

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
- regulation of protein kinase activity, GO:0045859
- DNA-directed 5'-3' RNA polymerase activity, GO:0003899
- histone acetyltransferase activity, GO:0004402
- mRNA polyadenylation, GO:0006378
- MLL1 complex, GO:0071339
- transcription from RNA polymerase III promoter, GO:0006383
- thyroid hormone receptor binding, GO:0046966
- DNA-templated transcription, GO:0006352
- transcription elongation factor complex, GO:0008023
- vitamin D receptor binding, GO:0042809
- negative regulation of myeloid cell differentiation, GO:0045638
- RNA polymerase binding, GO:0070063
- transcriptionally active chromatin, GO:0035327
- nucleosome binding, GO:0031491
- positive regulation of B cell differentiation, GO:0045579
- negative regulation of transcription elongation from RNA polymerase II promoter, GO:0034244
- transcription factor activity, GO:0001076
- mRNA cleavage, GO:0006379
- transcription coactivator binding, GO:0001223
- positive regulation of transcription elongation from RNA polymerase II promoter, GO:0032968
- DNA-directed RNA polymerase III complex, GO:0005666
- histone H2B ubiquitination, GO:0033523
- DNA-directed RNA polymerase I complex, GO:0005736
- RNA polymerase III activity, GO:0001056
- aryl hydrocarbon receptor binding, GO:0017162
- RNA polymerase I activity, GO:0001054
- RNA polymerase II activity, GO:0001055
- pre-mRNA cleavage required for polyadenylation, GO:0098789