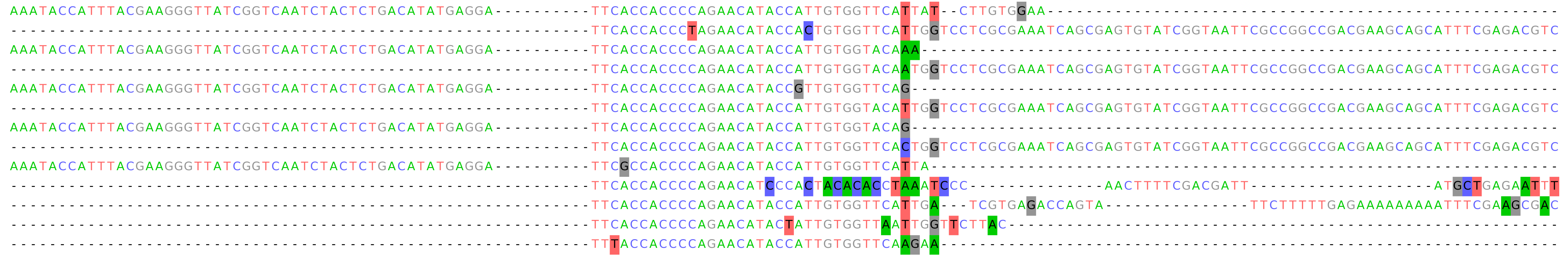
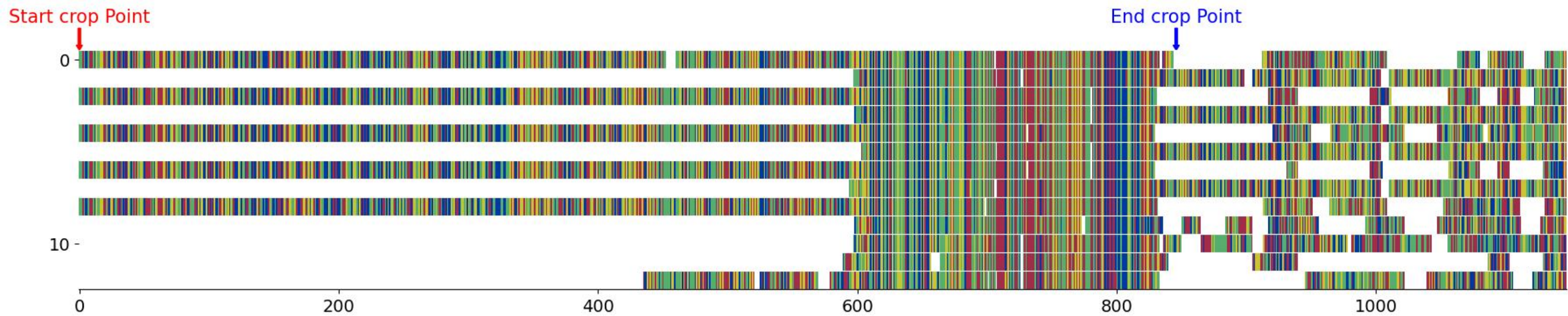


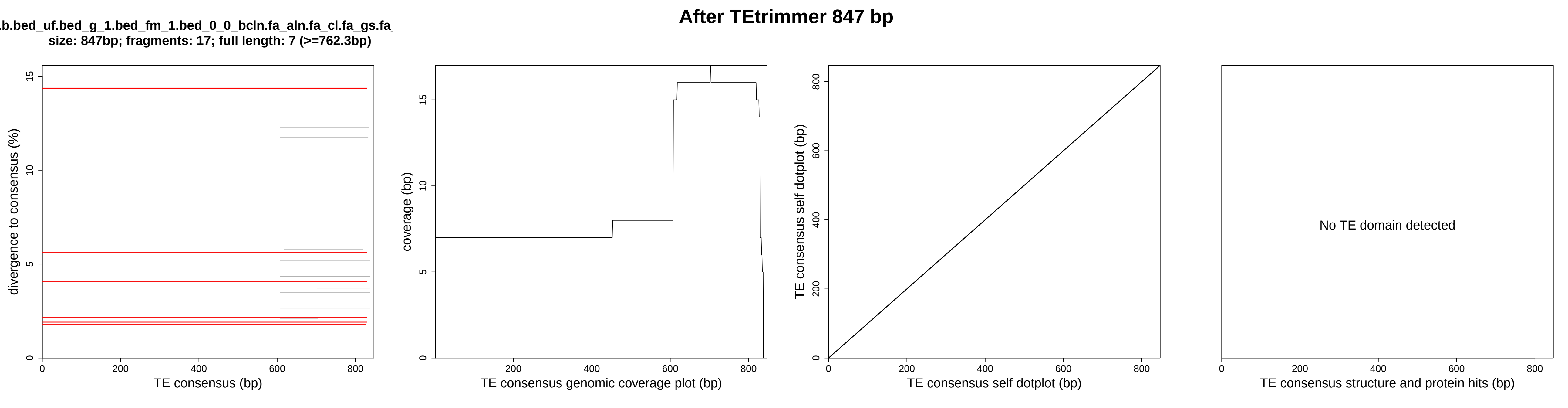
