

transcription elongation from RNA polymerase II promoter, GO:0006368

- snRNA transcription from RNA polymerase II promoter, GO:0042795
- positive regulation of gene expression, GO:0045815
- histone acetyltransferase activity, GO:0004402
- regulation of protein kinase activity, GO:0045859
- RNA polymerase II core promoter sequence-specific DNA binding, GO:0000979
- DNA-directed 5'-3' RNA polymerase activity, GO:0003899
- MLL1 complex, GO:0071339
- DNA-templated transcription, GO:0006352
- somatic stem cell population maintenance, GO:0035019
- histone acetyltransferase complex, GO:0000123
- transcription from RNA polymerase III promoter, GO:0006383
- positive regulation of viral transcription, GO:0050434
- transcription factor TFTC complex, GO:0033276
- mRNA polyadenylation, GO:0006378
- DNA-directed RNA polymerase II, GO:0005665
- RNA polymerase III activity, GO:0001056
- TBP-class protein binding, GO:0017025
- transcription elongation factor complex, GO:0008023
- nucleosome binding, GO:0031491
- transcriptionally active chromatin, GO:0035327
- STAGA complex, GO:0030914
- translation elongation factor activity, GO:0003746
- DNA-directed RNA polymerase I complex, GO:0005736
- SAGA complex, GO:0000124
- RNA polymerase binding, GO:0070063
- RNA polymerase I activity, GO:0001054
- histone deubiquitination, GO:0016578
- proteasome binding, GO:0070628
- RNA polymerase II transcriptional preinitiation complex assembly, GO:0051123
- RNA polymerase II activity, GO:0001055
- DNA-directed RNA polymerase III complex, GO:0005666
- vitamin D receptor binding, GO:0042809
- mRNA cleavage, GO:0006379
- positive regulation of endopeptidase activity, GO:0010950
- stem cell differentiation, GO:0048863
- proteasome regulatory particle, GO:0008541
- regulation of mRNA processing, GO:0050684
- segment specification, GO:0007379
- negative regulation of epithelial cell proliferation involved in prostate gland development, GO:0060770
- aryl hydrocarbon receptor binding, GO:0017162
- endopeptidase activator activity, GO:0061133
- phosphorylation of RNA polymerase II C-terminal domain, GO:0070816
- LRR domain binding, GO:0030275
- neural crest cell migration, GO:0001755
- negative regulation of myeloid cell differentiation, GO:0045638