

Start crop Point



End crop Point

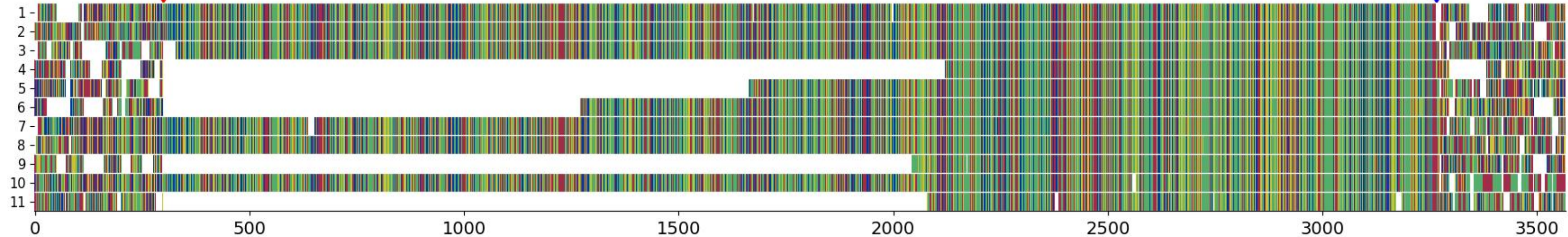


MSA length = 2967



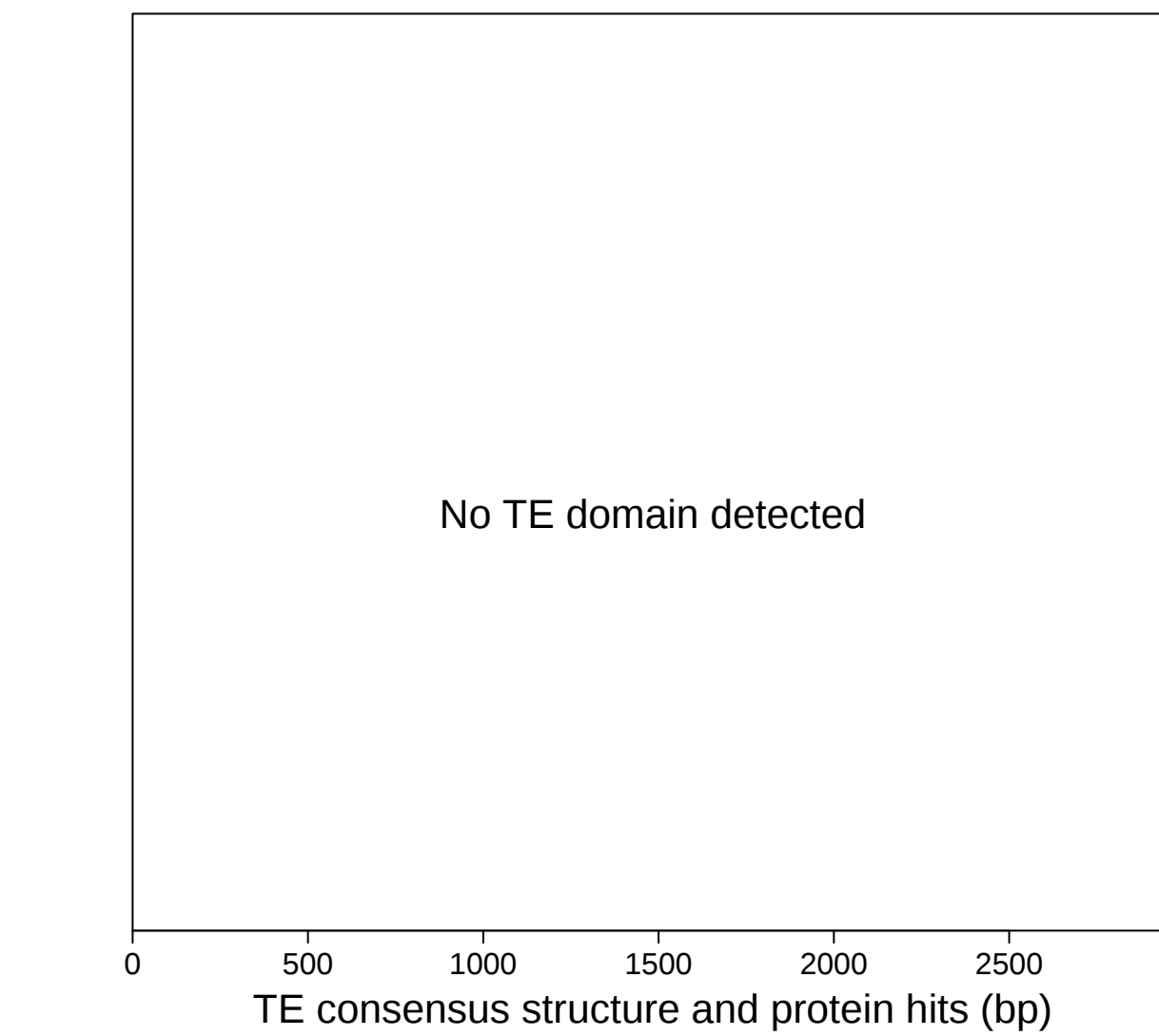
