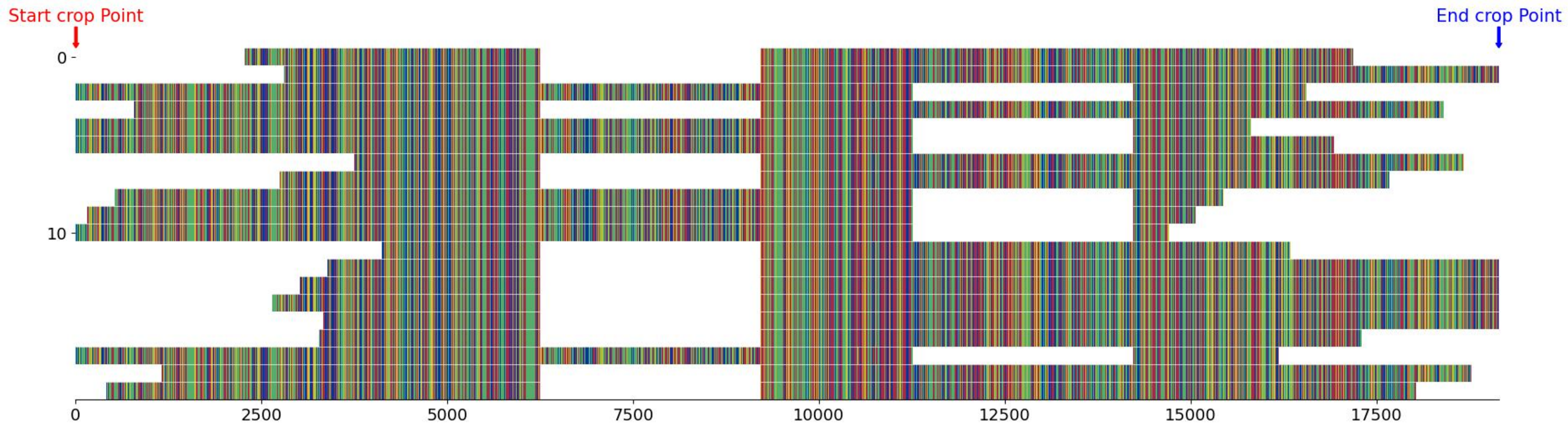


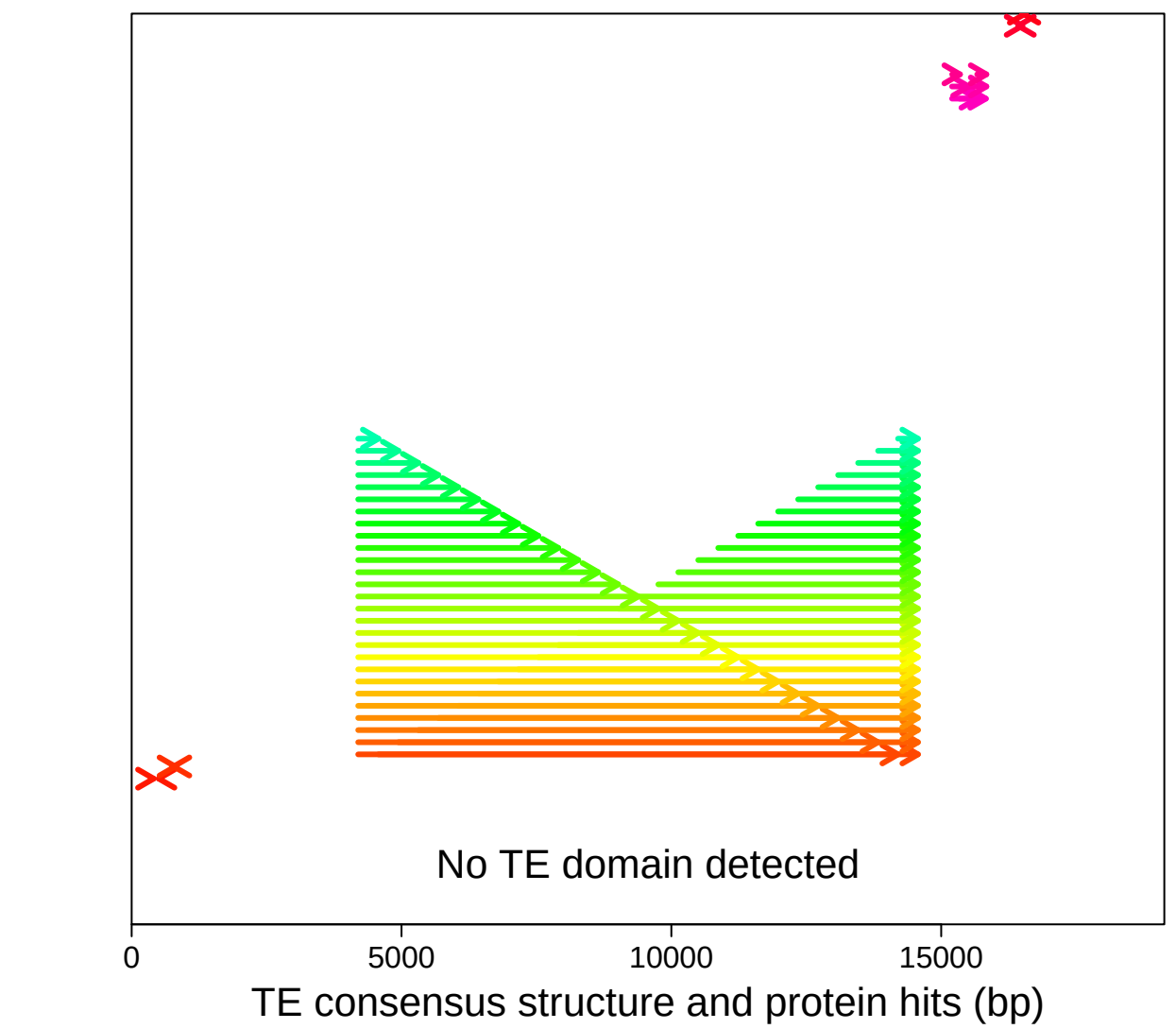
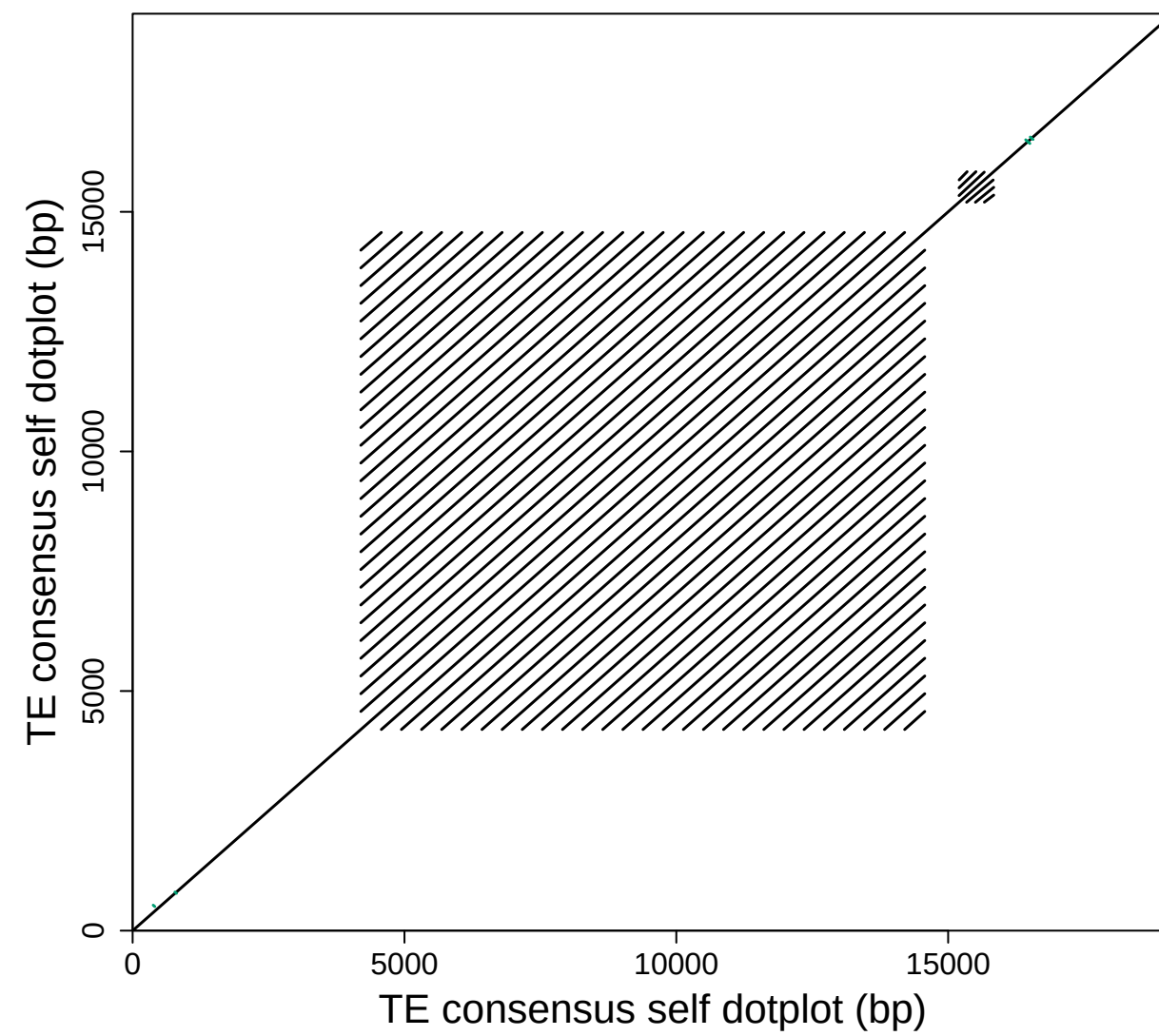
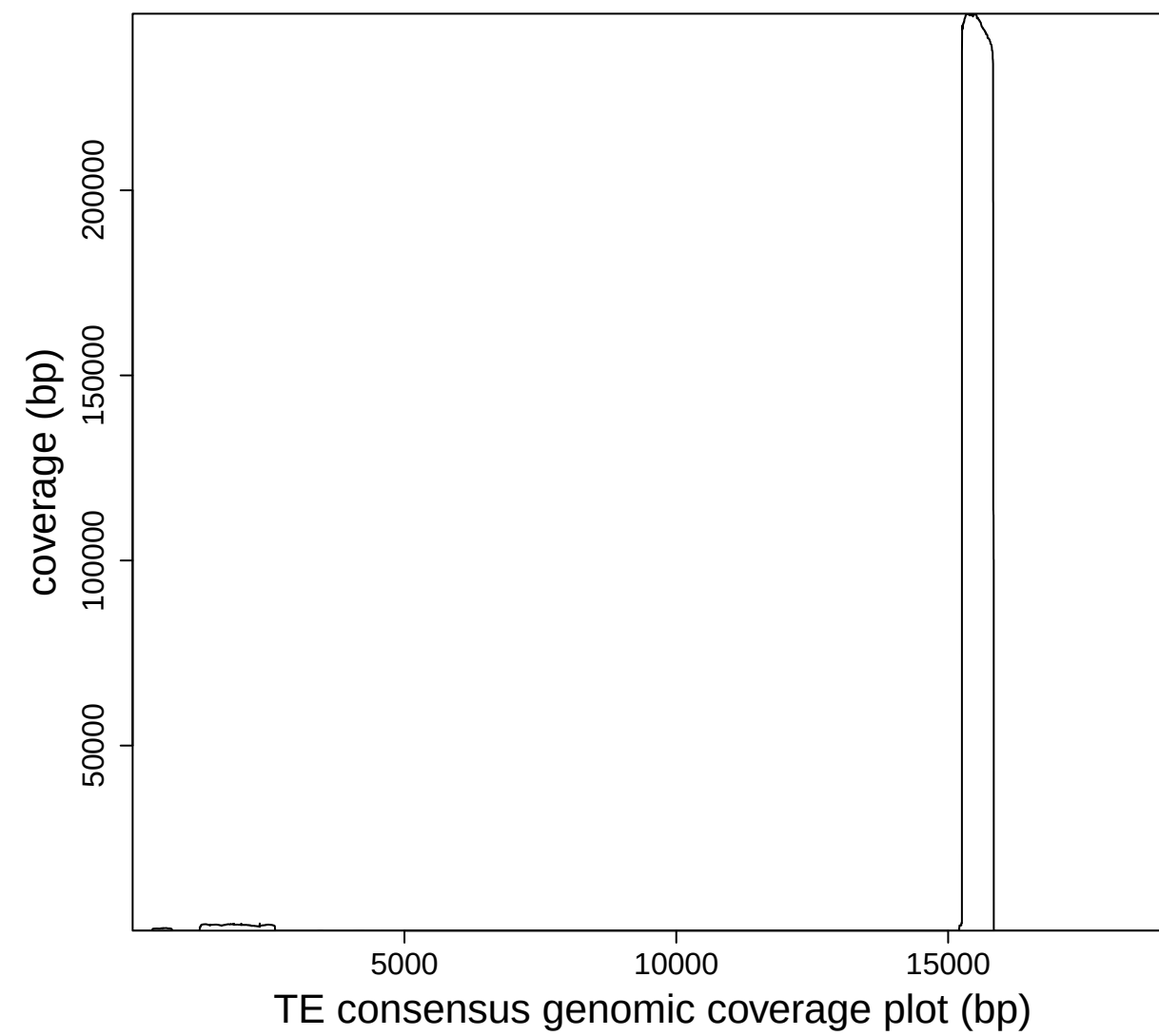