Start crop Point

End crop Point

MSA length = 847

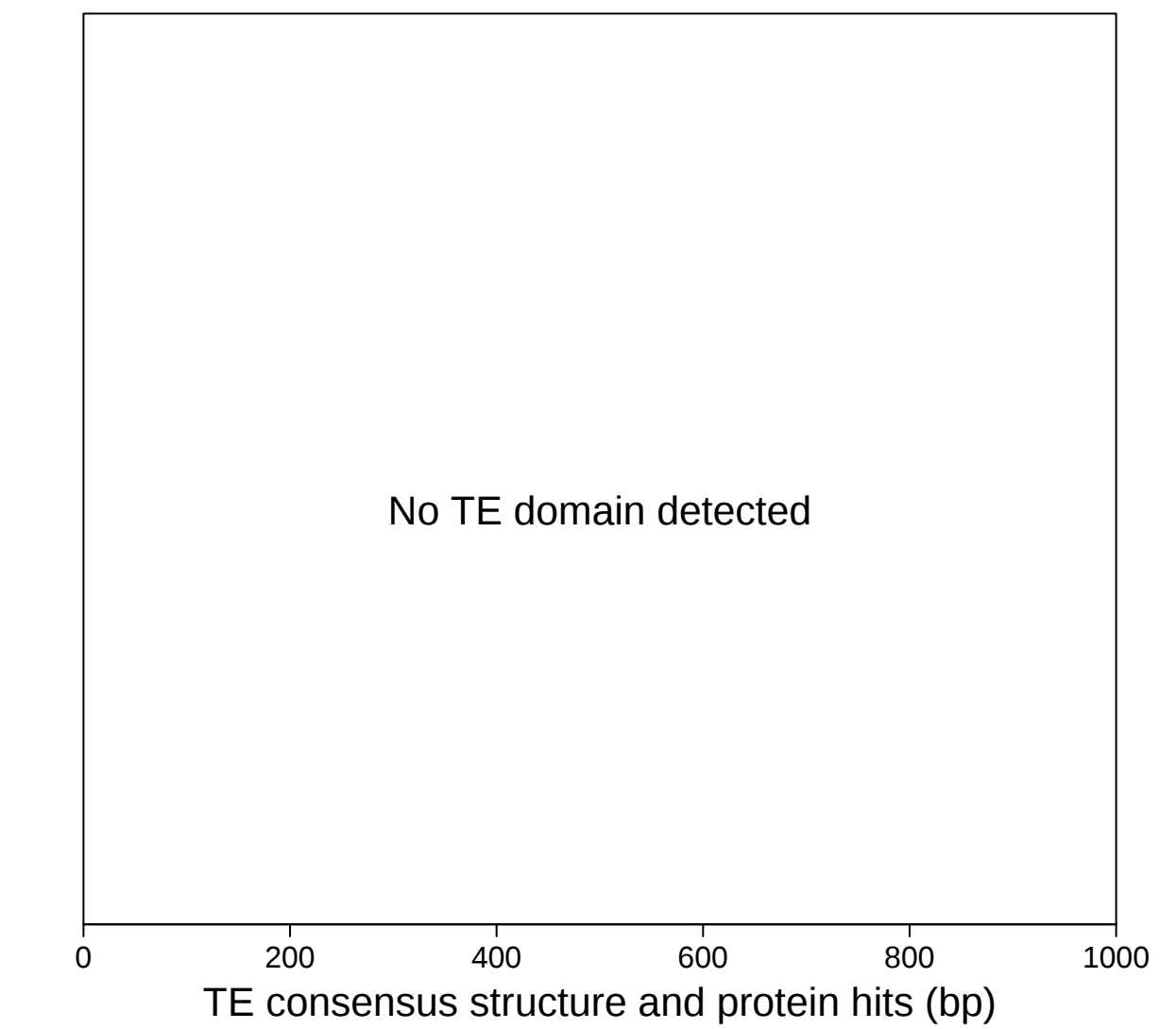
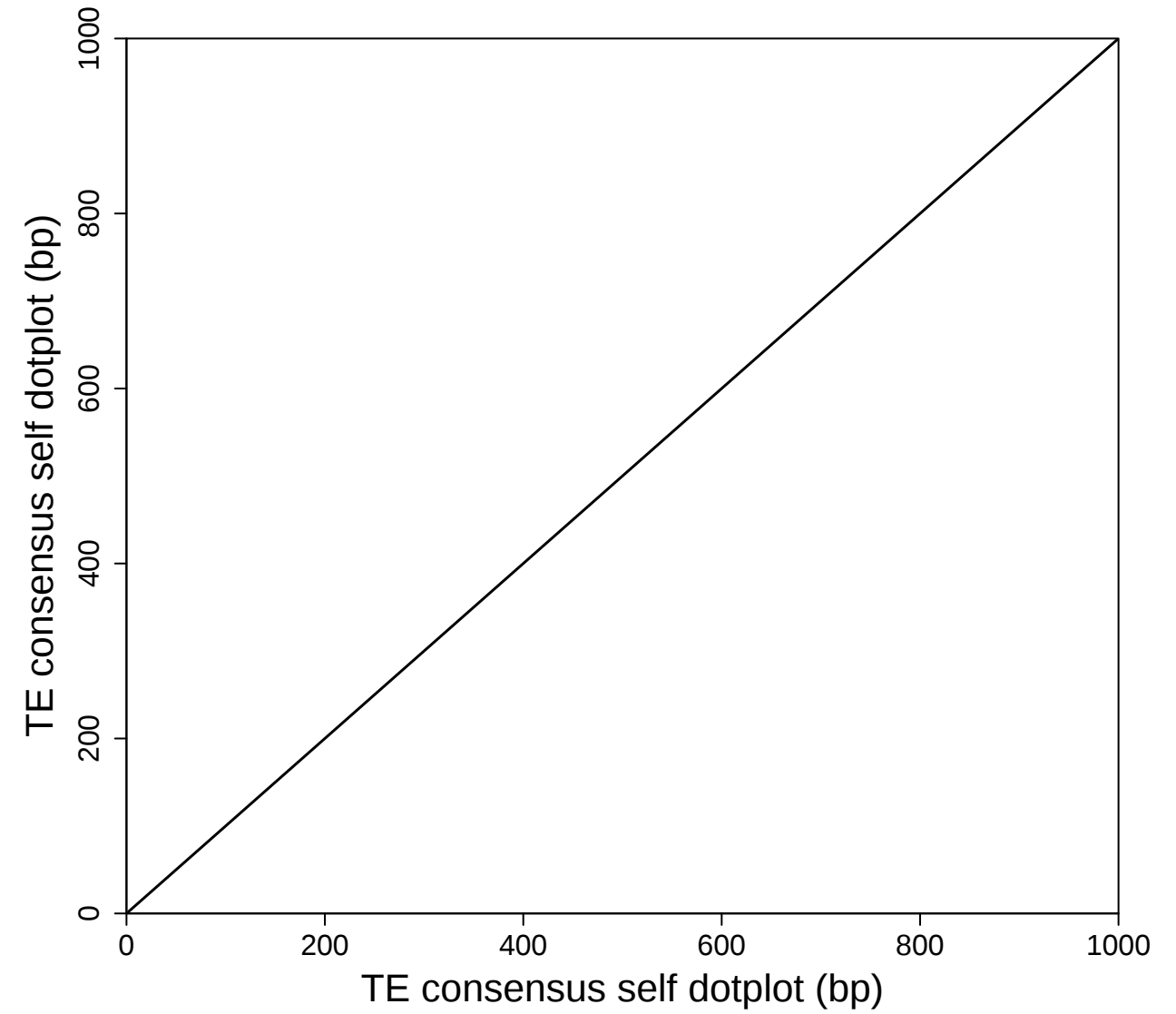
