-log10(P) 0 234 6 10 R-HSA-2990846: SUMOylation ko04630: Jak-STAT signaling pathway GO:0051235: maintenance of location GO:0042537: benzene-containing compound metabolic process GO:0040017: positive regulation of locomotion hsa00982: Drug metabolism - cytochrome P450 GO:0001667: ameboidal-type cell migration GO:0006957: complement activation, alternative pathway GO:0070266: necroptotic process GO:0002031: G protein-coupled receptor internalization GO:0002224: toll-like receptor signaling pathway GO:0060353: regulation of cell adhesion molecule production R-HSA-5633007: Regulation of TP53 Activity GO:0061450: trophoblast cell migration R-HSA-211958: Miscellaneous substrates R-HSA-5689901: Metalloprotease DUBs ko05206: MicroRNAs in cancer ko00620: Pyruvate metabolism M5193: SIG CHEMOTAXIS R-HSA-1679131: Trafficking and processing of endosomal TLR GO:0045596: negative regulation of cell differentiation R-HSA-114508: Effects of PIP2 hydrolysis R-HSA-442742: CREB1 phosphorylation through NMDA receptor-mediated activation of RAS signaling R-HSA-1169092: Activation of RAS in B cells GO:0006805: xenobiotic metabolic process GO:0006069: ethanol oxidation GO:0009101: glycoprotein biosynthetic process GO:0015721: bile acid and bile salt transport GO:0046854: phosphatidylinositol phosphorylation R-HSA-68884: Mitotic Telophase/Cytokinesis GO:0071539: protein localization to centrosome M63: PID AVB3 OPN PATHWAY GO:0050732: negative regulation of peptidyl-tyrosine phosphorylation GO:0032071: regulation of endodeoxyribonuclease activity CORUM:5268: TNF-alpha/NF-kappa B signaling complex 7 GO:2000114: regulation of establishment of cell polarity GO:0018209: peptidyl-serine modification GO:0051493: regulation of cytoskeleton organization R-HSA-9006931: Signaling by Nuclear Receptors GO:0150012: positive regulation of neuron projection arborization GO:0009074: aromatic amino acid family catabolic process R-HSA-373760: L1CAM interactions ko04960: Aldosterone-regulated sodium reabsorption GO:0044282: small molecule catabolic process GO:0006839: mitochondrial transport GO:1903827: regulation of cellular protein localization ko04151: PI3K-Akt signaling pathway R-HSA-3700989: Transcriptional Regulation by TP53 GO:0031589: cell-substrate adhesion GO:1900117: regulation of execution phase of apoptosis ko04120: Ubiquitin mediated proteolysis R-HSA-1640170: Cell Cycle GO:0030307: positive regulation of cell growth GO:0042060: wound healing R-HSA-5688426: Deubiquitination R-HSA-211897: Cytochrome P450 - arranged by substrate type ko05222: Small cell lung cancer GO:0032988: ribonucleoprotein complex disassembly R-HSA-168898: Toll-like Receptor Cascades CORUM:1332: Large Drosha complex GO:0035405: histone-threonine phosphorylation R-HSA-187037: Signaling by NTRK1 (TRKA) GO:0006338: chromatin remodeling M151: PID AR TF PATHWAY GO:0031647: regulation of protein stability GO:0022407: regulation of cell-cell adhesion GO:0006611: protein export from nucleus GO:0097190: apoptotic signaling pathway GO:0042770: signal transduction in response to DNA damage GO:0051186: cofactor metabolic process M286: PID TGFBR PATHWAY GO:0044257: cellular protein catabolic process GO:0098927: vesicle-mediated transport between endosomal compartments R-HSA-983705: Signaling by the B Cell Receptor (BCR) GO:0071826: ribonucleoprotein complex subunit organization R-HSA-3858494: Beta-catenin independent WNT signaling GO:0006623: protein targeting to vacuole GO:0043281: regulation of cysteine-type endopeptidase activity involved in apoptotic process R-HSA-389359: CD28 dependent Vav1 pathway GO:0016569: covalent chromatin modification R-HSA-180024: DARPP-32 events M53: PID INTEGRIN3 PATHWAY hsa05169: Epstein-Barr virus infection GO:0003307: regulation of Wnt signaling pathway involved in heart development GO:0034655: nucleobase-containing compound catabolic process GO:0015718: monocarboxylic acid transport R-HSA-77305: Beta oxidation of palmitoyl-CoA to myristoyl-CoA M255: PID HIF1 TFPATHWAY GO:0010638: positive regulation of organelle organization GO:0042791: 5S class rRNA transcription by RNA polymerase III GO:0042180: cellular ketone metabolic process ko05205: Proteoglycans in cancer R-HSA-2454202: Fc epsilon receptor (FCERI) signaling hsa05165: human papillomavirus infection R-HSA-8953854: Metabolism of RNA GO:0022613: ribonucleoprotein complex biogenesis GO:0019216: regulation of lipid metabolic process GO:0032787: monocarboxylic acid metabolic process R-HSA-556833: Metabolism of lipids R-HSA-211859: Biological oxidations 10 uM 100