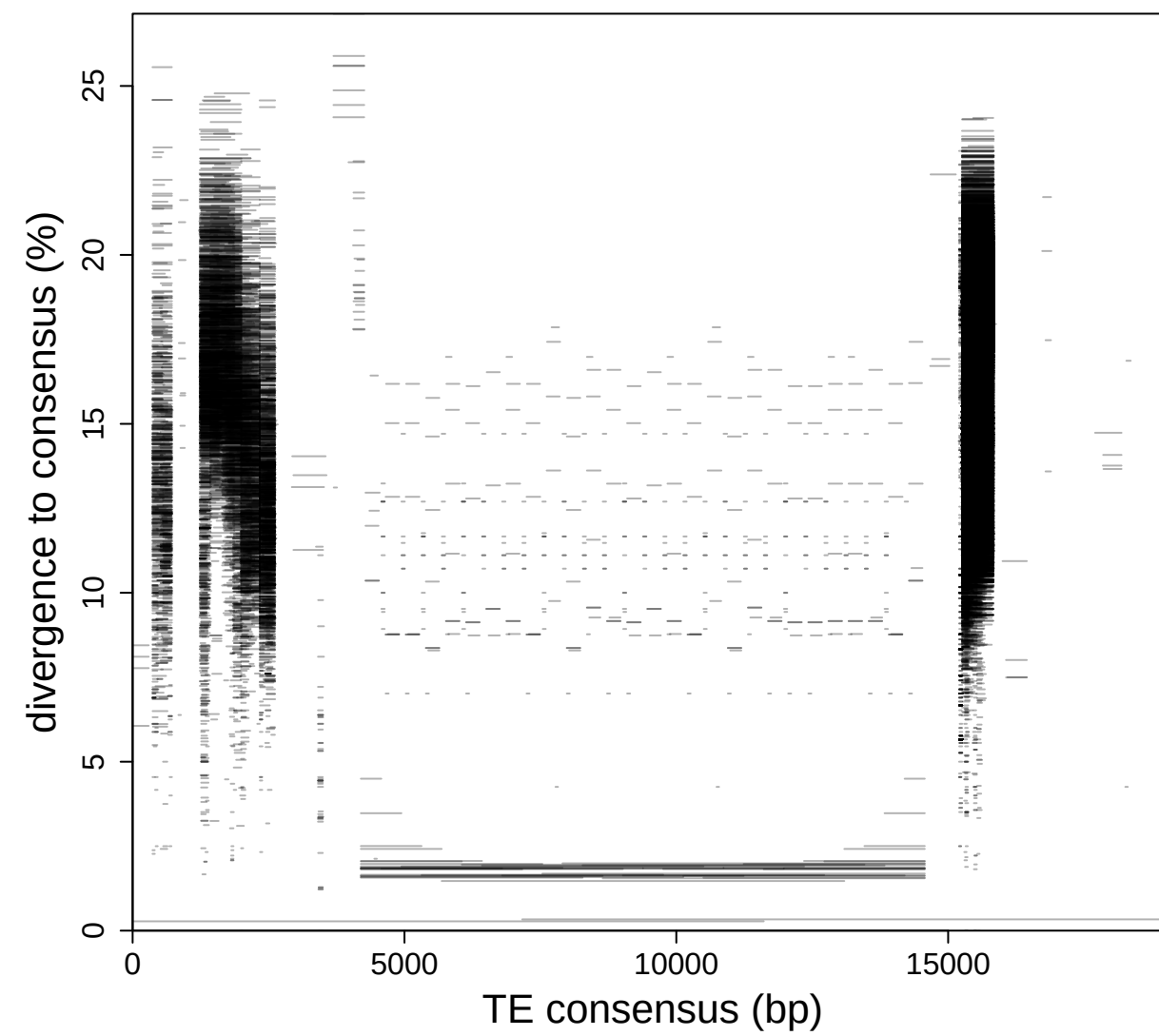


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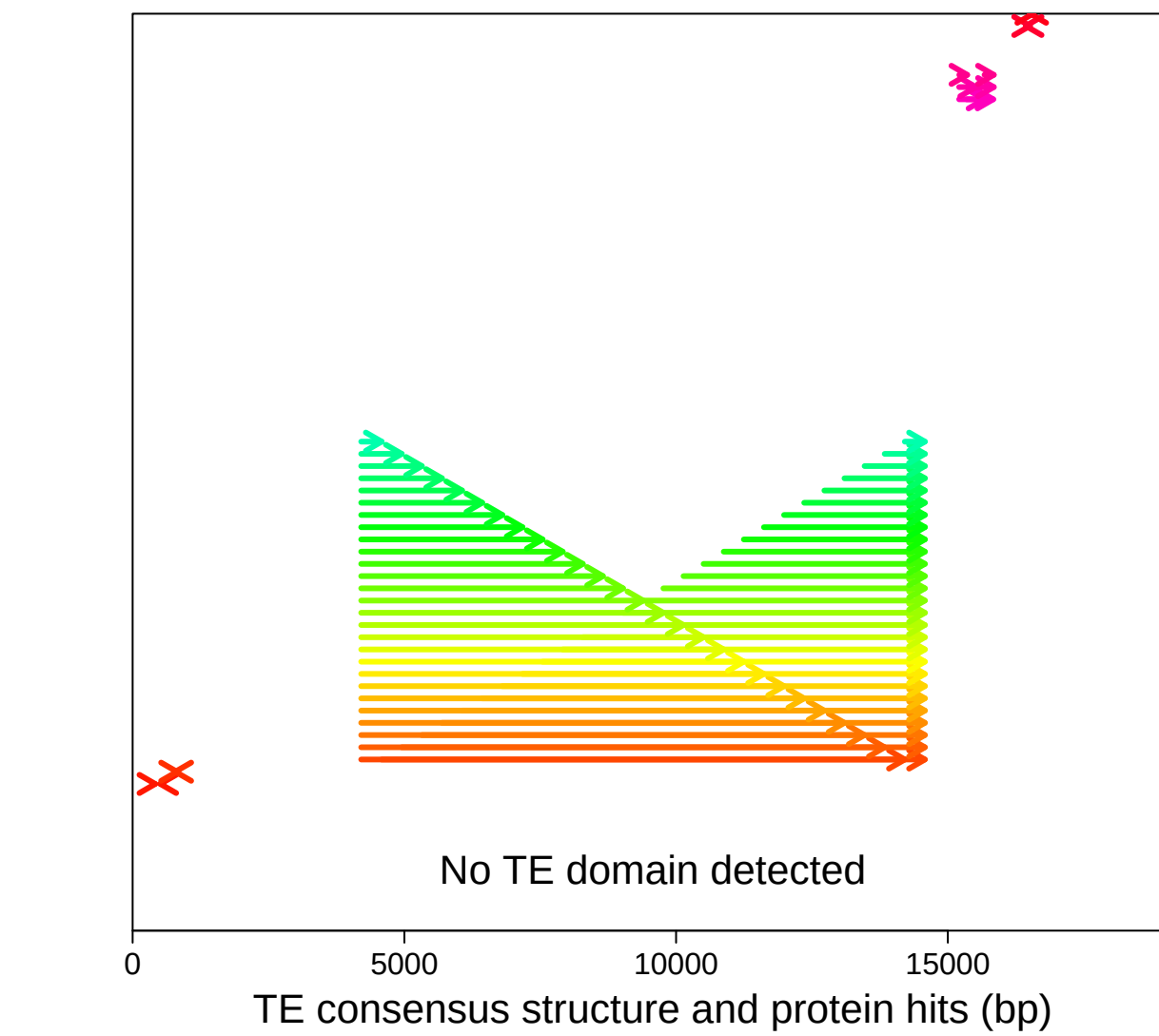
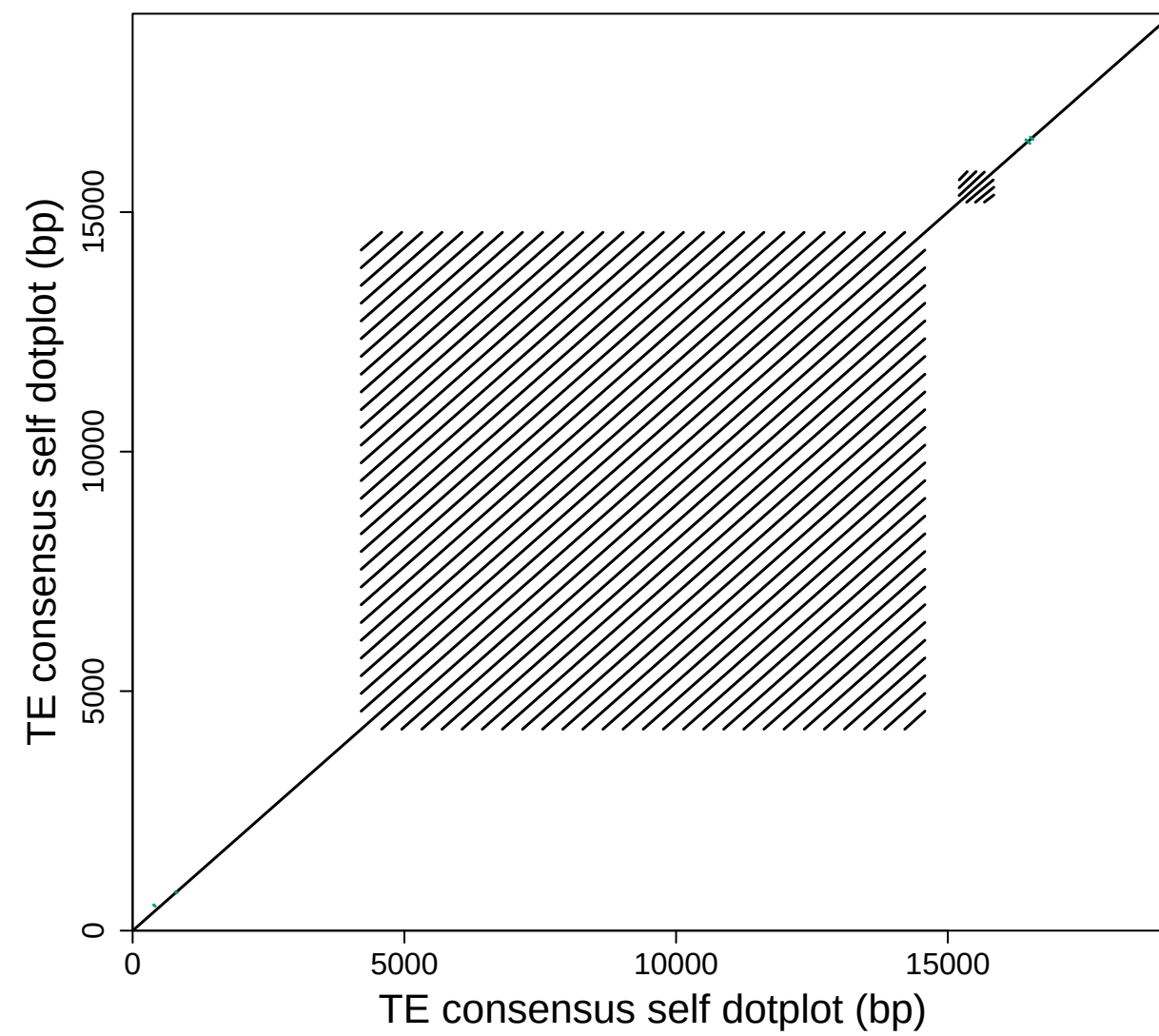
