```
R-HSA-382551: Transport of small molecules
                                           R-HSA-1483206: Glycerophospholipid biosynthesis
                                           GO:0030855: epithelial cell differentiation
                                           GO:0043009: chordate embryonic development
                                           GO:0035239: tube morphogenesis
                                           GO:0046394: carboxylic acid biosynthetic process
                                           R-HSA-75205: Dissolution of Fibrin Clot
                                           hsa05418: Fluid shear stress and atherosclerosis
                                           M3468: NABA ECM REGULATORS
                                           GO:0048608: reproductive structure development
                                           WP4754: IL-18 signaling pathway
                                           GO:0010951: negative regulation of endopeptidase activity
                                           GO:0009636: response to toxic substance
                                           GO:0001655: urogenital system development
                                           GO:0006575: cellular modified amino acid metabolic process
                                           GO:0050801: ion homeostasis
                                           GO:0006820: anion transport
                                           hsa05200: Pathways in cancer
                                           WP98: Prostaglandin synthesis and regulation
                                           GO:0071214: cellular response to abiotic stimulus
                                           GO:1901615: organic hydroxy compound metabolic process
                                           GO:0090066: regulation of anatomical structure size
                                           GO:0050678: regulation of epithelial cell proliferation
                                           GO:0009725: response to hormone
                                           M3008: NABA ECM GLYCOPROTEINS
                                           GO:0060541: respiratory system development
                                           GO:0044282: small molecule catabolic process
                                           hsa00561: Glycerolipid metabolism
                                           M53: PID INTEGRIN3 PATHWAY
                                           GO:0006790: sulfur compound metabolic process
                                           GO:0016485: protein processing
                                           WP5065: SARS-CoV-2 altering angiogenesis via NRP1
                                           GO:0030111: regulation of Wnt signaling pathway
                                           R-HSA-983712: Ion channel transport
                                           GO:0010035: response to inorganic substance
                                           GO:0040008: regulation of growth
                                           GO:0010575: positive regulation of vascular endothelial growth factor production
                                           GO:0097435: supramolecular fiber organization
                                           GO:0007568: aging
                                           R-HSA-15869: Metabolism of nucleotides
                                           GO:1901605: alpha-amino acid metabolic process
                                           GO:0003013: circulatory system process
                                           GO:0071470: cellular response to osmotic stress
                                           R-HSA-425366: Transport of bile salts and organic acids, metal ions and amine compounds
                                           R-HSA-5083635: Defective B3GALTL causes PpS
                                           hsa04960: Aldosterone-regulated sodium reabsorption
                                           WP5053: Development of ureteric collection system
                                           GO:0070294: renal sodium ion absorption
                                           GO:0003018: vascular process in circulatory system
                                           GO:0000278: mitotic cell cycle
                                           GO:0070372: regulation of ERK1 and ERK2 cascade
                                           GO:1901989: positive regulation of cell cycle phase transition
                                           GO:0090270: regulation of fibroblast growth factor production
                                           GO:0048871: multicellular organismal homeostasis
                                           GO:0022407: regulation of cell-cell adhesion
                                           GO:1902600: proton transmembrane transport
                                           GO:0048589: developmental growth
                                           GO:0016042: lipid catabolic process
                                           GO:2000048: negative regulation of cell-cell adhesion mediated by cadherin
                                           GO:0032147: activation of protein kinase activity
                                           M239: PID A6B1 A6B4 INTEGRIN PATHWAY
                                           M183: PID IL6 7 PATHWAY
                                           GO:1901890: positive regulation of cell junction assembly
                                           WP2873: Aryl hydrocarbon receptor pathway
                                           GO:0045606: positive regulation of epidermal cell differentiation
                                           GO:0046903: secretion
                                           GO:0033280: response to vitamin D
                                           WP3888: VEGFA-VEGFR2 signaling pathway
                                           R-HSA-425393: Transport of inorganic cations/anions and amino acids/oligopeptides
                                           R-HSA-6785807: Interleukin-4 and Interleukin-13 signaling
                                           GO:2000179: positive regulation of neural precursor cell proliferation
                                           GO:0006468: protein phosphorylation
                                           GO:0048246: macrophage chemotaxis
                                           GO:0001991: regulation of systemic arterial blood pressure by circulatory renin-angiotensin
                                           GO:0008643: carbohydrate transport
                                           GO:0045661: regulation of myoblast differentiation
                                           GO:0008360: regulation of cell shape
                                           GO:0042359: vitamin D metabolic process
                                           R-HSA-68962: Activation of the pre-replicative complex
                                           hsa04919: Thyroid hormone signaling pathway
                                           GO:0060341: regulation of cellular localization
                                           GO:0030900: forebrain development
                                           R-HSA-983189: Kinesins
                                           R-HSA-381426: Regulation of Insulin-like Growth Factor (IGF) transport and uptake by Insulin-like Growth Factor Binding Proteins (IGFBPs)
                                           GO:0032233: positive regulation of actin filament bundle assembly
                                           GO:0035641: locomotory exploration behavior
                                           GO:0031175: neuron projection development
                                           hsa05323: Rheumatoid arthritis
                                           WP3624: Luna fibrosis
                                           R-HSA-2129379: Molecules associated with elastic fibres
                                           R-HSA-194315: Signaling by Rho GTPases
                                           WP5033: Genes associated with the development of rheumatoid arthritis
                                           GO:0099518: vesicle cytoskeletal trafficking
                                           GO:0006641: triglyceride metabolic process
                                           GO:0070482: response to oxygen levels
                                           M167: PID AP1 PATHWAY
                                           WP2880: Glucocorticoid receptor pathway
                                           GO:0002520: immune system development
                                           GO:0019722: calcium-mediated signaling
                                           hsa01230: Biosynthesis of amino acids
                     8
                               10
4
      -log10(P)
```