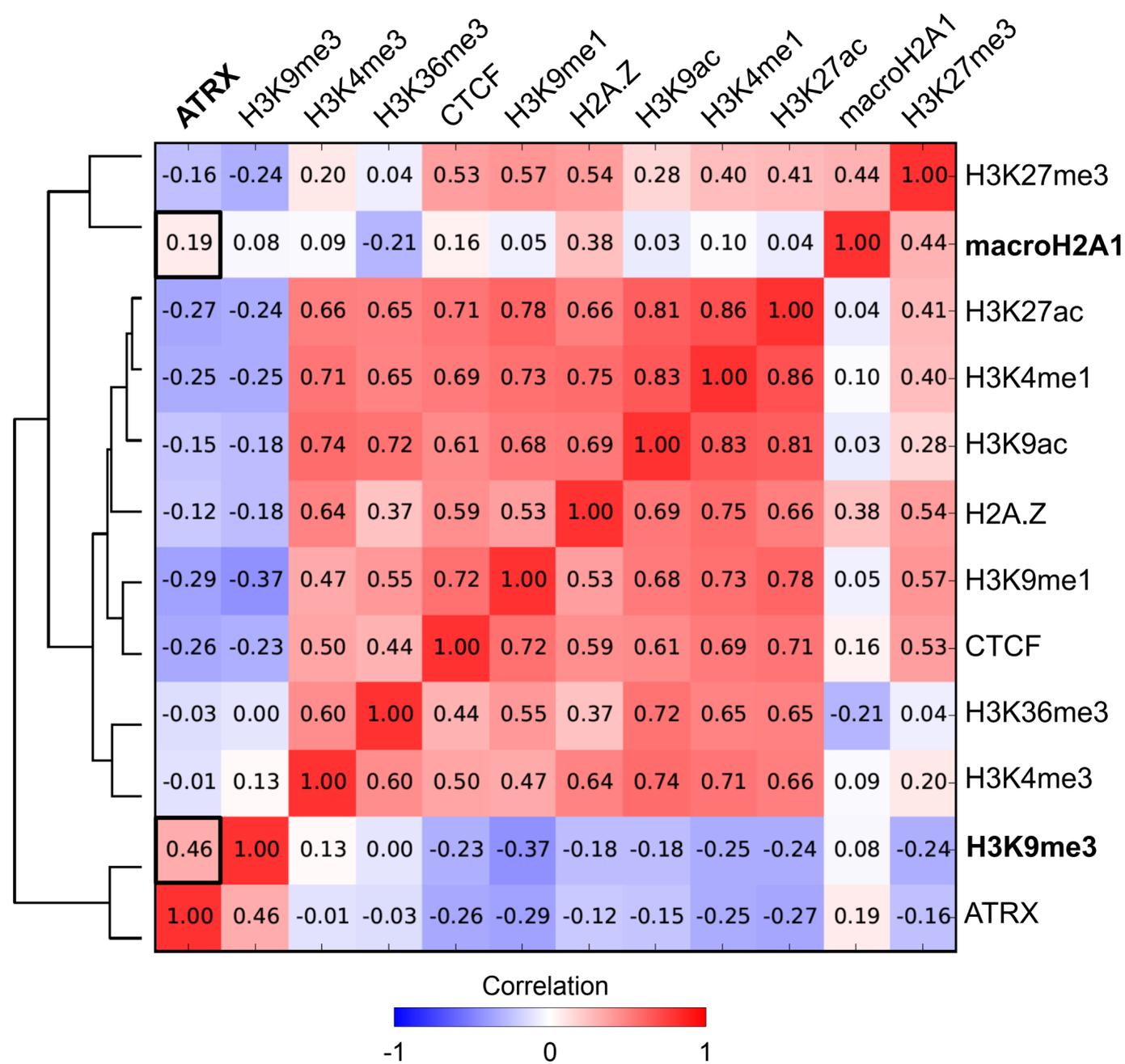


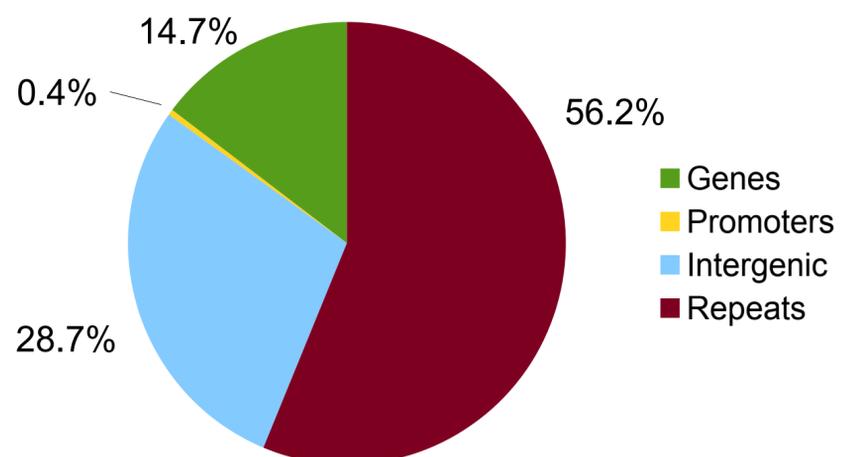
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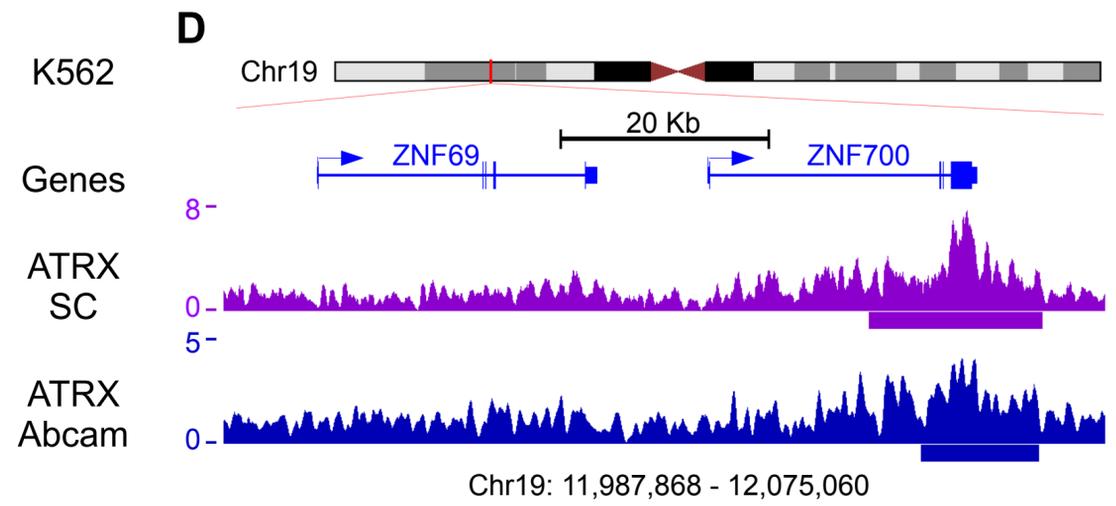
