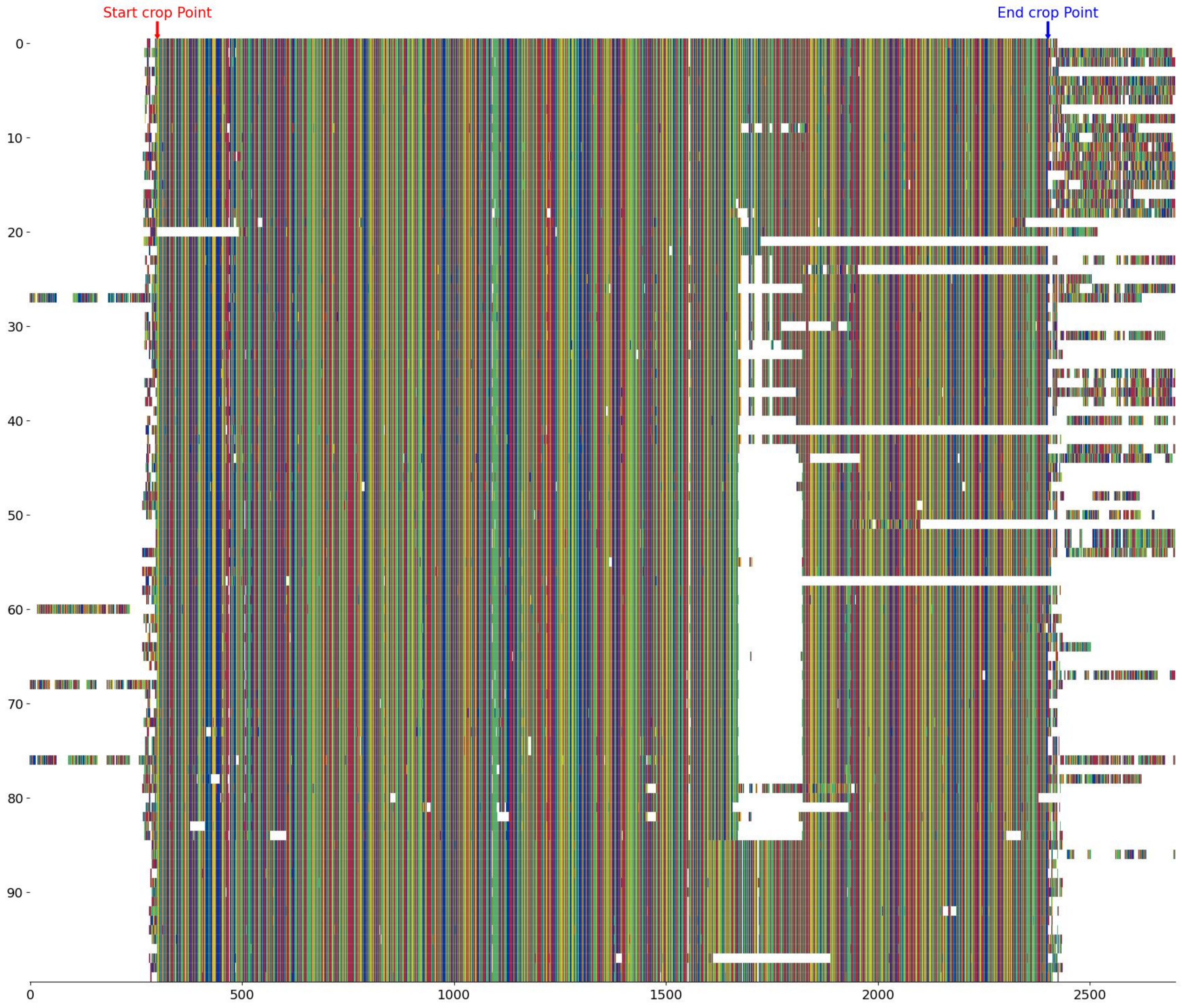






Start crop Point

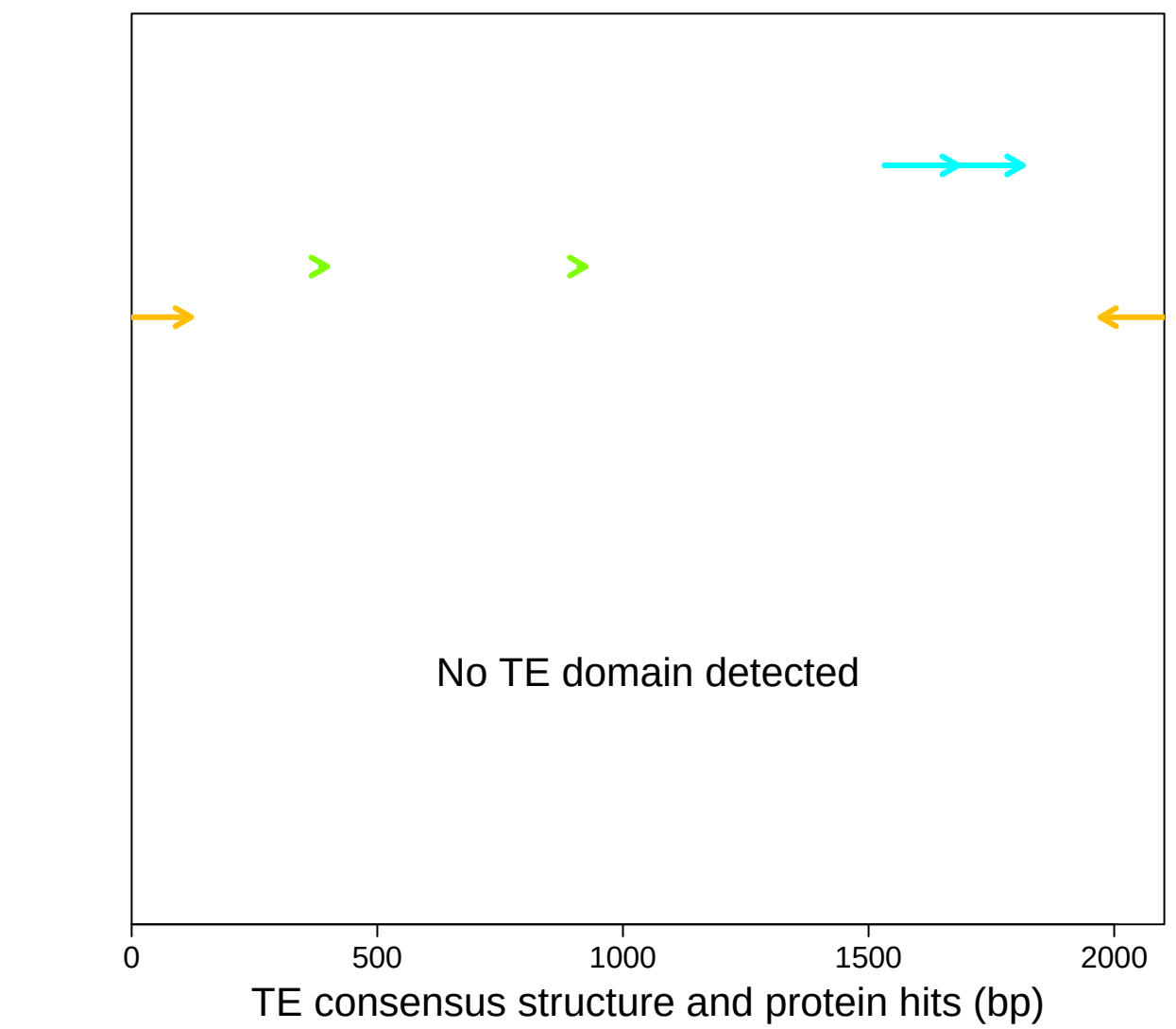
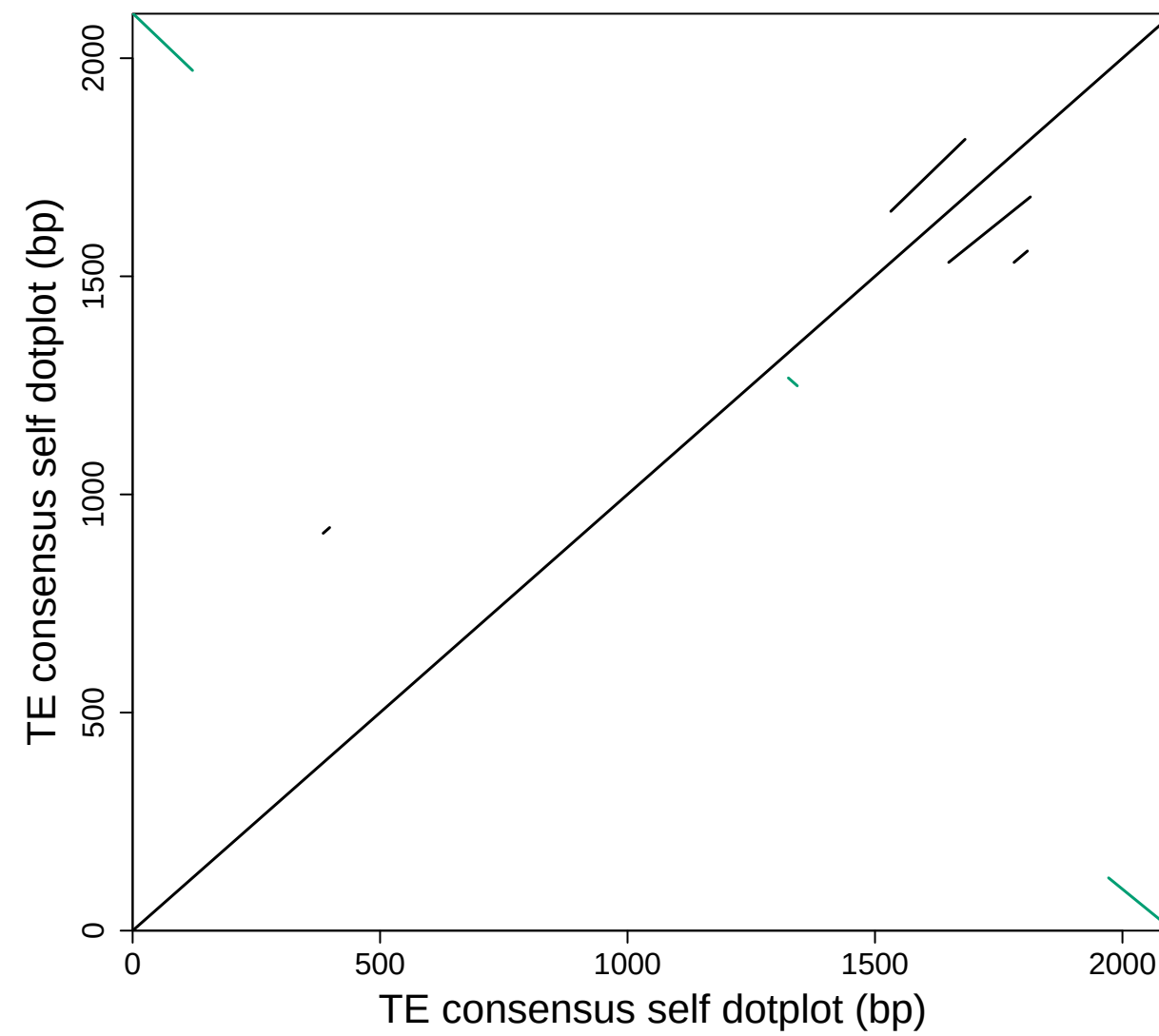
