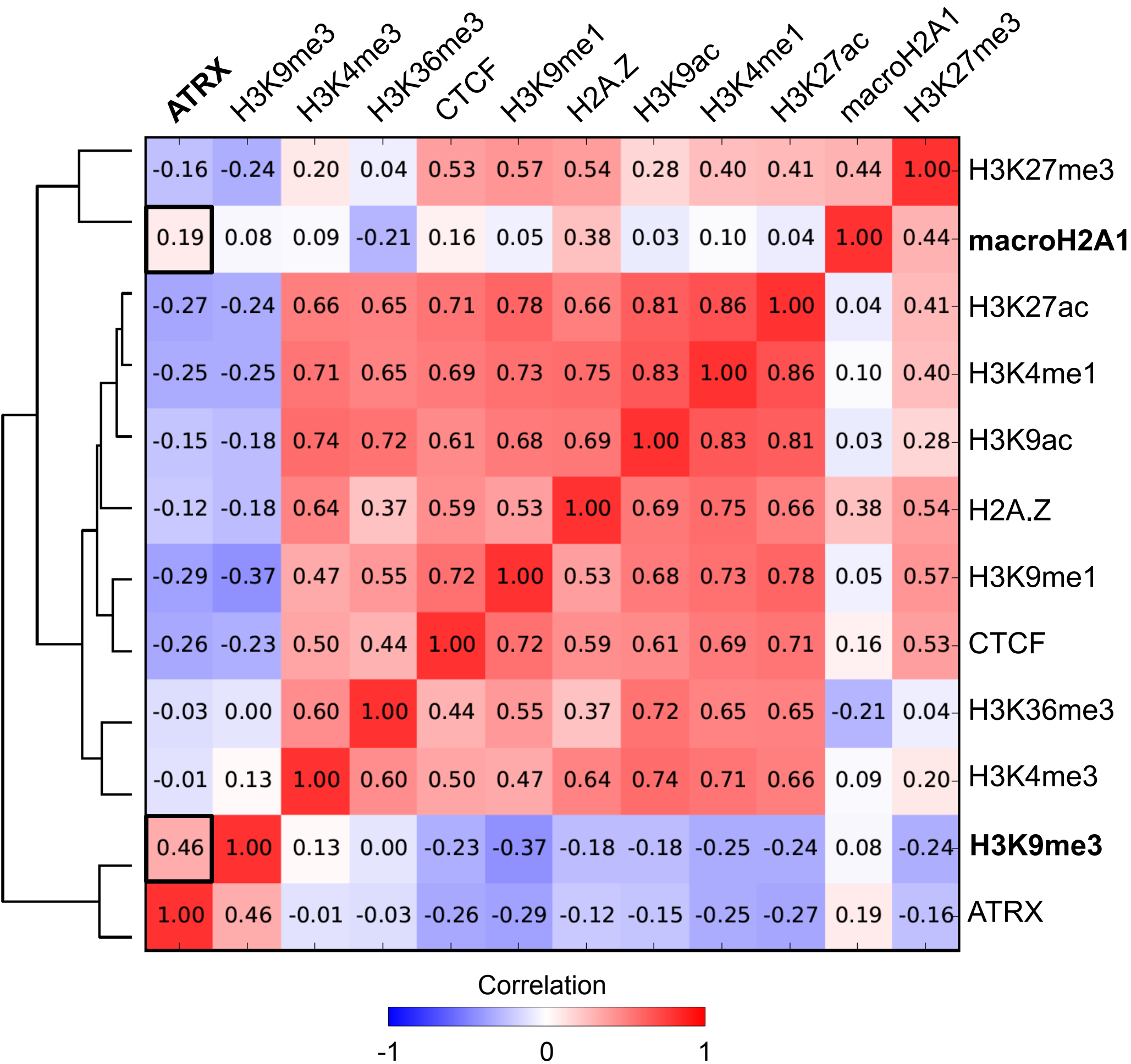


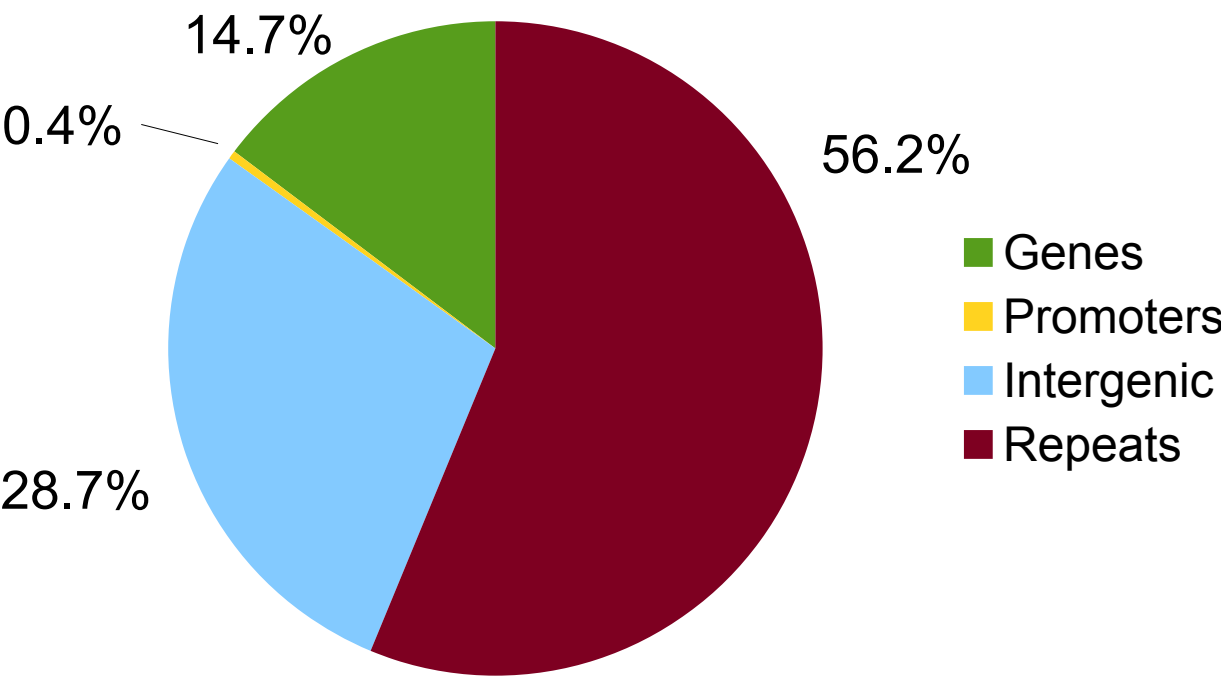
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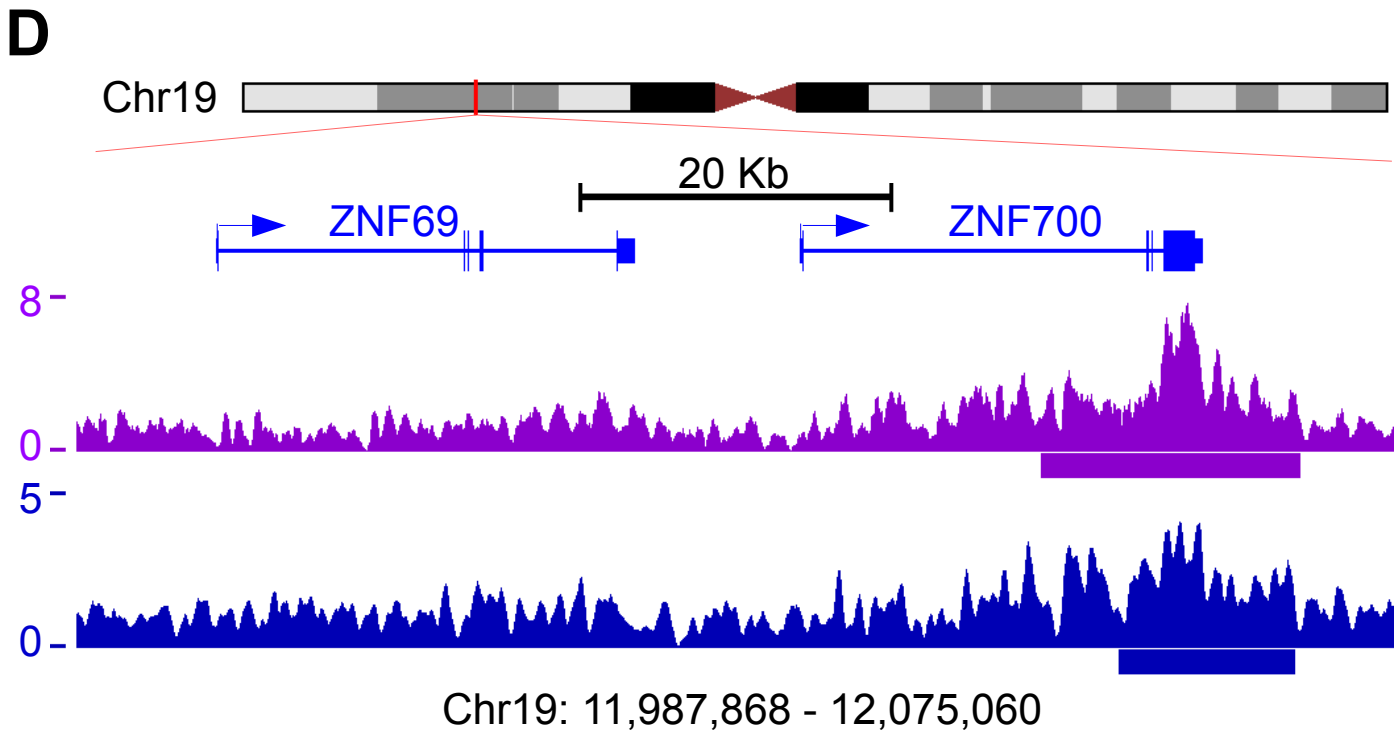
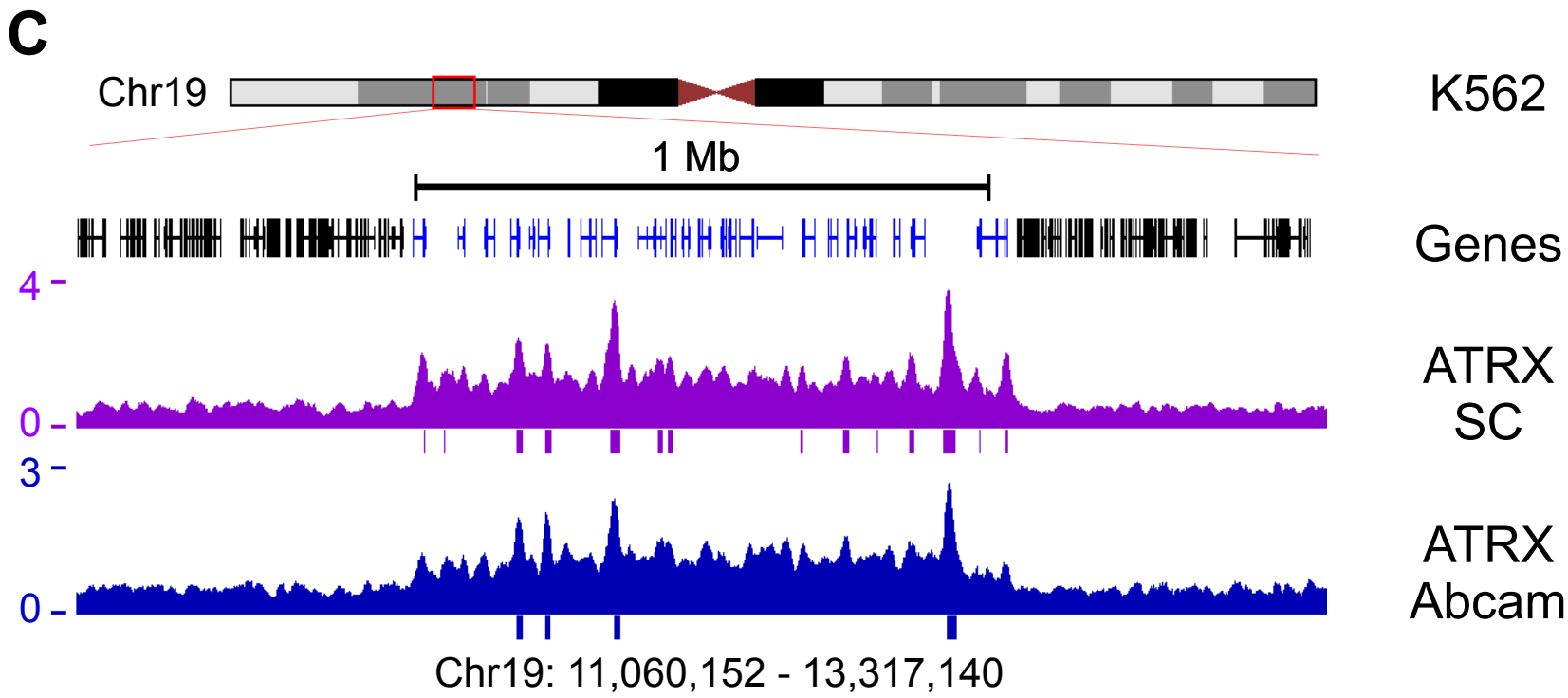
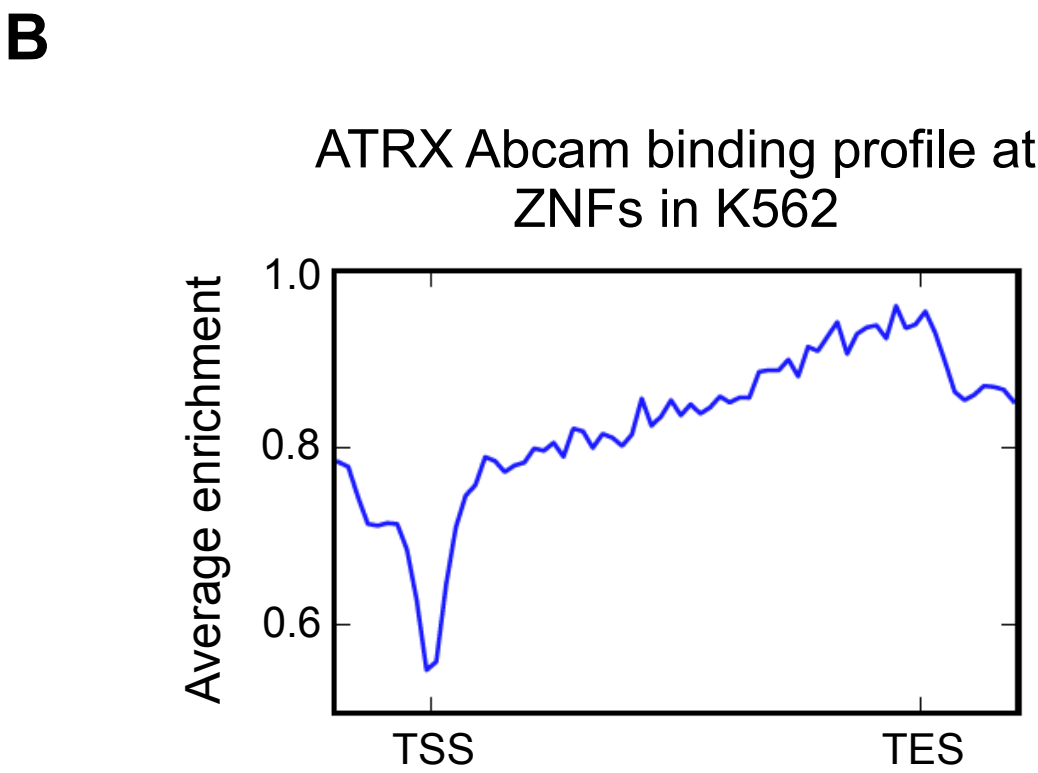
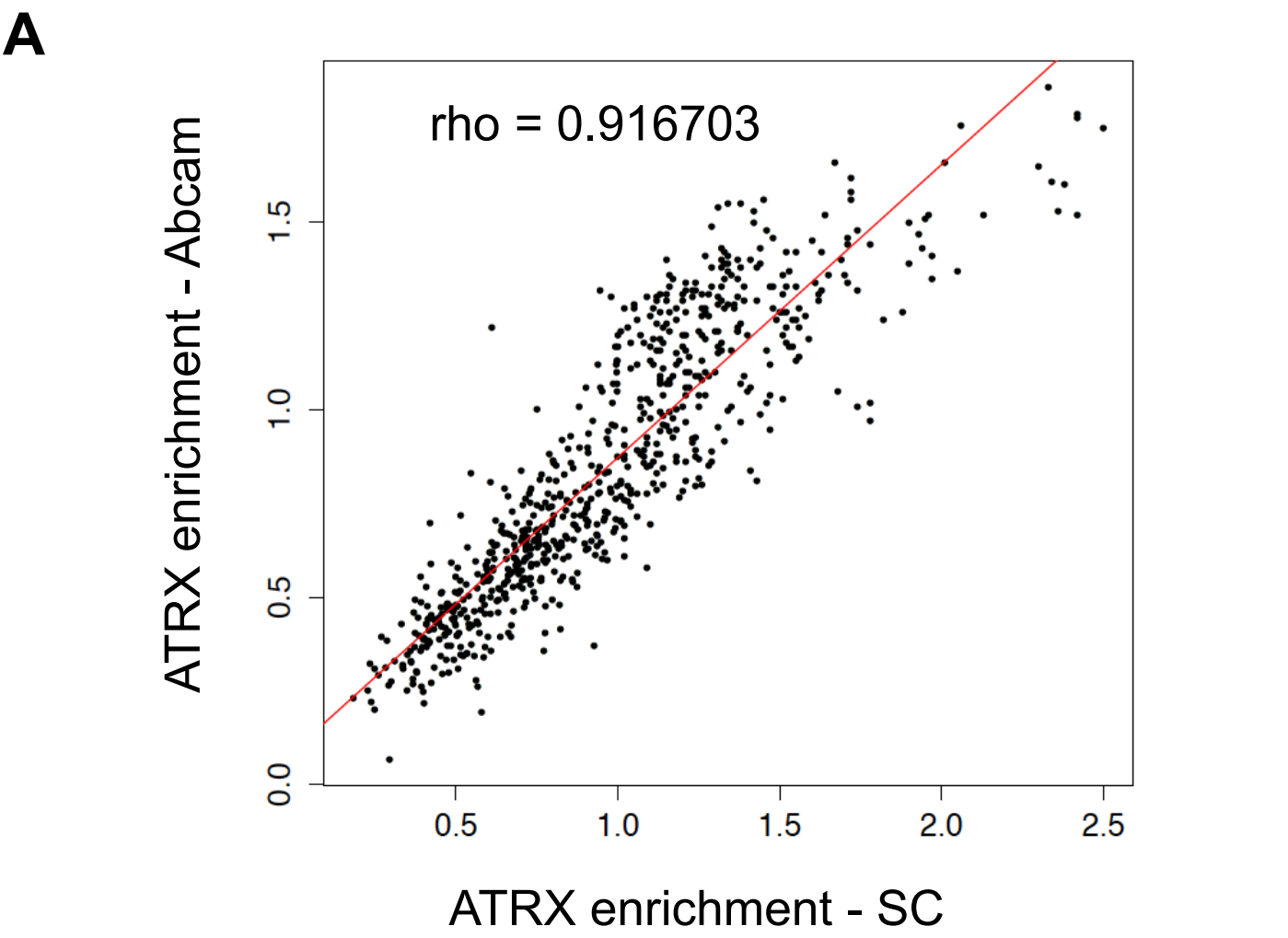
Genome wide correlation of ChIP-seq profiles in K562

