

transcription elongation from RNA polymerase II promoter, GO:0006368

- positive regulation of gene expression, GO:0045815
- snRNA transcription from RNA polymerase II promoter, GO:0042795
- DNA-directed 5'-3' RNA polymerase activity, GO:0003899
- DNA-templated transcription, GO:0006352
- transcription from RNA polymerase III promoter, GO:0006383
- transcription initiation from RNA polymerase I promoter, GO:0006361
- MLL1 complex, GO:0071339
- histone acetyltransferase activity, GO:0004402
- histone acetyltransferase complex, GO:0000123
- thyroid hormone receptor binding, GO:0046966
- positive regulation of viral transcription, GO:0050434
- translational elongation, GO:0006414
- somatic stem cell population maintenance, GO:0035019
- mRNA polyadenylation, GO:0006378
- TBP-class protein binding, GO:0017025
- positive regulation of cell growth, GO:0030307
- DNA-directed RNA polymerase III complex, GO:0005666
- translation elongation factor activity, GO:0003746
- RNA polymerase III activity, GO:0001056
- transcription elongation factor complex, GO:0008023
- DNA-directed RNA polymerase II, GO:0005665
- DNA-directed RNA polymerase I complex, GO:0005736
- vitamin D receptor binding, GO:0042809
- RNA polymerase I activity, GO:0001054
- transcriptionally active chromatin, GO:0035327
- RNA polymerase II transcriptional preinitiation complex assembly, GO:0051123
- proteasome regulatory particle, GO:0008541
- proteasome assembly, GO:0043248
- RNA polymerase II activity, GO:0001055
- transcription factor TFTC complex, GO:0033276
- negative regulation of myeloid cell differentiation, GO:0045638
- STAGA complex, GO:0030914
- nucleosome binding, GO:0031491
- regulation of transcription from RNA polymerase III promoter, GO:0006359
- SAGA complex, GO:0000124
- histone deubiquitination, GO:0016578
- positive regulation of transcription elongation from RNA polymerase II promoter, GO:0032968
- nuclear-transcribed mRNA catabolic process, GO:0000288
- elongin complex, GO:0070449
- positive regulation of endopeptidase activity, GO:0010950
- regulation of mRNA processing, GO:0050684
- negative regulation of mRNA polyadenylation, GO:1900364
- negative regulation of epithelial cell proliferation involved in prostate gland development, GO:0060770
- cellular response to ATP, GO:0071318
- negative regulation of stem cell differentiation, GO:2000737
- transcription from RNA polymerase I promoter, GO:0006360
- aryl hydrocarbon receptor binding, GO:0017162
- mRNA cleavage, GO:0006379
- poly(A)+ mRNA export from nucleus, GO:0016973
- Elongator holoenzyme complex, GO:0033588
- endopeptidase activator activity, GO:0061133
- pre-mRNA cleavage required for polyadenylation, GO:0098789
- histone H2B ubiquitination, GO:0033523
- negative regulation of cysteine-type endopeptidase activity, GO:2000117
- LRR domain binding, GO:0030275
- snRNA transcription from RNA polymerase III promoter, GO:0042796
- central nervous system myelination, GO:0022010
- positive regulation of B cell differentiation, GO:0045579