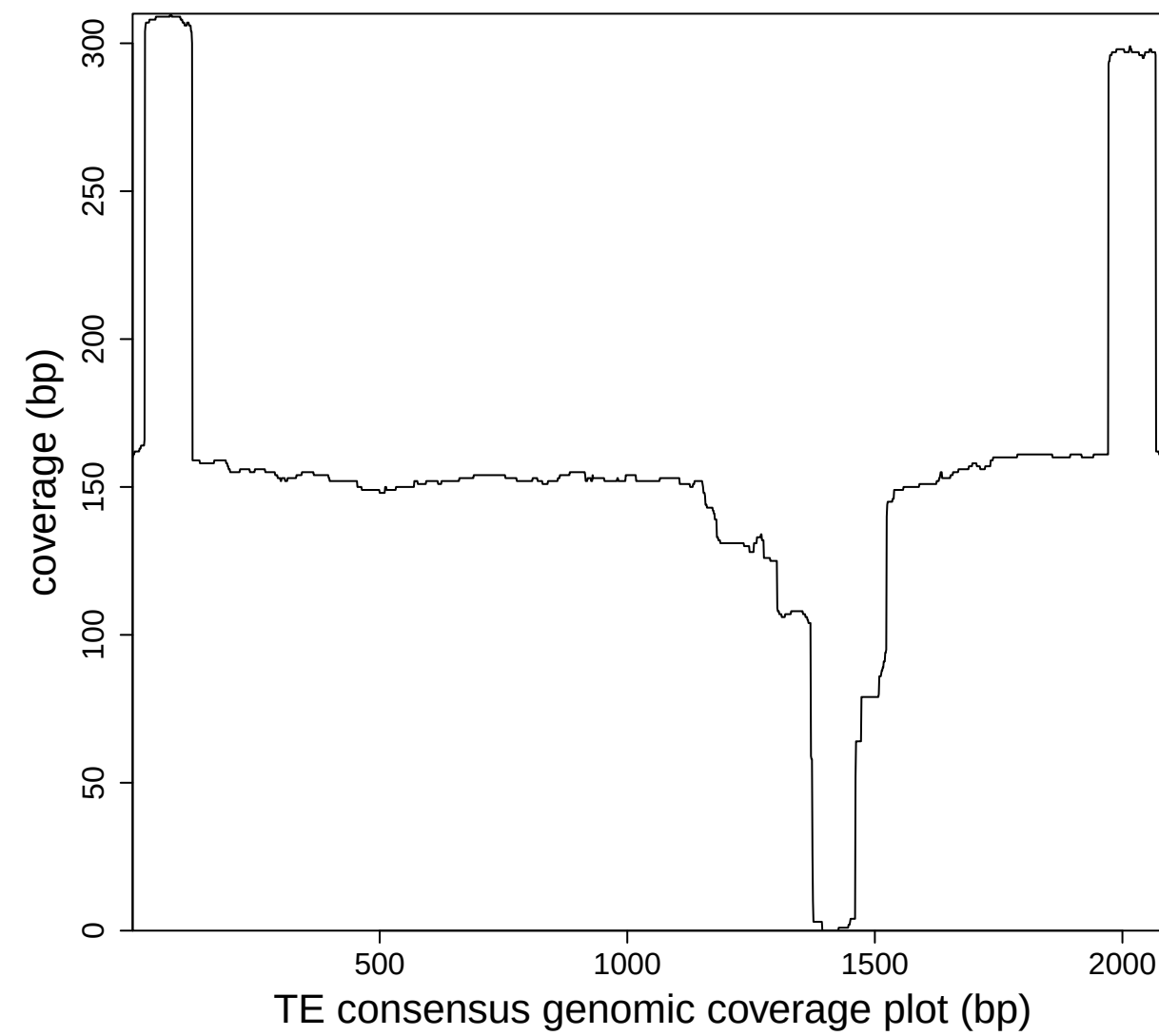
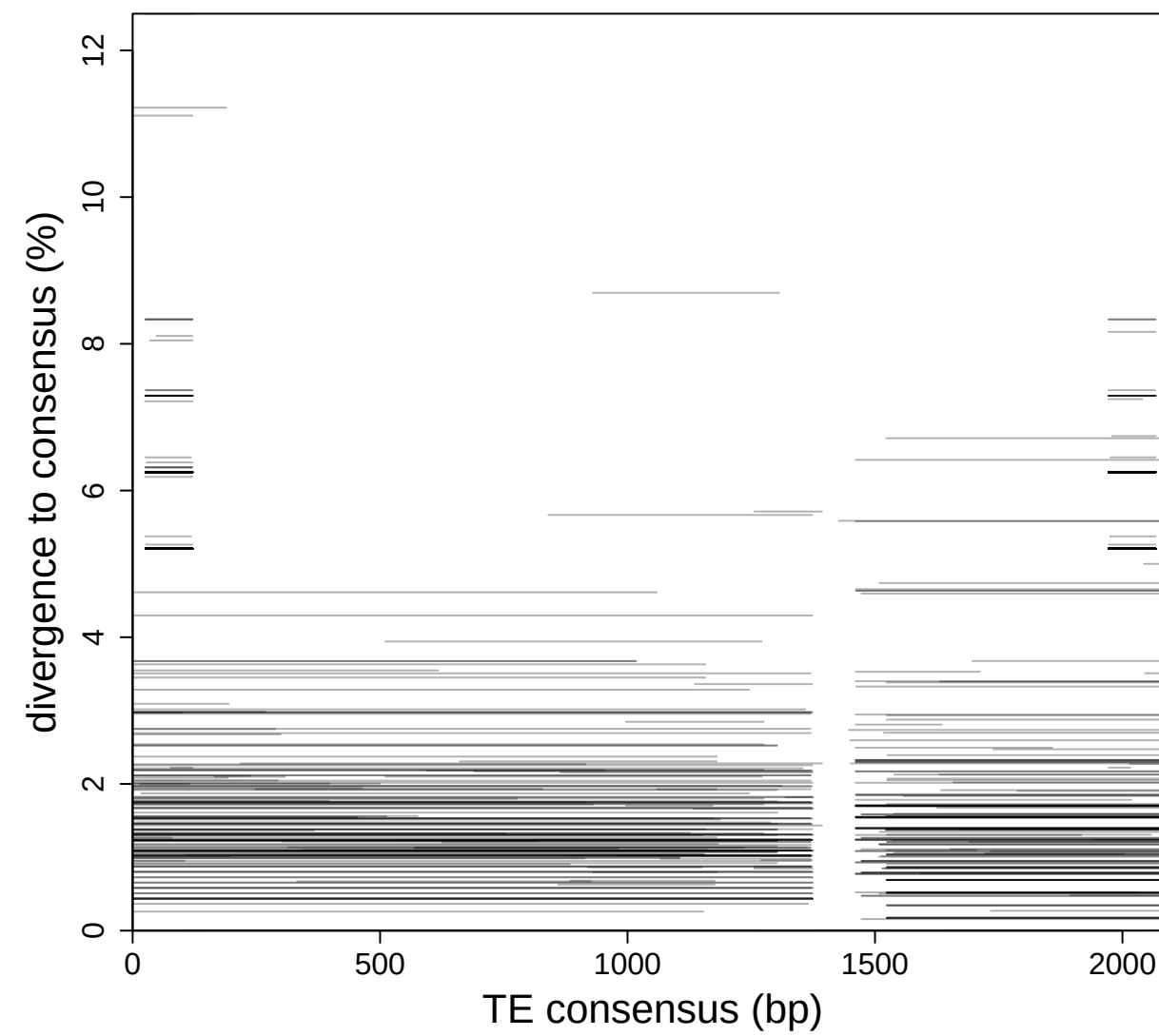
End crop Point





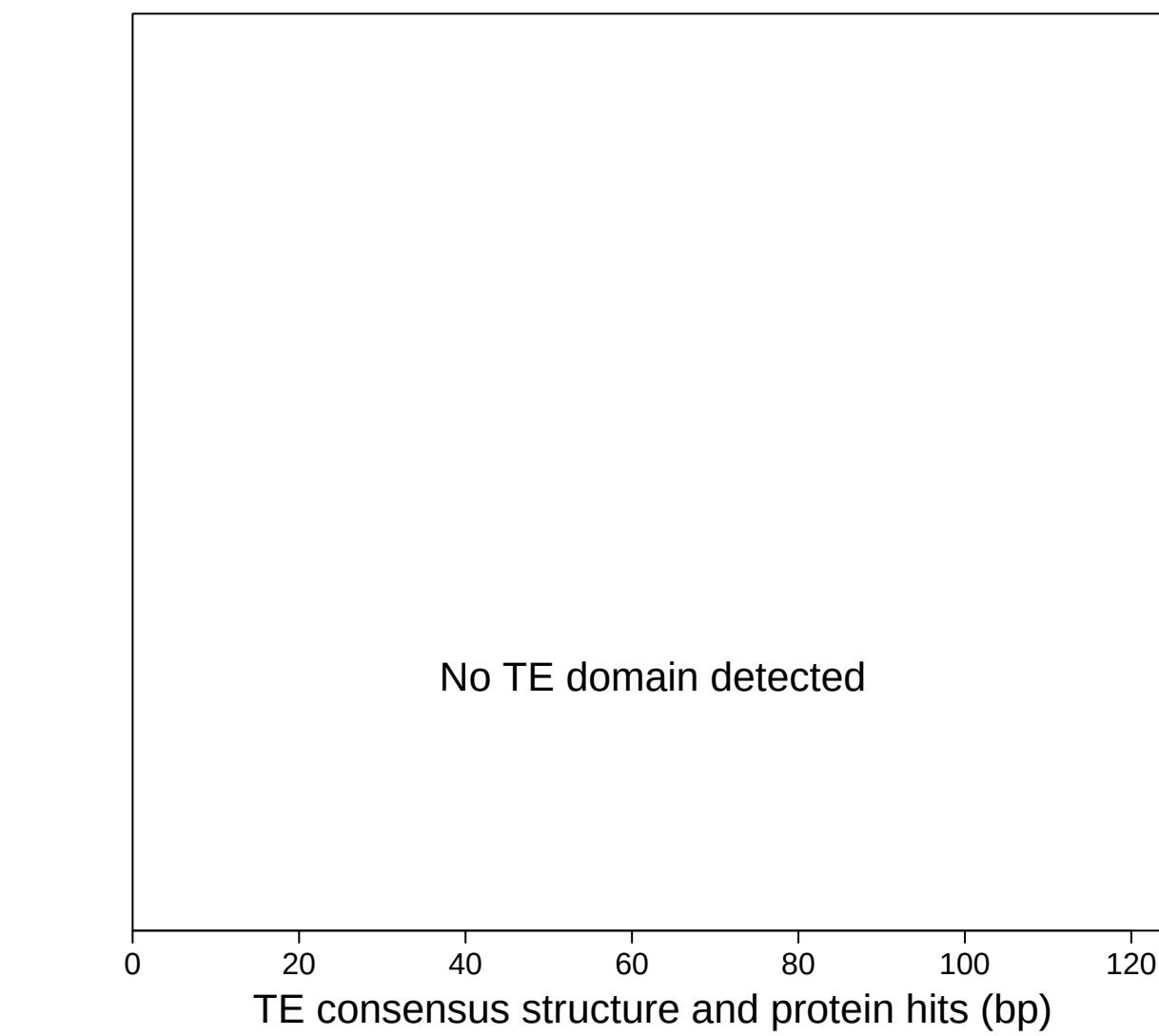
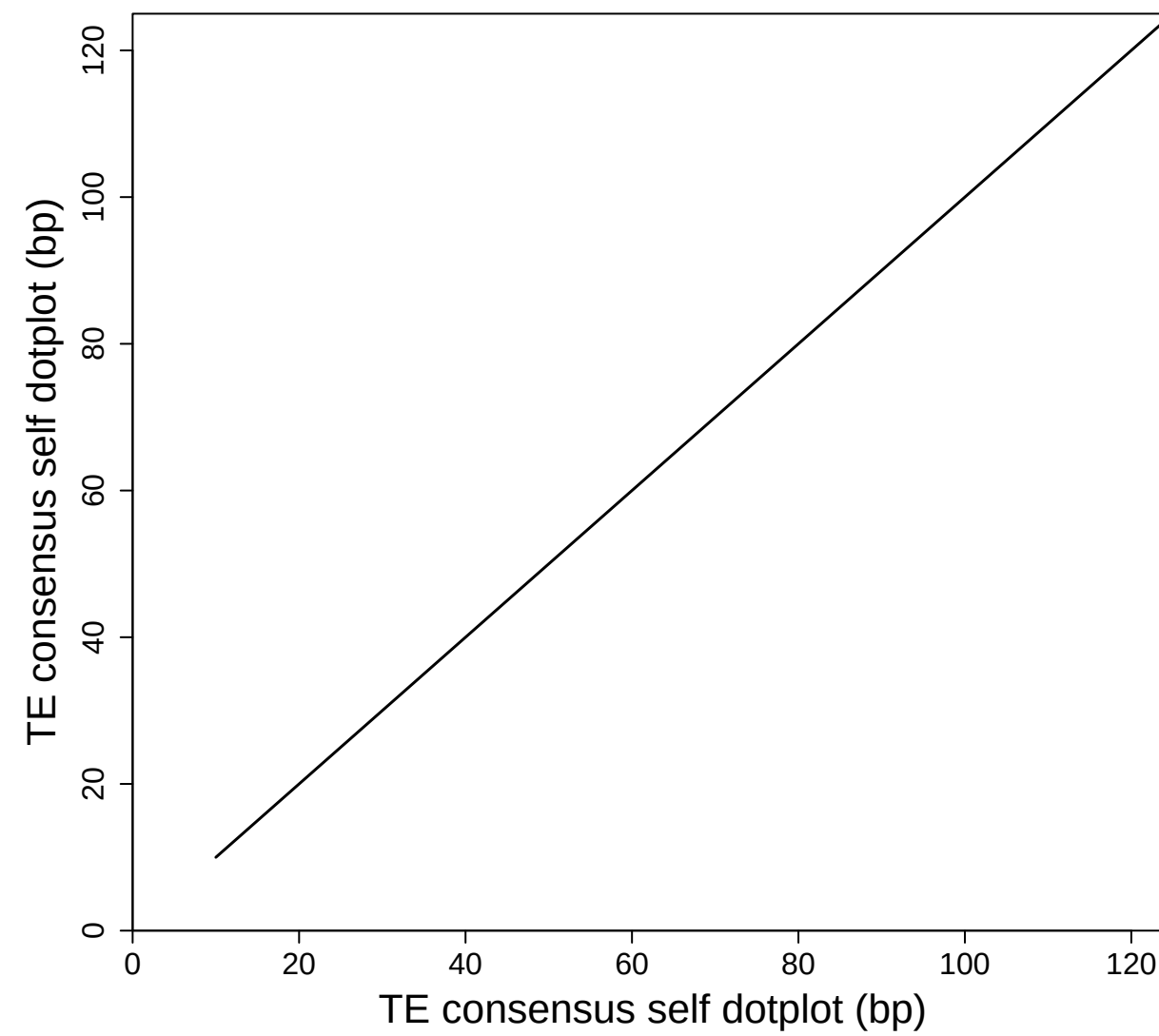