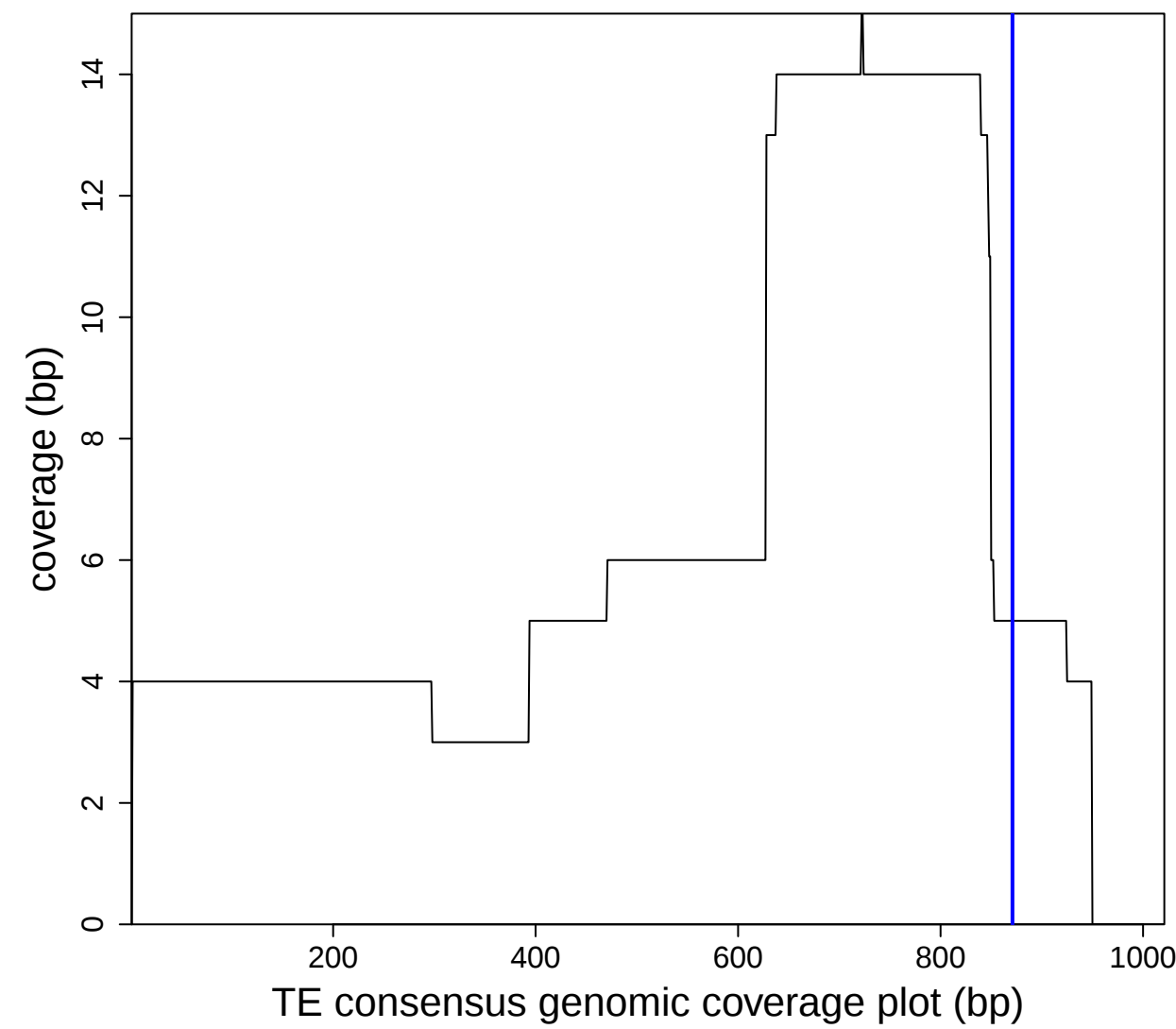
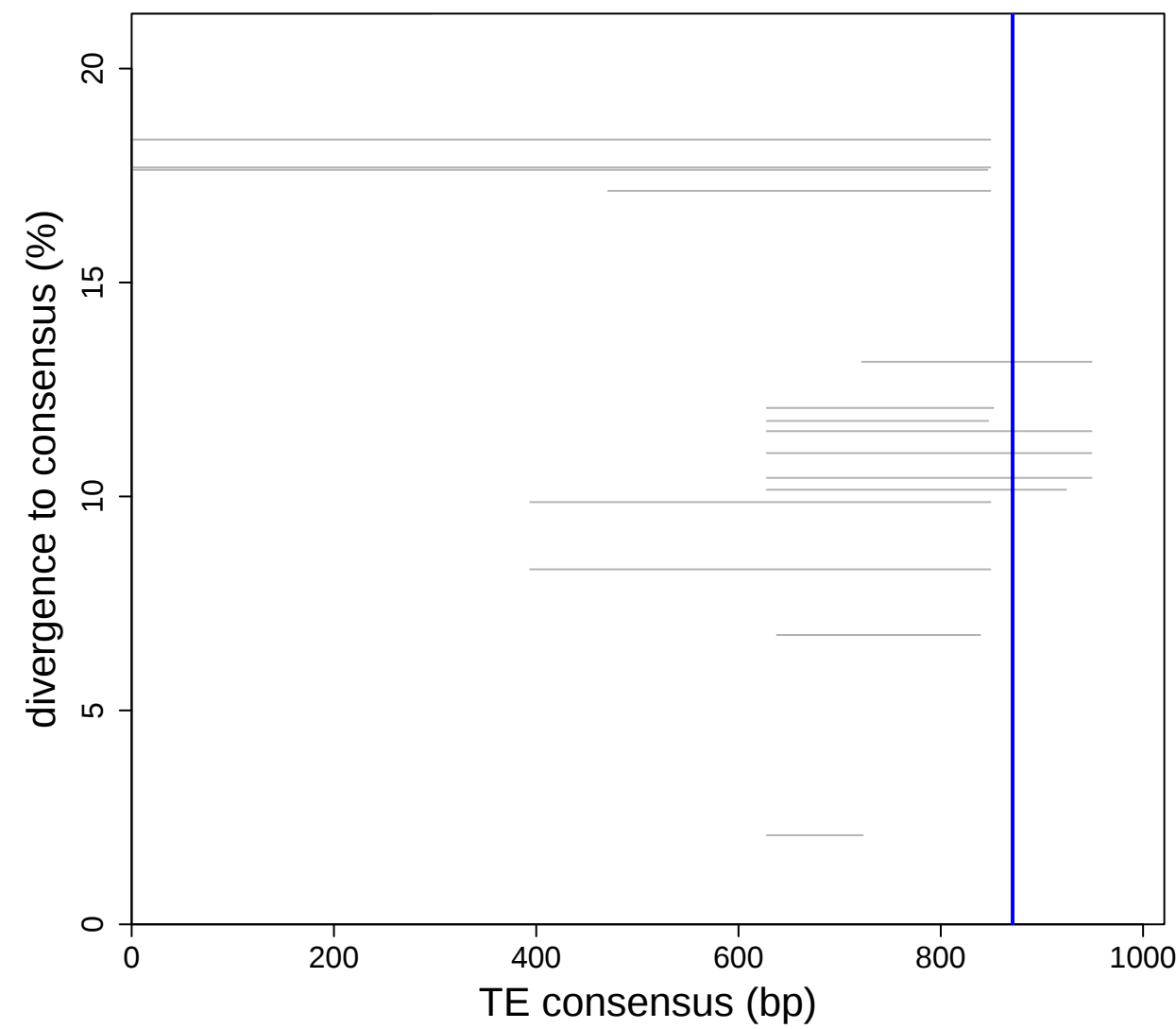




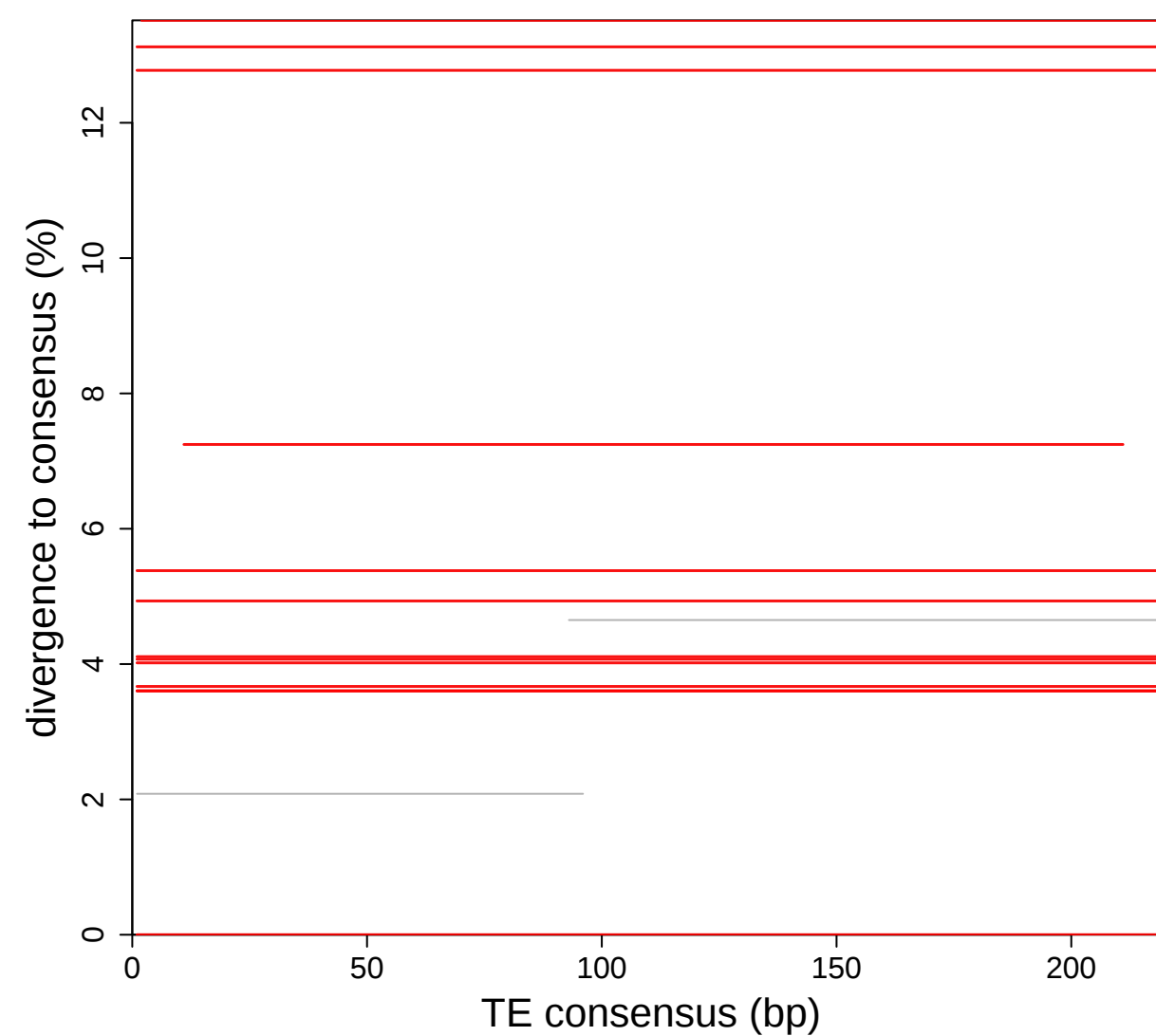