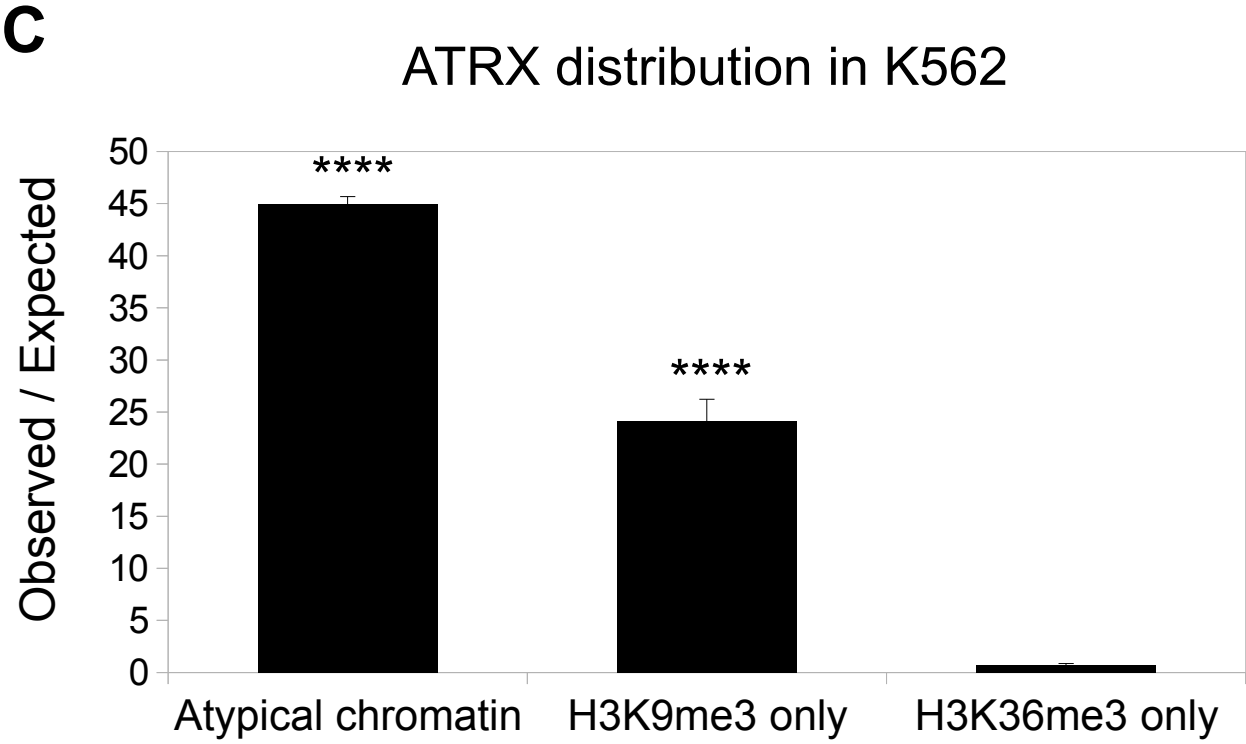
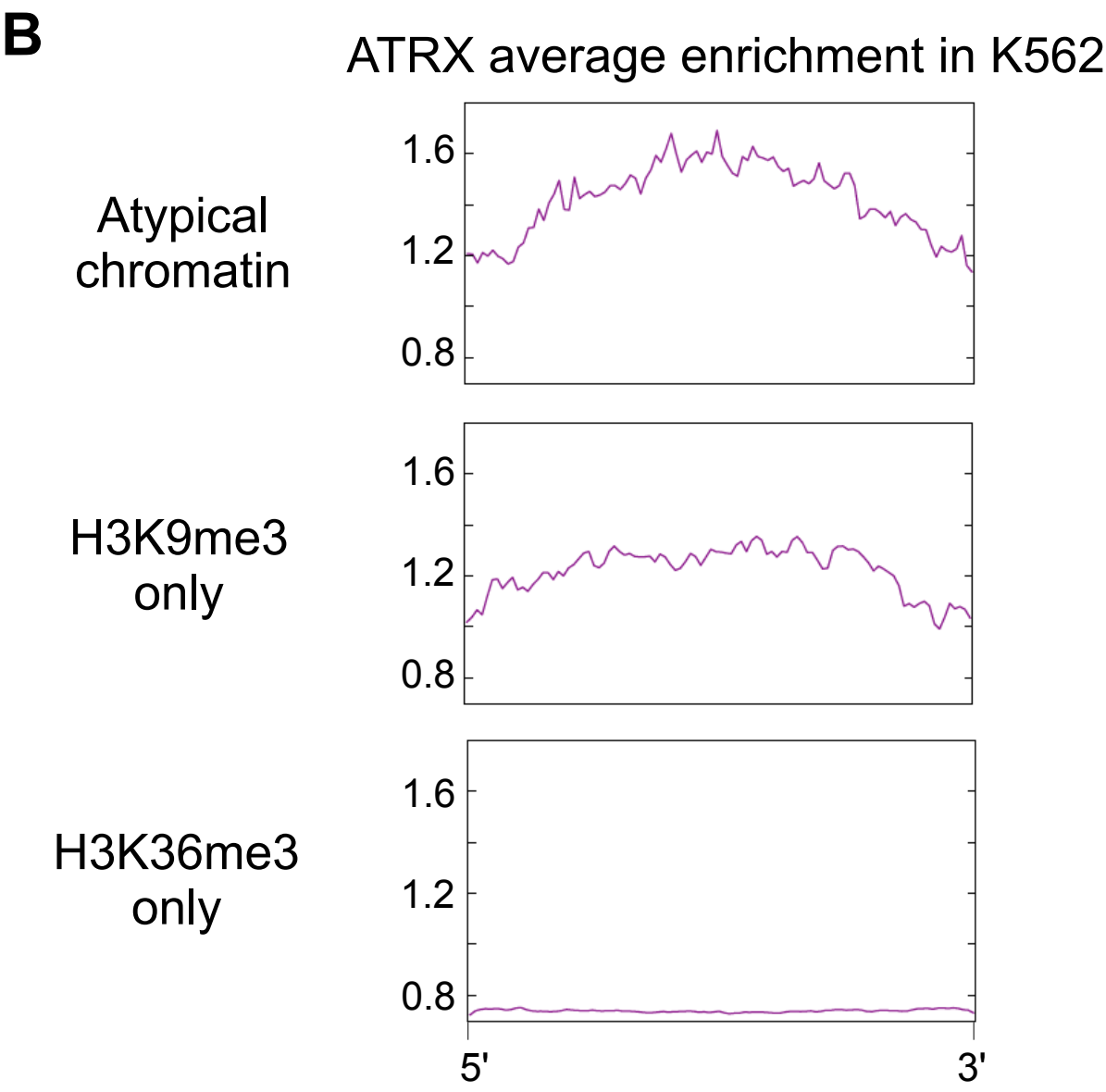
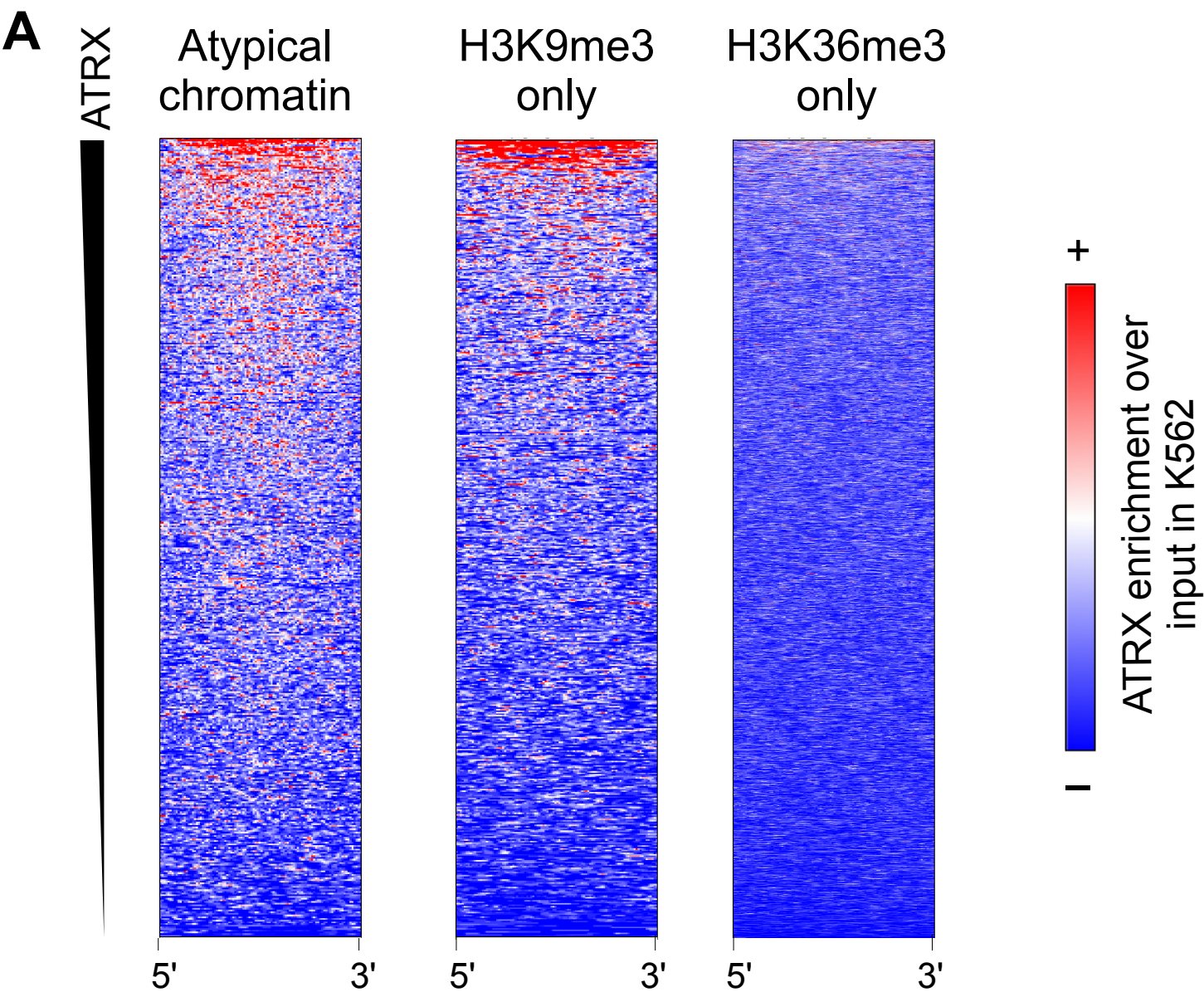