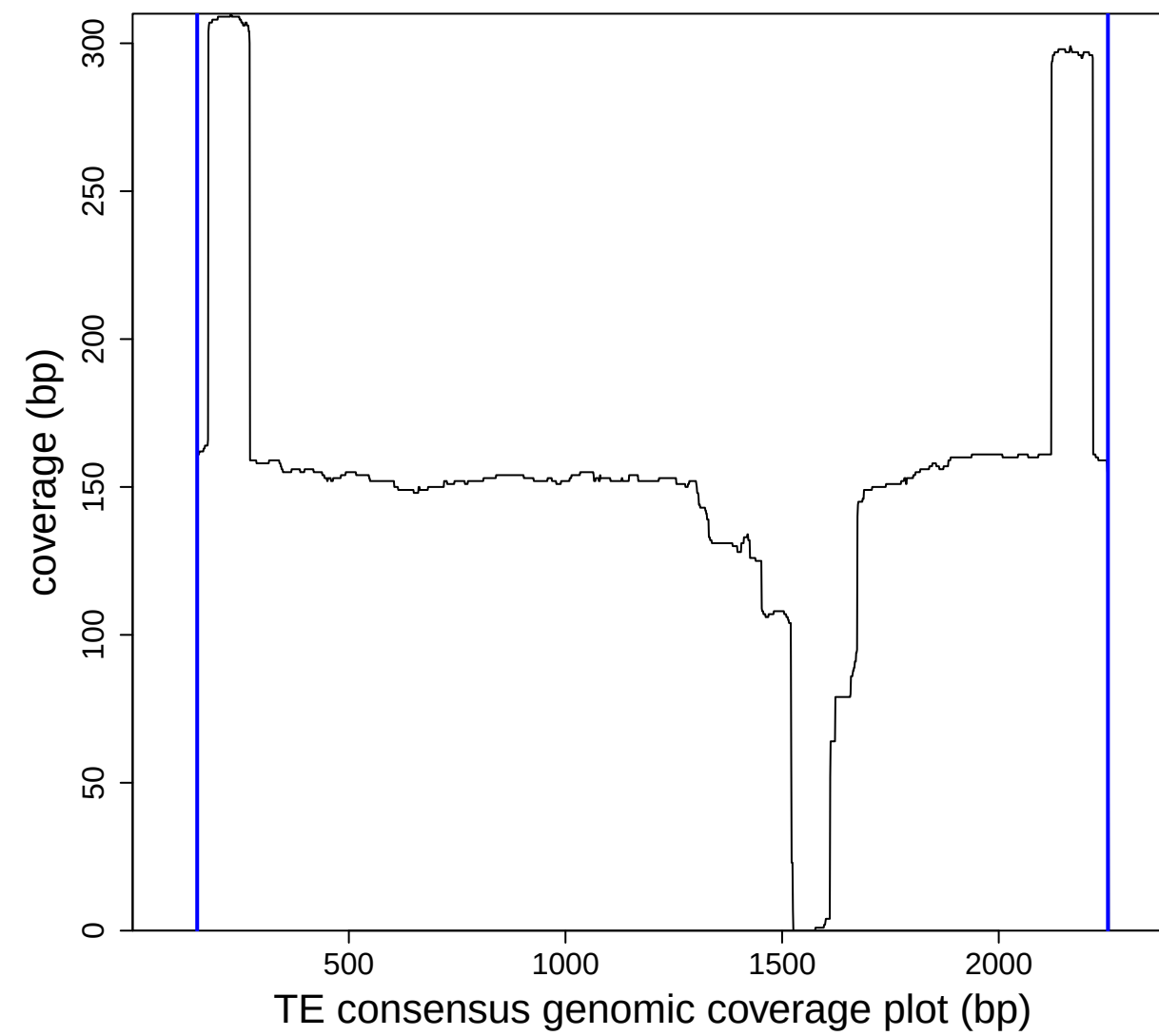
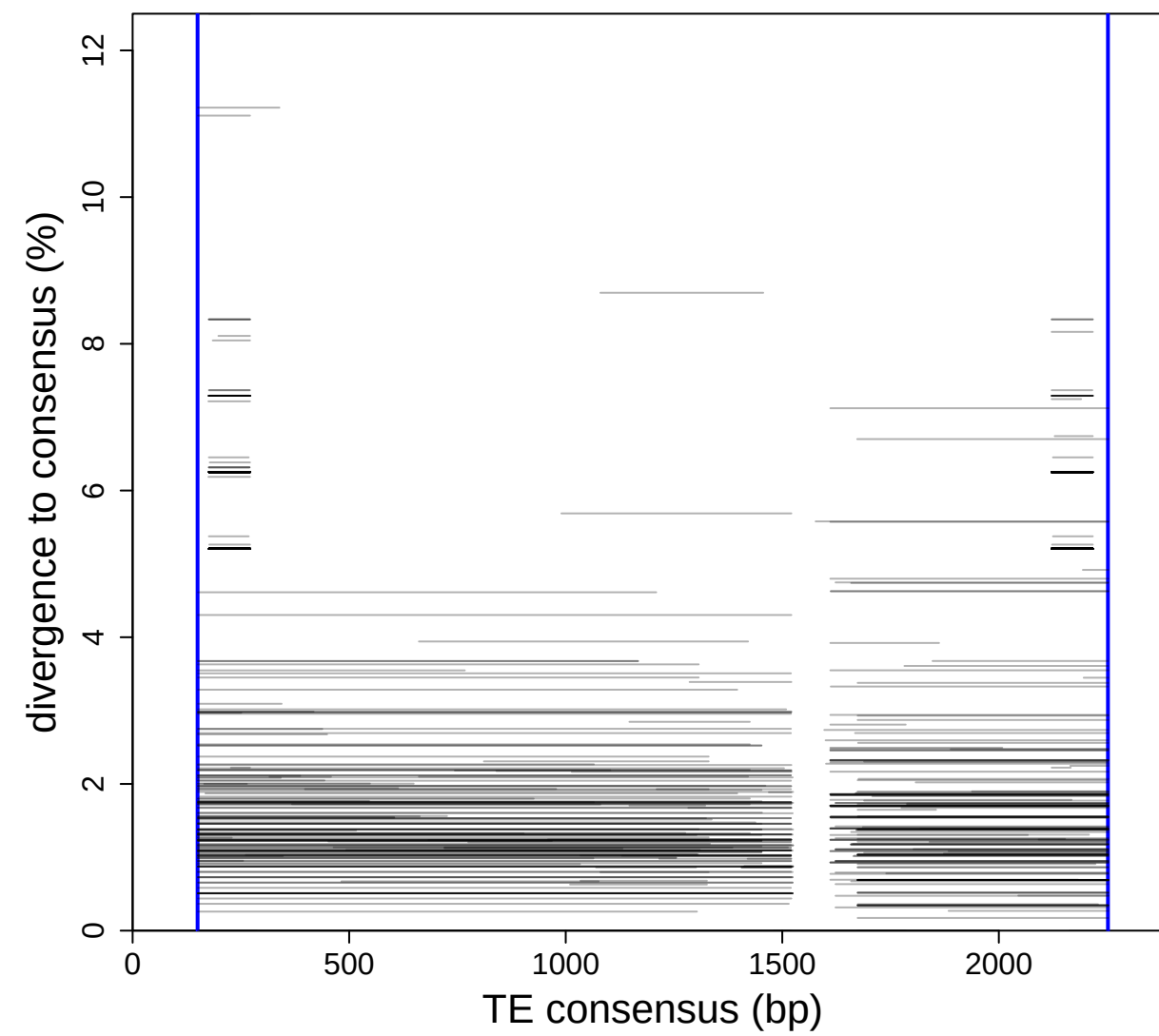
a.b.bed\_uf.bed\_g\_1.bed\_fm\_1.bed\_0\_0\_bcln.fa\_aln.fa\_cl.fa\_gs.f  