.fasta.b.bed\_uf.bed\_g\_1.bed\_fm\_1.bed\_0\_0\_bcln.fa\_aln.fa\_cl.fa  
size: 1021bp; fragments: 16; full length: 0 (>=918.9bp)

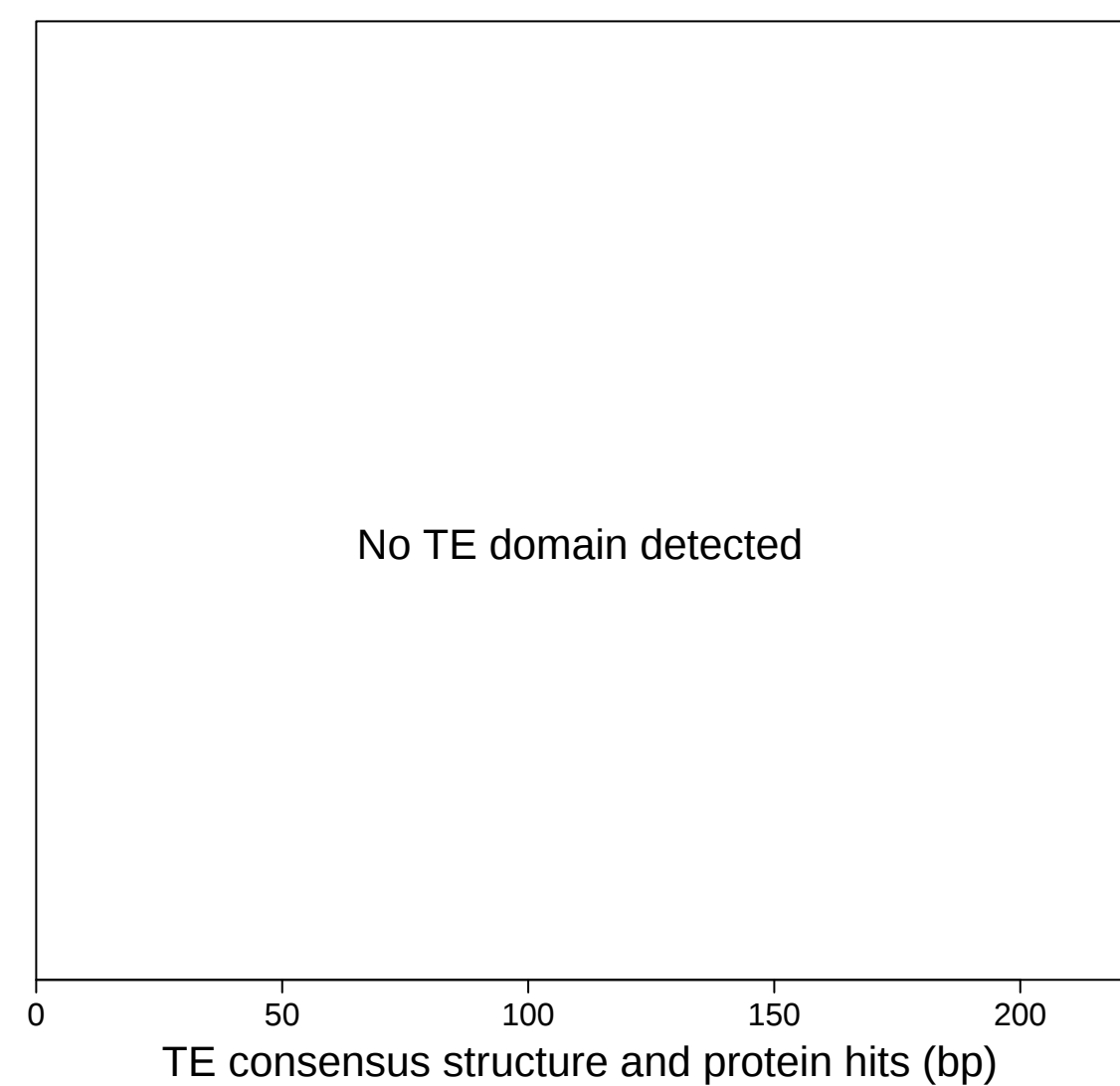
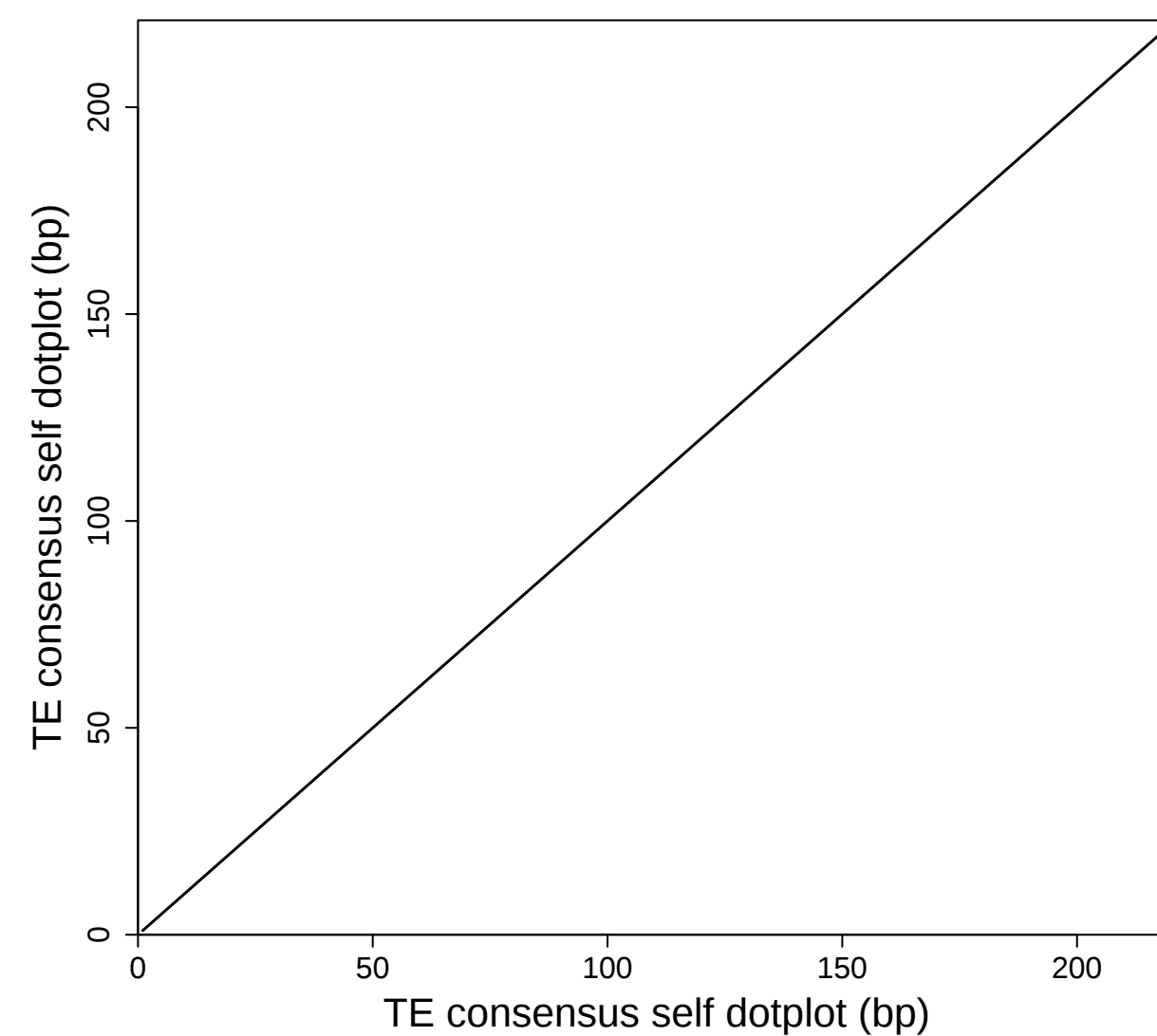
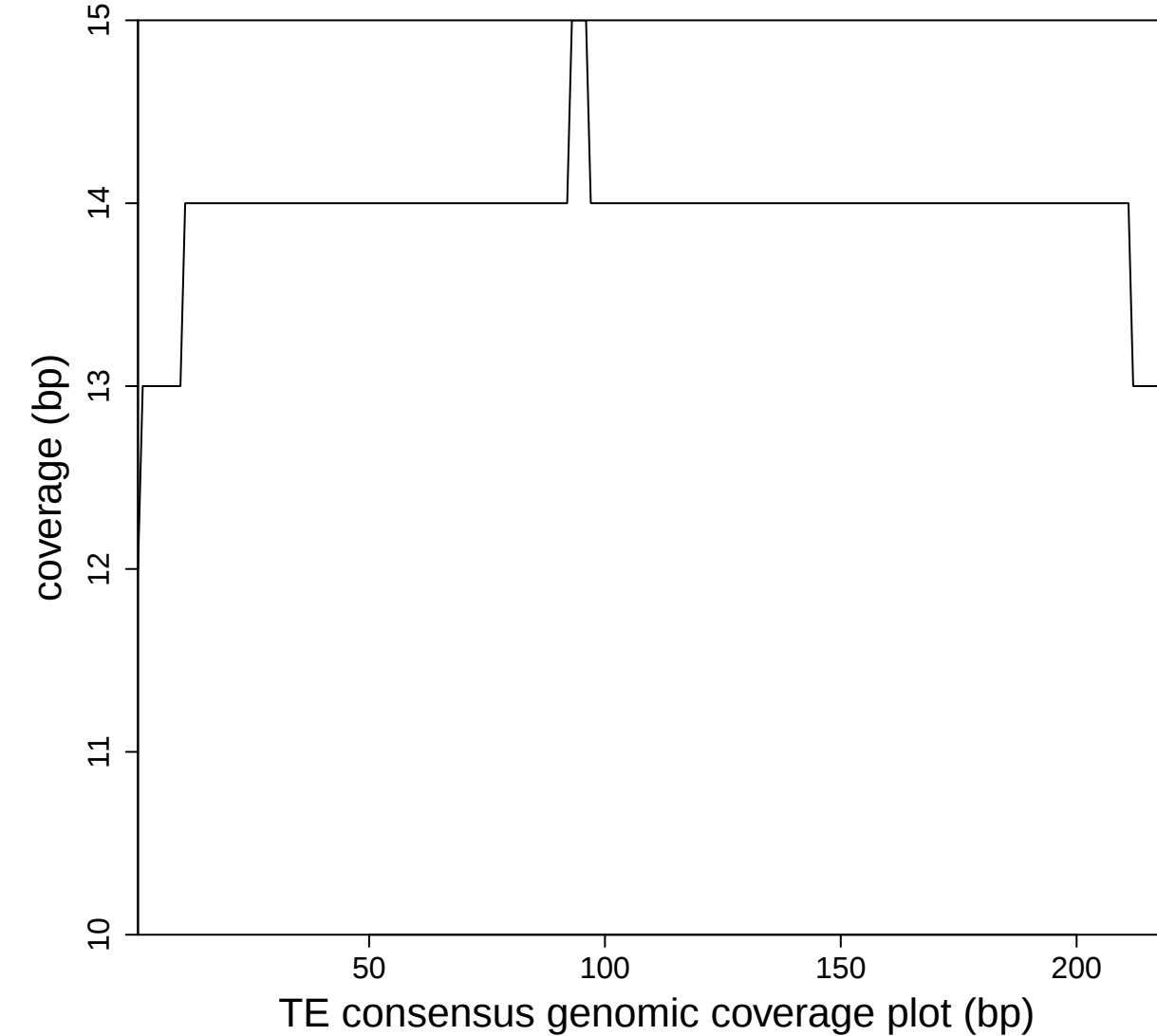