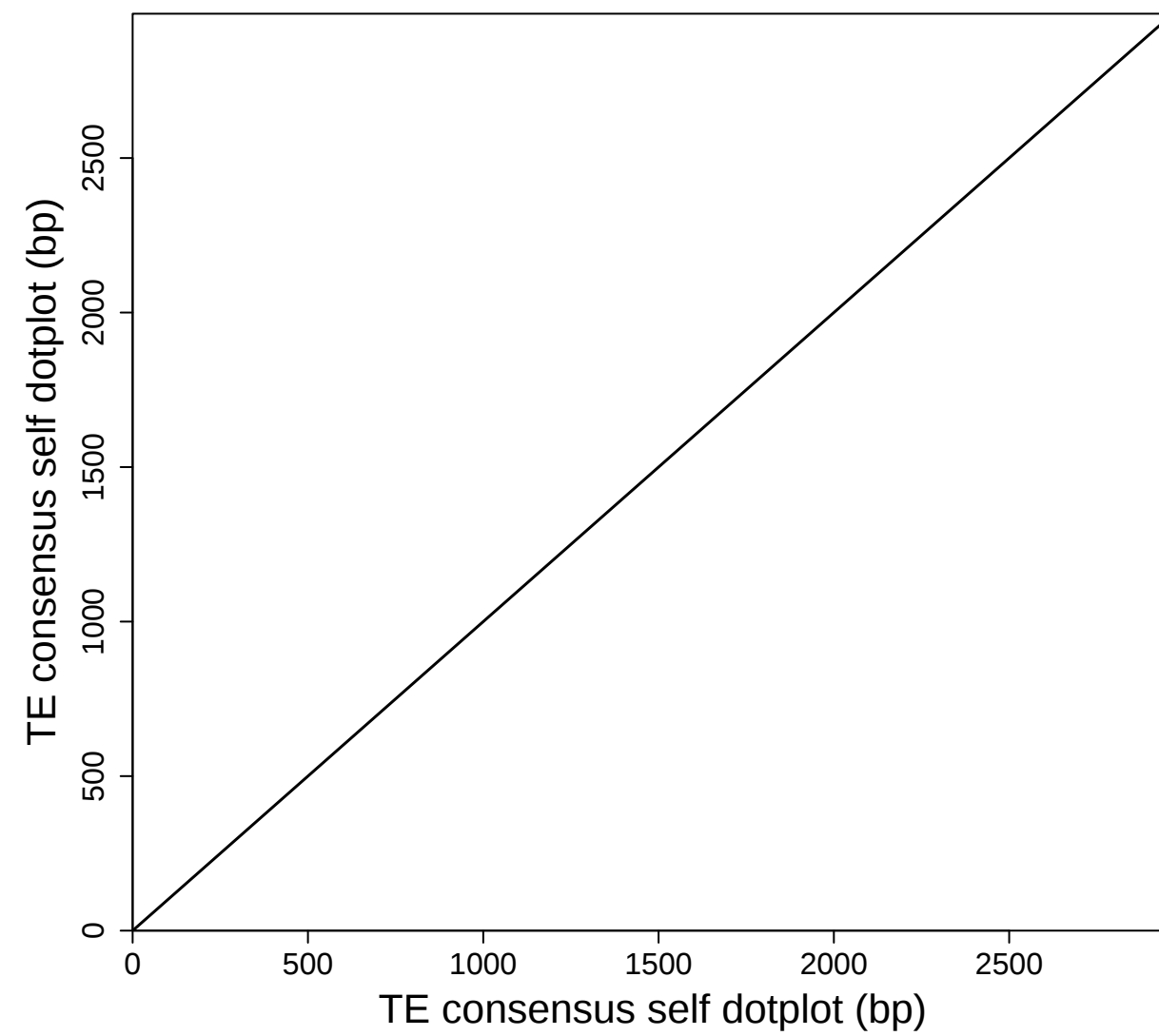
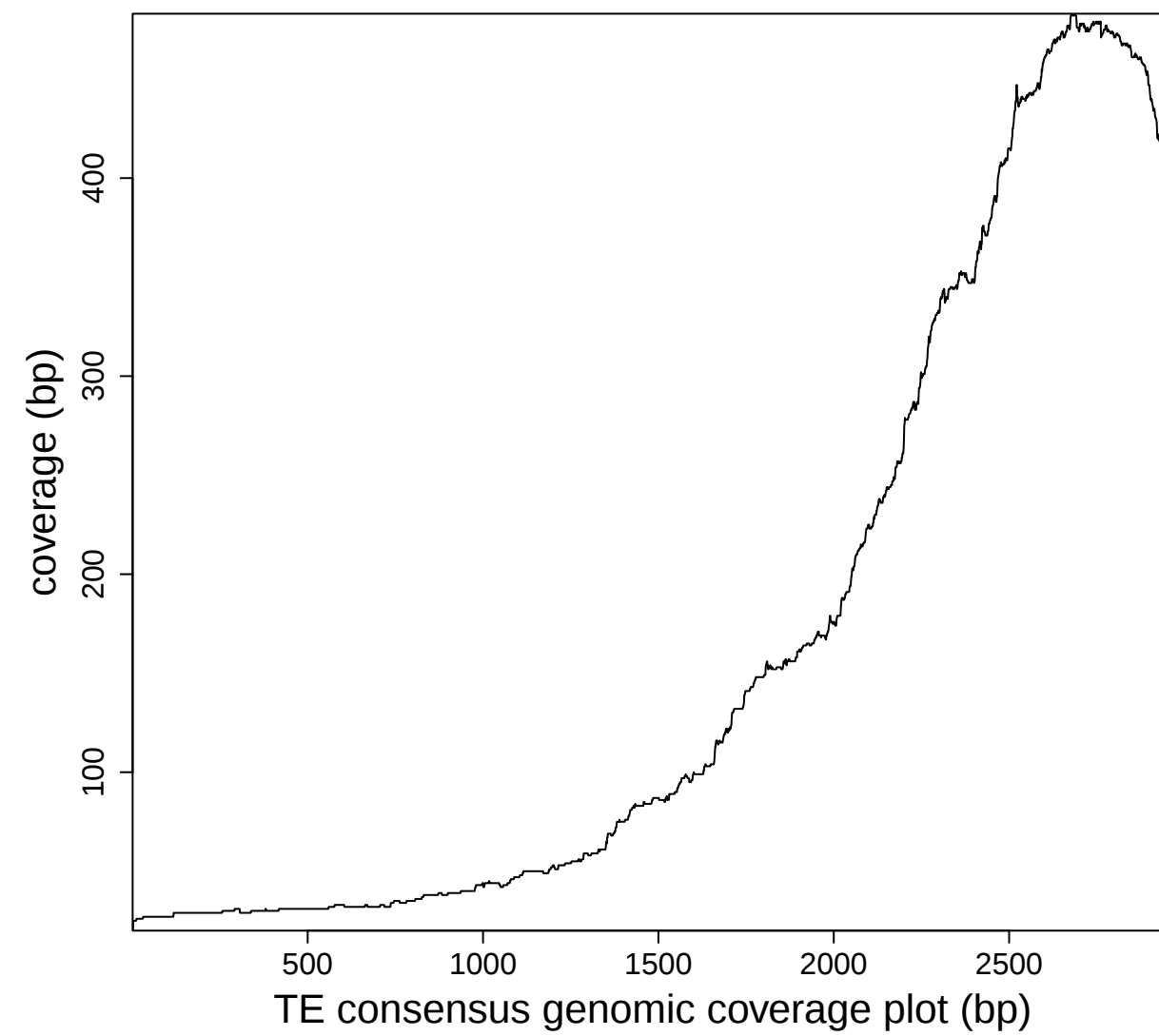
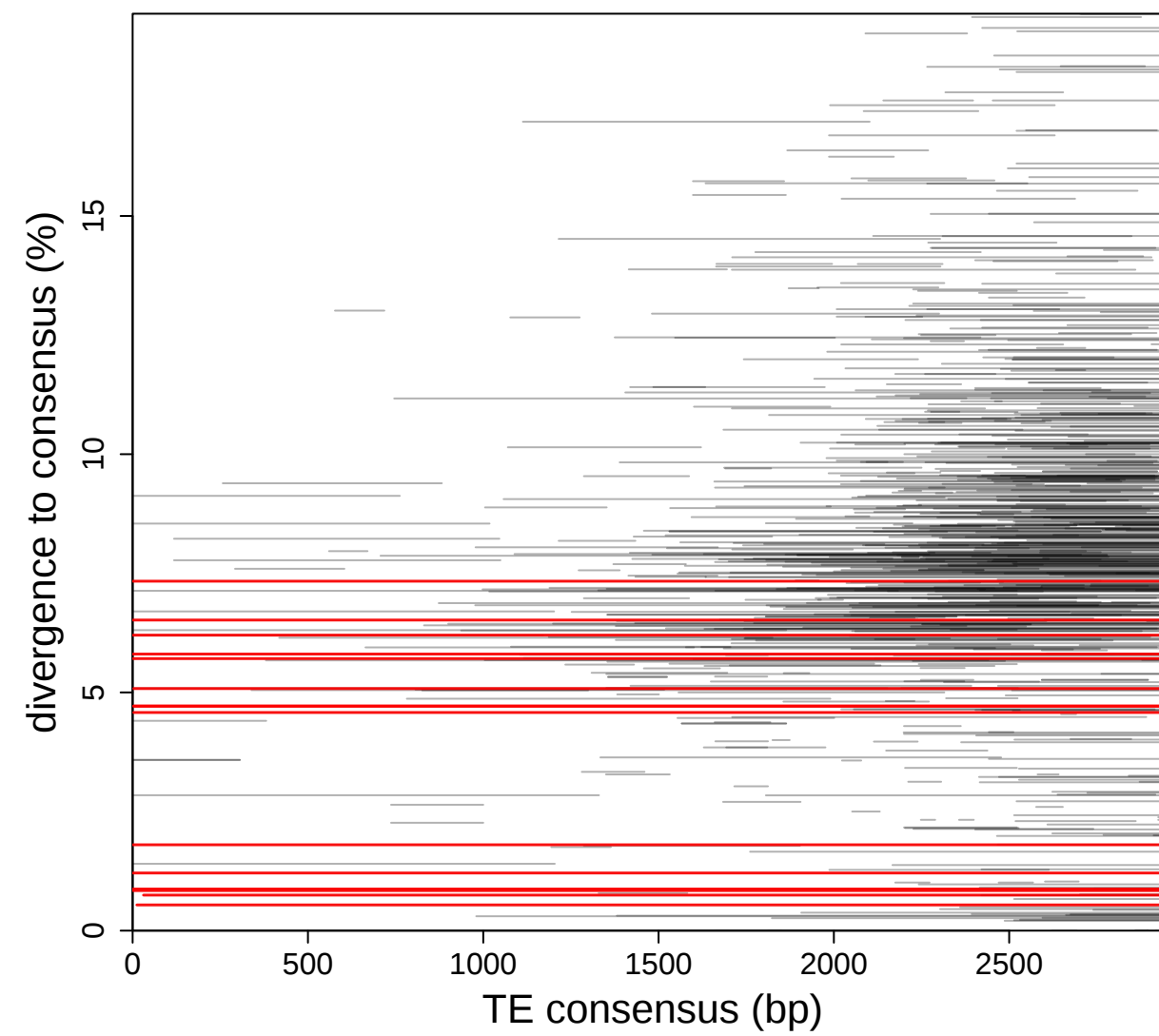
Start crop Point