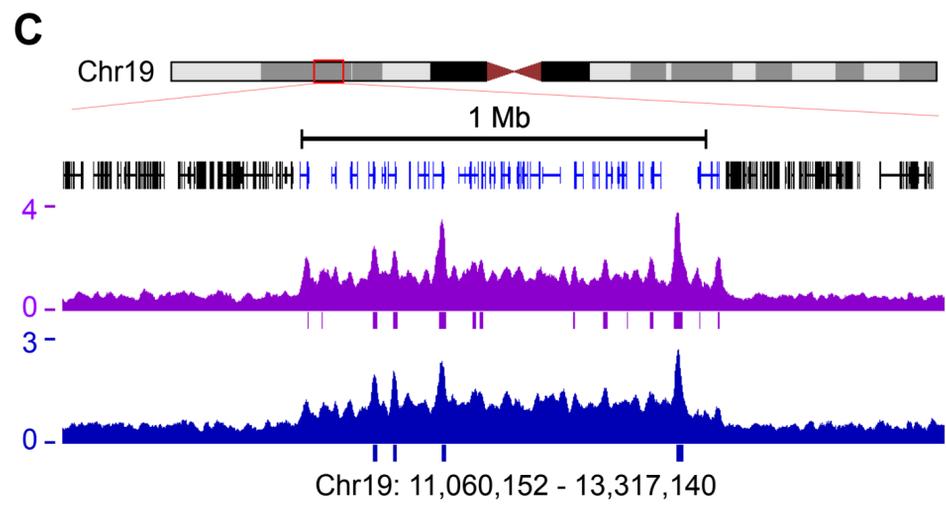
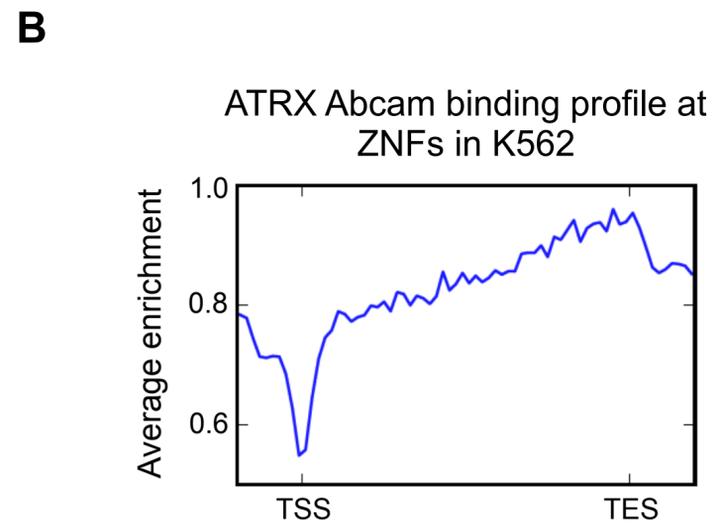
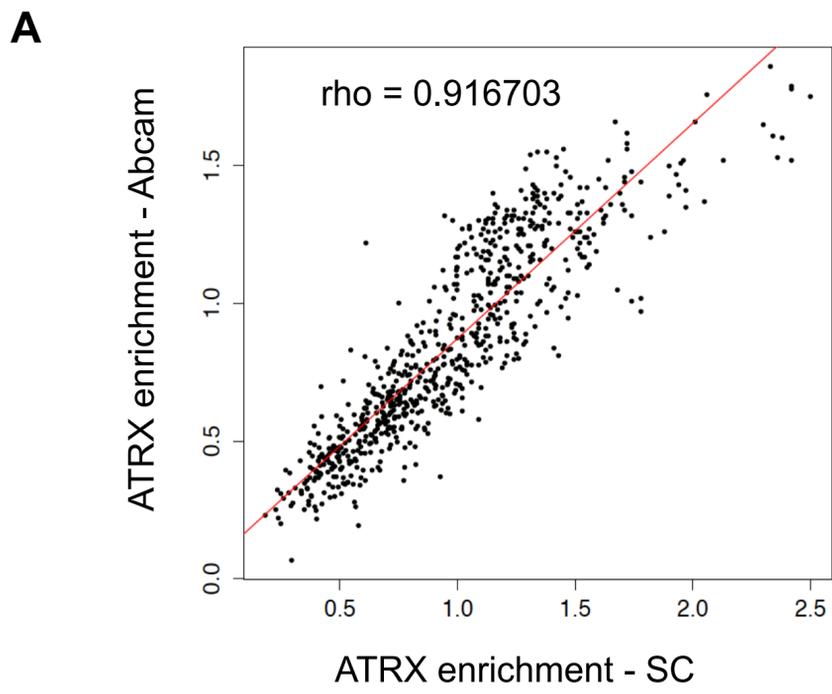
## Genome wide correlation of ChIP-seq profiles in K562