# After TEtrimmer Extended plot Blue lines are boundaries



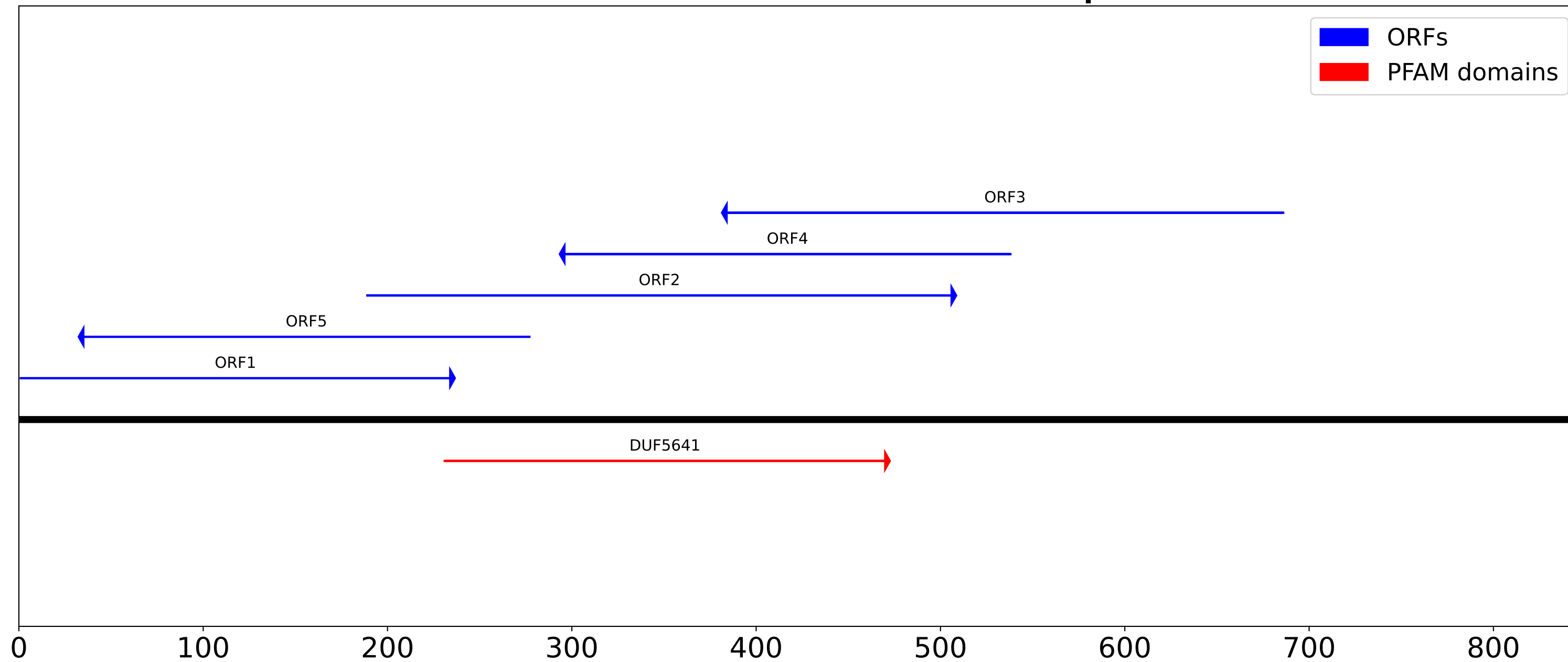