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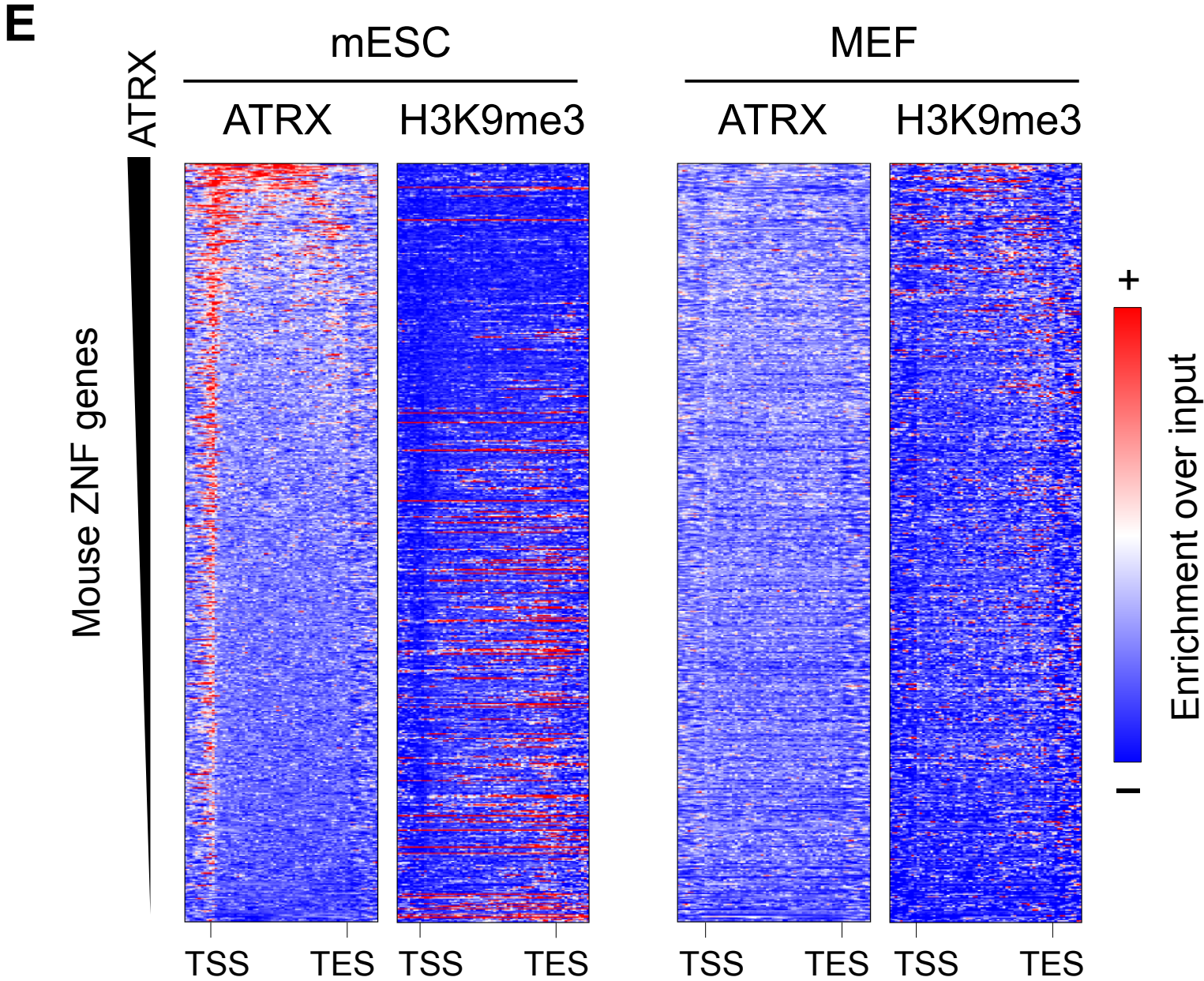
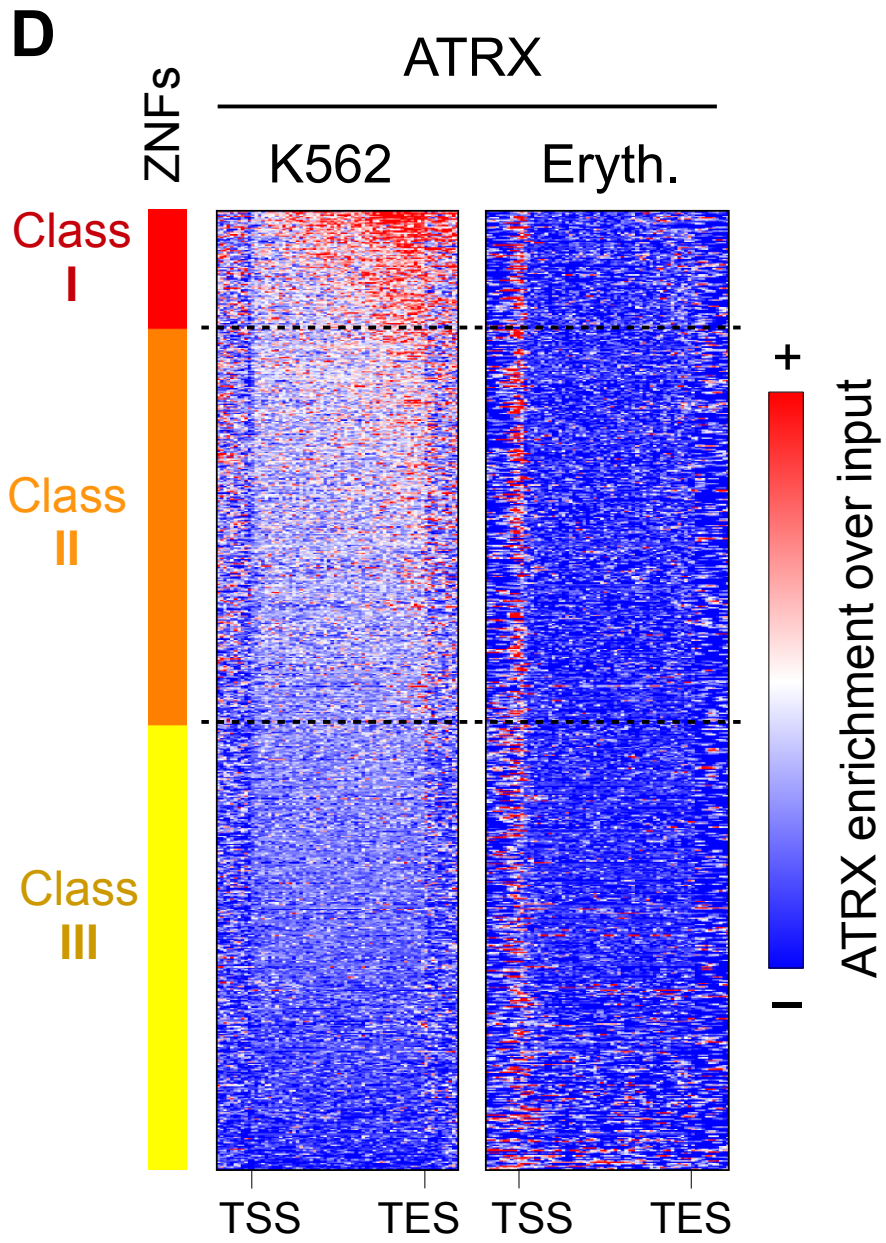
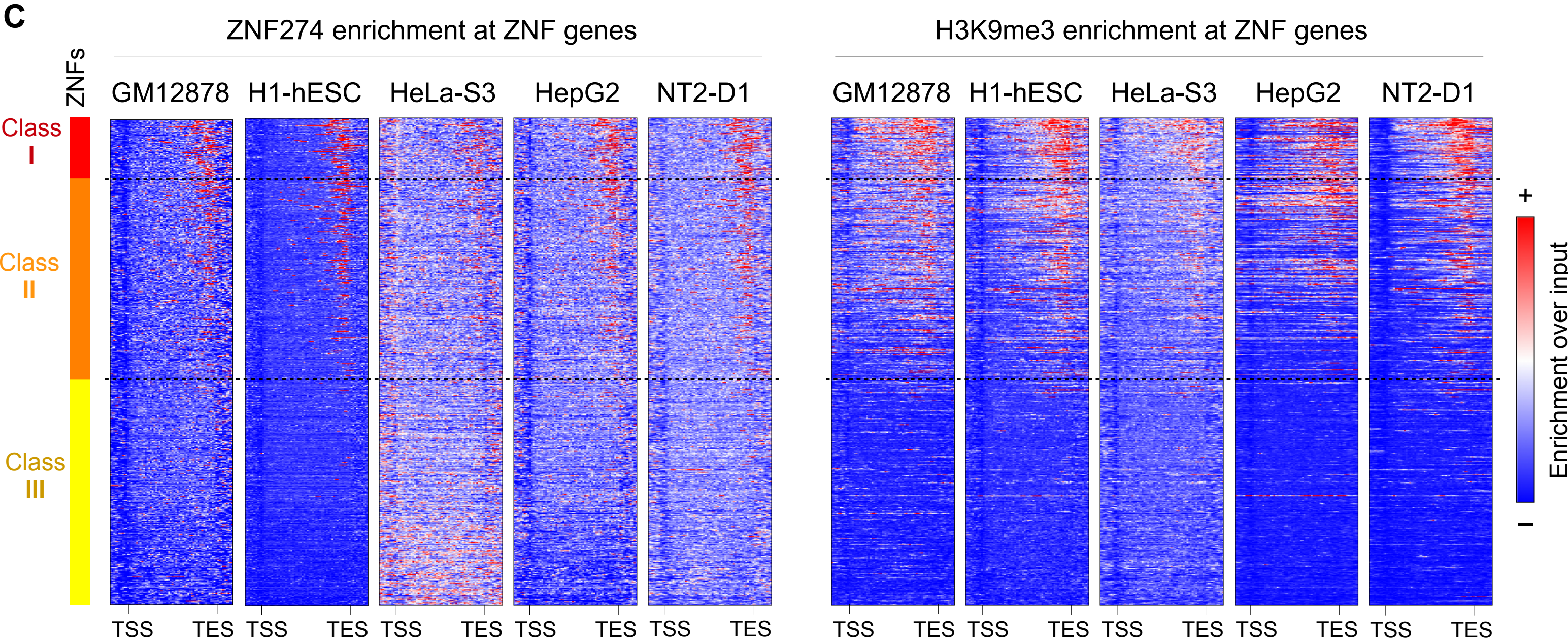
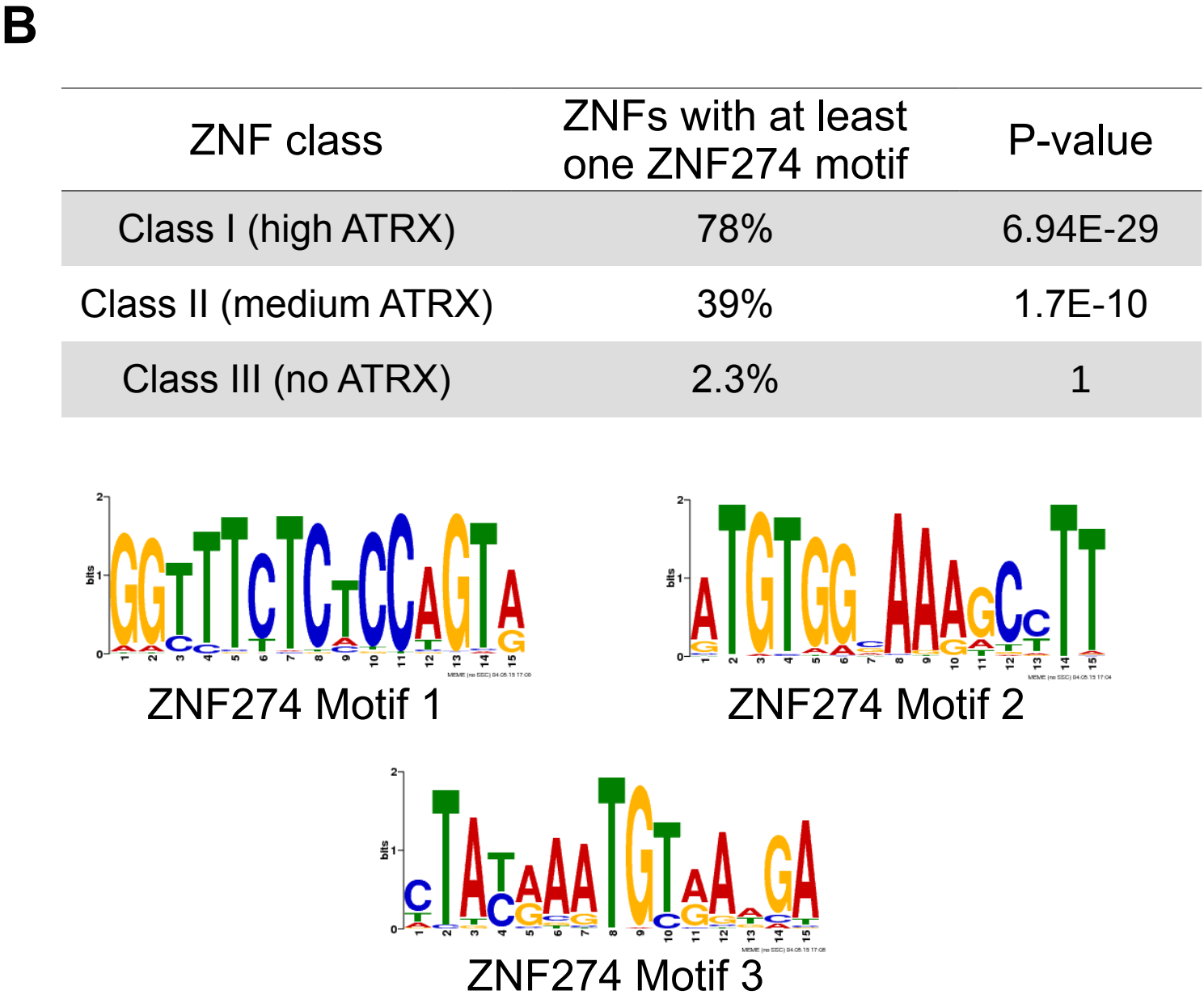
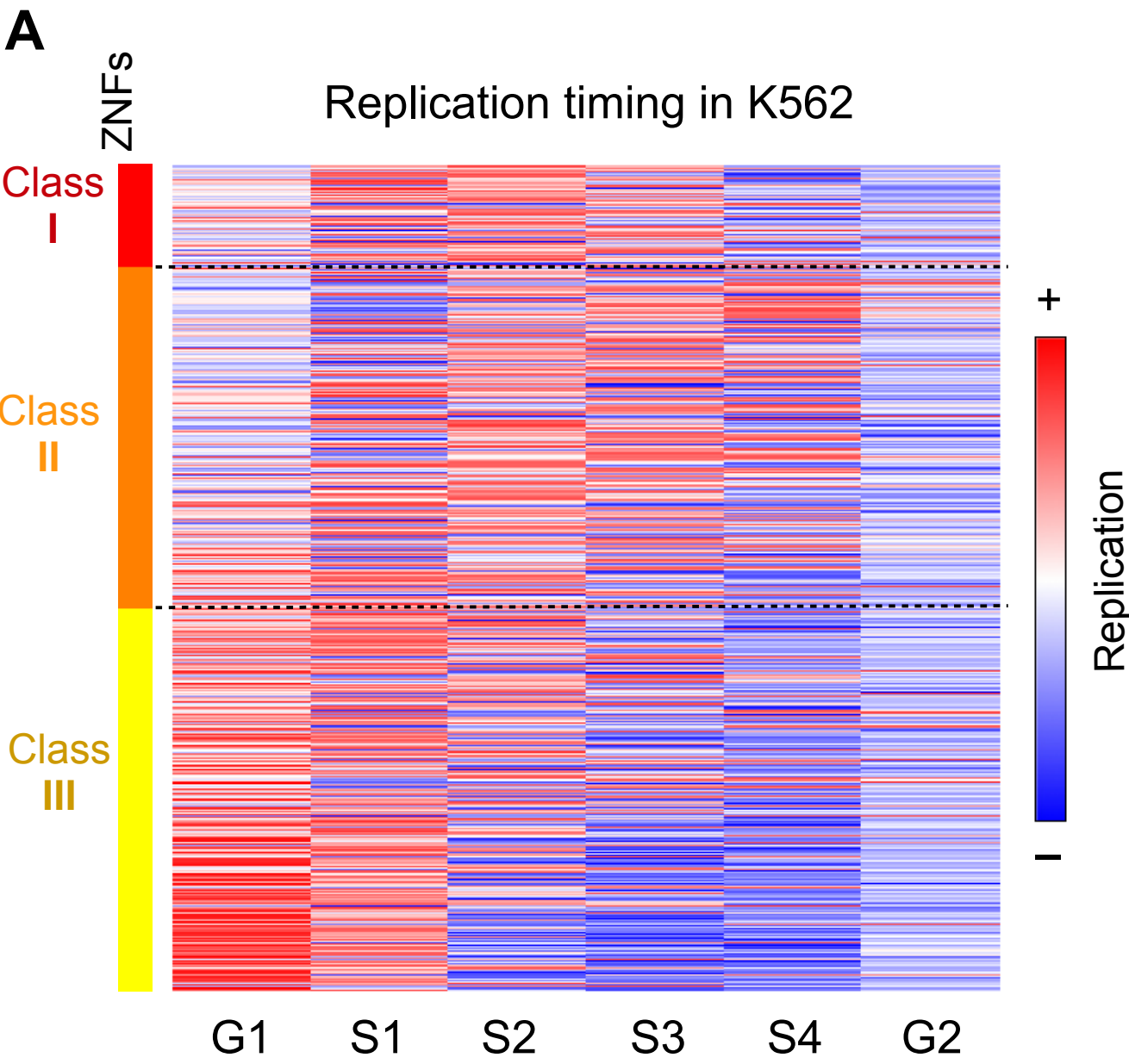
ATRX peak distribution in K562