size: 2102bp; fragments: 660; full length: 0 (>=1891.8bp)

# After TEtrimmer 2102 bp



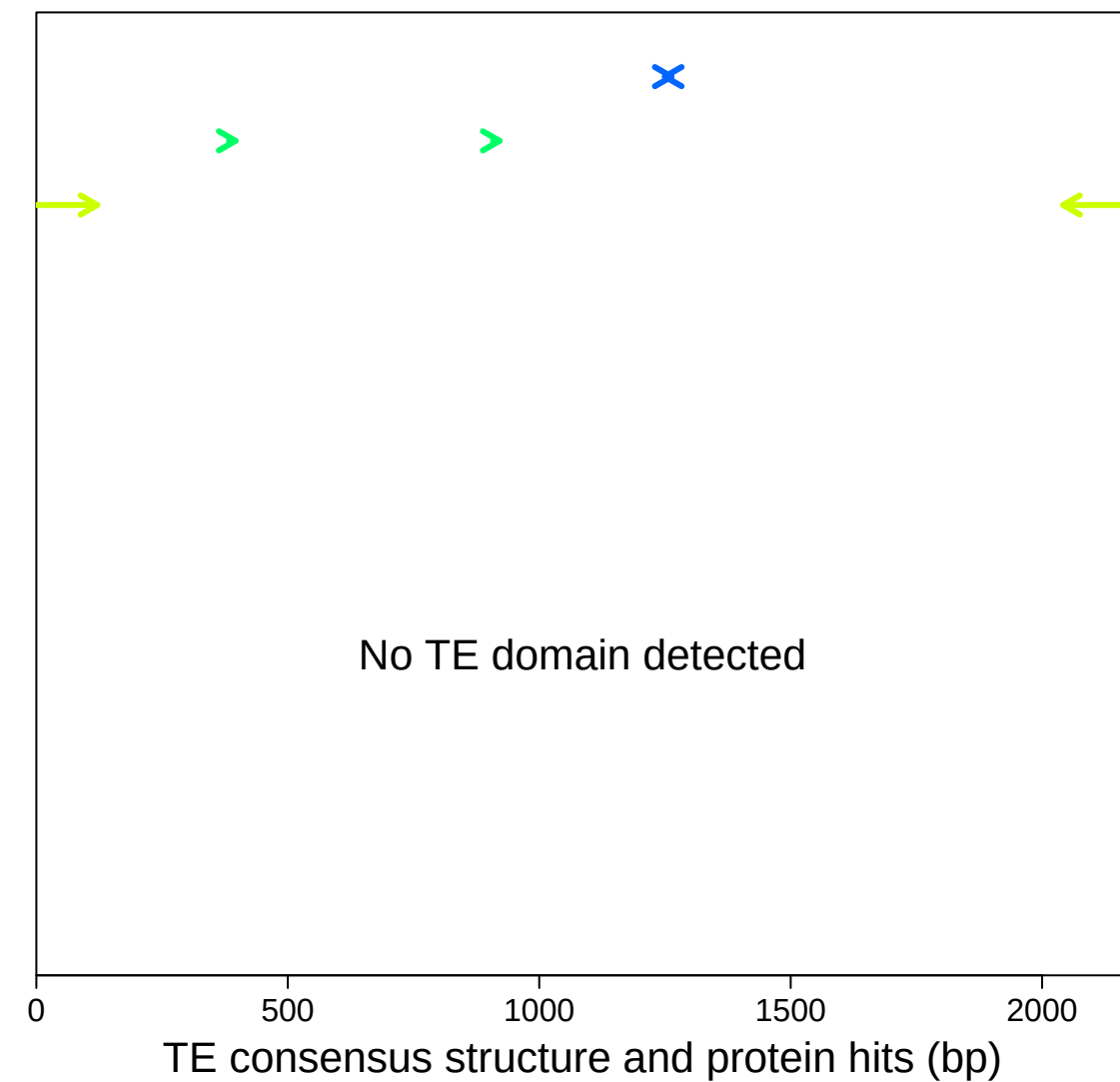
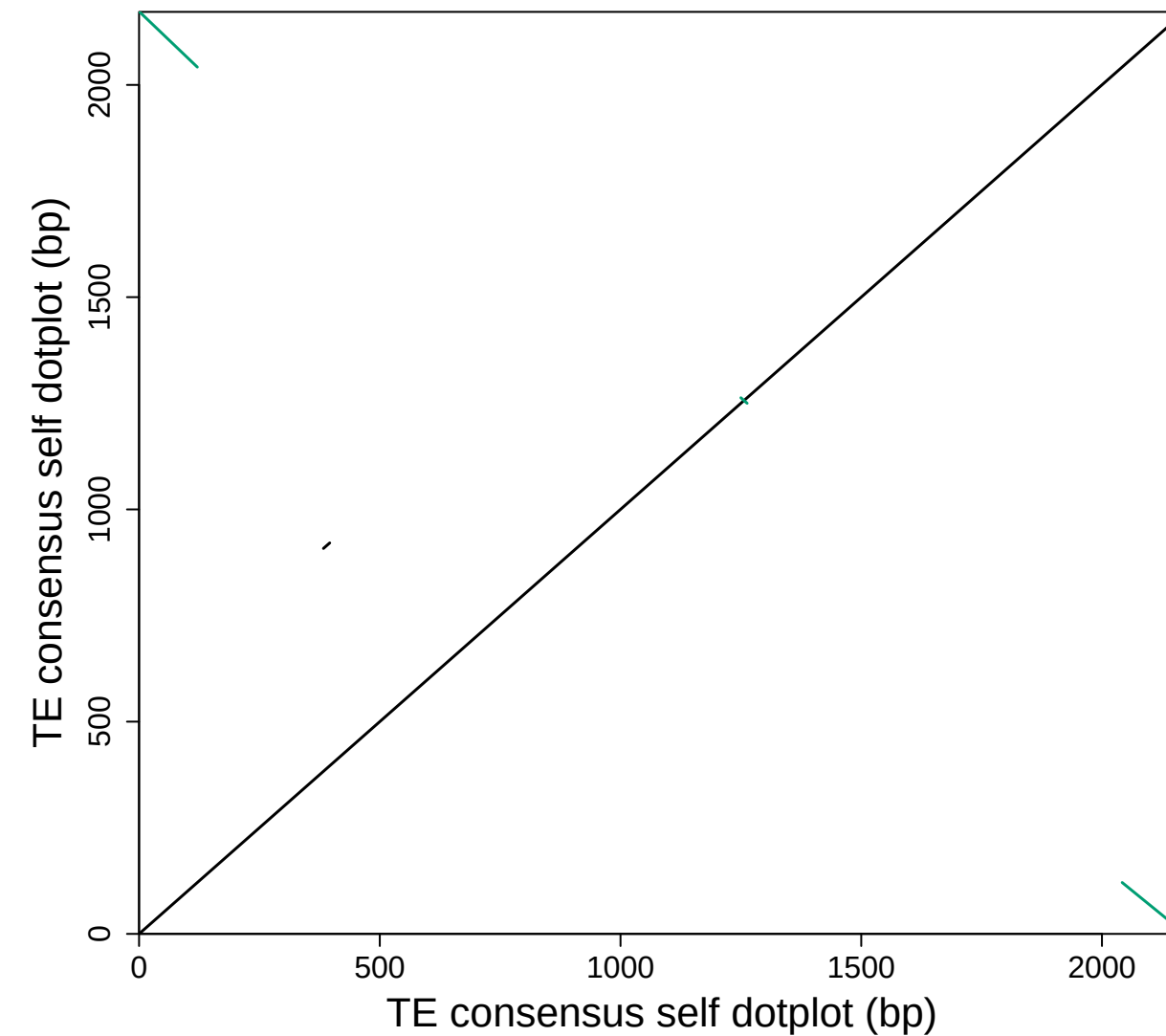
