

**chromatin remodeling, GO:0006338**

- regulation of growth, GO:0040008
- Ada2/Gcn5/Ada3 transcription activator complex, GO:0005671
- transcription regulatory region sequence-specific DNA binding, GO:0000976
- ligand-dependent nuclear receptor binding, GO:0016922
- nucleotide metabolic process, GO:0009117
- histone deacetylase complex, GO:0000118
- negative regulation of osteoblast differentiation, GO:0045668
- ATP-dependent chromatin remodeling, GO:0043044
- anatomical structure morphogenesis, GO:0009653
- SWI/SNF complex, GO:0016514
- N-acetyltransferase activity, GO:0008080
- NuA4 histone acetyltransferase complex, GO:0035267
- nBAF complex, GO:0071565
- smooth muscle tissue development, GO:0048745
- embryonic limb morphogenesis, GO:0030326
- seminiferous tubule development, GO:0072520
- BAF-type complex, GO:0090544
- myoblast differentiation, GO:0045445
- Ino80 complex, GO:0031011
- nuclear replication fork, GO:0043596
- histone H2A acetylation, GO:0043968
- neural retina development, GO:0003407
- positive regulation of interleukin-4 production, GO:0032753
- npBAF complex, GO:0071564
- negative regulation by host of viral transcription, GO:0043922
- negative regulation of cytokine secretion, GO:0050710
- actomyosin, GO:0042641
- anterior/posterior pattern specification, GO:0009952
- positive regulation of striated muscle cell differentiation, GO:0051155
- anatomical structure formation involved in morphogenesis, GO:0048646