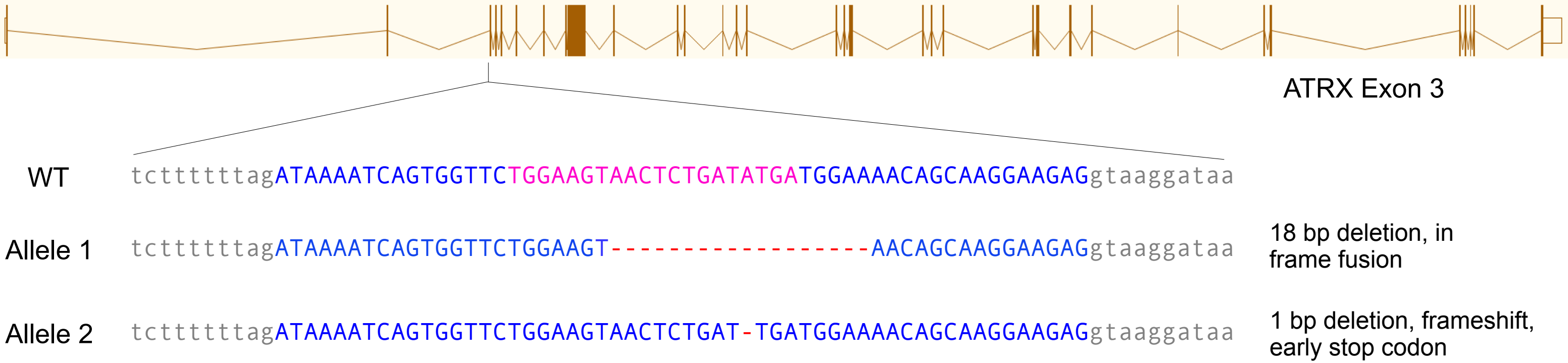




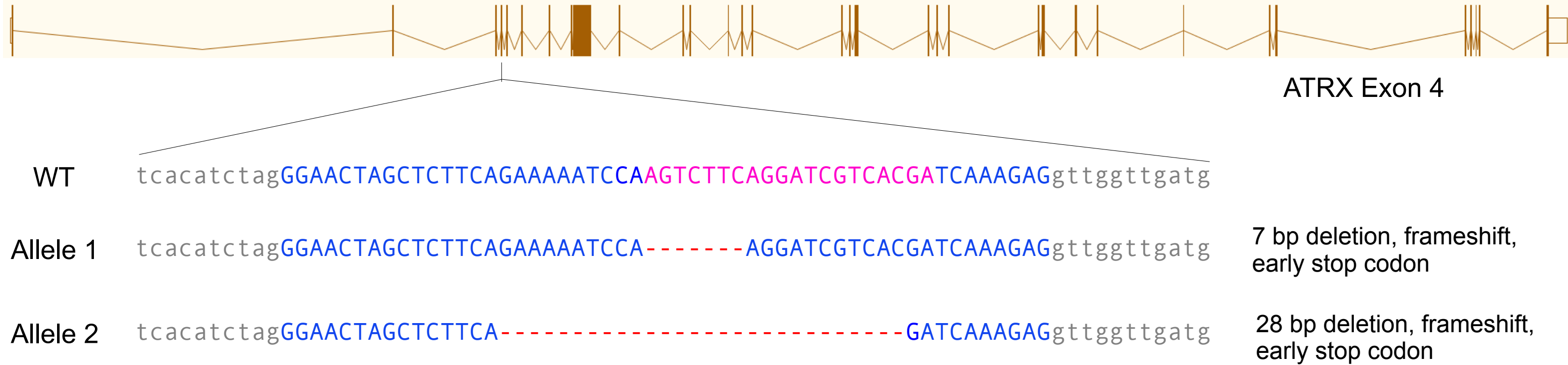


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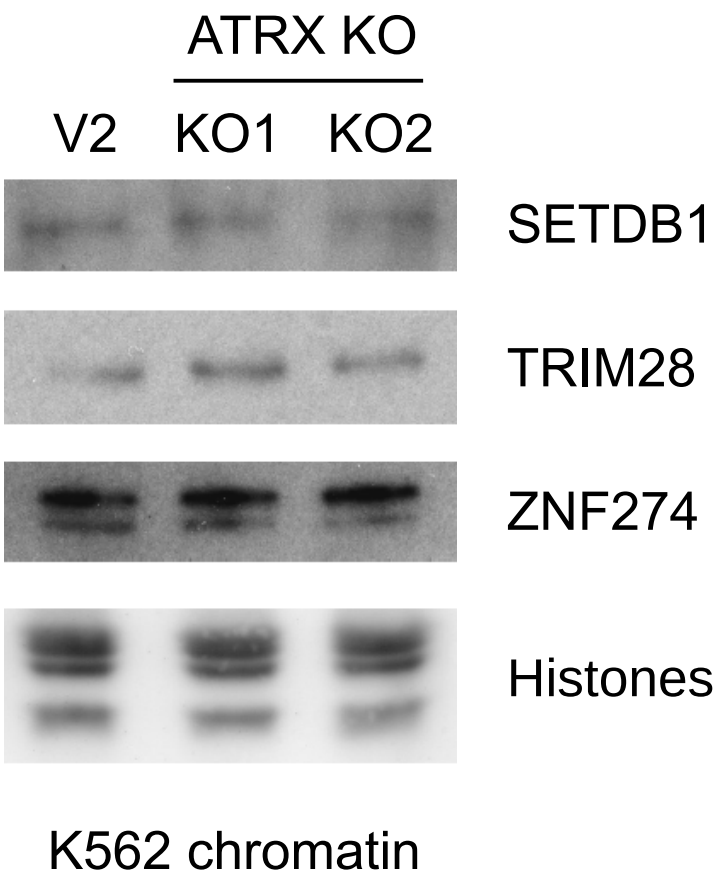
ATRX KO1, K562