B

## ATRX peak distribution in K562

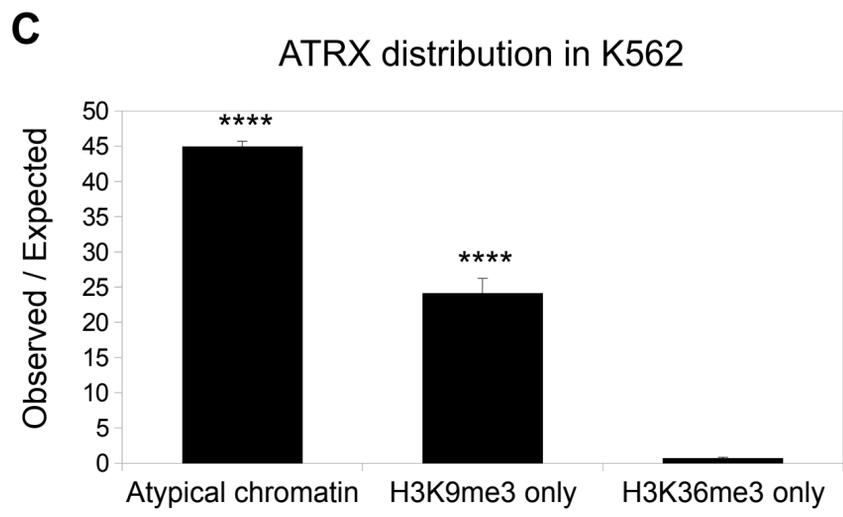
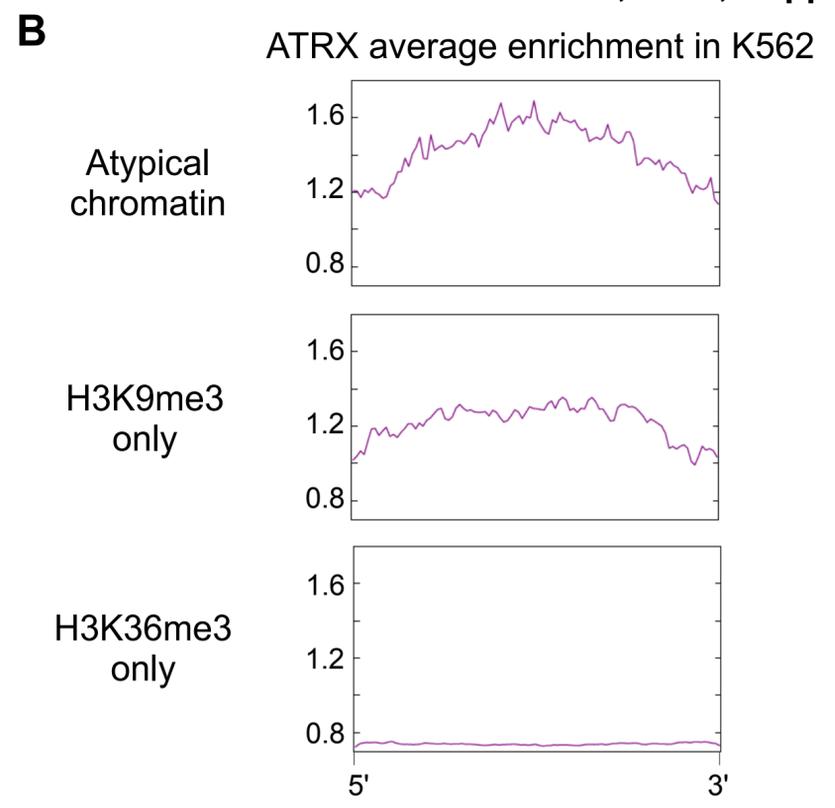
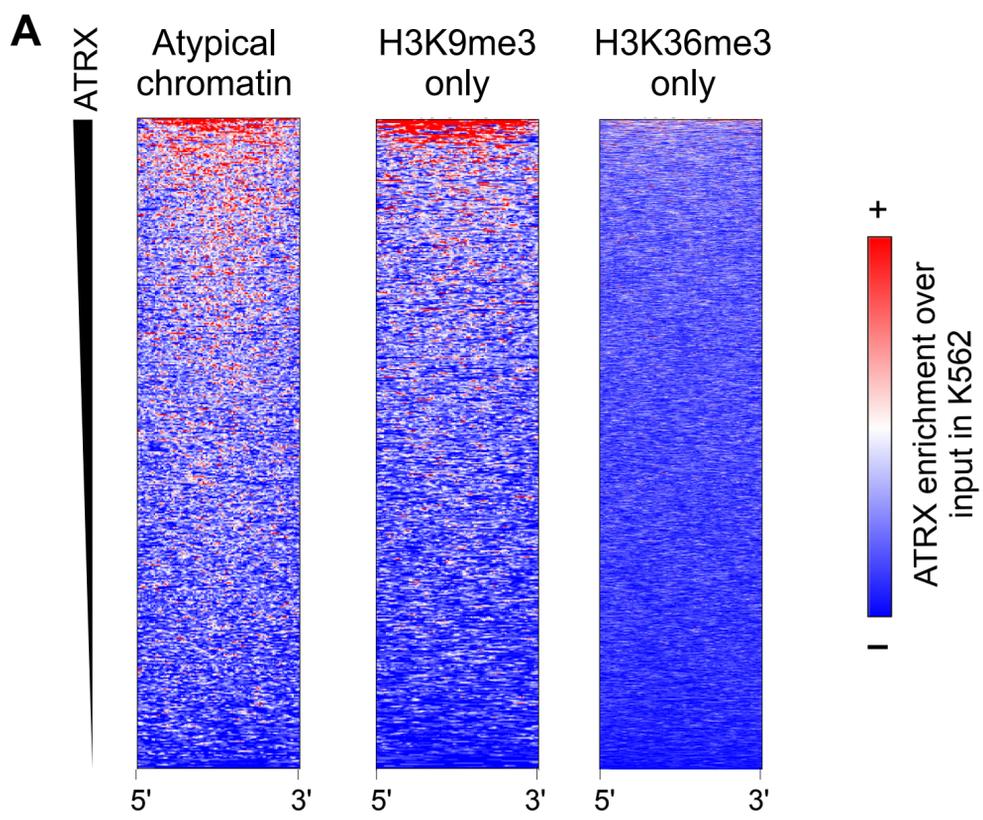


