

**chromatin remodeling, GO:0006338**

- RNA polymerase II transcription factor activity, GO:0000981
- transcription regulatory region sequence-specific DNA binding, GO:0000976
- histone deacetylase complex, GO:0000118
- ATP-dependent chromatin remodeling, GO:0043044
- NuA4 histone acetyltransferase complex, GO:0035267
- regulation of growth, GO:0040008
- Ada2/Gcn5/Ada3 transcription activator complex, GO:0005671
- ligand-dependent nuclear receptor binding, GO:0016922
- skeletal system development, GO:0001501
- intermediate filament cytoskeleton, GO:0045111
- transcriptional activator activity, GO:0001228
- histone H2A acetylation, GO:0043968
- negative regulation of cell cycle, GO:0045786
- cardiac muscle cell differentiation, GO:0055007
- npBAF complex, GO:0071564
- N-acetyltransferase activity, GO:0008080
- Ino80 complex, GO:0031011
- SWI/SNF complex, GO:0016514
- positive regulation of interleukin-4 production, GO:0032753
- nBAF complex, GO:0071565
- cellular response to estradiol stimulus, GO:0071392
- cartilage development, GO:0051216
- BAF-type complex, GO:0090544
- embryonic limb morphogenesis, GO:0030326
- actomyosin, GO:0042641
- histone H4-K16 acetylation, GO:0043984
- aorta morphogenesis, GO:0035909
- regulation of gene expression, GO:0040029
- positive regulation of collagen biosynthetic process, GO:0032967
- histone H3 deacetylation, GO:0070932
- core promoter proximal region sequence-specific DNA binding, GO:0000987
- ventricular trabecula myocardium morphogenesis, GO:0003222
- embryonic placenta development, GO:0001892
- protein-arginine deiminase activity, GO:0004668
- protein citrullination, GO:0018101
- negative regulation of chondrocyte differentiation, GO:0032331
- skeletal muscle cell differentiation, GO:0035914
- positive regulation of cartilage development, GO:0061036
- negative regulation of lymphocyte proliferation, GO:0050672
- positive regulation of interleukin-5 production, GO:0032754
- regulation of lipid kinase activity, GO:0043550
- oxygen transport, GO:0015671
- I band, GO:0031674
- telencephalon development, GO:0021537
- NAD-dependent histone deacetylase activity (H3-K14 specific), GO:0032041
- neuron maturation, GO:0042551