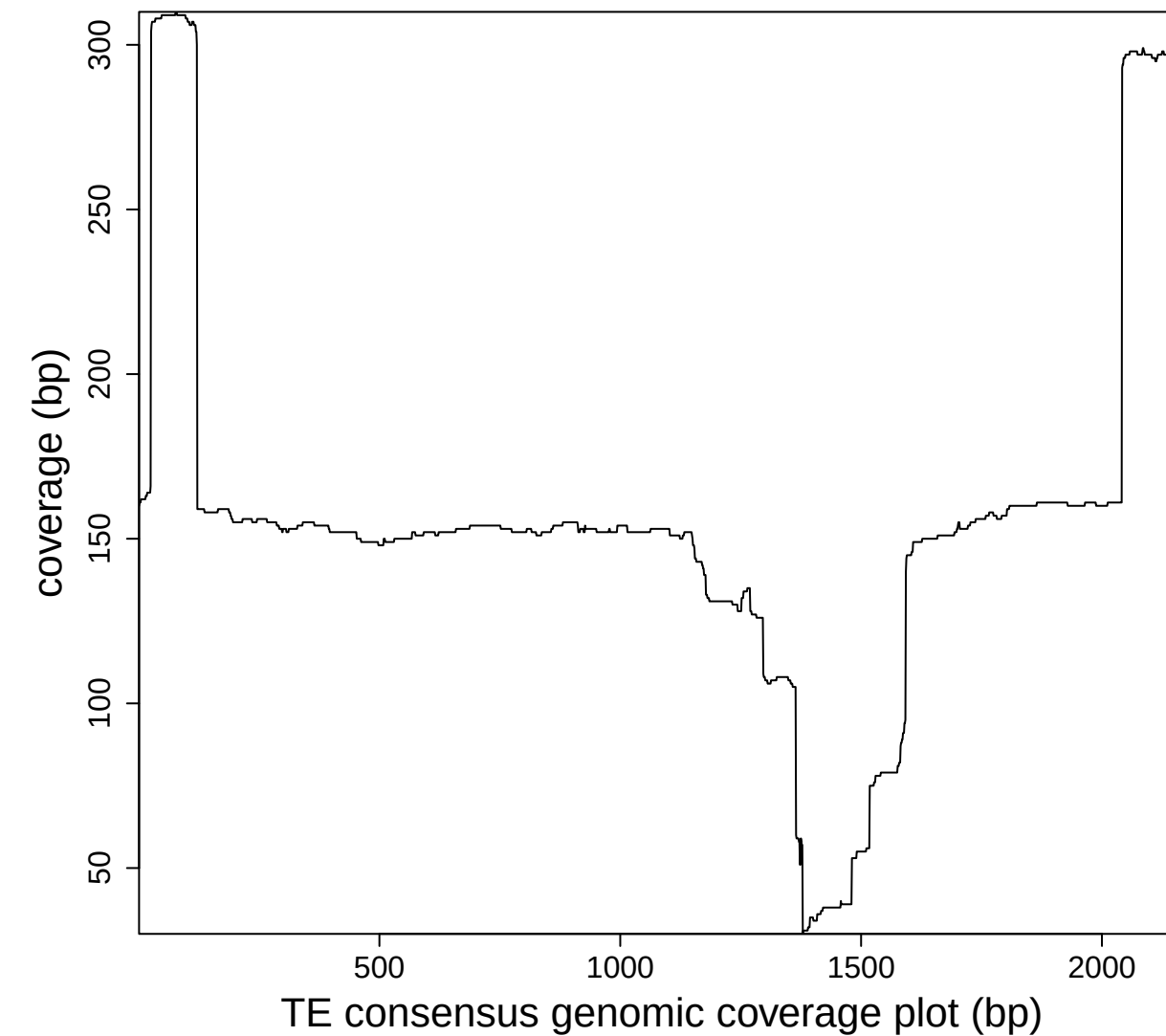
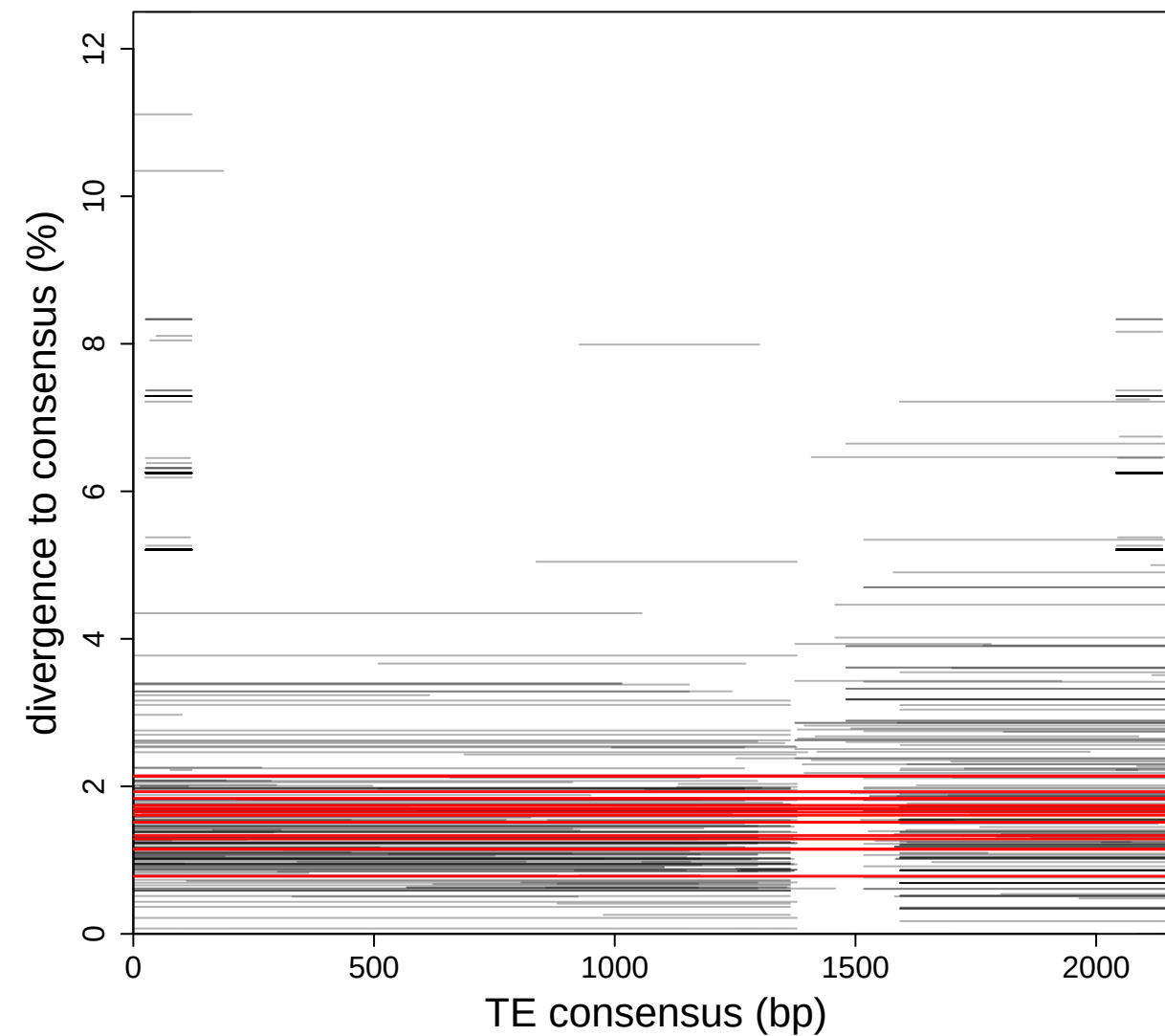
0.fasta.b.bed uf.bed g\_1.bed fm\_1.bed 0\_0\_bcln.fa aln.fa cl.f  