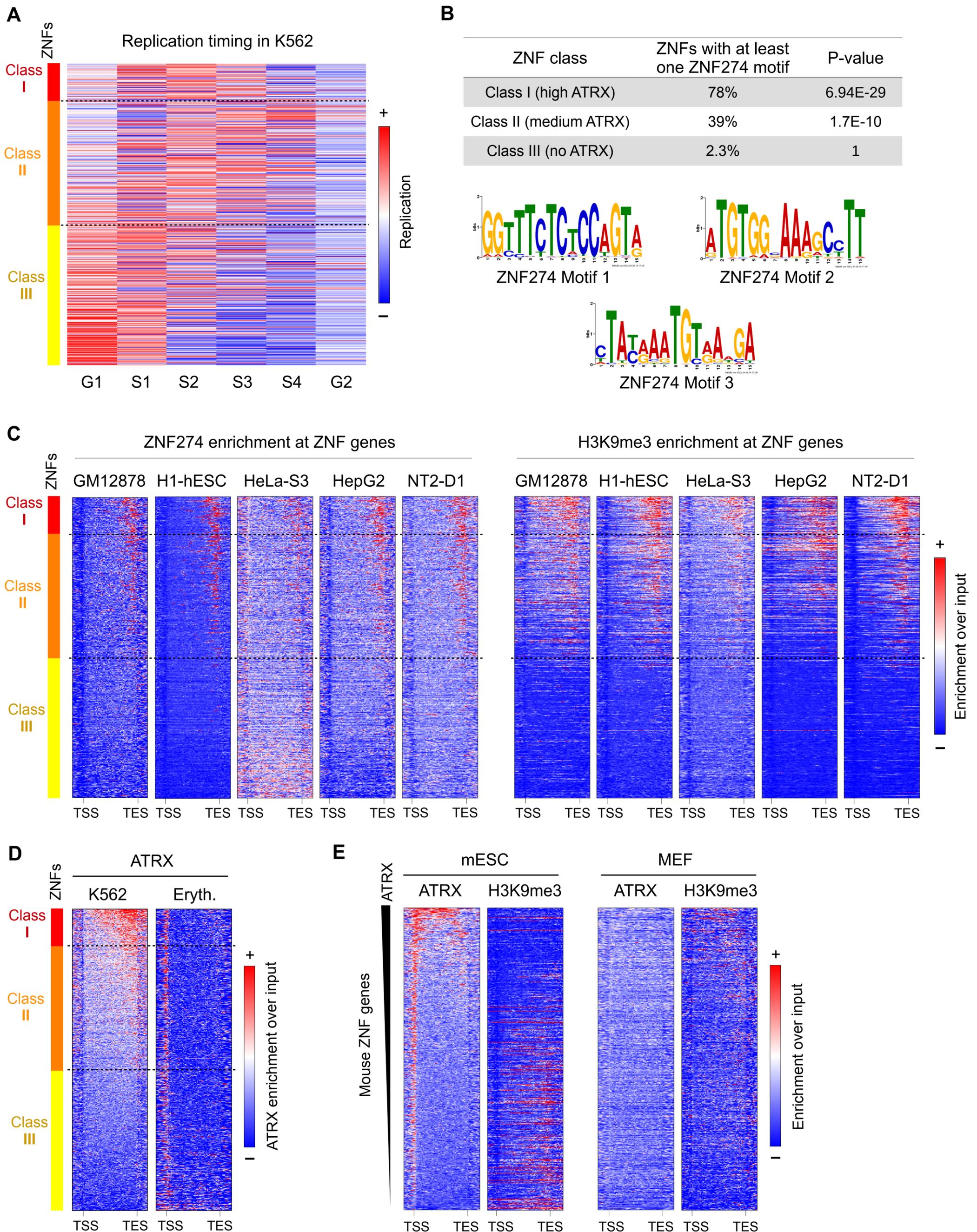


K562

Genes

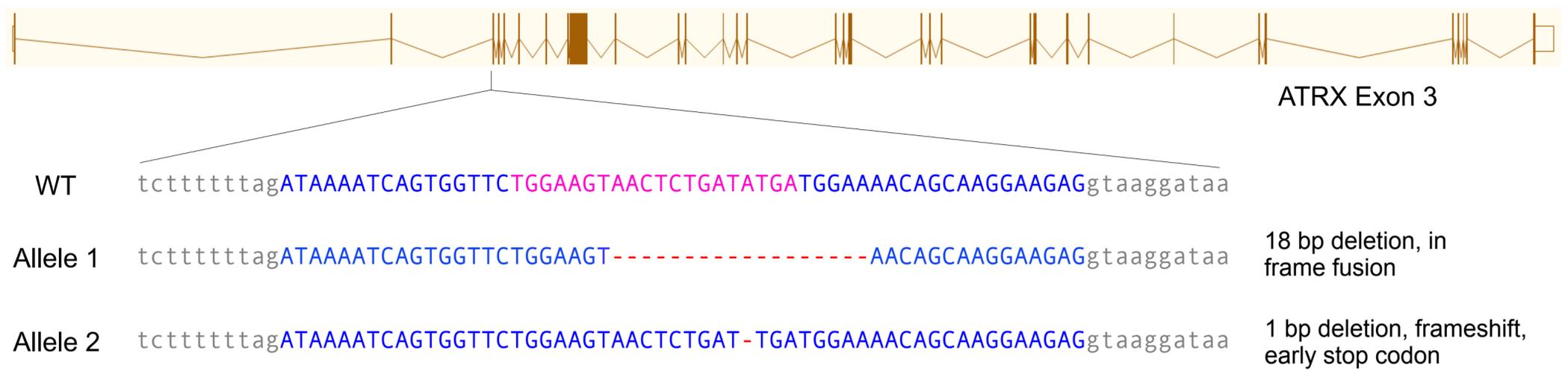
ATRAX  
SCATRAX  
Abcam



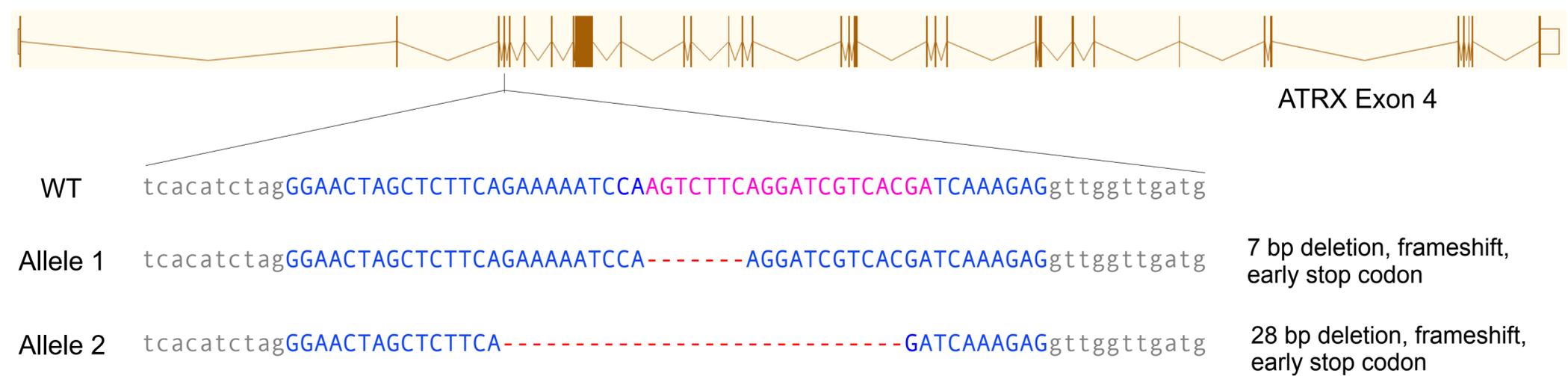
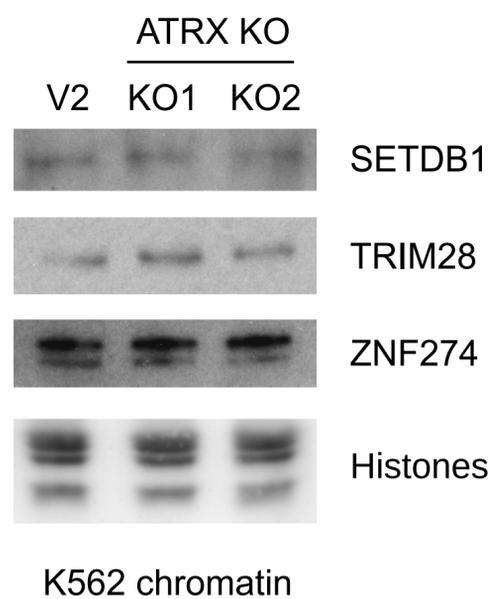