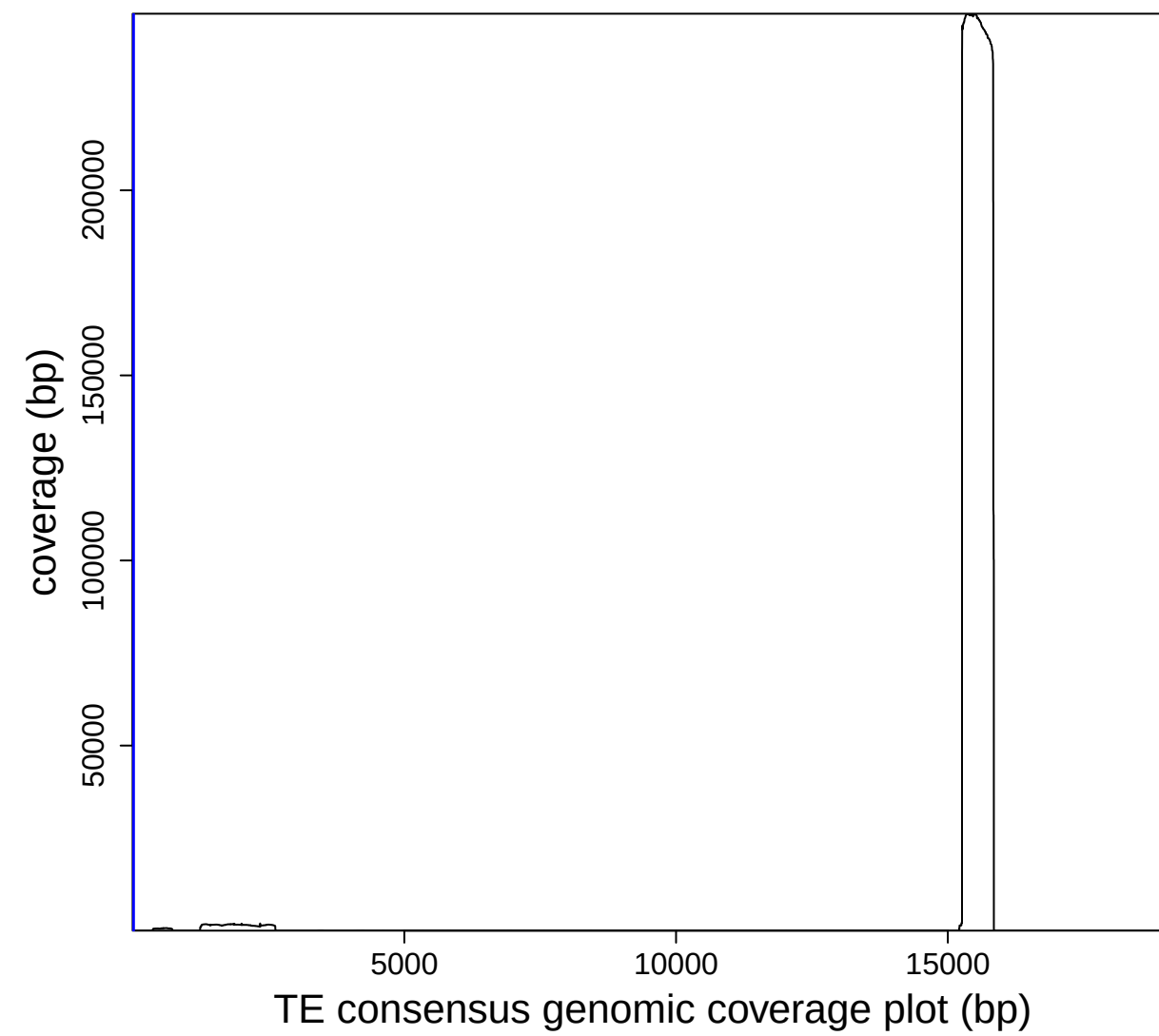
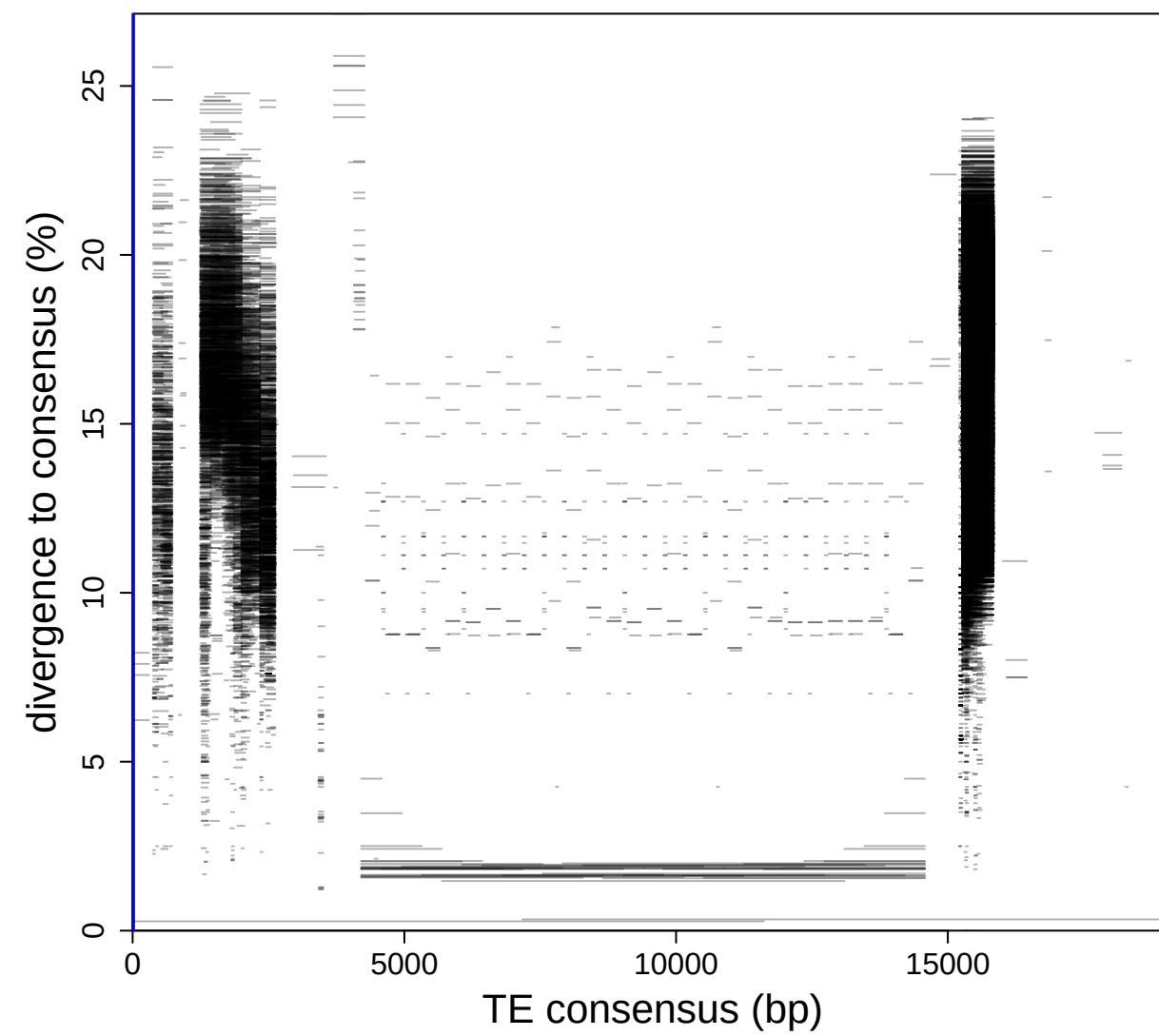
b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa  
