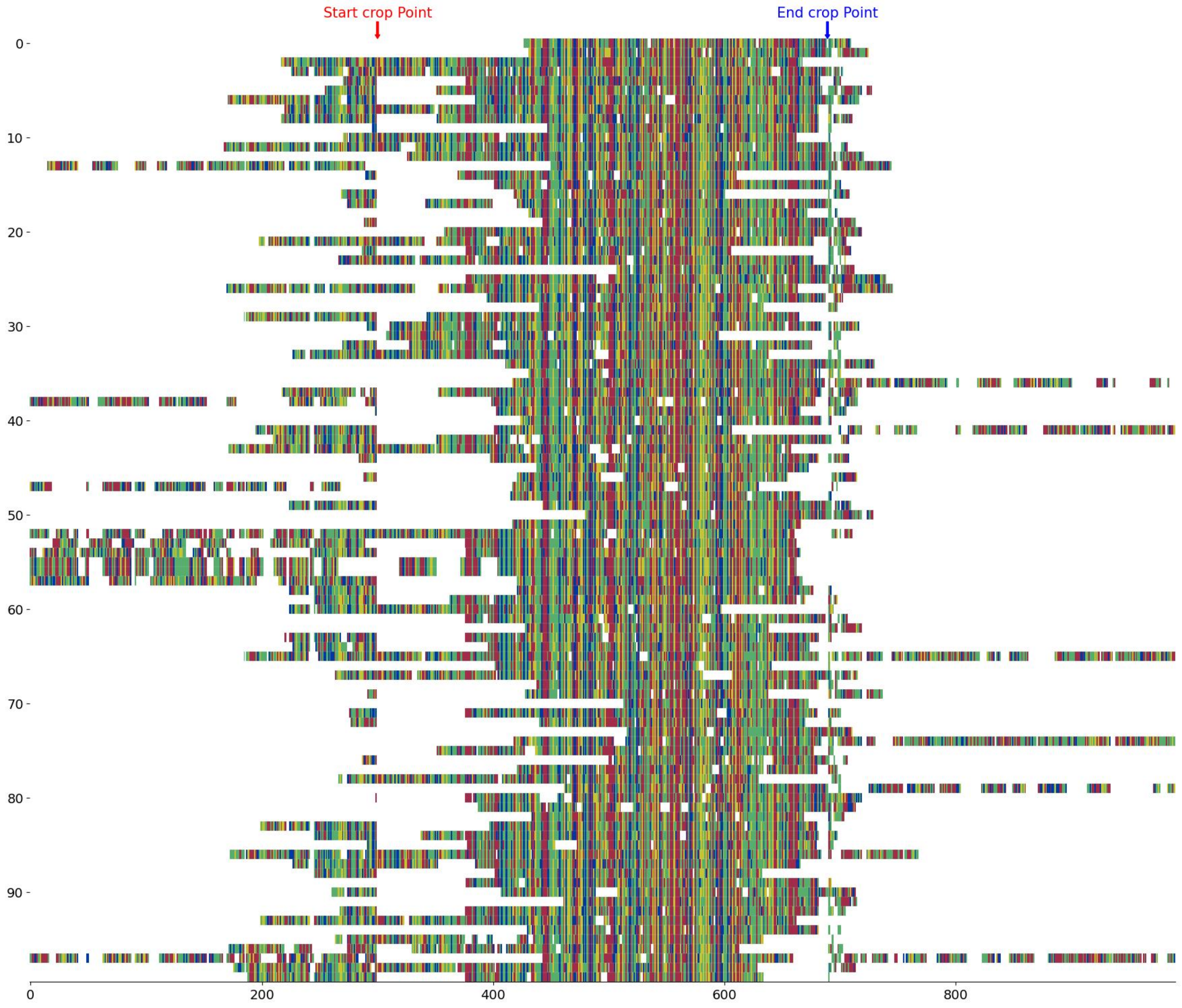
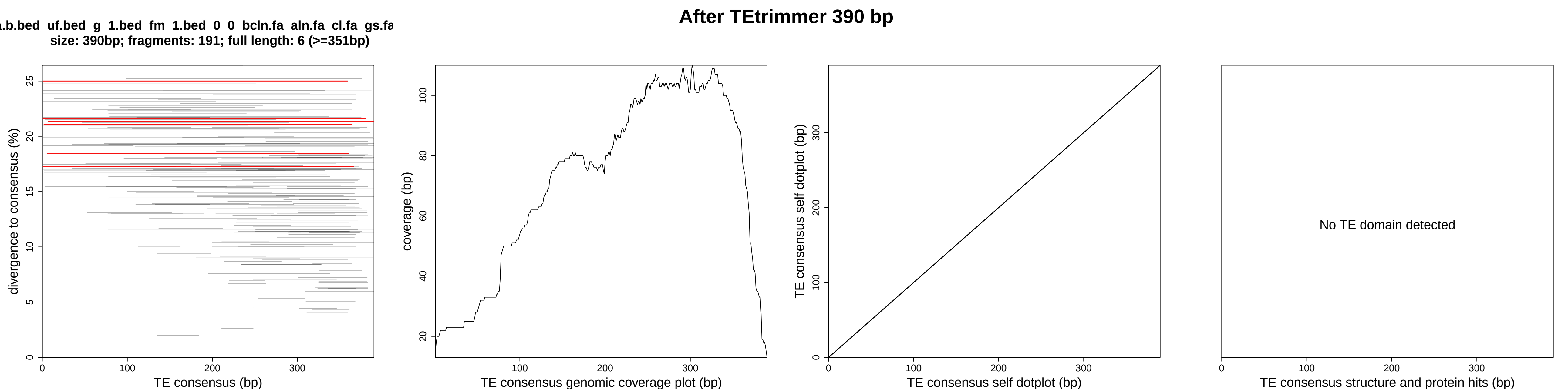






