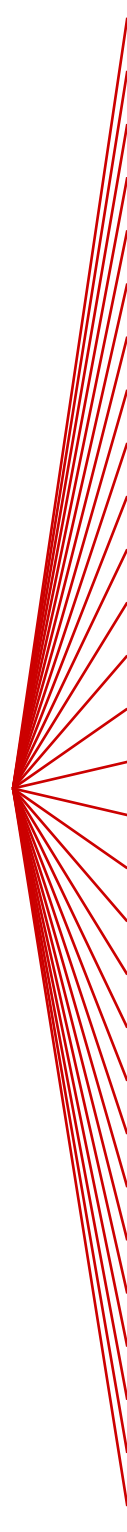


**core promoter binding, GO:0001047**



- circadian regulation of gene expression, GO:0032922
- ligand-dependent nuclear receptor transcription coactivator activity, GO:0030374
- RNA polymerase II transcription factor activity, GO:0000981
- histone deacetylase complex, GO:0000118
- transcriptional repressor activity, GO:0001227
- transcription regulatory region sequence-specific DNA binding, GO:0000976
- neural tube development, GO:0021915
- ATP-dependent chromatin remodeling, GO:0043044
- negative regulation of nucleic acid-templated transcription, GO:1903507
- RNA polymerase II transcription factor complex, GO:0090575
- DNA binding, GO:0008301
- skeletal muscle cell differentiation, GO:0035914
- entrainment of circadian clock by photoperiod, GO:0043153
- regulation of G1/S transition of mitotic cell cycle, GO:2000045
- transdifferentiation, GO:0060290
- monocyte differentiation, GO:0030224
- NAD-dependent histone deacetylase activity (H3-K14 specific), GO:0032041
- left/right axis specification, GO:0070986
- response to redox state, GO:0051775
- chondrocyte differentiation, GO:0002062
- transcription factor activity, GO:0000989
- lymphocyte differentiation, GO:0030098
- Notch binding, GO:0005112
- transcription factor activity, GO:0000982
- negative regulation of neurogenesis, GO:0050768
- cardiac muscle cell proliferation, GO:0060038
- steroid binding, GO:0005496
- response to metal ion, GO:0010038
- integrator complex, GO:0032039