size: 2402bp; fragments: 659; full length: 0 (>=2161.8bp)

# After TEtrimmer Extended plot Blue lines are boundaries

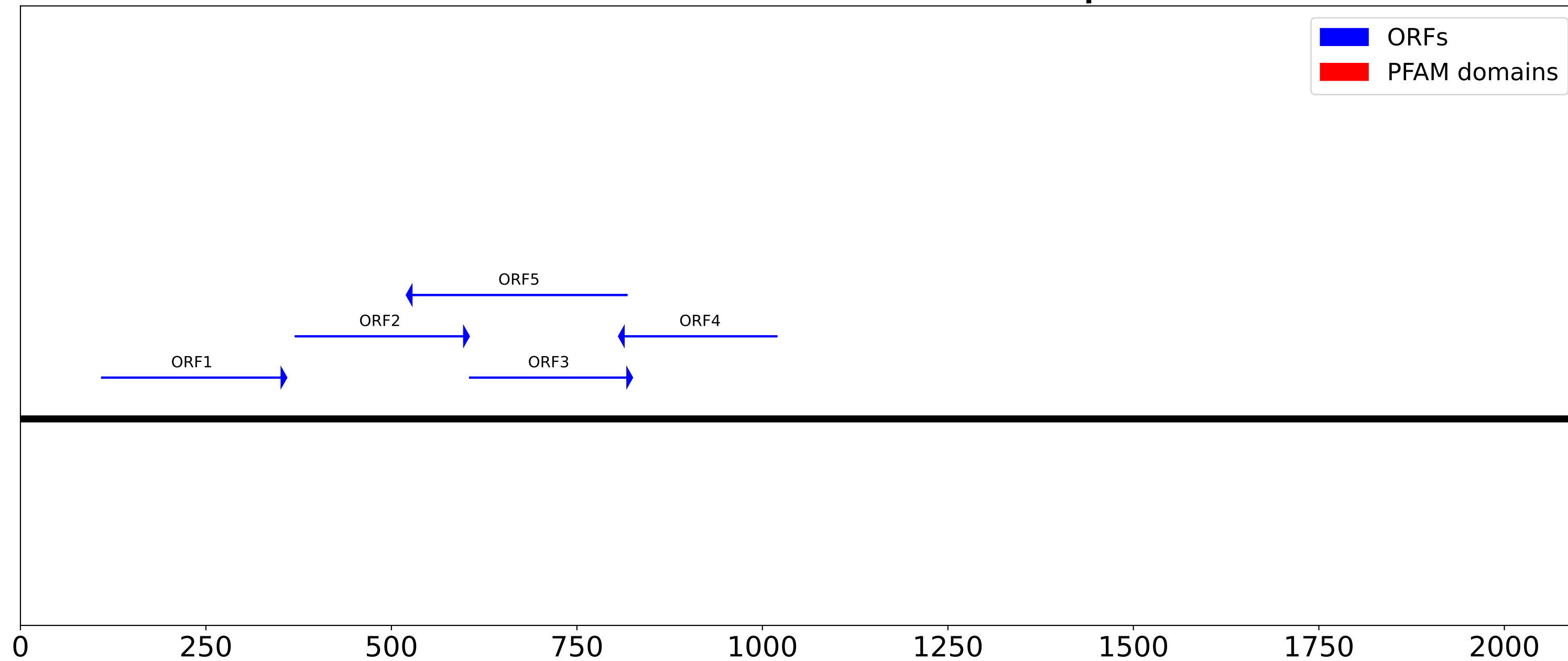