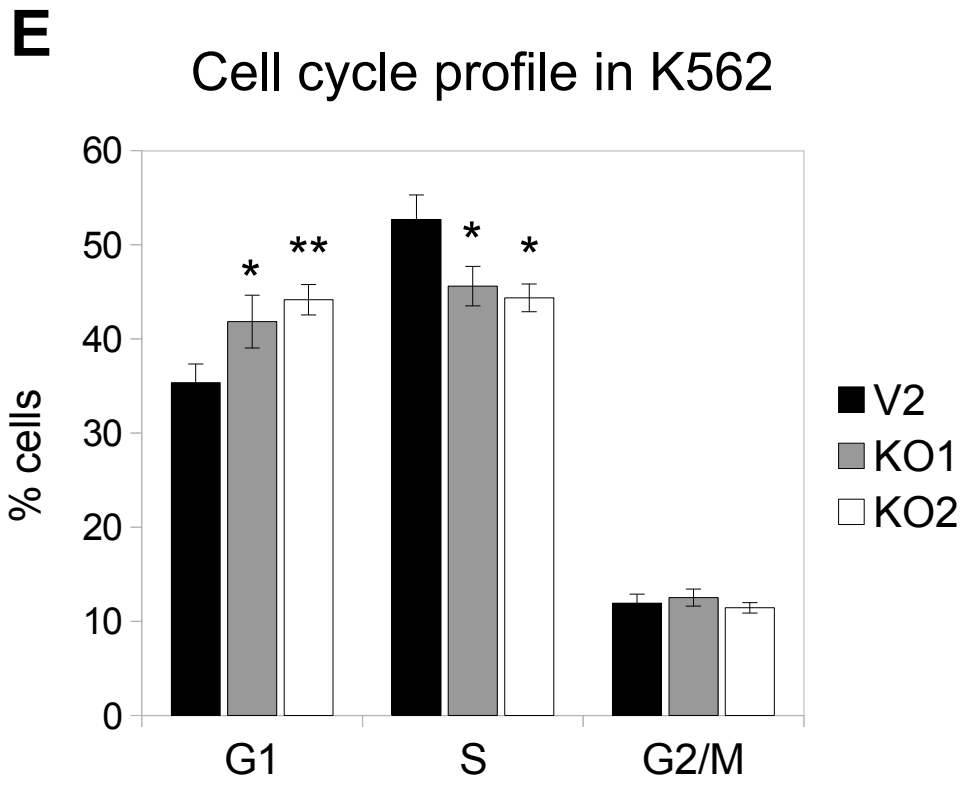
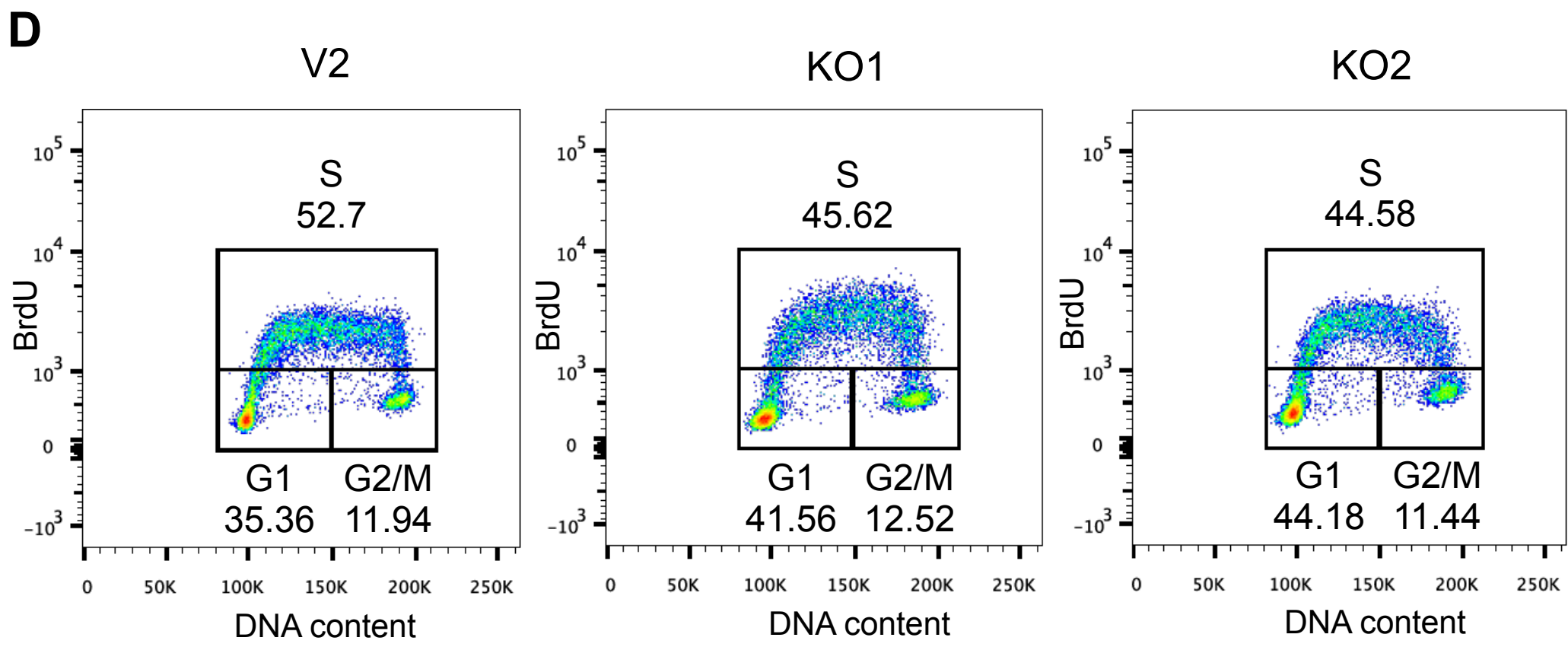
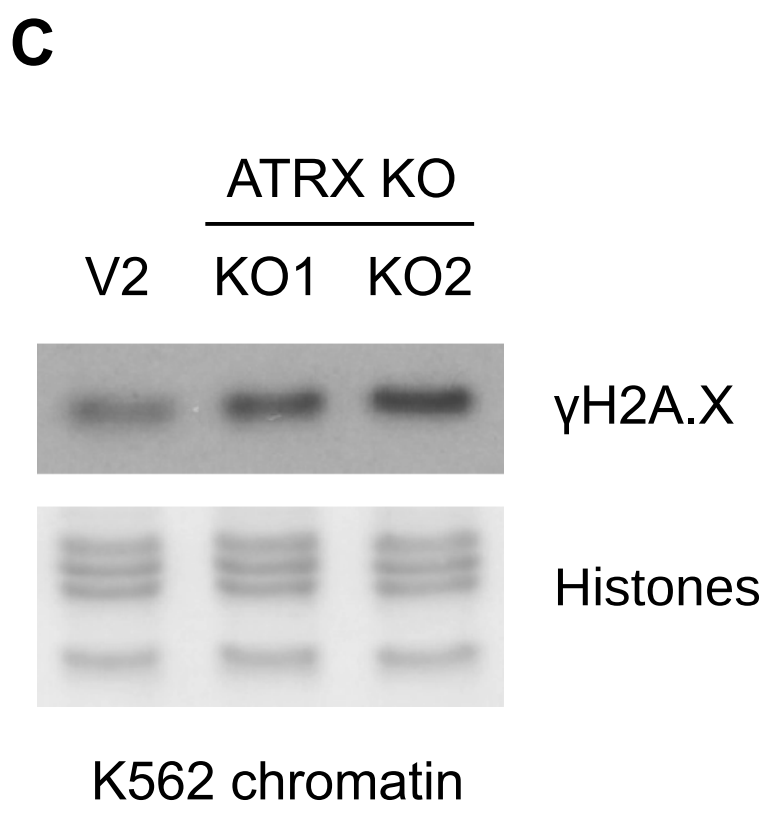
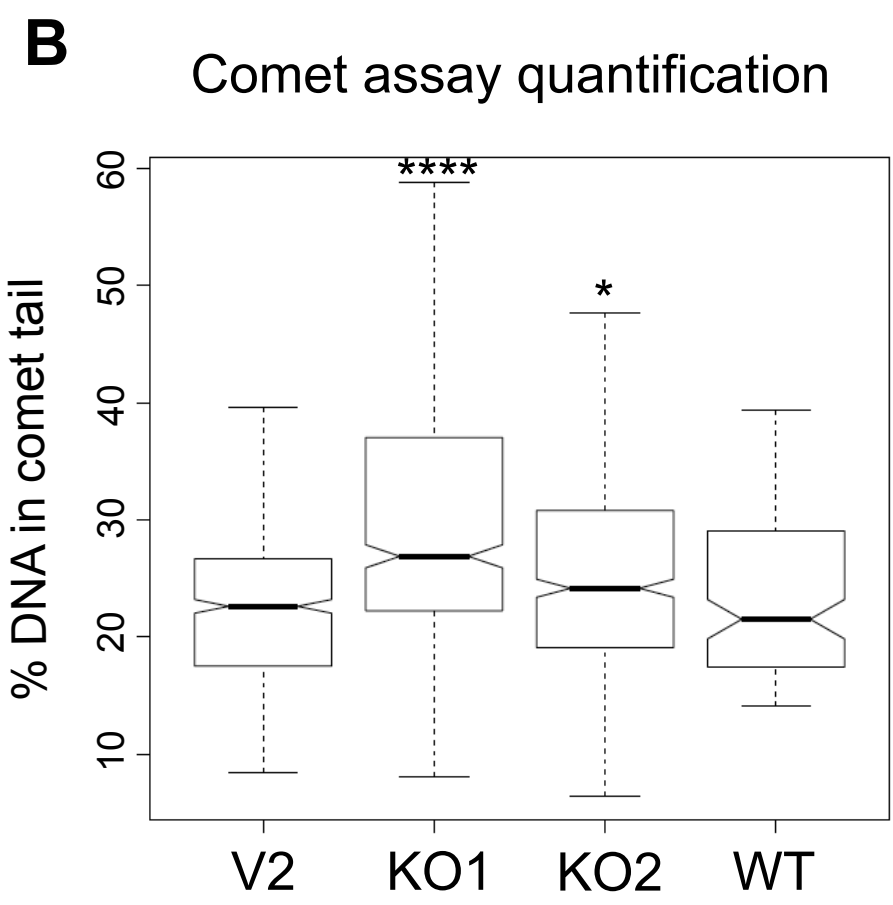
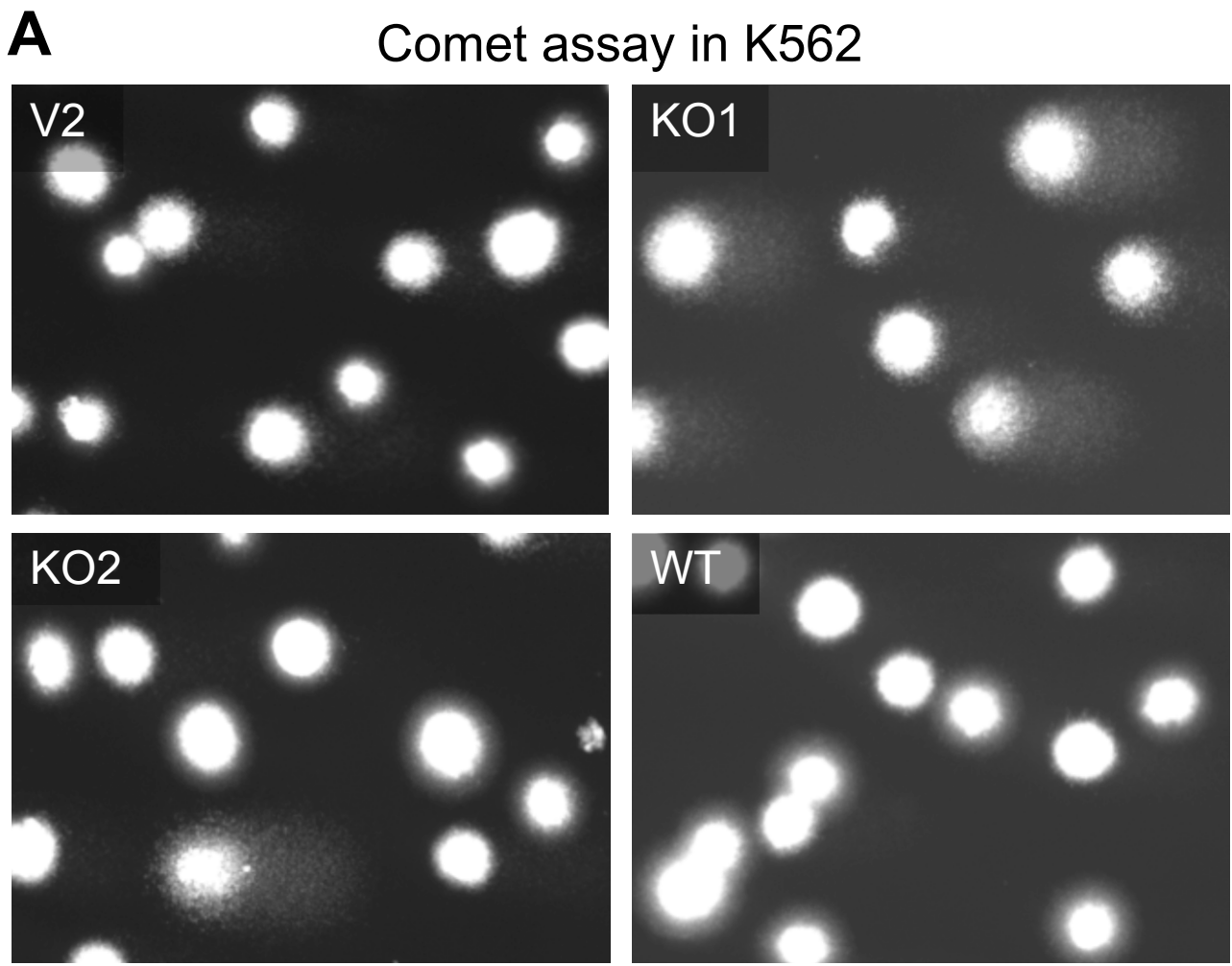