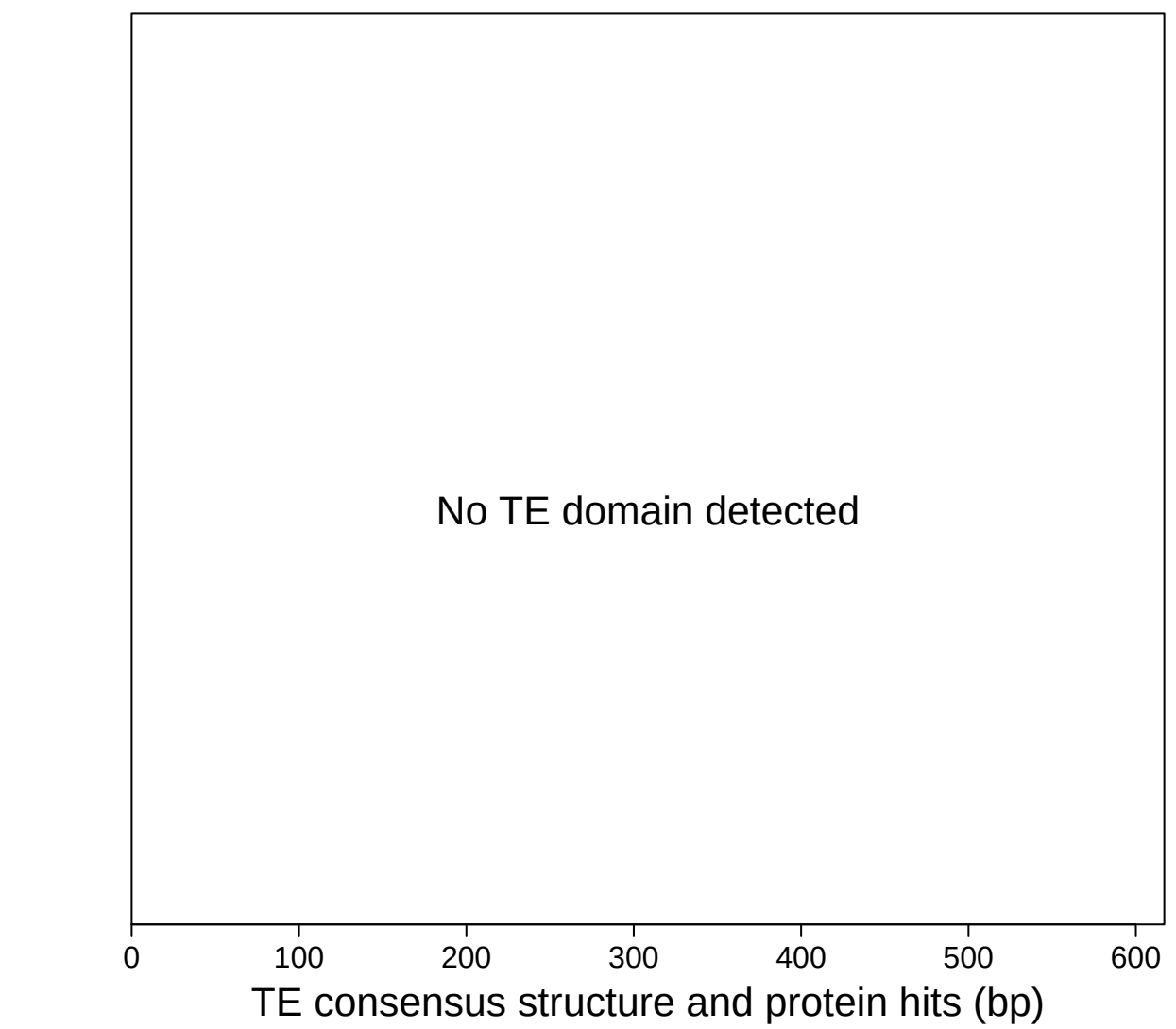
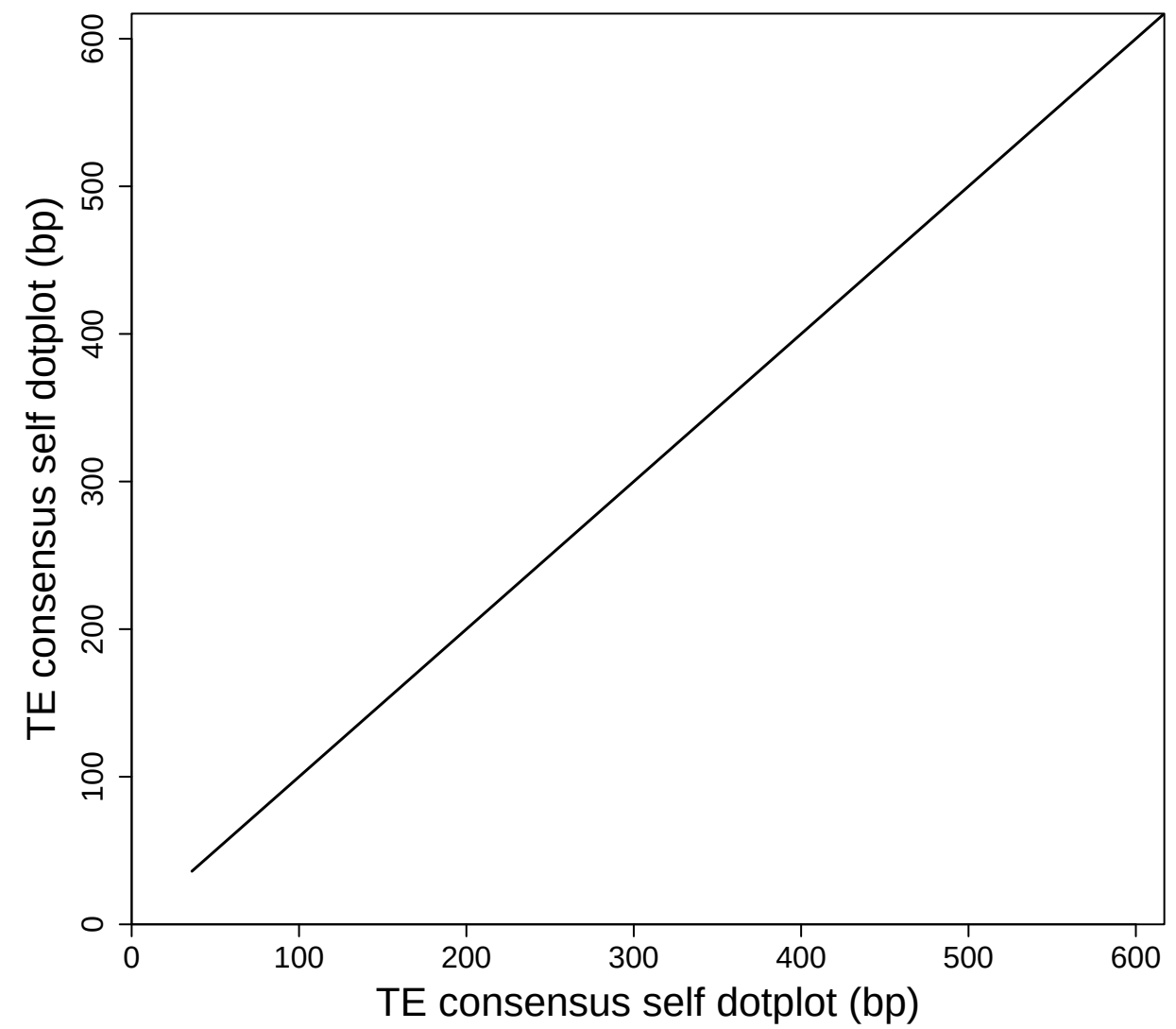