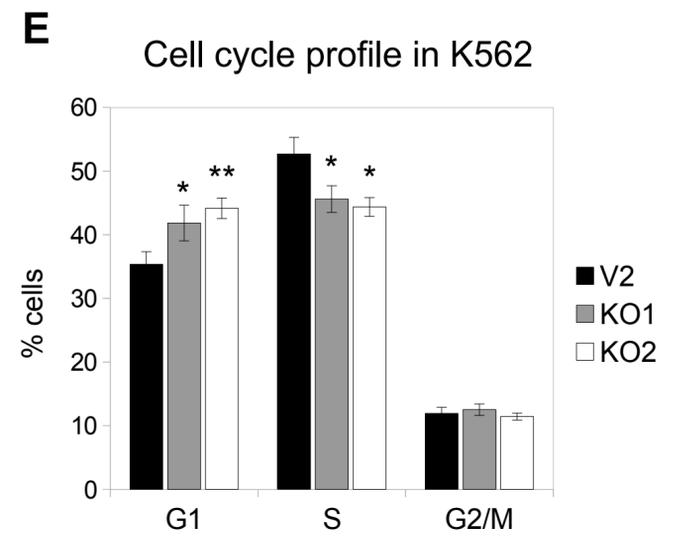
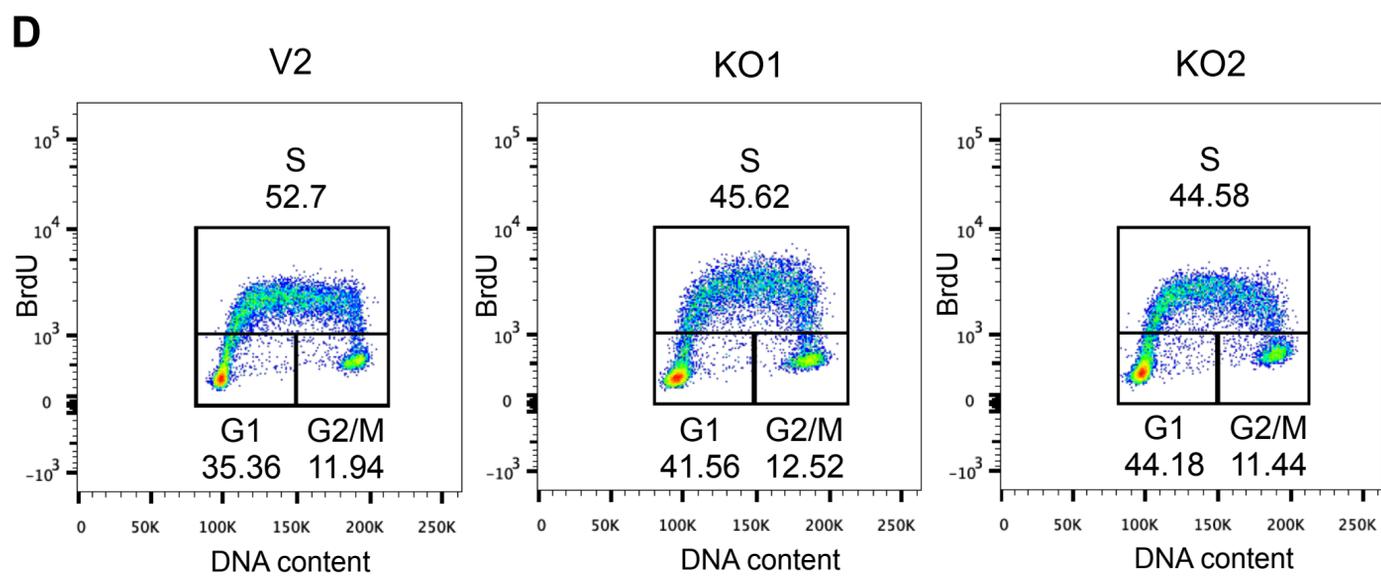
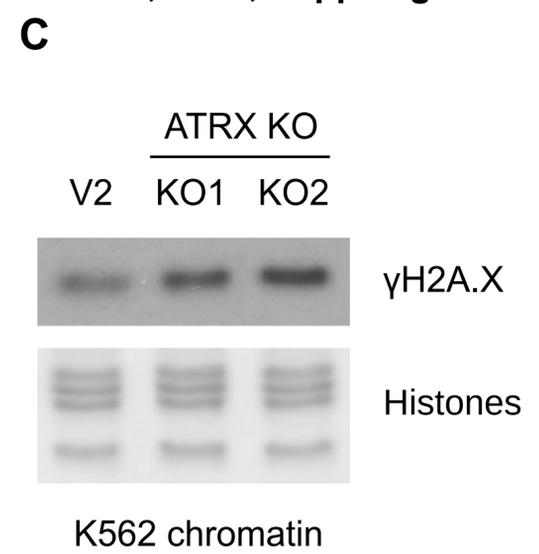
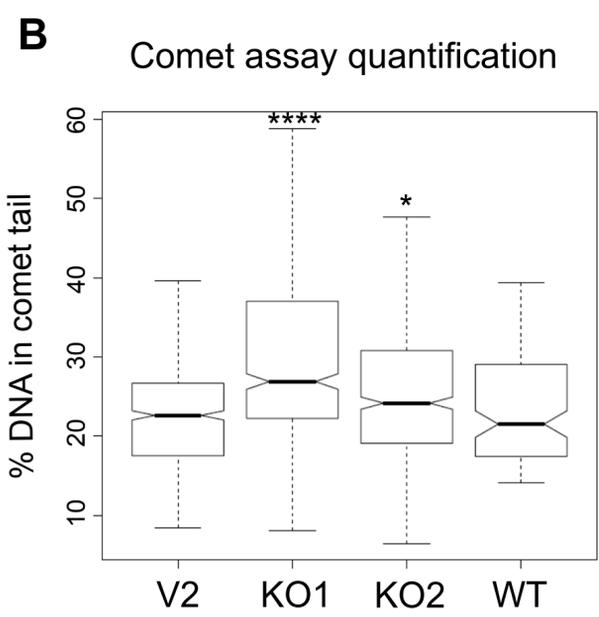
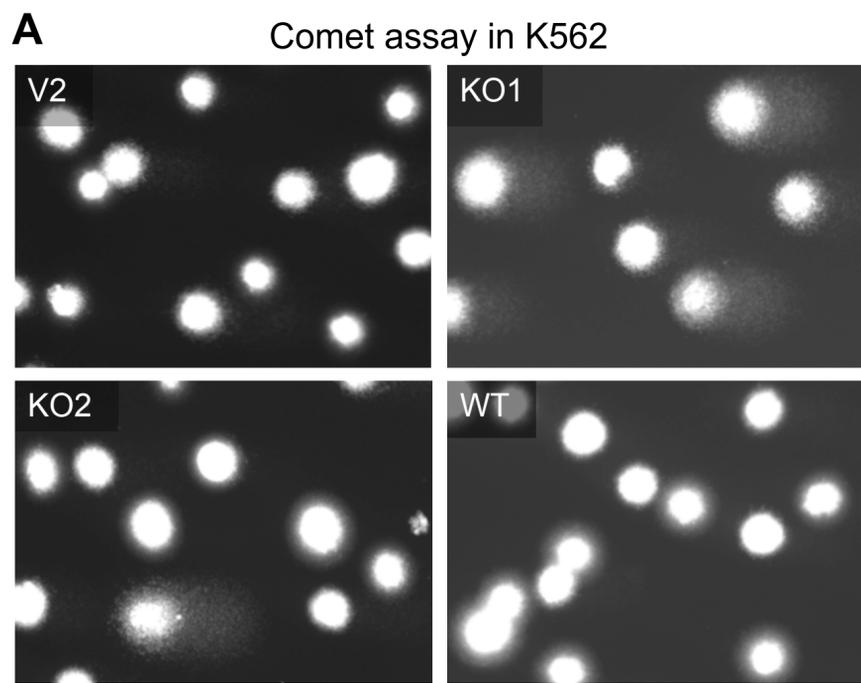
**A**