size: 19134bp; fragments: 259227; full length: 0 (>=17220.6bp)

# After TEtrimmer 19134 bp



3.fasta.b.bed\_uf.bed\_g\_2.bed\_fm\_1.bed\_0\_0\_bcln.fa\_aln.fa\_cl.fa  
size: 19142bp; fragments: 259227; full length: 0 (>=17227.8bp)

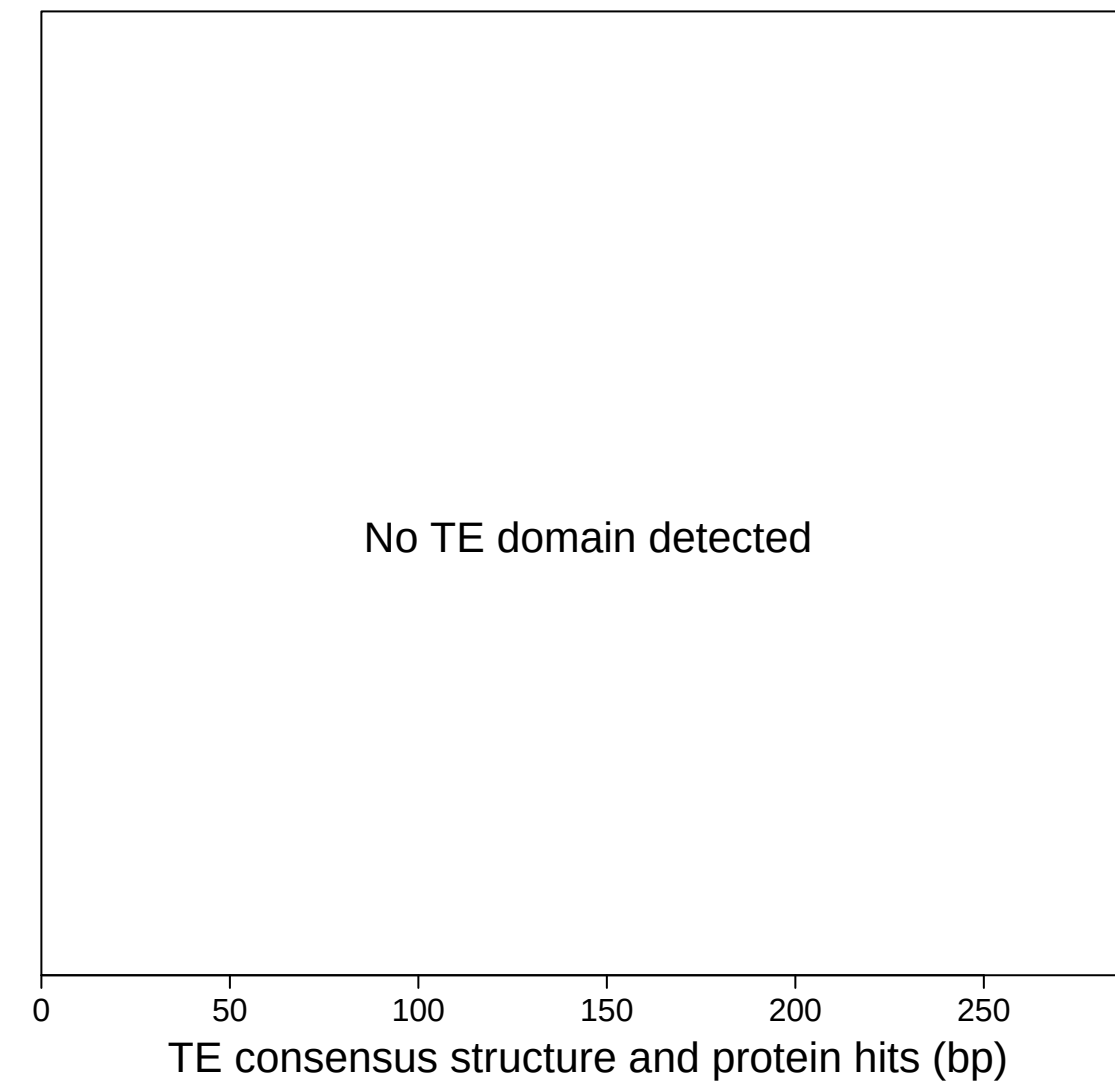
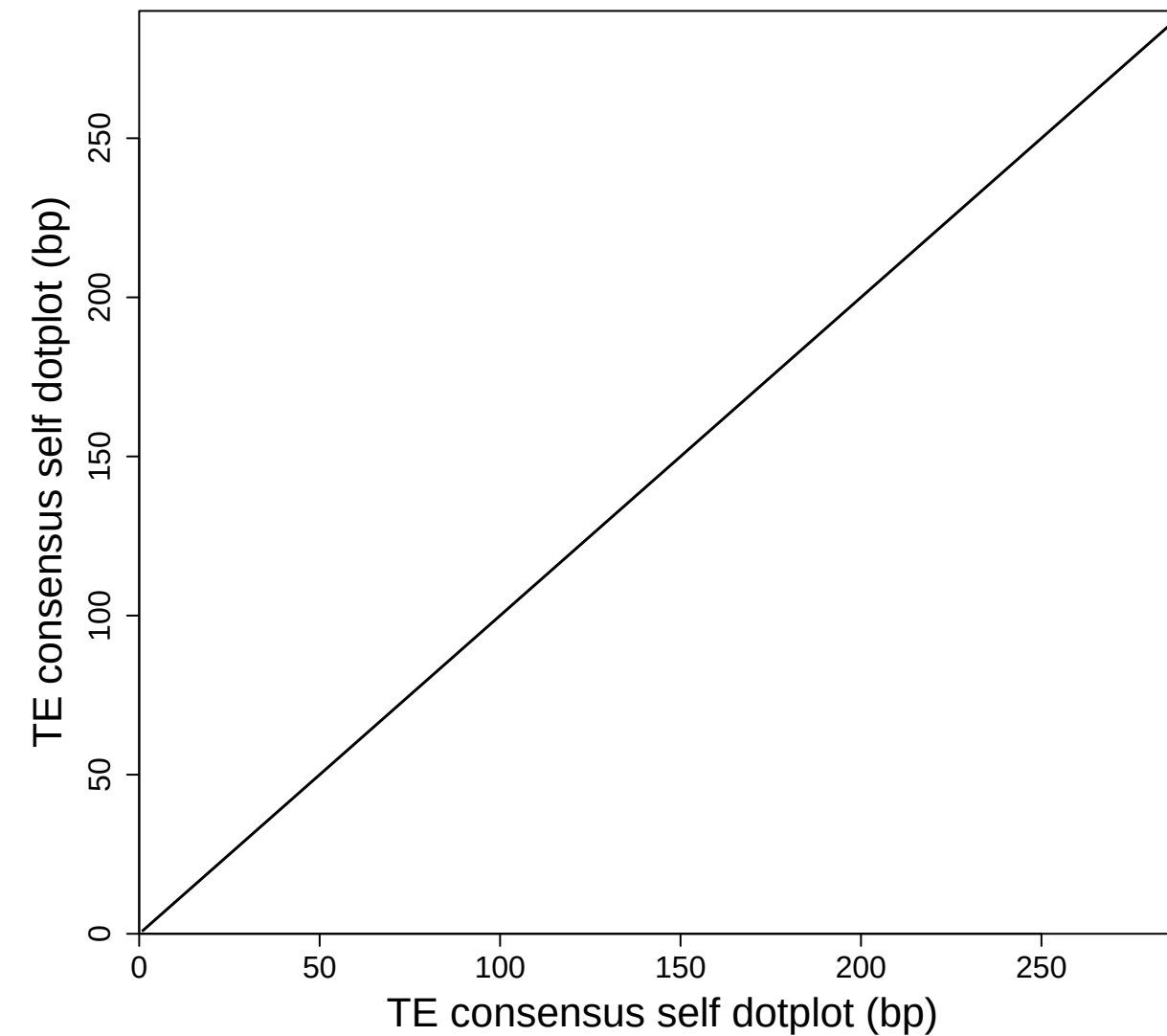