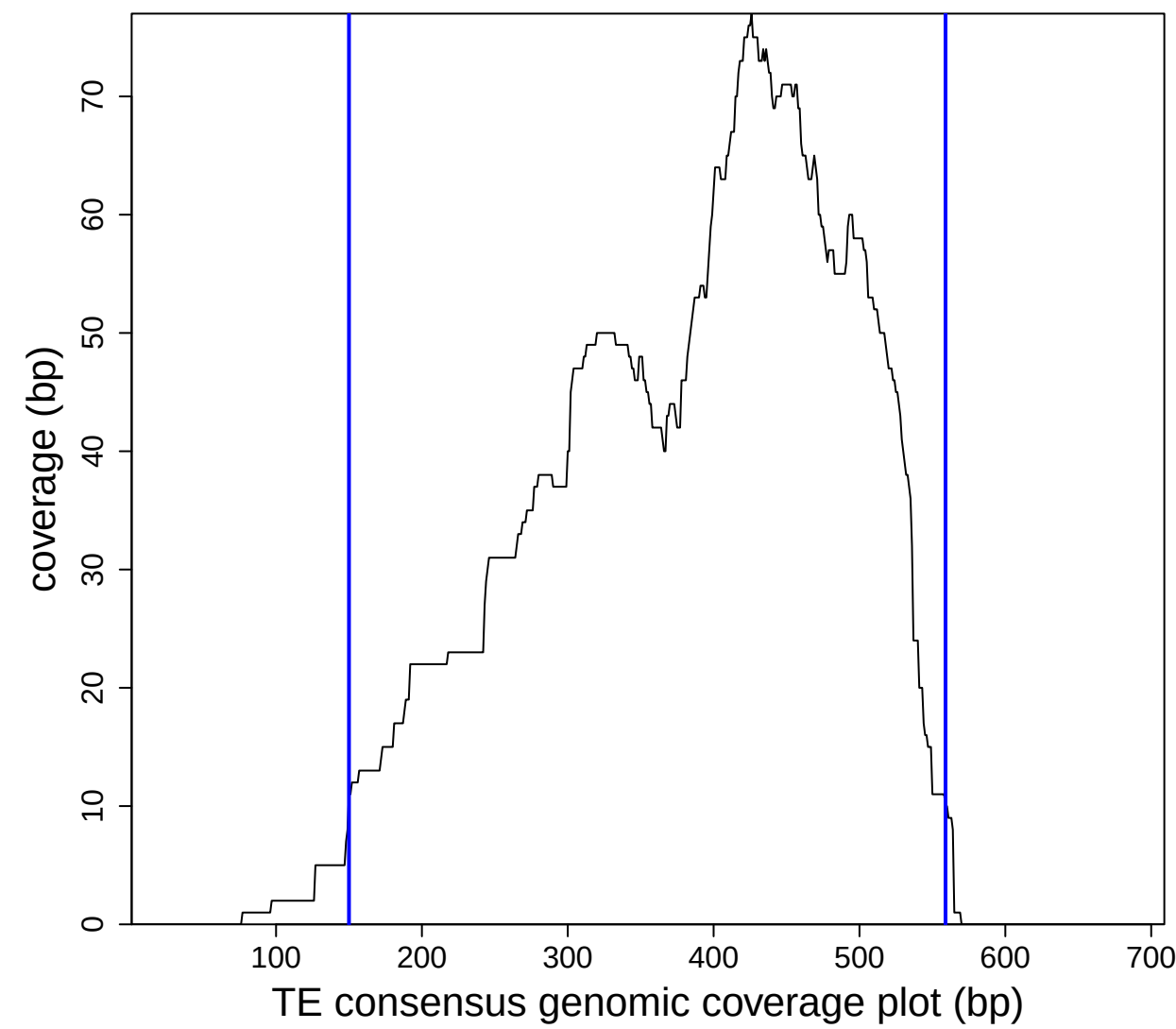