End crop Point



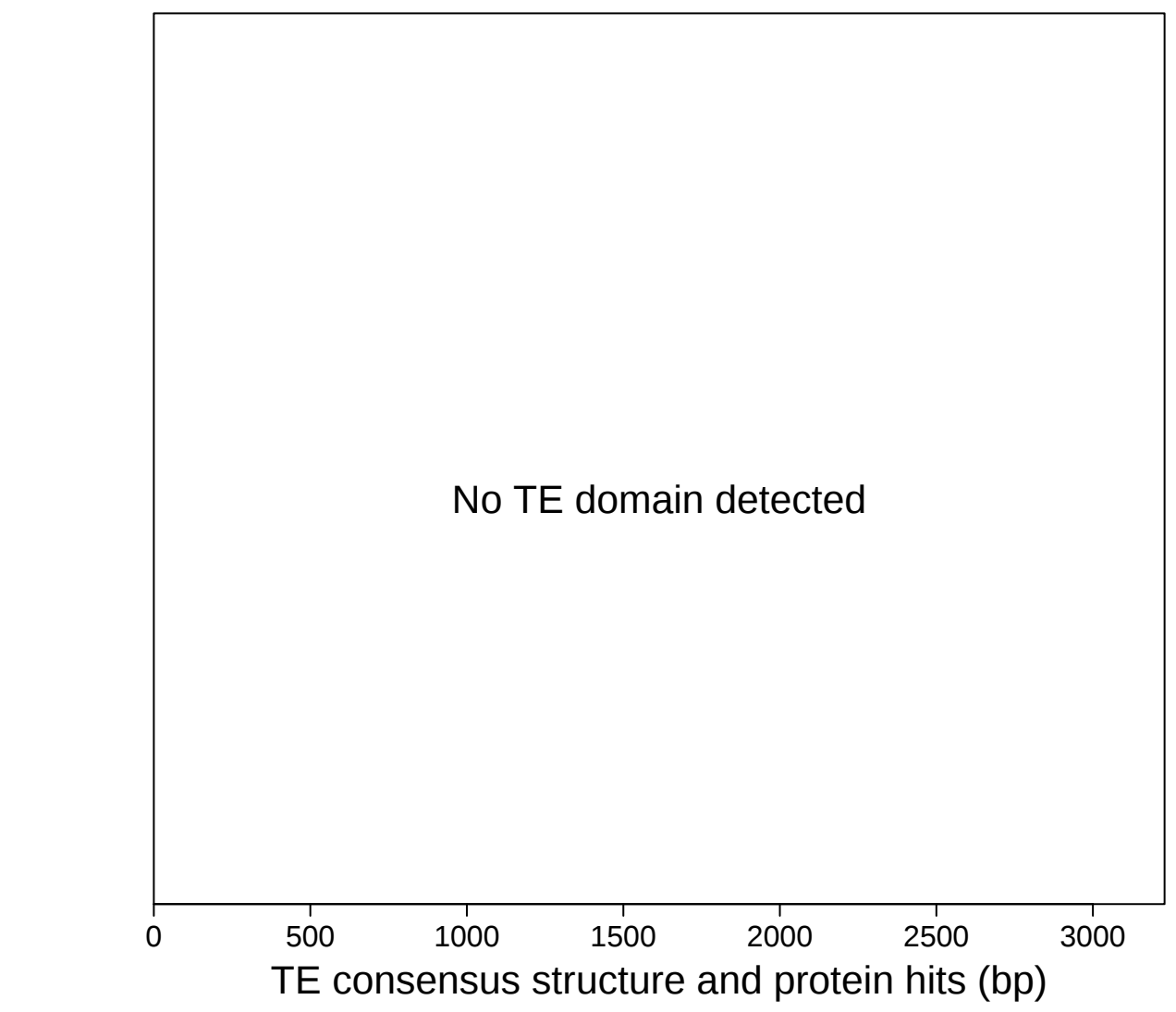
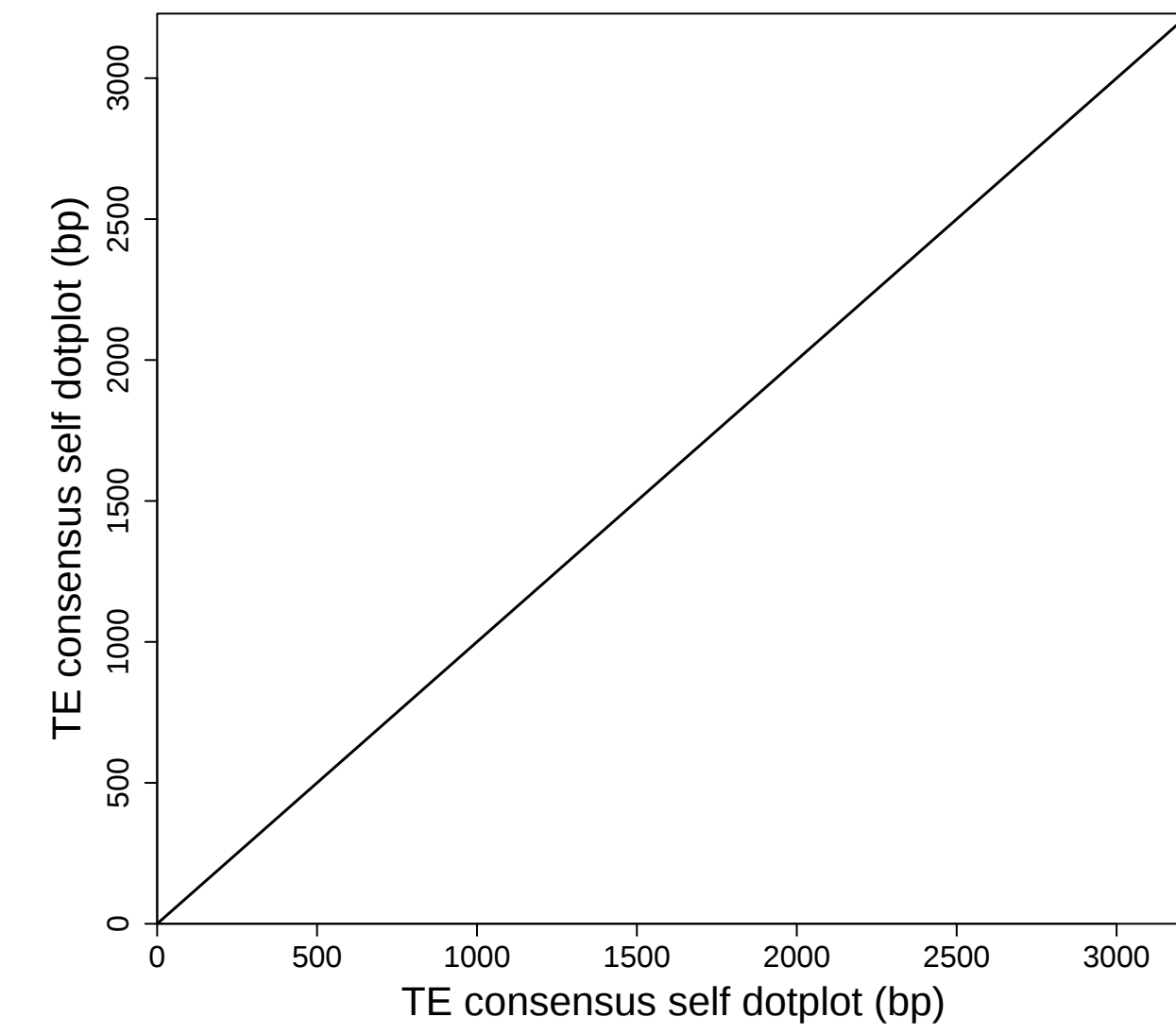
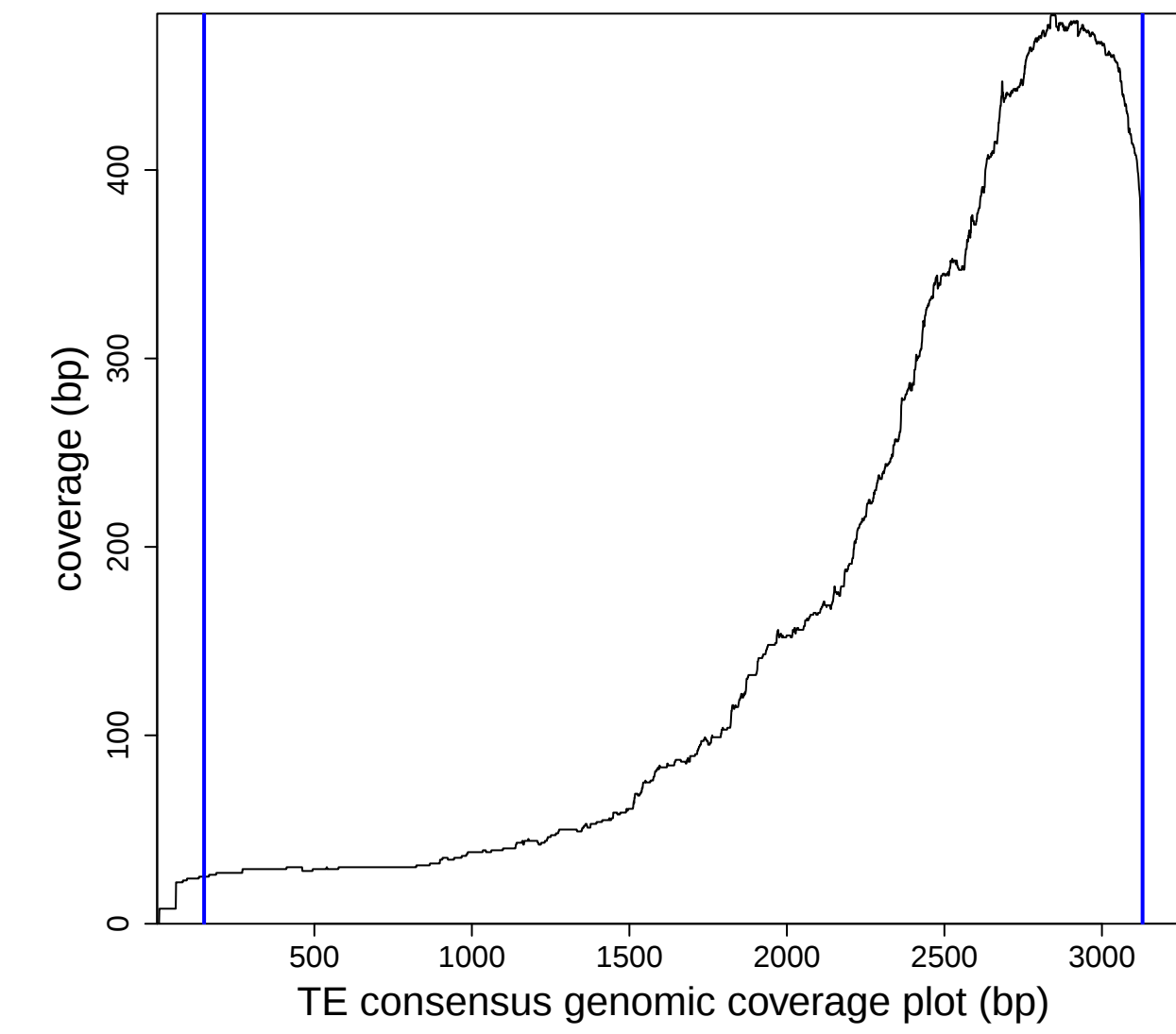