TE: rnd\_5\_family\_4620  
size: 2172bp; fragments: 642; full length: 18 ( $\geq 1954.8$ bp)

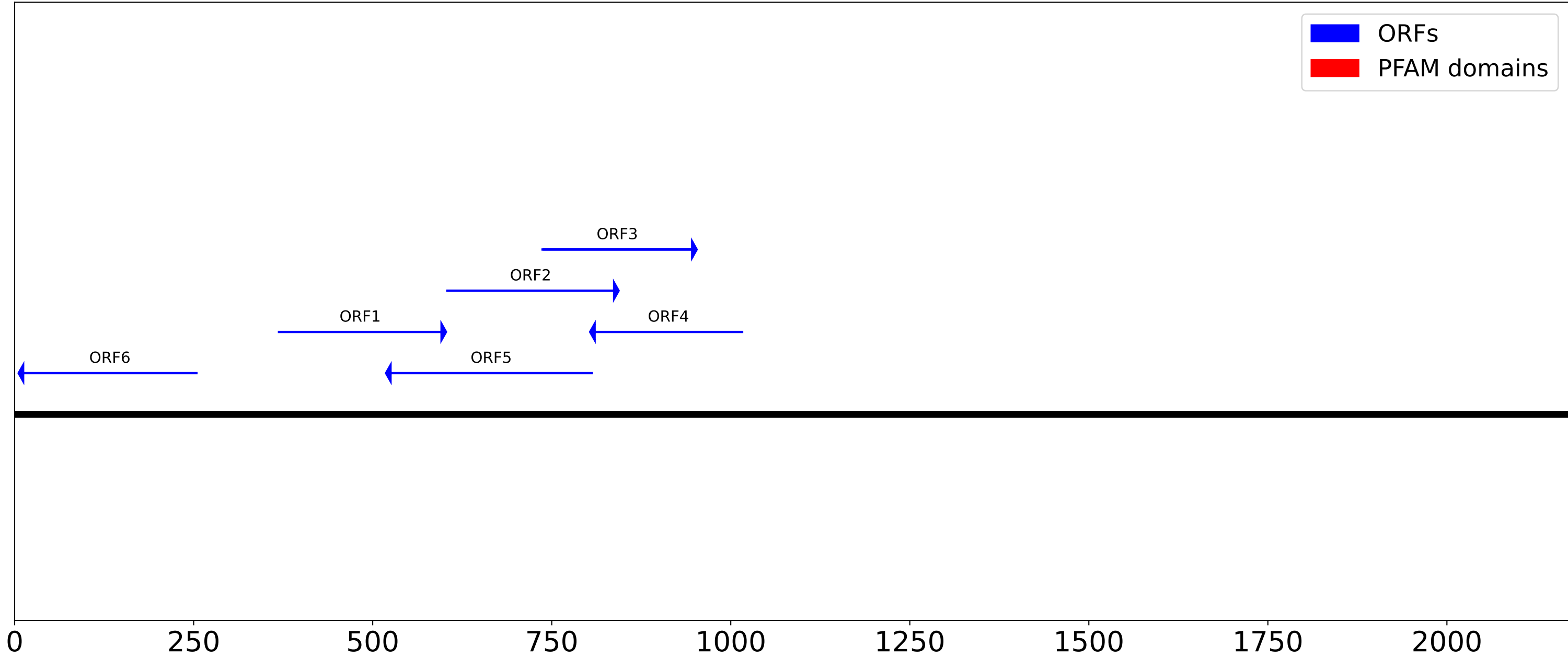
## Before TEtrimmer 2172 bp



## After TEtrimmer ORF and PFAM domain plot



**Before TEtrimmer ORF and PFAM domain plot**





# Dotplot

(windowsize = 25, threshold = 50.00 10/10/25)