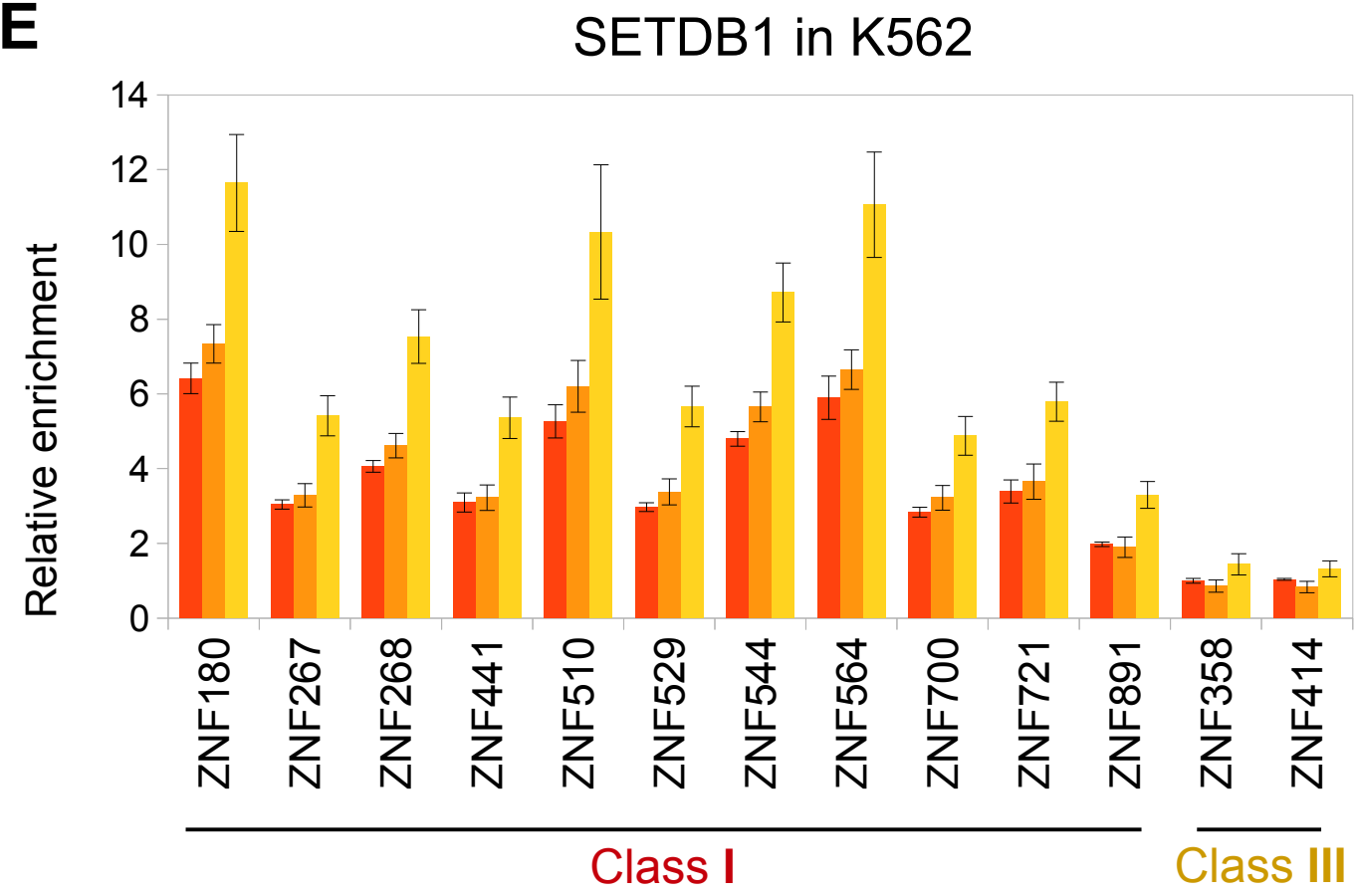
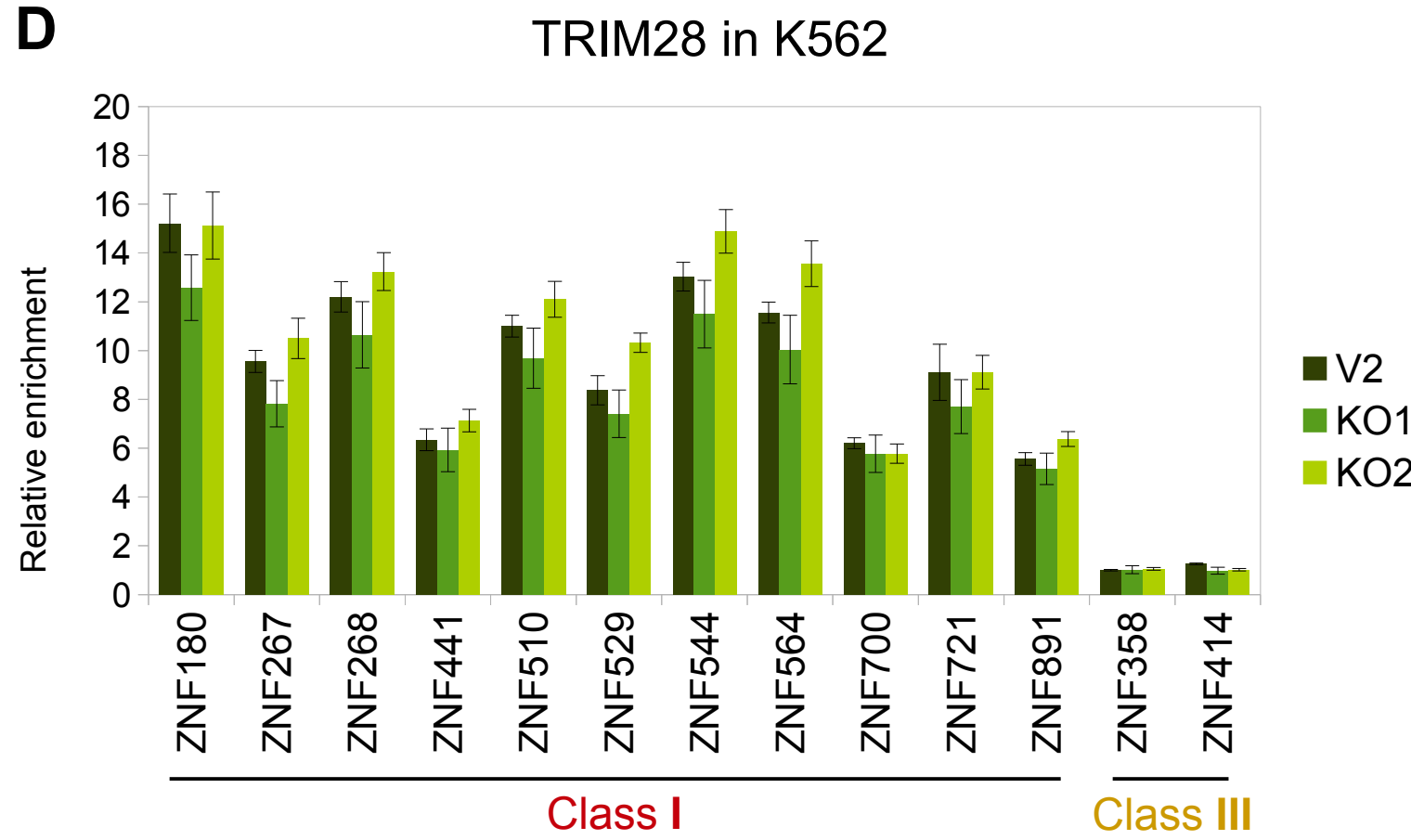
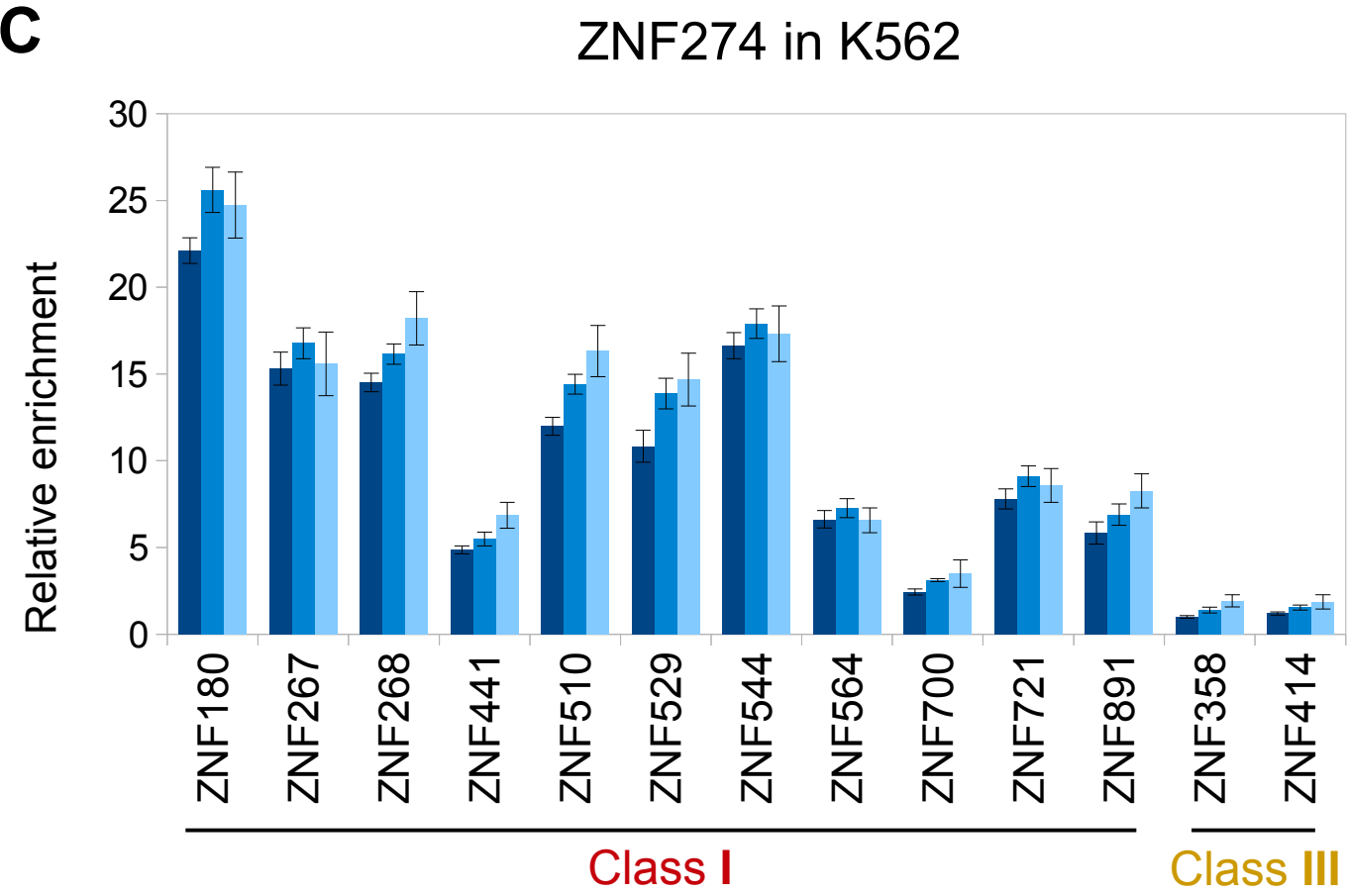
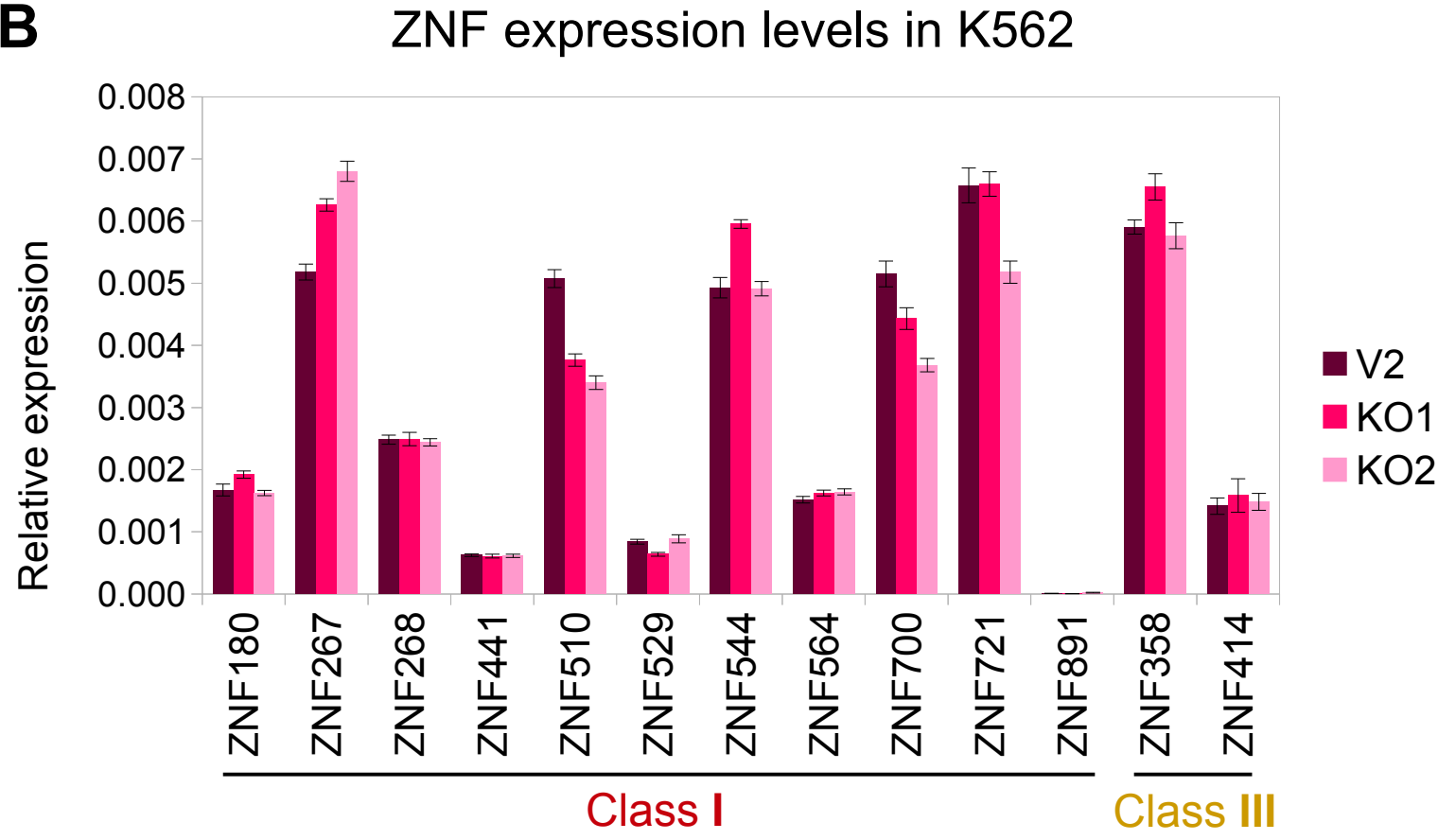
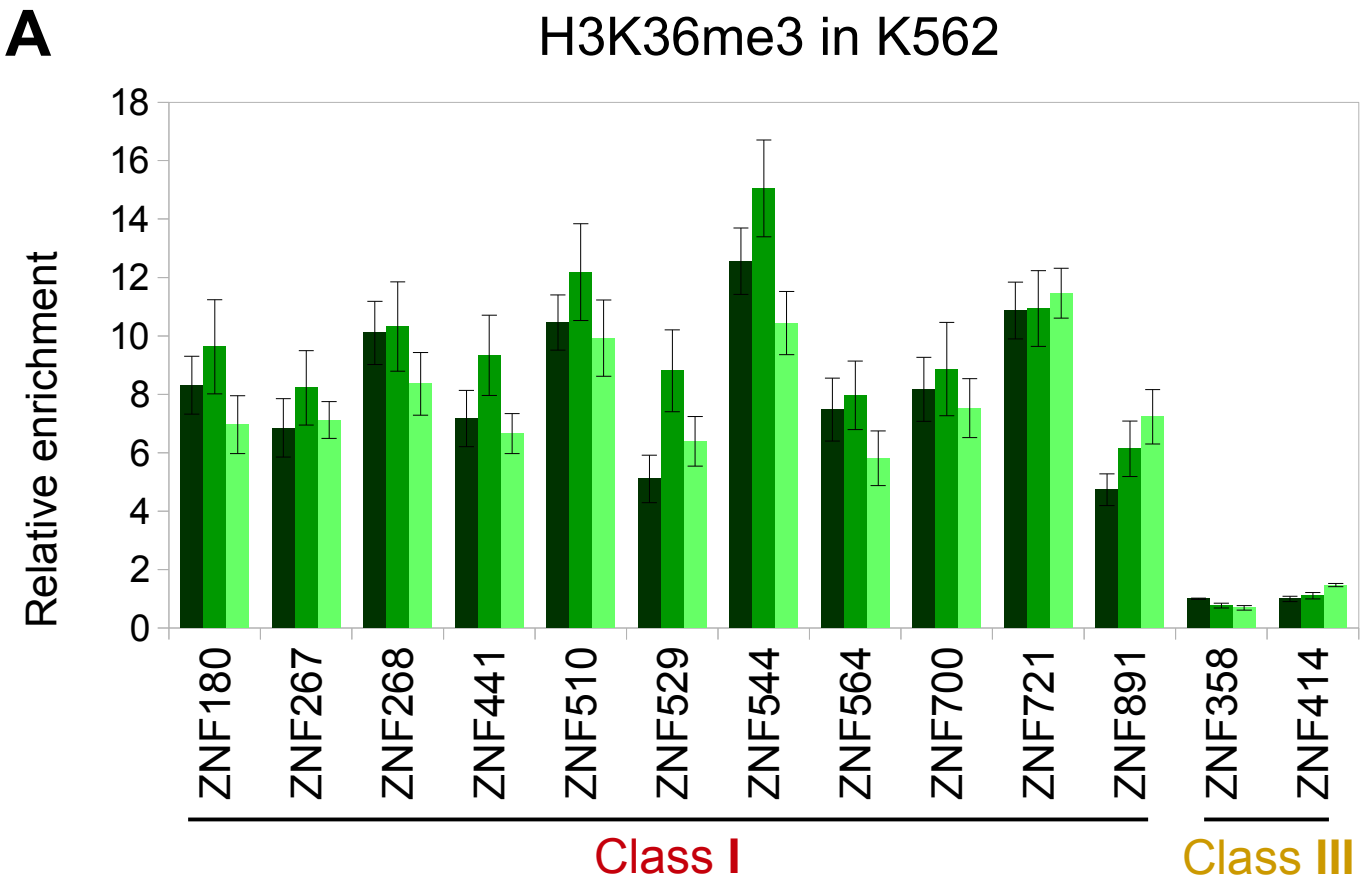
ATRX KO2, K562