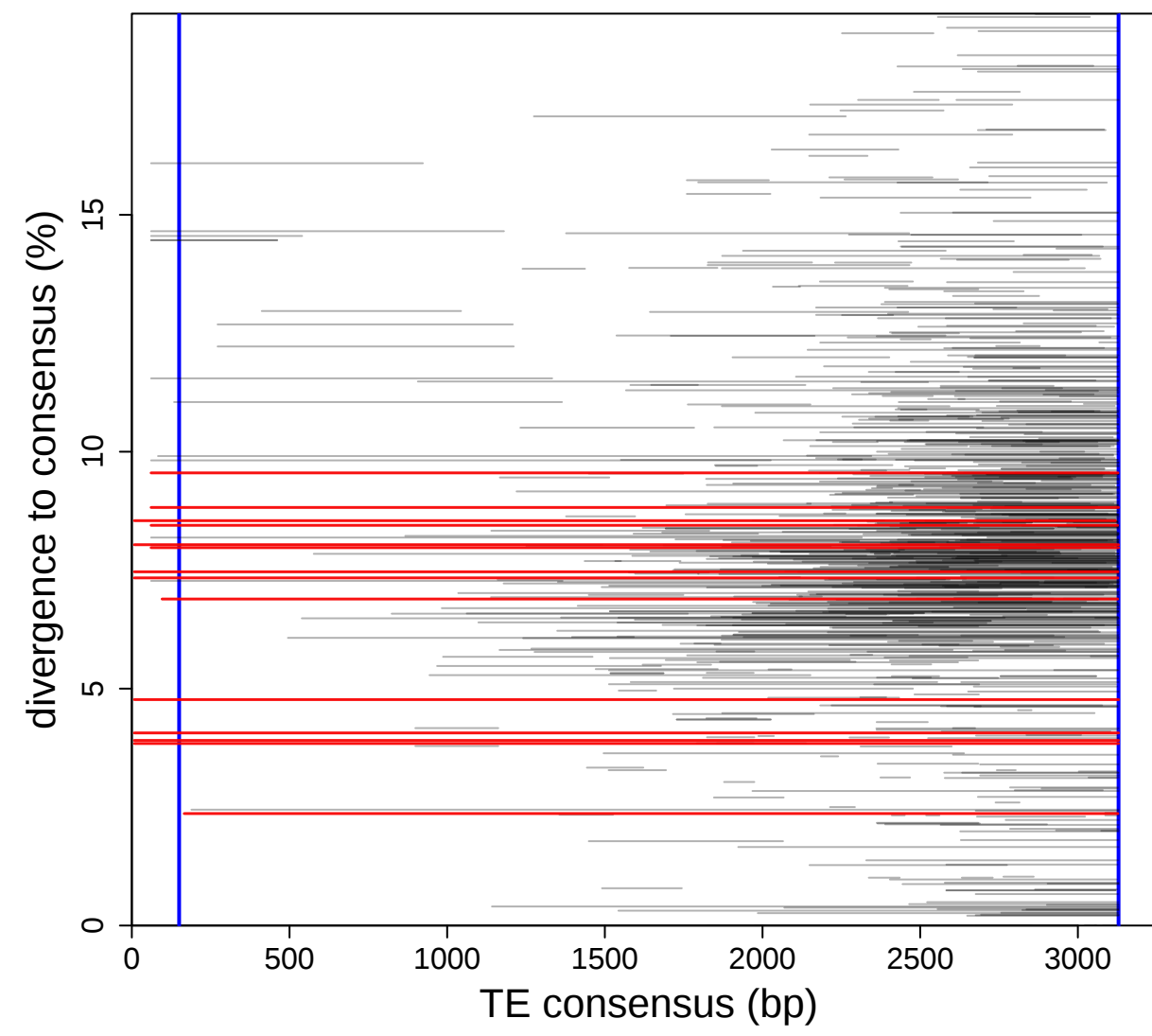
bed\_fm\_1.bed\_0\_0\_n.bed\_g\_1.bed\_fm\_2.bed\_0\_0\_bcln.fa\_aln.fa  
size: 2967bp; fragments: 766; full length: 15 ( $\geq 2670.3$ bp)

