

chromatin remodeling, GO:0006338

- regulation of growth, GO:0040008
- core promoter binding, GO:0001047
- skeletal system development, GO:0001501
- NF-kappaB binding, GO:0051059
- chromatin silencing, GO:0006342
- Ada2/Gcn5/Ada3 transcription activator complex, GO:0005671
- histone deacetylase complex, GO:0000118
- ATP-dependent chromatin remodeling, GO:0043044
- NuA4 histone acetyltransferase complex, GO:0035267
- intermediate filament cytoskeleton, GO:0045111
- negative regulation of osteoblast differentiation, GO:0045668
- Ino80 complex, GO:0031011
- histone H2A acetylation, GO:0043968
- histone H3 deacetylation, GO:0070932
- ligand-dependent nuclear receptor binding, GO:0016922
- N-acetyltransferase activity, GO:0008080
- NAD-dependent histone deacetylase activity (H3-K14 specific), GO:0032041
- actomyosin, GO:0042641
- npBAF complex, GO:0071564
- BAF-type complex, GO:0090544
- anatomical structure formation involved in morphogenesis, GO:0048646
- positive regulation of interleukin-4 production, GO:0032753
- neural retina development, GO:0003407
- nBAF complex, GO:0071565
- positive regulation of phospholipase C activity, GO:0010863
- eyelid development in camera-type eye, GO:0061029
- negative regulation of activated T cell proliferation, GO:0046007
- negative regulation of cytokine secretion, GO:0050710
- proximal/distal pattern formation, GO:0009954
- I band, GO:0031674
- negative regulation of immune response, GO:0050777
- negative regulation of interleukin-10 production, GO:0032693
- regulation of DNA recombination, GO:0000018
- negative regulation of myotube differentiation, GO:0010832