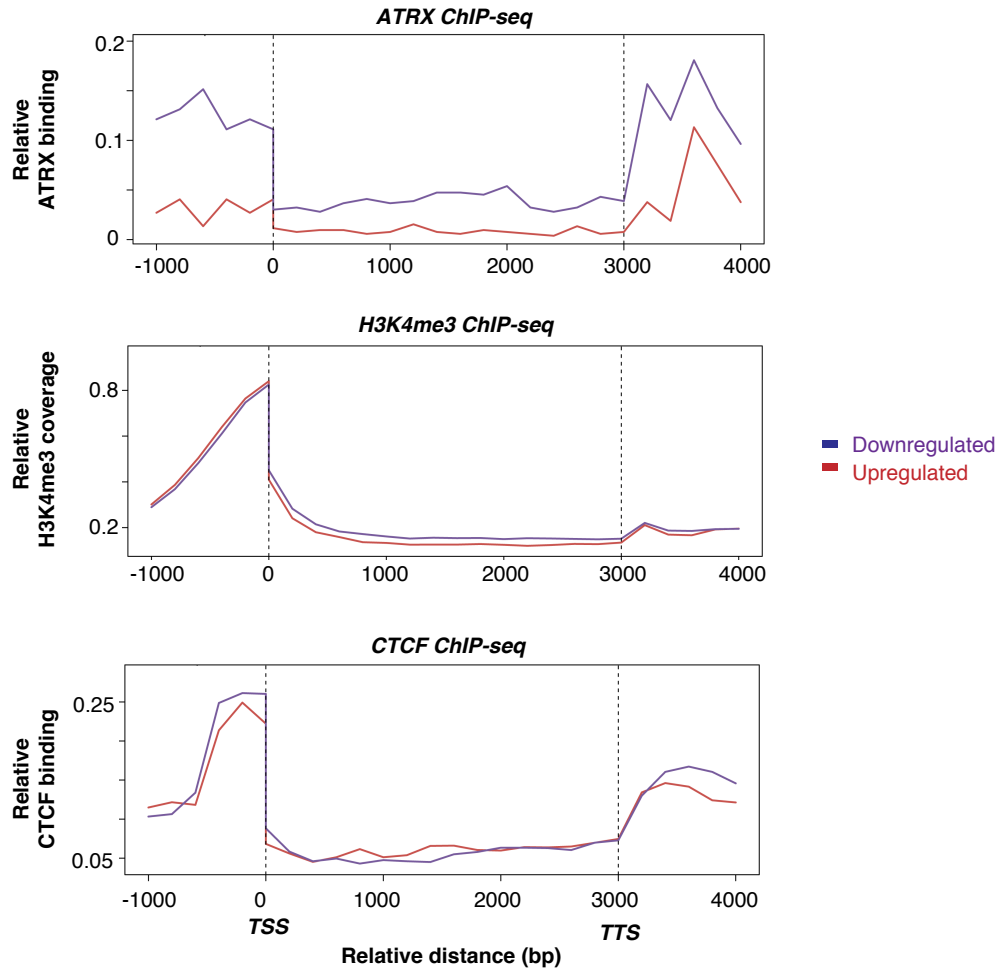
ATRX ChIP-seq
Relative ATRX binding
0.2
0.1
0
-1000
0
1000
2000
3000
4000
H3K4me3 ChIP-seq
Relative H3K4me3 coverage
0.8
0.2
CTCF ChIP-seq
Relative CTCF binding
0.25
0.05
TSS
TTS
Relative distance (bp)
Downregulated
Upregulated

Figure S6