# After TEtrimmer 2967 bp



f.bed\_g\_3.bed\_fm\_1.bed\_0\_0\_n.bed\_g\_1.bed\_fm\_2.bed\_0\_0\_bcl  
size: 3279bp; fragments: 763; full length: 14 (>=2951.1bp)

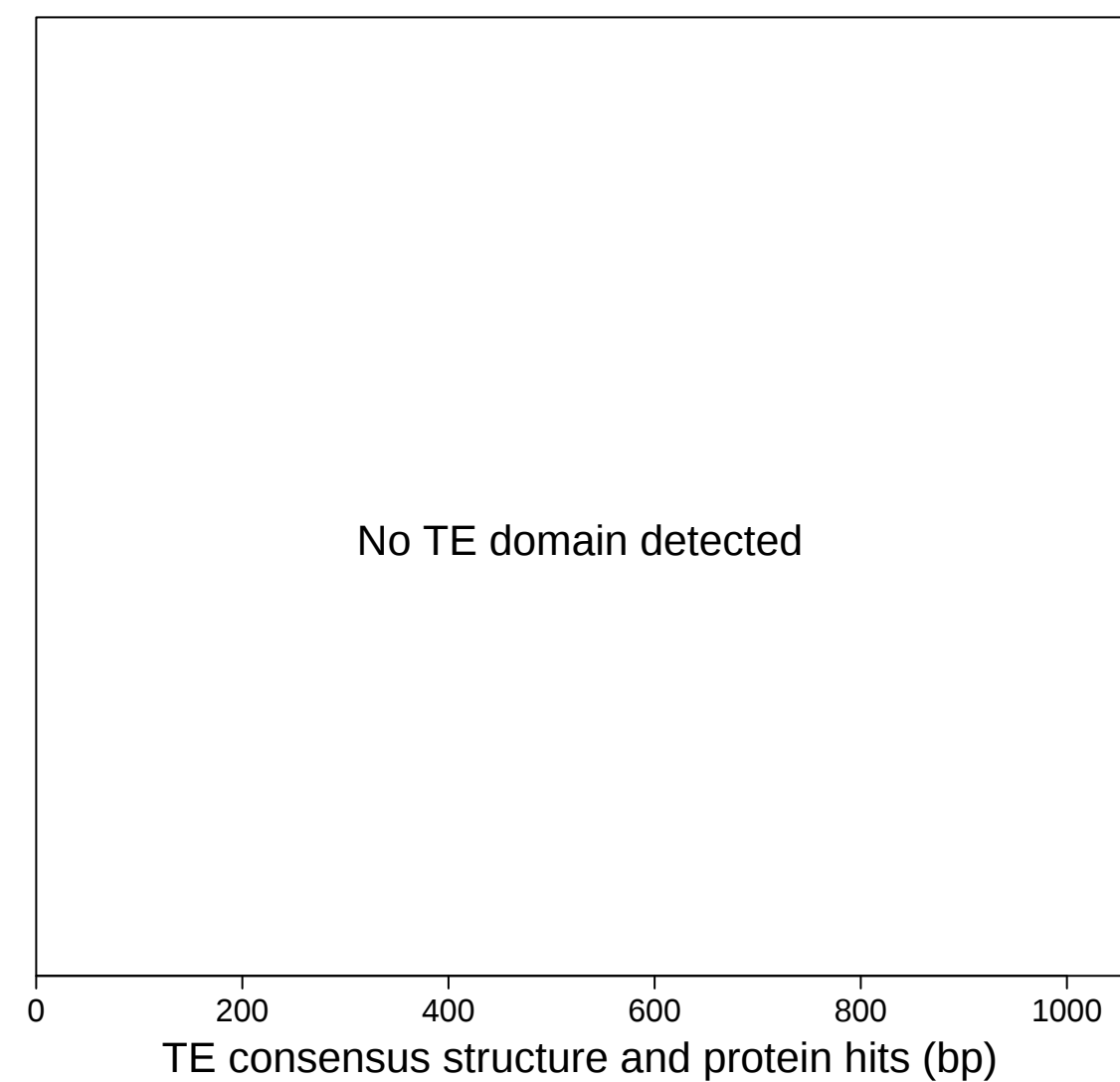