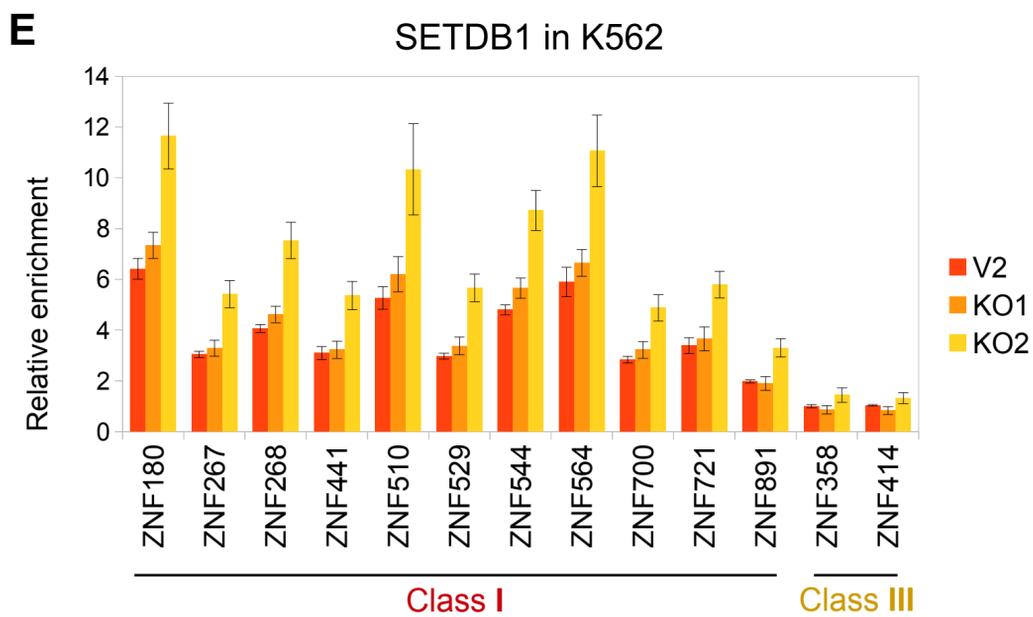
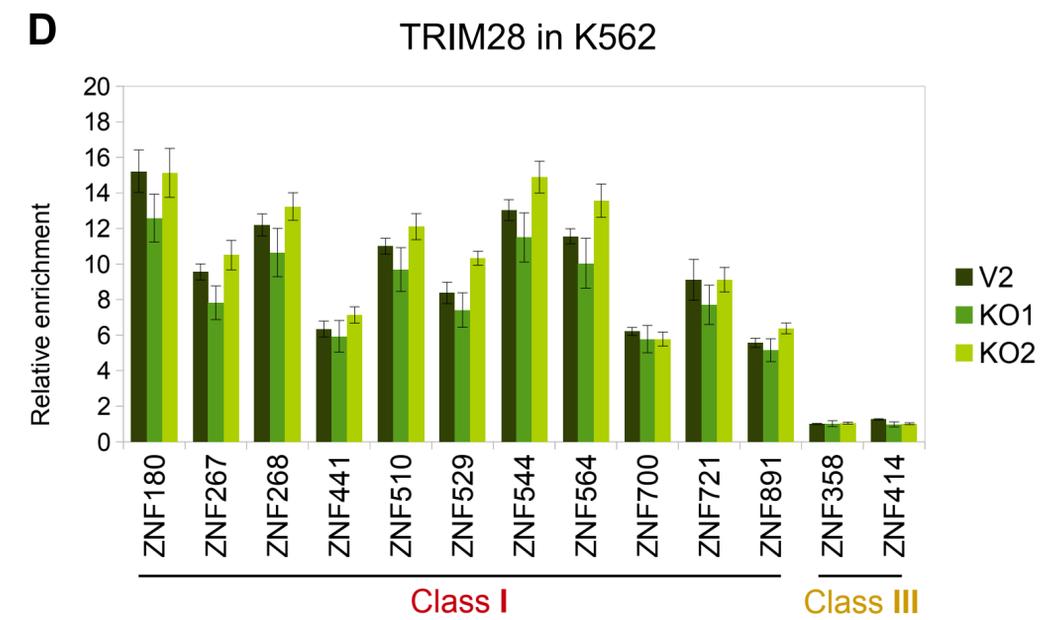
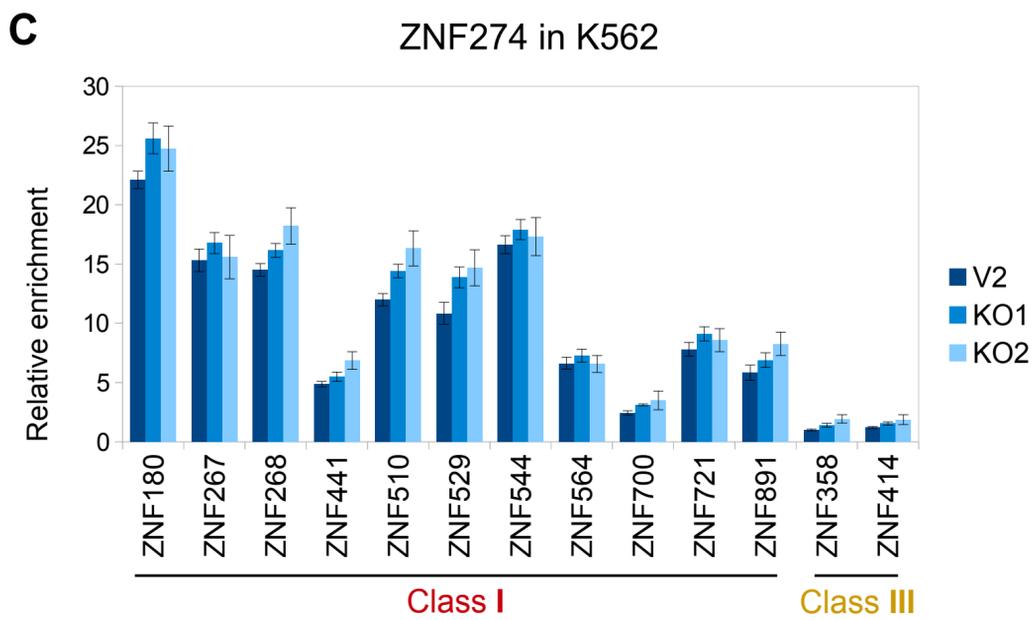