TE: ltr\_1\_family\_270  
size: 221bp; fragments: 15; full length: 13 ( $\geq 198.9$ bp)



## Before TEtrimmer 221 bp



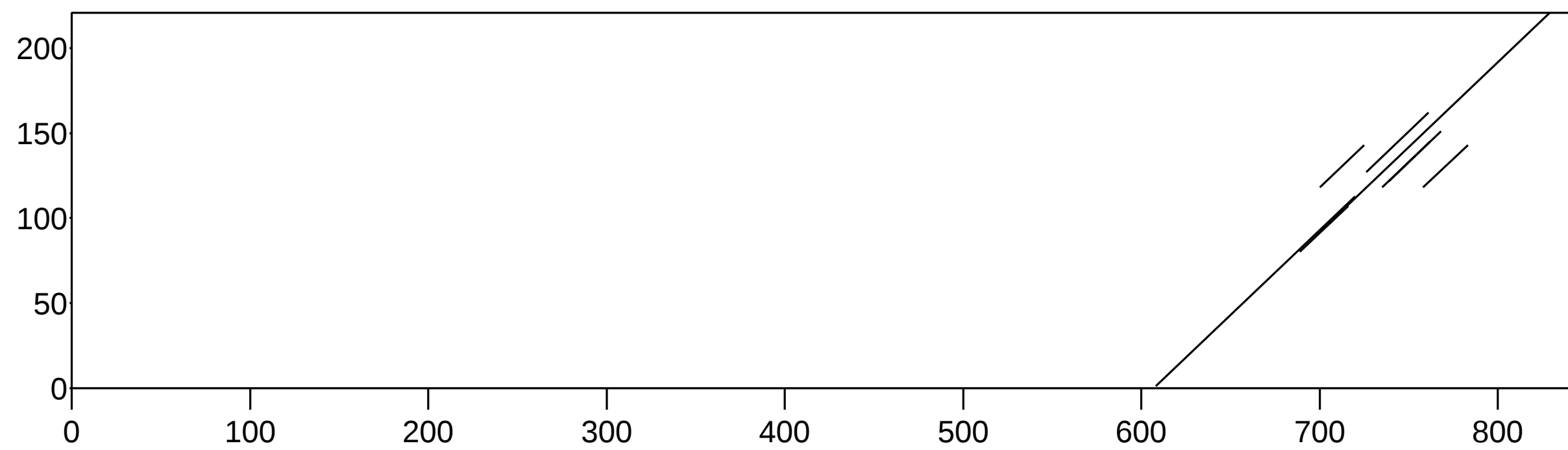
## After TEtrimmer ORF and PFAM domain plot



# Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)