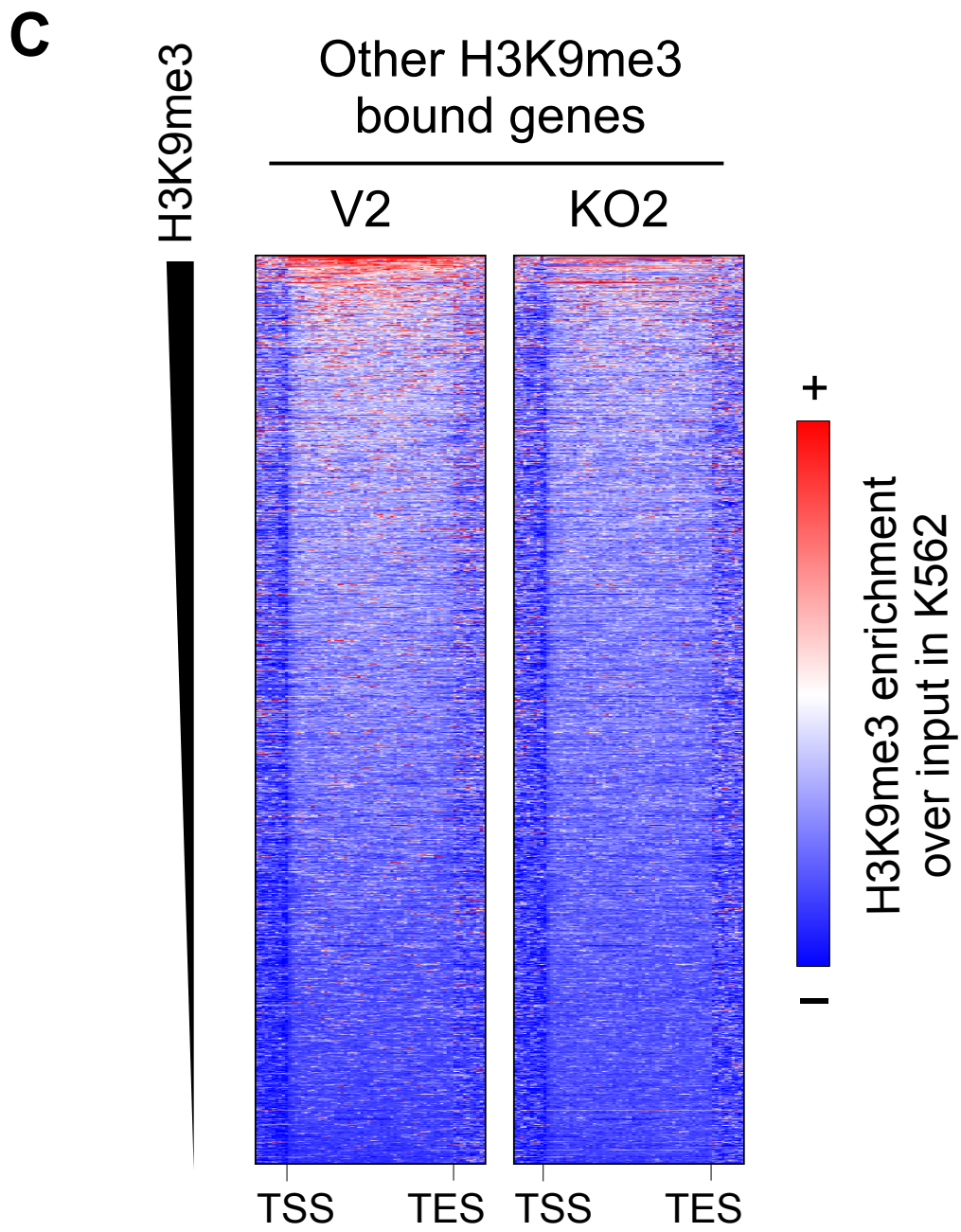
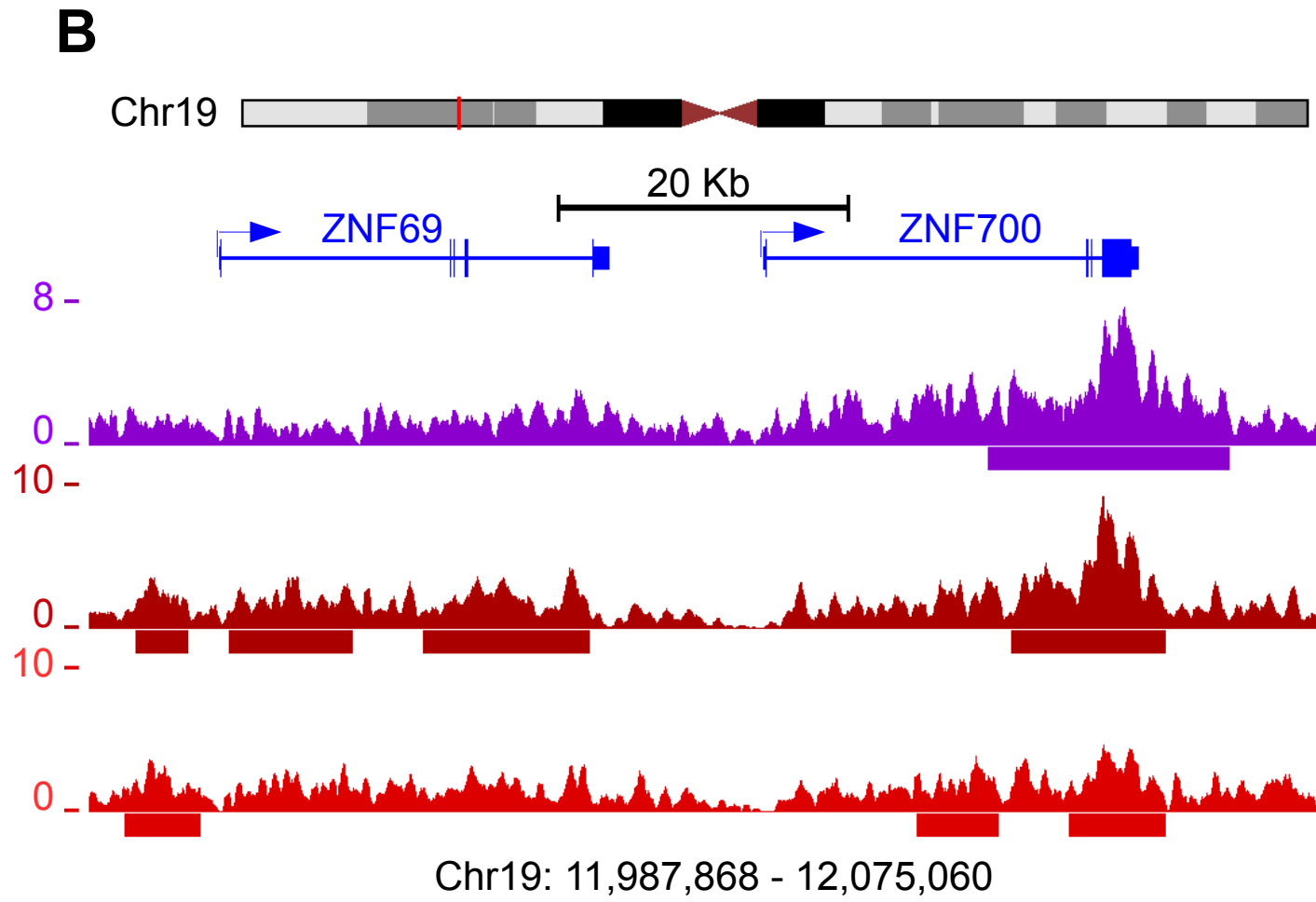
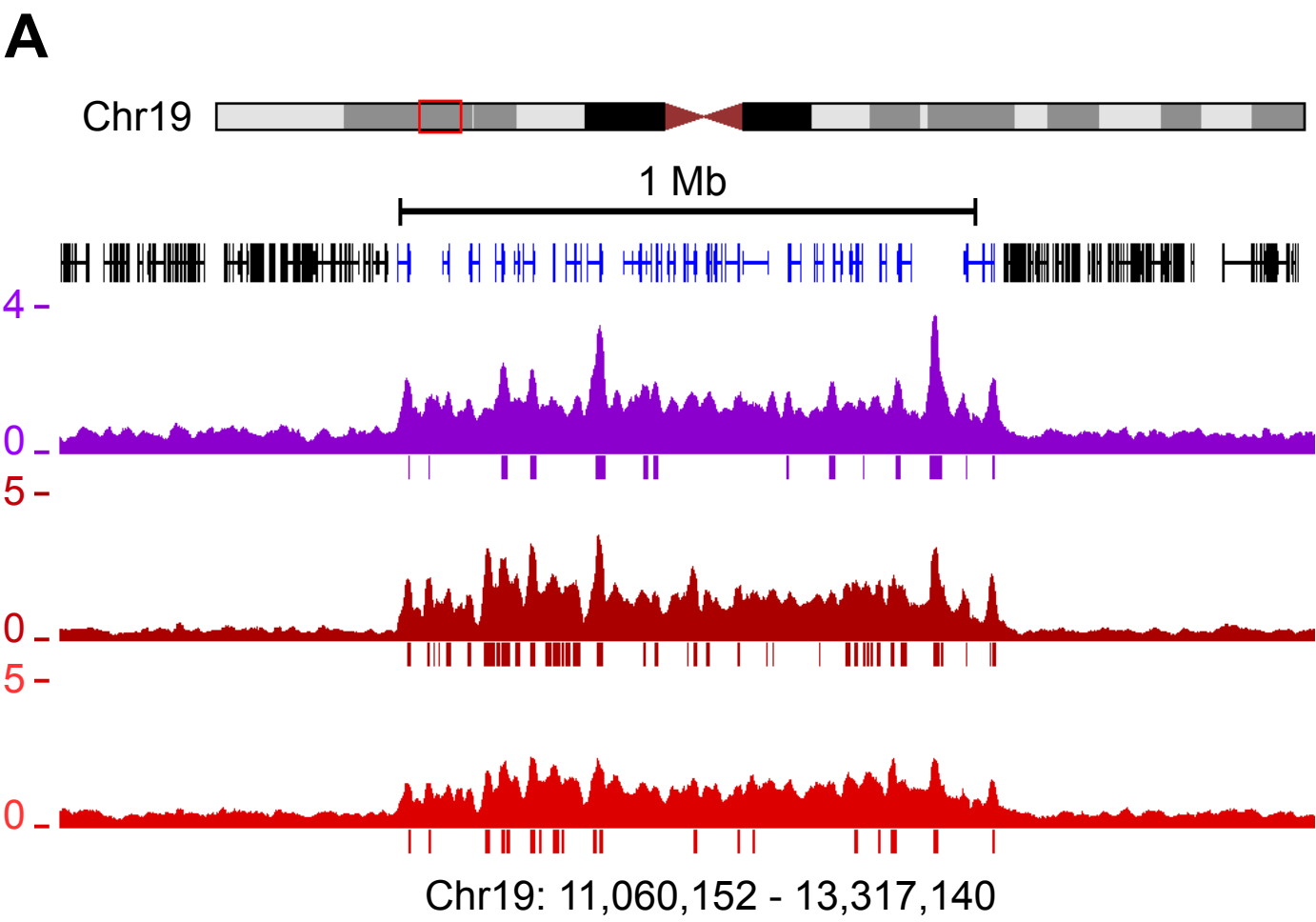