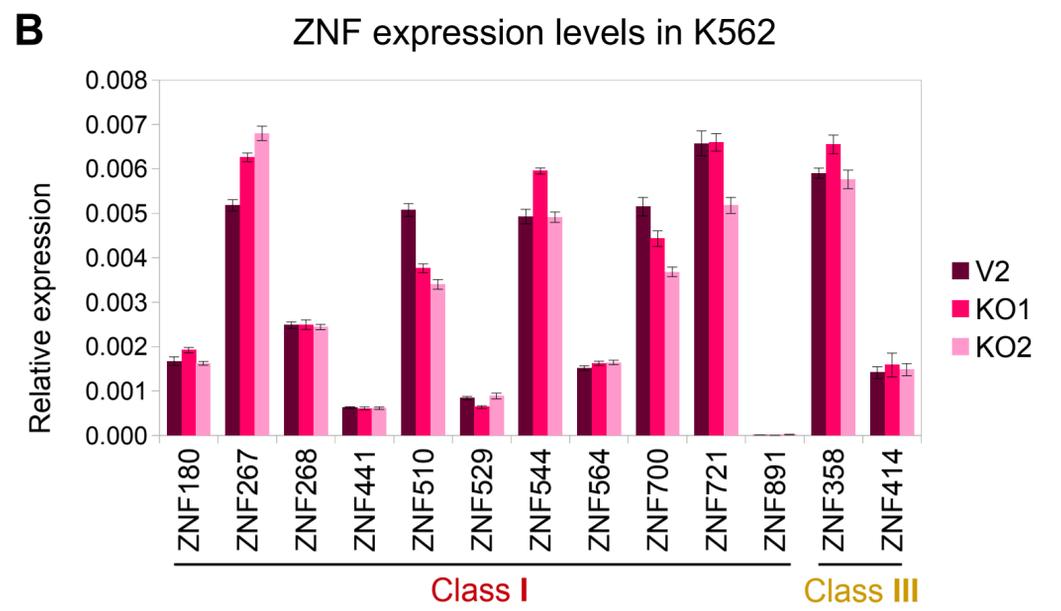
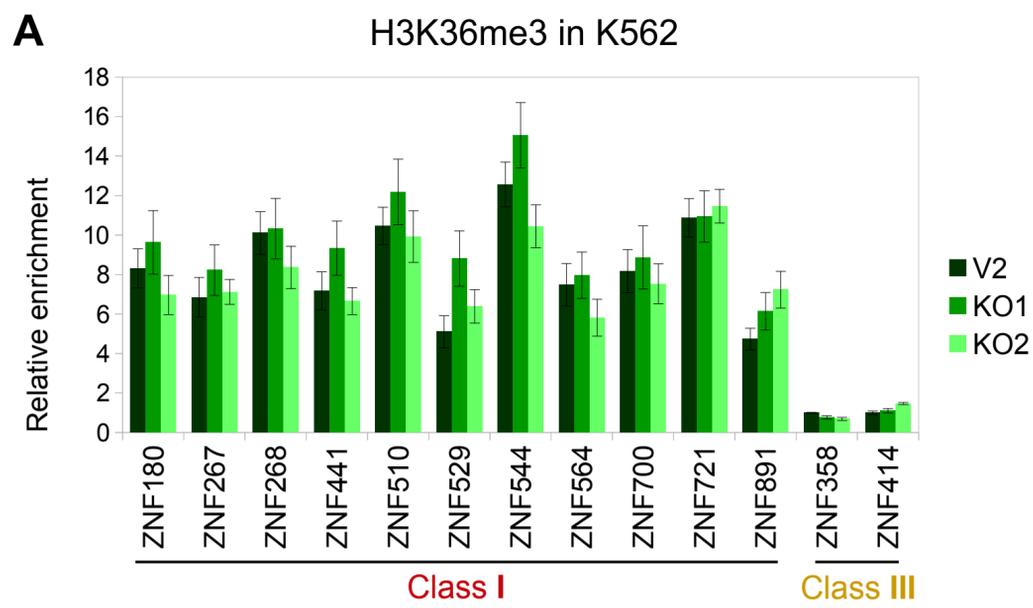
ATRX KO1, K562