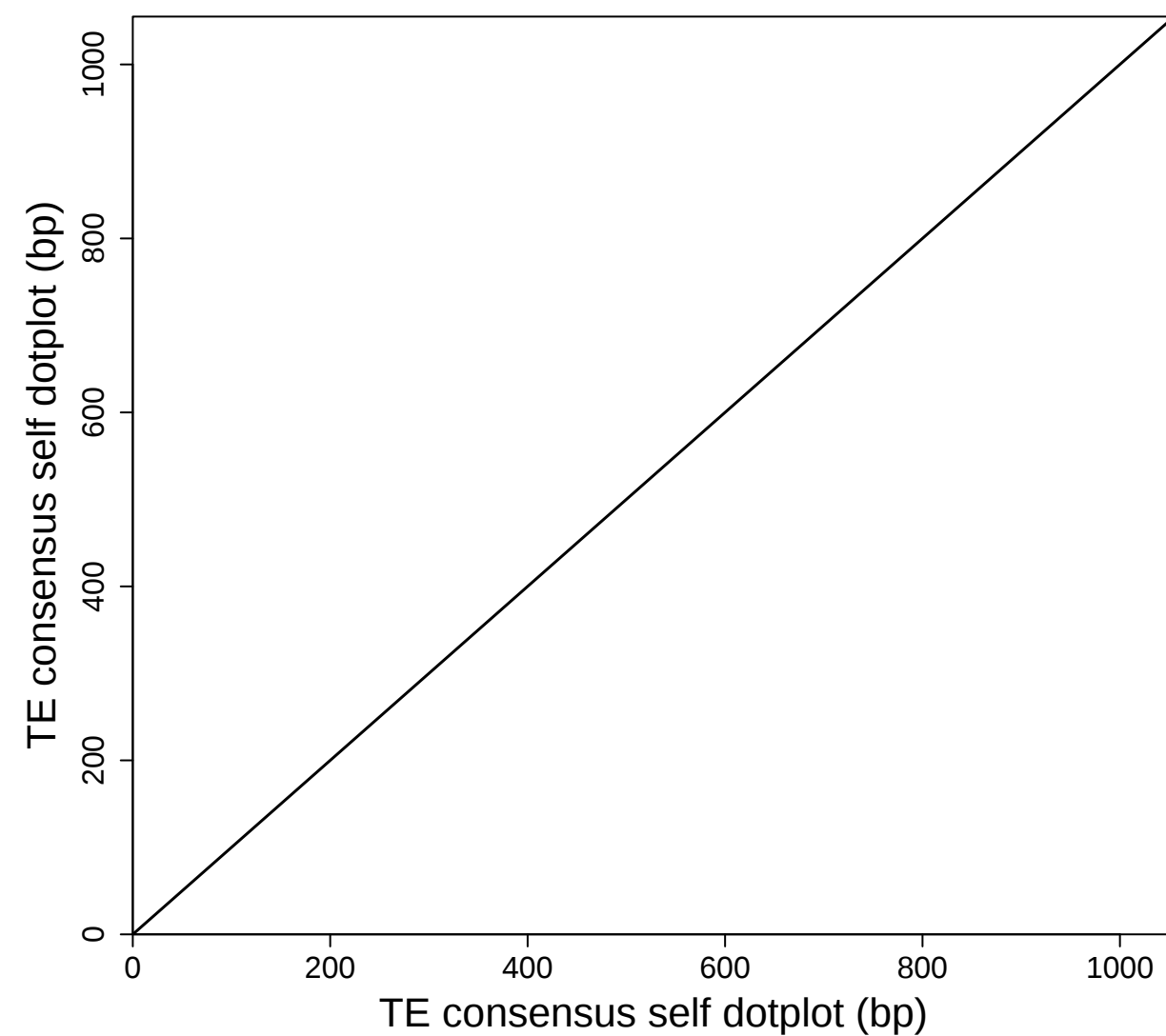
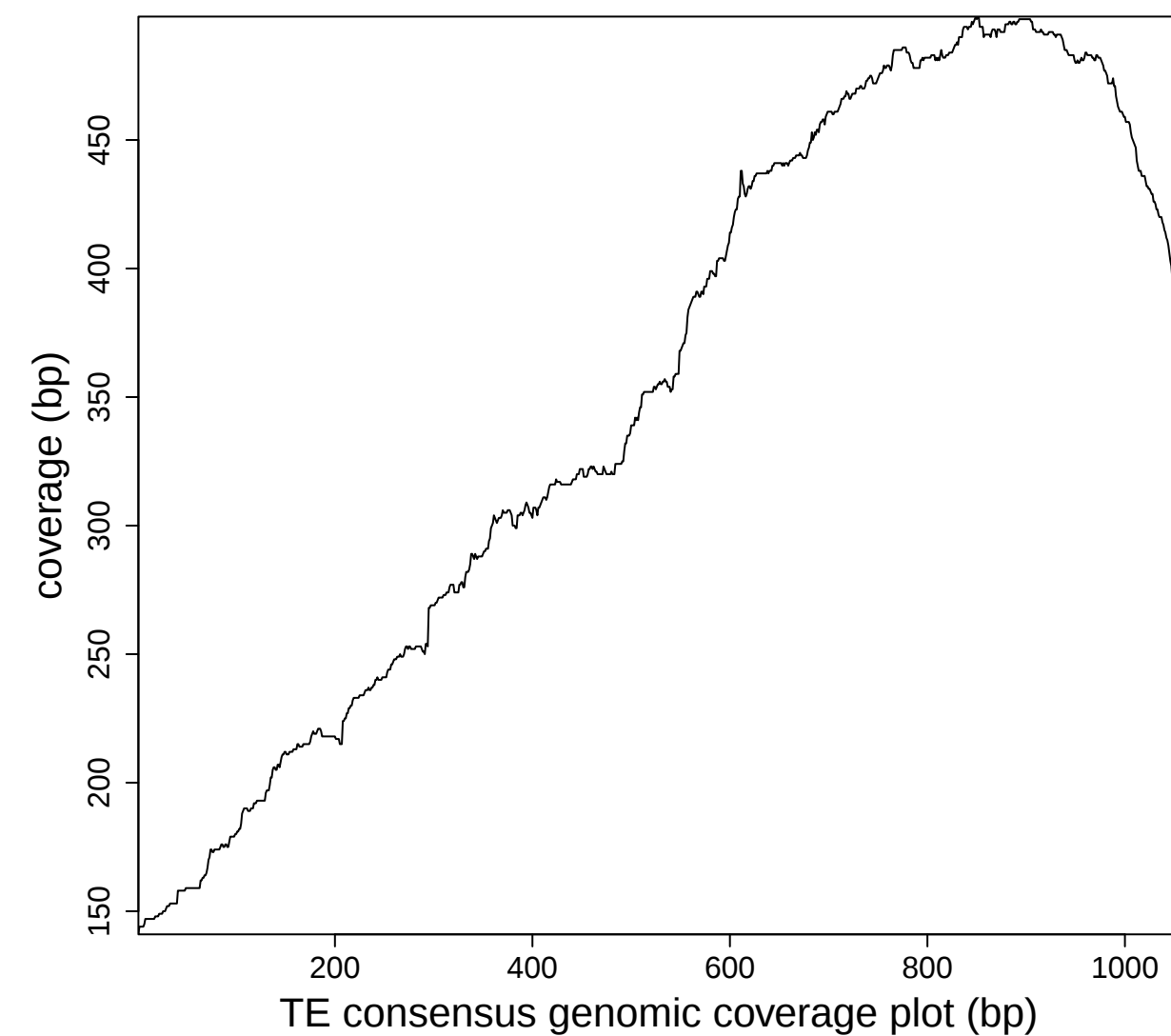
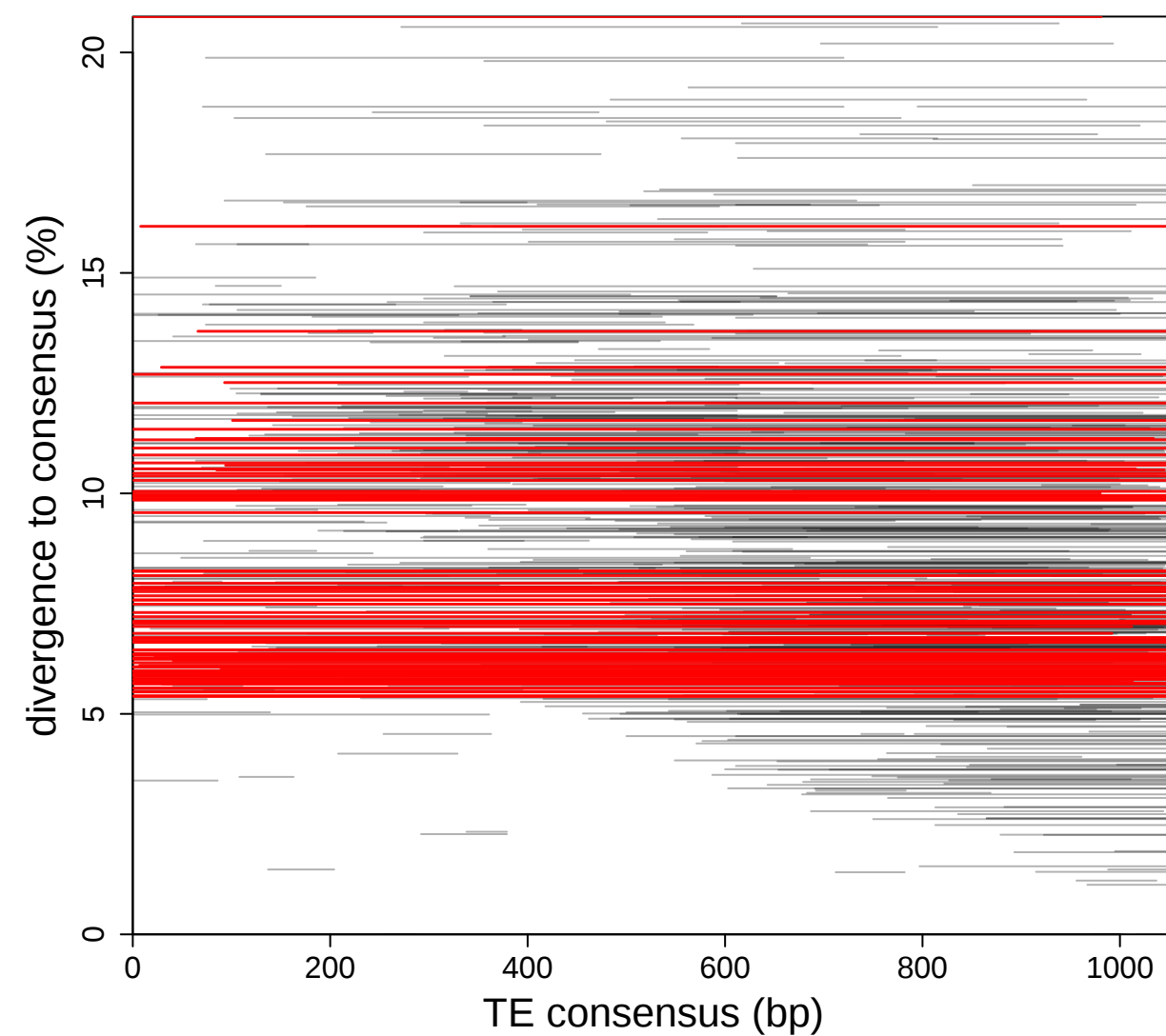
# After TEtrimmer Extended plot Blue lines are boundaries