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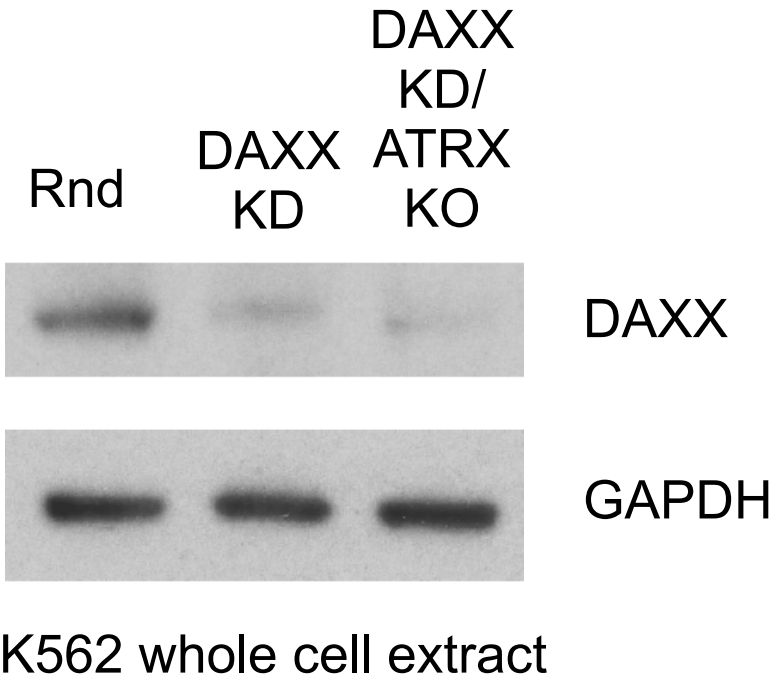




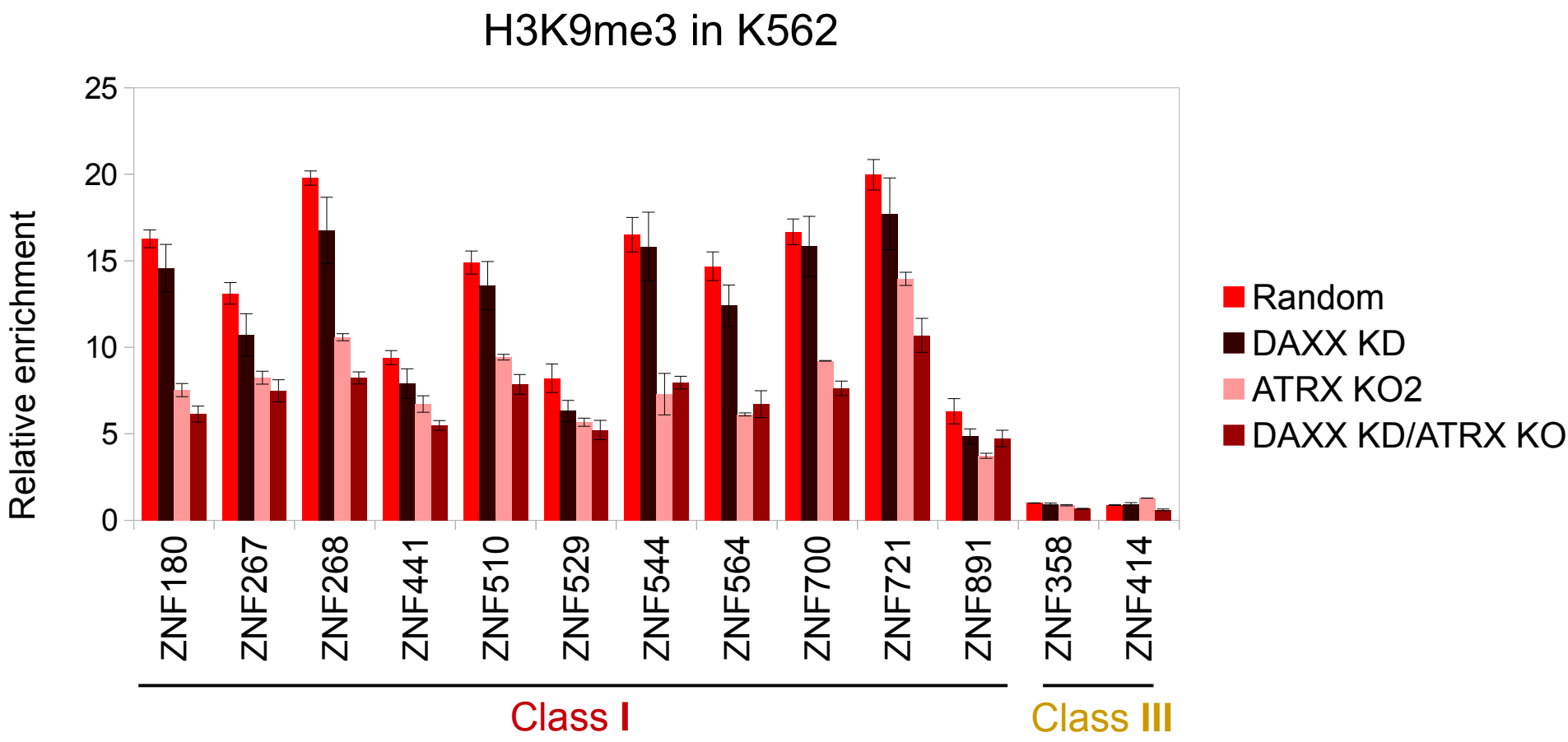




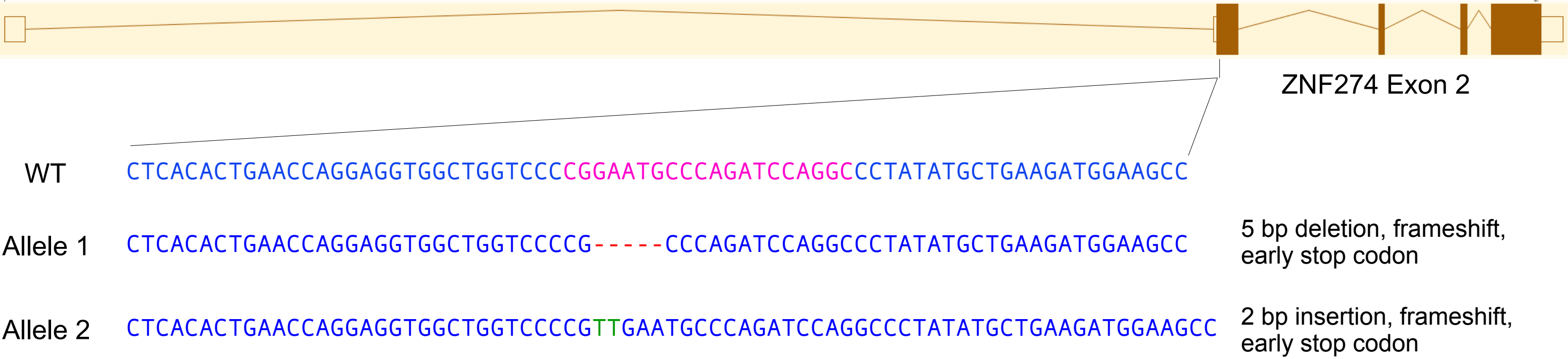
A



B



Z274 KO, K562



Z274/ATRX KO, K562