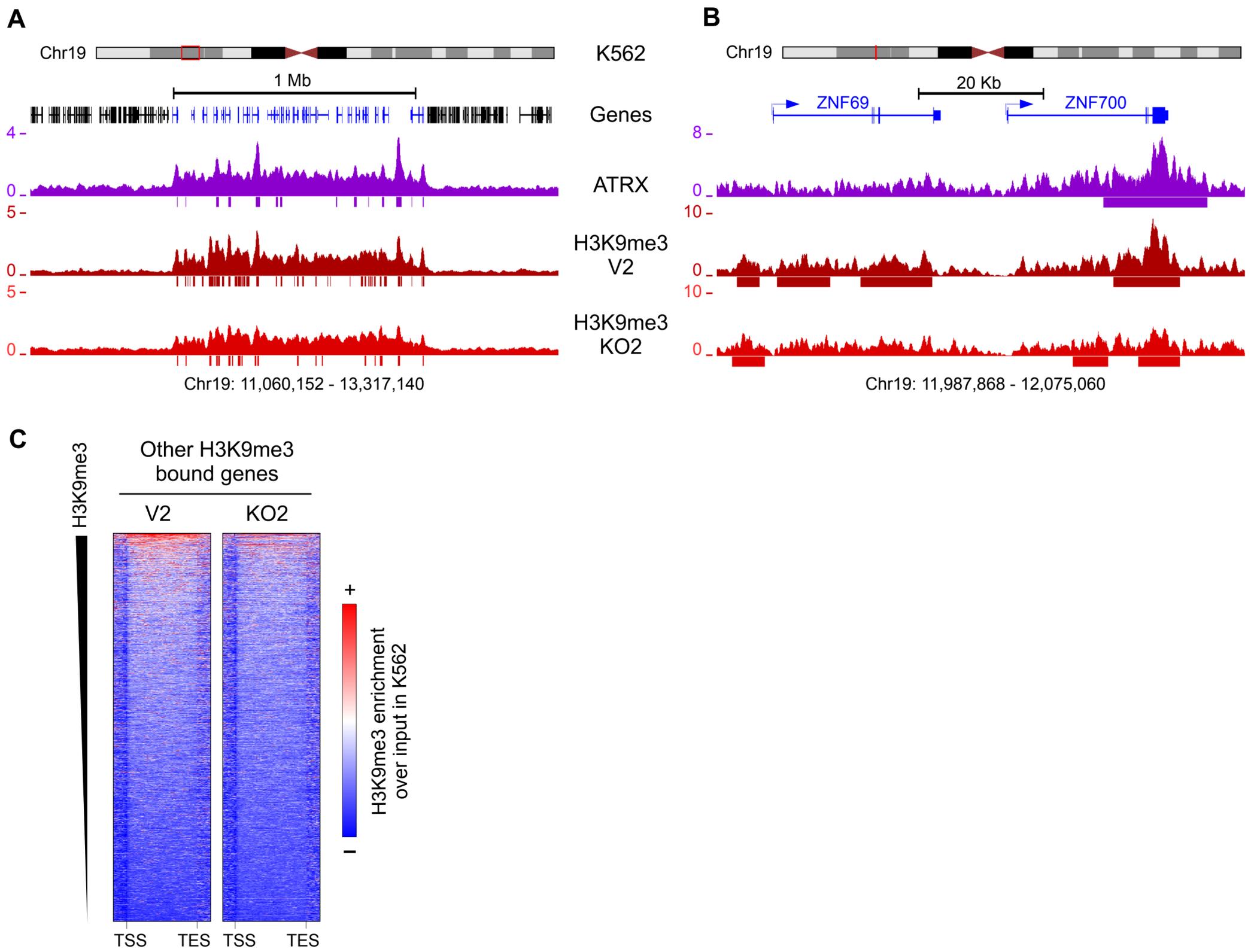


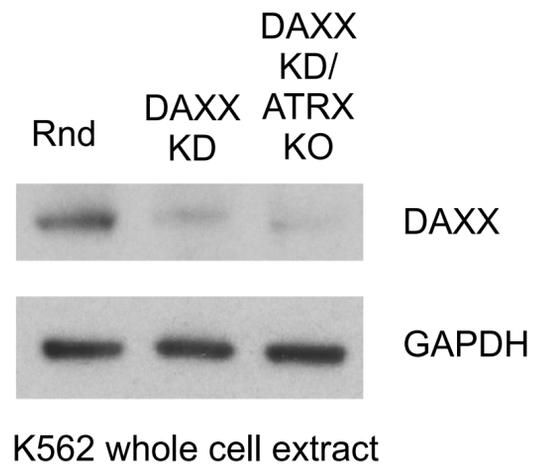
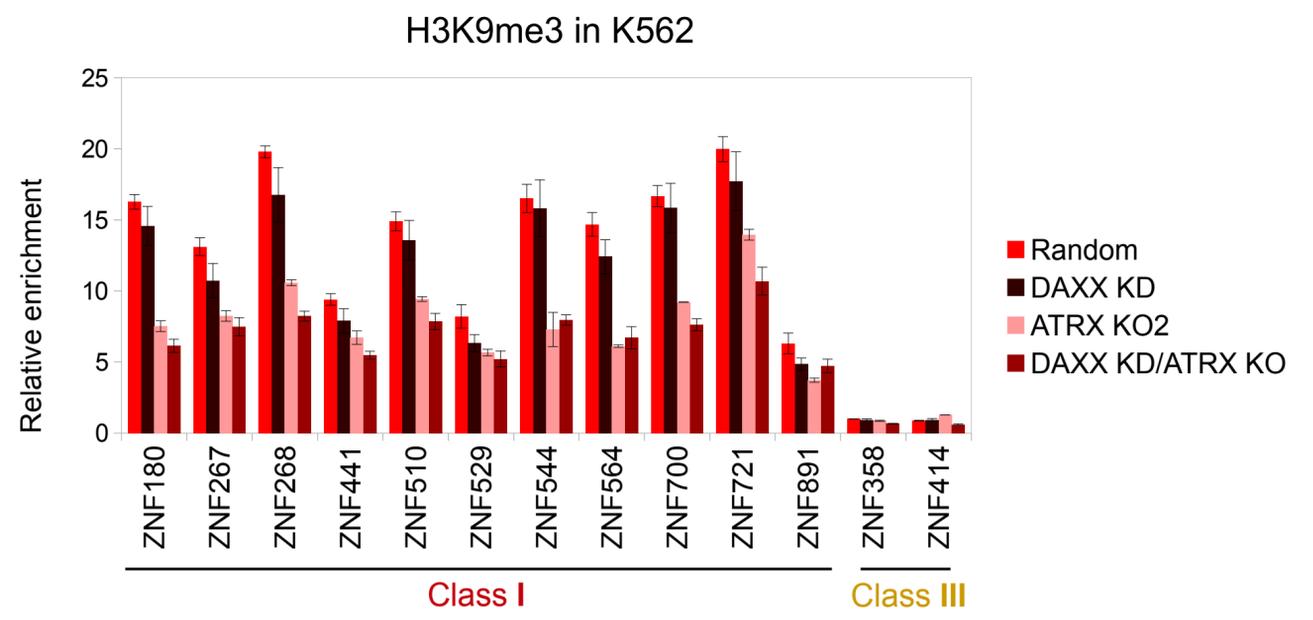
ATRX KO2, K562

**B**

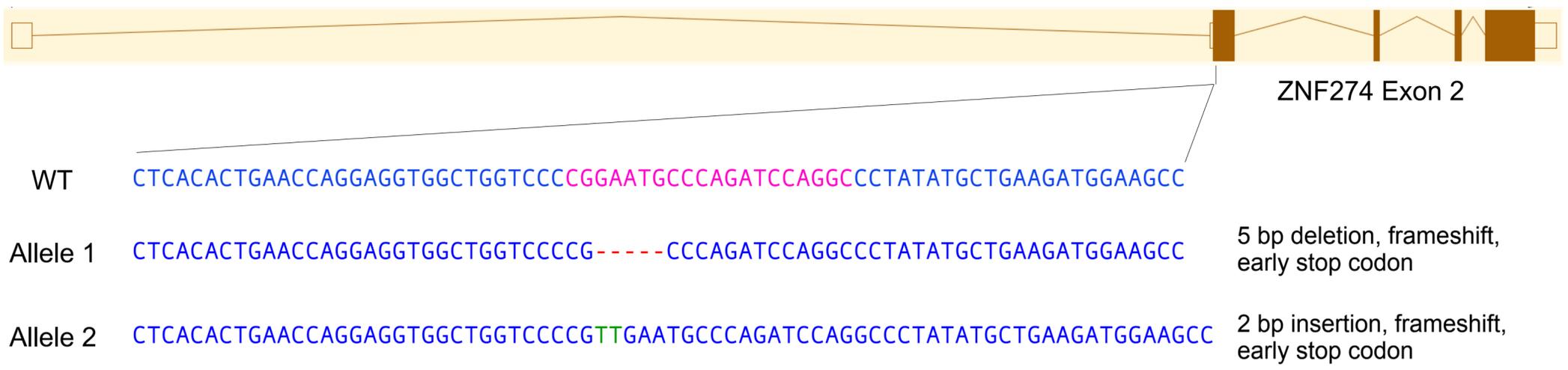






**A****B**

Z274 KO, K562



Z274/ATRX KO, K562