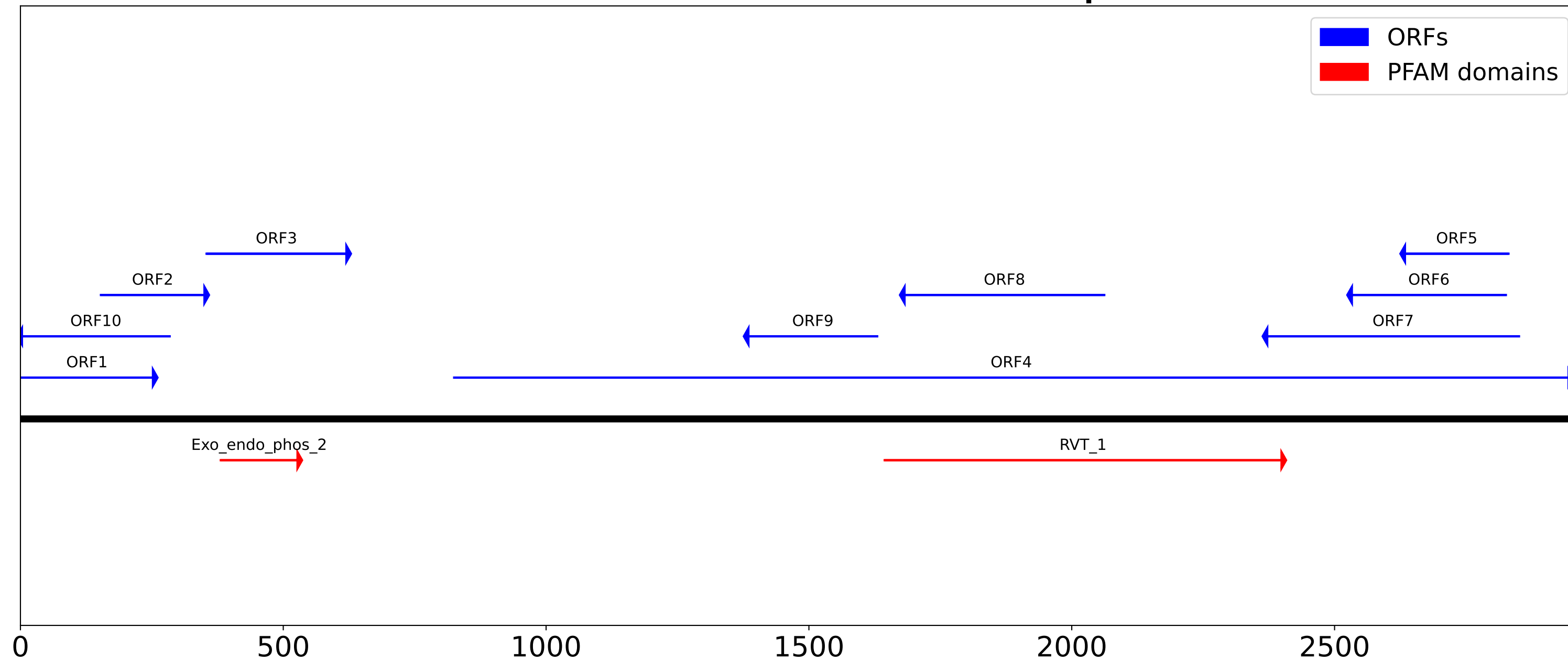
## Before TEtrimmer 1055 bp

TE: rnd\_4\_family\_846

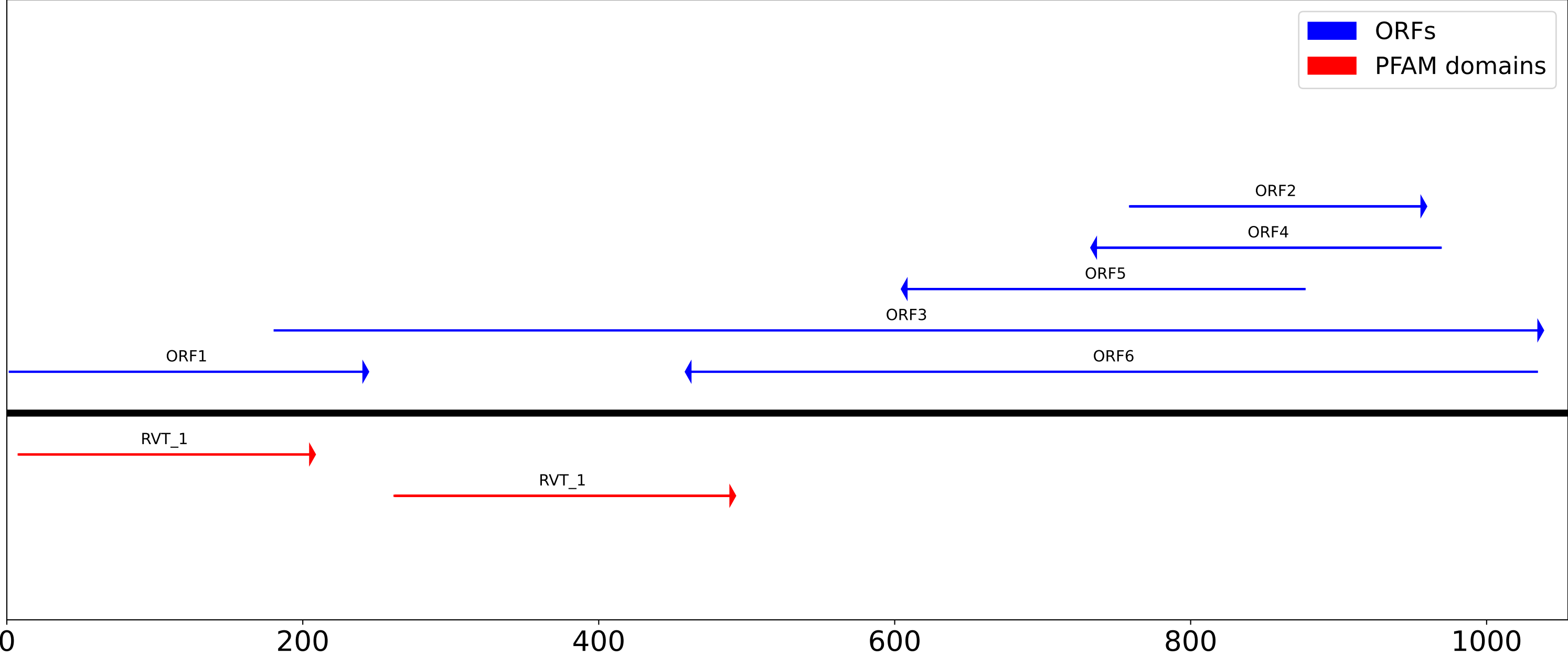
size: 1055bp; fragments: 710; full length: 109 ( $\geq 949.5$ bp)



