

regulation of hematopoietic stem cell differentiation, GO:1902036

- tumor necrosis factor-mediated signaling pathway, GO:0033209
- negative regulation of canonical Wnt signaling pathway, GO:0090090
- proteasome complex, GO:0000502
- antigen processing and presentation of exogenous peptide antigen via MHC class I, GO:0002479
- nucleosome, GO:0000786
- hemopoiesis, GO:0030097
- response to virus, GO:0009615
- chromatin organization, GO:0006325
- regulation of cellular amino acid metabolic process, GO:0006521
- nucleosome assembly, GO:0006334
- endopeptidase activity, GO:0004175
- RNA polymerase II transcription factor activity, GO:0000981
- nucleosomal DNA binding, GO:0031492
- RNA polymerase II distal enhancer sequence-specific DNA binding, GO:0000980
- protein heterotetramerization, GO:0051290
- chromatin silencing at rDNA, GO:0000183
- proteolysis involved in cellular protein catabolic process, GO:0051603
- telomere organization, GO:0032200
- transcriptional activator activity, GO:0001228
- positive regulation of neuron differentiation, GO:0045666
- proteasome accessory complex, GO:0022624
- transcription regulatory region sequence-specific DNA binding, GO:0000976
- RNA polymerase II transcription factor complex, GO:0090575
- telomere capping, GO:0016233
- embryo implantation, GO:0007566
- erythrocyte differentiation, GO:0030218
- nuclear nucleosome, GO:0000788
- regulation of G1/S transition of mitotic cell cycle, GO:2000045
- proteasome core complex, GO:0005839
- inner ear morphogenesis, GO:0042472
- cytosolic proteasome complex, GO:0031597
- negative regulation of megakaryocyte differentiation, GO:0045653
- negative regulation of fat cell differentiation, GO:0045599
- threonine-type endopeptidase activity, GO:0004298
- proteasome regulatory particle, GO:0005838
- proteasome regulatory particle, GO:0008540
- cell fate determination, GO:0001709
- negative regulation of programmed cell death, GO:0043069
- gastrulation with mouth forming second, GO:0001702
- regulation of intracellular estrogen receptor signaling pathway, GO:0033146
- muscle cell differentiation, GO:0042692
- positive regulation of cytokine production, GO:0001819
- HMG box domain binding, GO:0071837
- regulation of B cell receptor signaling pathway, GO:0050855
- regulation of keratinocyte differentiation, GO:0045616
- MDM2/MDM4 family protein binding, GO:0097371
- spinal cord association neuron differentiation, GO:0021527
- embryonic placenta development, GO:0001892
- positive regulation of stress-activated MAPK cascade, GO:0032874
- enhancer binding, GO:0035326
- transcriptional activator activity, GO:0001205
- positive regulation of cell-matrix adhesion, GO:0001954
- uterus development, GO:0060065
- enzyme regulator activity, GO:0030234
- granulocyte differentiation, GO:0030851