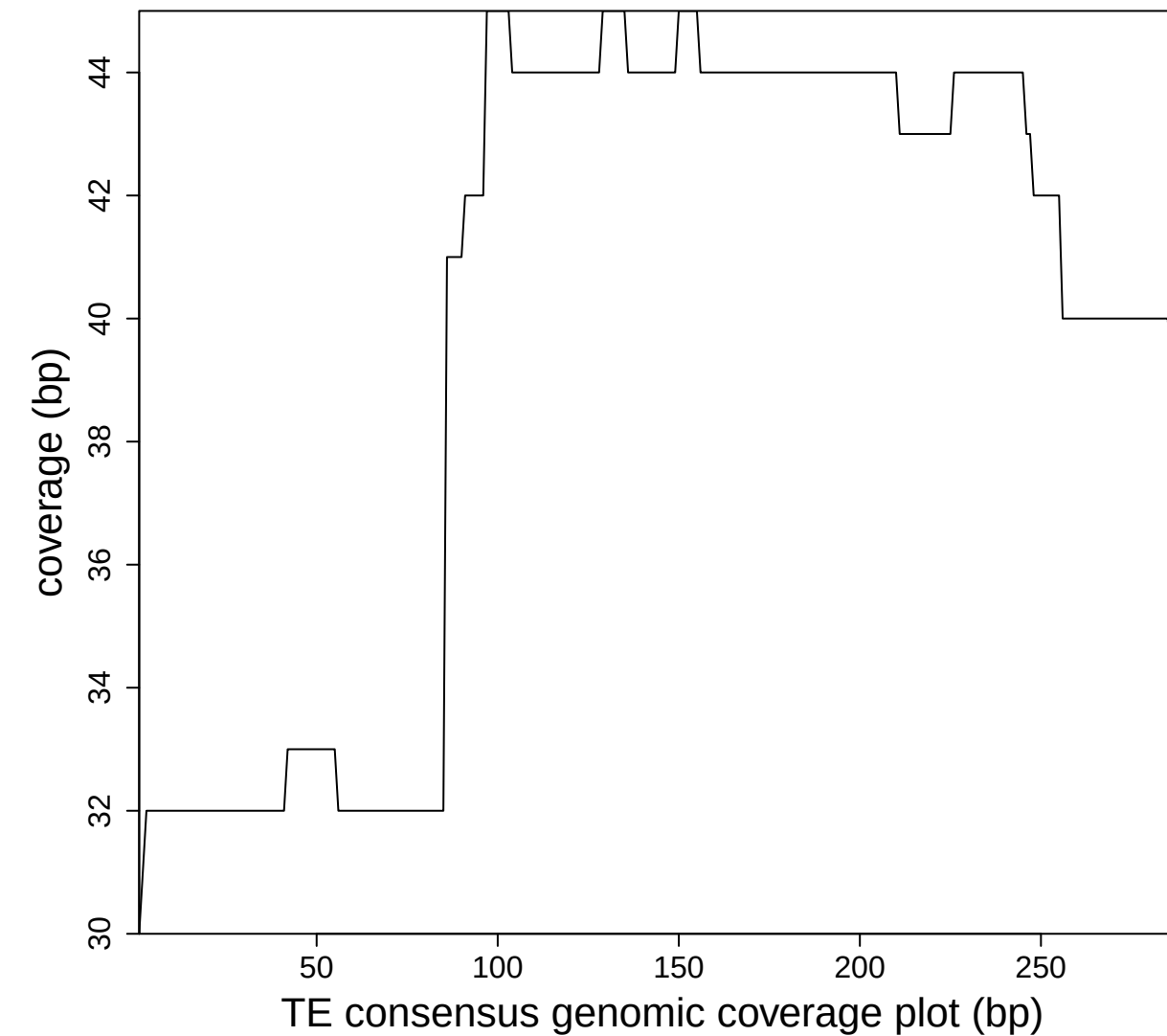
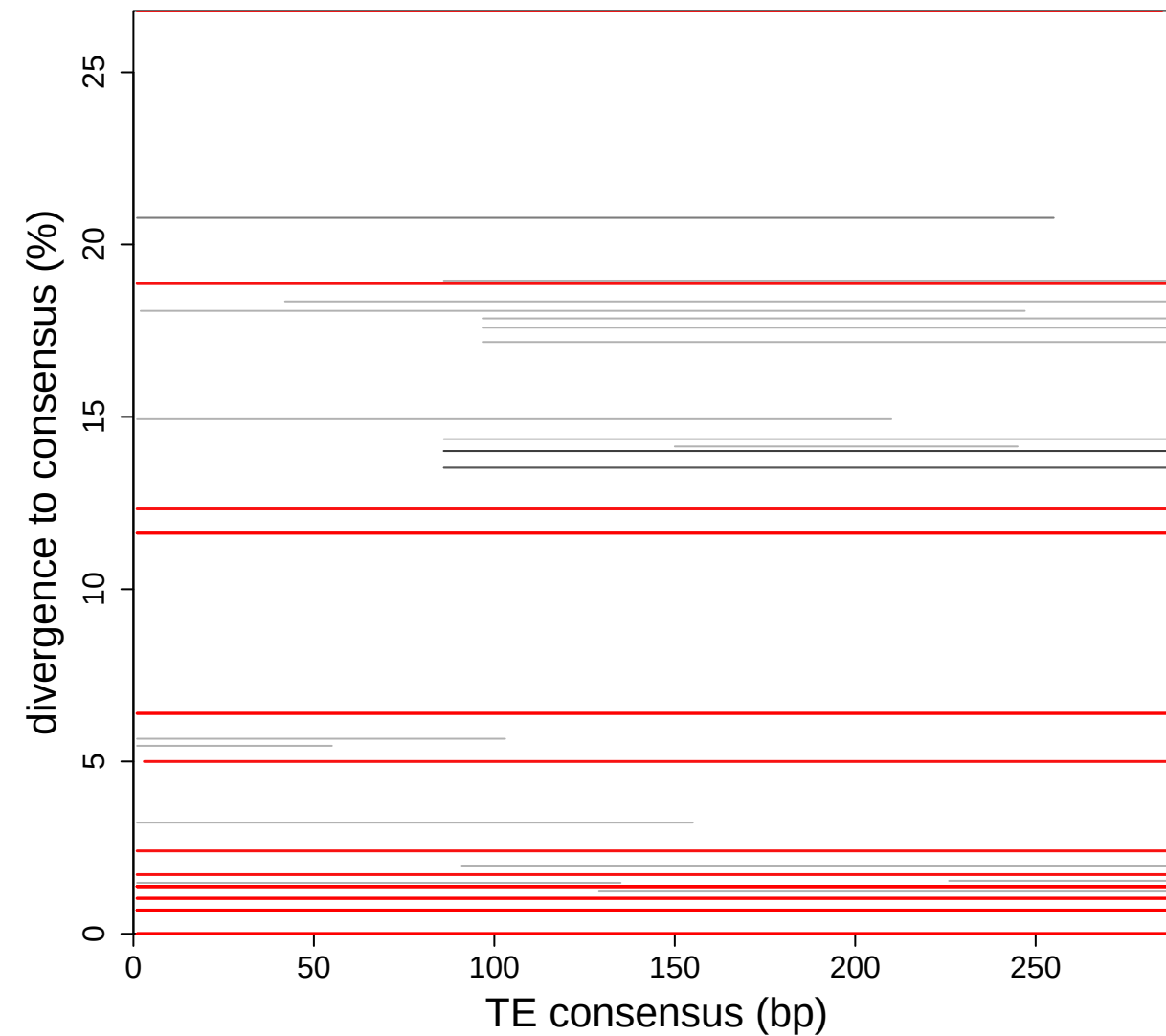
# After TEtrimmer Extended plot Blue lines are boundaries



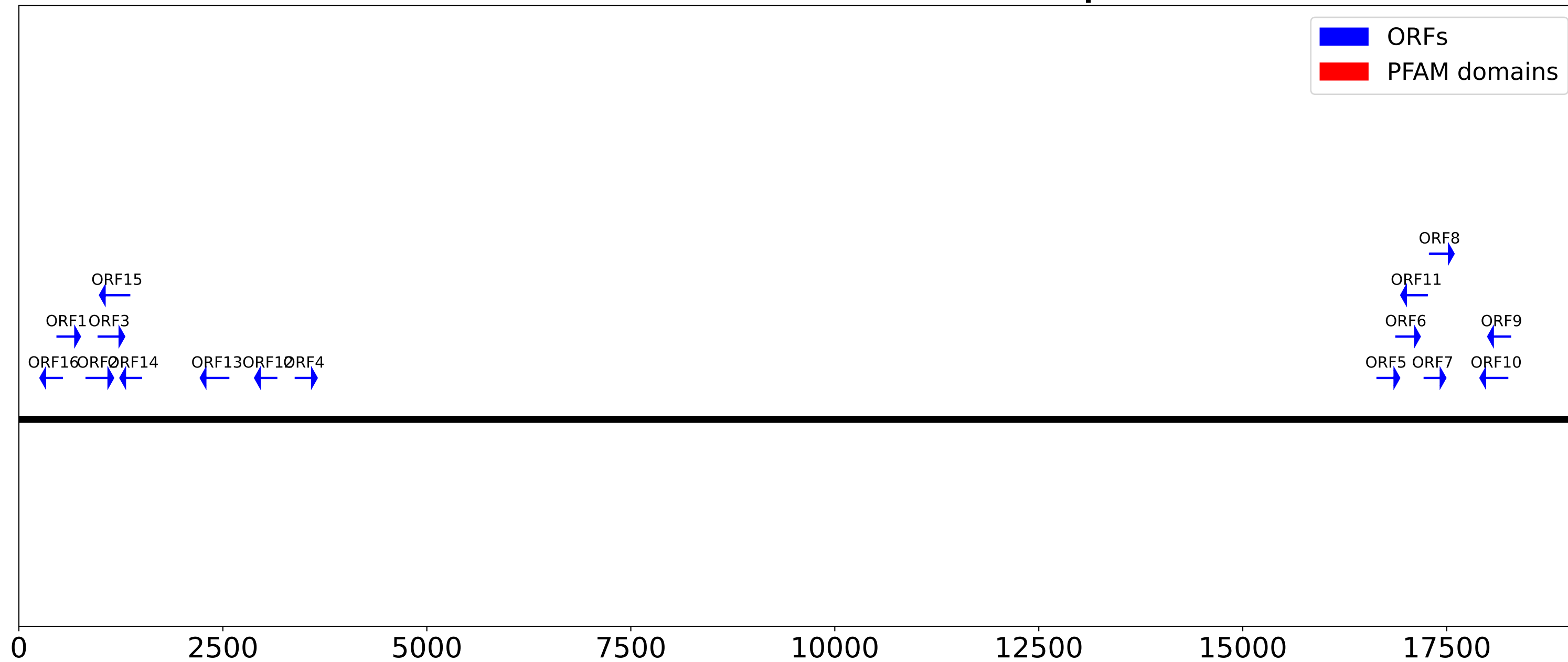
# Before TEtrimmer 290 bp

TE: ltr\_1\_family\_368

size: 290bp; fragments: 49; full length: 24 ( $\geq 261$ bp)



## After TEtrimmer ORF and PFAM domain plot

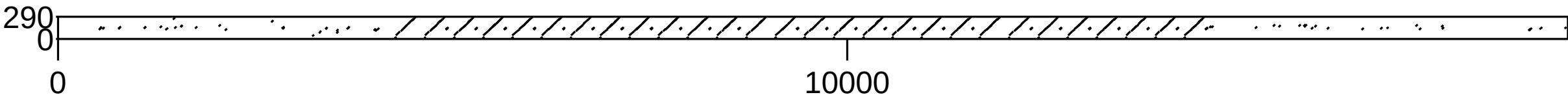




# Dotplot

(windowsize = 25, threshold = 50.00 09/10/25)

1sus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)