organelle fusion
                                          GO:1990778: protein localization to cell periphery
                                          R-HSA-199992: trans-Golgi Network Vesicle Budding
                                          GO:0001501: skeletal system development
                                           M72: PID NECTIN PATHWAY
                                          GO:1990440: positive regulation of transcription from RNA polymerase II promoter in response to endoplasmic reti
                                          GO:0030335: positive regulation of cell migration
                                          GO:0031589: cell-substrate adhesion
                                          GO:0046686: response to cadmium ion
                                          GO:0014074: response to purine-containing compound
                                          GO:0040008: regulation of growth
                                          GO:0006890: retrograde vesicle-mediated transport, Golgi to ER
                                          GO:0071407: cellular response to organic cyclic compound
                                          GO:0051345: positive regulation of hydrolase activity
                                          R-HSA-75205: Dissolution of Fibrin Clot
                                          GO:0051098: regulation of binding
                                          GO:2001233: regulation of apoptotic signaling pathway
                                          M229: PID P38 ALPHA BETA DOWNSTREAM PATHWAY
                                          GO:0030155: regulation of cell adhesion
                                          GO:0010720: positive regulation of cell development
                                          R-HSA-69205: G1/S-Specific Transcription
                                          GO:0048251: elastic fiber assembly
                                          GO:0007033: vacuole organization
                                          M145: PID P53 DOWNSTREAM PATHWAY
                                          GO:0065005: protein-lipid complex assembly
                                          R-HSA-9022692: Regulation of MECP2 expression and activity
                                           GO:0031647: regulation of protein stability
                                           M210: PID IL8 CXCR2 PATHWAY
                                          GO:1902292: cell cycle DNA replication initiation
                                           GO:0032964: collagen biosynthetic process
                                           GO:0034446: substrate adhesion-dependent cell spreading
                                          GO:0030212: hyaluronan metabolic process
                                          GO:0051235: maintenance of location
                                          GO:0001704: formation of primary germ layer
                                          GO:0007610: behavior
                                          GO:0099003: vesicle-mediated transport in synapse
                                          R-HSA-76009: Platelet Aggregation (Plug Formation)
                                          GO:0051051: negative regulation of transport
                                          GO:0061029: eyelid development in camera-type eye
                                          GO:0032077: positive regulation of deoxyribonuclease activity
                                          GO:1901873: regulation of post-translational protein modification
                                          GO:0060359: response to ammonium ion
                                          GO:0051602: response to electrical stimulus
                                          R-HSA-3700989: Transcriptional Regulation by TP53
                                          GO:0046688: response to copper ion
                                          GO:0048308: organelle inheritance
                                          GO:0045861: negative regulation of proteolysis
                                          GO:0001503: ossification
                                          GO:0070848: response to growth factor
                                          M5193: SIG CHEMOTAXIS
                                          M120: PID ARF6 DOWNSTREAM PATHWAY
10
                      20
                                 25
           15
      -log10(P)
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