



nucleus	16	72.7%
negative regulation of transcription by RNA polymerase II	10	45.5%
cytoplasm	9	40.9%
negative regulation of DNA-templated transcription	9	40.9%
cytosol	7	31.8%
chromatin remodeling	6	27.3%
positive regulation of DNA-templated transcription	6	27.3%
chromatin	5	22.7%
DNA binding	5	22.7%
Goterm	Count	%
negative regulation of transcription by RNA polymerase II	10	45.5%
negative regulation of DNA-templated transcription	9	40.9%
chromatin remodeling	6	27.3%
positive regulation of DNA-templated transcription	6	27.3%
positive regulation of stem cell population maintenance	5	22.7%
regulation of cell fate specification	5	22.7%
regulation of DNA-templated transcription	5	22.7%
regulation of stem cell differentiation	5	22.7%
negative regulation of cell migration	4	18.2%
negative regulation of stem cell population maintenance	4	18.2%
Goterm	Count	%
DNA binding	5	22.7%
RNA binding	5	22.7%
transcription cis-regulatory region binding	5	22.7%
zinc ion binding	5	22.7%
ATP binding	4	18.2%
chromatin binding	4	18.2%
histone deacetylase binding	4	18.2%
metal ion binding	4	18.2%