## After TEtrimmer ORF and PFAM domain plot



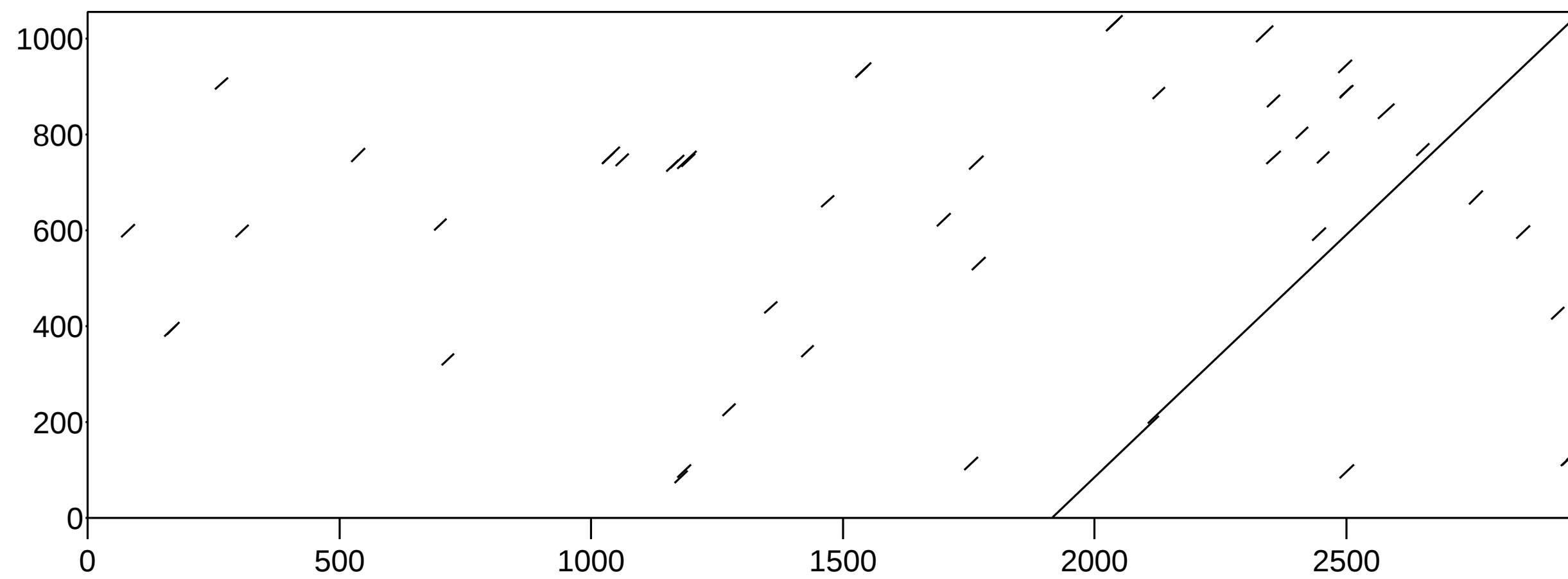
Before TEtrimmer ORF and PFAM domain plot



# Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)