

transcription elongation from RNA polymerase II promoter, GO:0006368

- snRNA transcription from RNA polymerase II promoter, GO:0042795
- histone acetyltransferase activity, GO:0004402
- DNA-directed 5'-3' RNA polymerase activity, GO:0003899
- transcription from RNA polymerase III promoter, GO:0006383
- positive regulation of viral transcription, GO:0050434
- thyroid hormone receptor binding, GO:0046966
- mRNA polyadenylation, GO:0006378
- MLL1 complex, GO:0071339
- TBP-class protein binding, GO:0017025
- RNA polymerase III activity, GO:0001056
- somatic stem cell population maintenance, GO:0035019
- transcription elongation factor complex, GO:0008023
- DNA-directed RNA polymerase III complex, GO:0005666
- vitamin D receptor binding, GO:0042809
- proteasome regulatory particle, GO:0008541
- DNA-directed RNA polymerase II, GO:0005665
- transcription factor TFTC complex, GO:0033276
- SAGA complex, GO:0000124
- RNA polymerase II activity, GO:0001055
- RNA polymerase II transcriptional preinitiation complex assembly, GO:0051123
- negative regulation of myeloid cell differentiation, GO:0045638
- regulation of transcription from RNA polymerase III promoter, GO:0006359
- transcription factor activity, GO:0001076
- DNA-directed RNA polymerase I complex, GO:0005736
- positive regulation of B cell differentiation, GO:0045579
- STAGA complex, GO:0030914
- Cdc73/Paf1 complex, GO:0016593
- poly(A)+ mRNA export from nucleus, GO:0016973
- RNA polymerase I activity, GO:0001054
- mRNA cleavage, GO:0006379
- negative regulation of cysteine-type endopeptidase activity, GO:2000117
- positive regulation of phosphorylation of RNA polymerase II C-terminal domain, GO:1901409
- negative regulation of insulin secretion involved in cellular response to glucose stimulus, GO:0061179
- snRNA transcription from RNA polymerase III promoter, GO:0042796
- positive regulation of endopeptidase activity, GO:0010950