

regulation of cell cycle, GO:0051726

- cell cycle arrest, GO:0007050
- core promoter binding, GO:0001047
- aging, GO:0007568
- negative regulation of protein kinase activity, GO:0006469
- liver development, GO:0001889
- fat cell differentiation, GO:0045444
- kidney development, GO:0001822
- transcriptional activator activity, GO:0001228
- RNA polymerase II transcription factor activity, GO:0000981
- positive regulation of JNK cascade, GO:0046330
- negative regulation of cell migration, GO:0030336
- positive regulation of neuron death, GO:1901216
- transcription regulatory region sequence-specific DNA binding, GO:0000976
- animal organ morphogenesis, GO:0009887
- animal organ regeneration, GO:0031100
- response to mechanical stimulus, GO:0009612
- regulation of mitochondrial membrane potential, GO:0051881
- response to insulin, GO:0032868
- positive regulation of neuron apoptotic process, GO:0043525
- R-SMAD binding, GO:0070412
- positive regulation of pri-miRNA transcription from RNA polymerase II promoter, GO:1902895
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- transcriptional repressor activity, GO:0001227
- negative regulation of intrinsic apoptotic signaling pathway, GO:2001243
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- intrinsic apoptotic signaling pathway by p53 class mediator, GO:0072332
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- negative regulation of cell cycle, GO:0045786
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- cellular response to organic substance, GO:0071310
- T cell homeostasis, GO:0043029
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- anatomical structure morphogenesis, GO:0009653
- regulation of reactive oxygen species metabolic process, GO:2000377
- regulation of insulin secretion, GO:0050796
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- positive regulation of cell cycle arrest, GO:0071158
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- megakaryocyte development, GO:0035855
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- activation of cysteine-type endopeptidase activity, GO:0097202
- positive regulation of osteoblast differentiation, GO:0045669
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- negative regulation of organ growth, GO:0046621
- endoplasmic reticulum calcium ion homeostasis, GO:0032469
- response to steroid hormone, GO:0048545
- positive regulation of smooth muscle cell migration, GO:0014911
- transcriptional activator activity, GO:0001190
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- dentate gyrus development, GO:0021542
- activation of MAPKKK activity, GO:0000185
- CD40 receptor complex, GO:0035631
- adenosine deaminase activity, GO:0004000
- G-protein coupled receptor kinase activity, GO:0004703
- regulation of adaptive immune response, GO:0002819
- protein insertion into mitochondrial membrane involved in apoptotic signaling pathway, GO:0001844
- regulation of transforming growth factor beta receptor signaling pathway, GO:0017015
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- positive regulation of reactive oxygen species metabolic process, GO:2000379
- positive regulation of calcium ion transport into cytosol, GO:0010524
- positive regulation of B cell proliferation, GO:0030890
- regulation of gluconeogenesis, GO:0006111
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- germ cell migration, GO:0008354
- definitive hemopoiesis, GO:0060216
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- regulation of synaptic transmission, GO:0032228
- positive regulation of tyrosine phosphorylation of STAT protein, GO:0042531
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