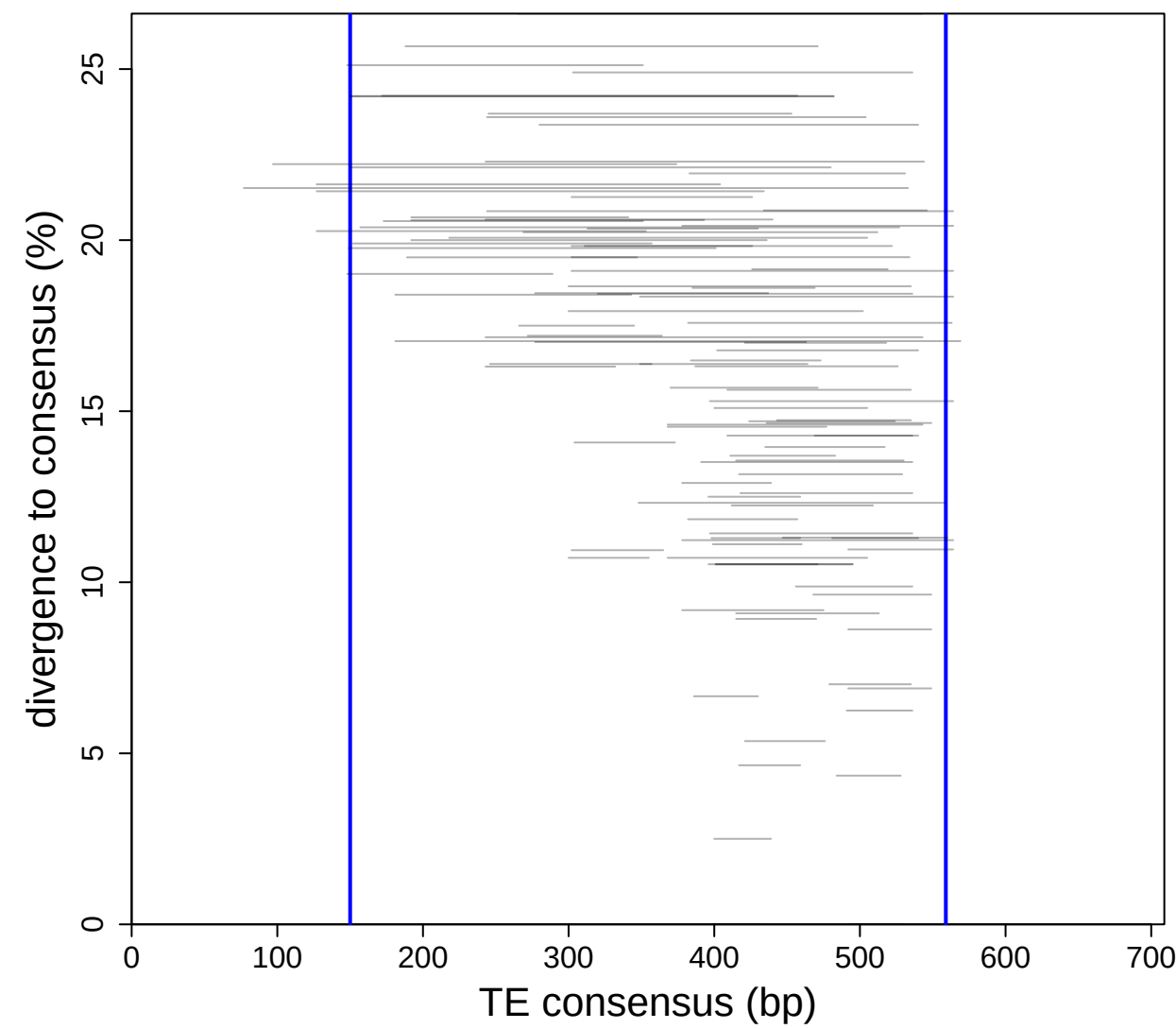






1.fasta.b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa  
size: 709bp; fragments: 112; full length: 0 (>=638.1bp)

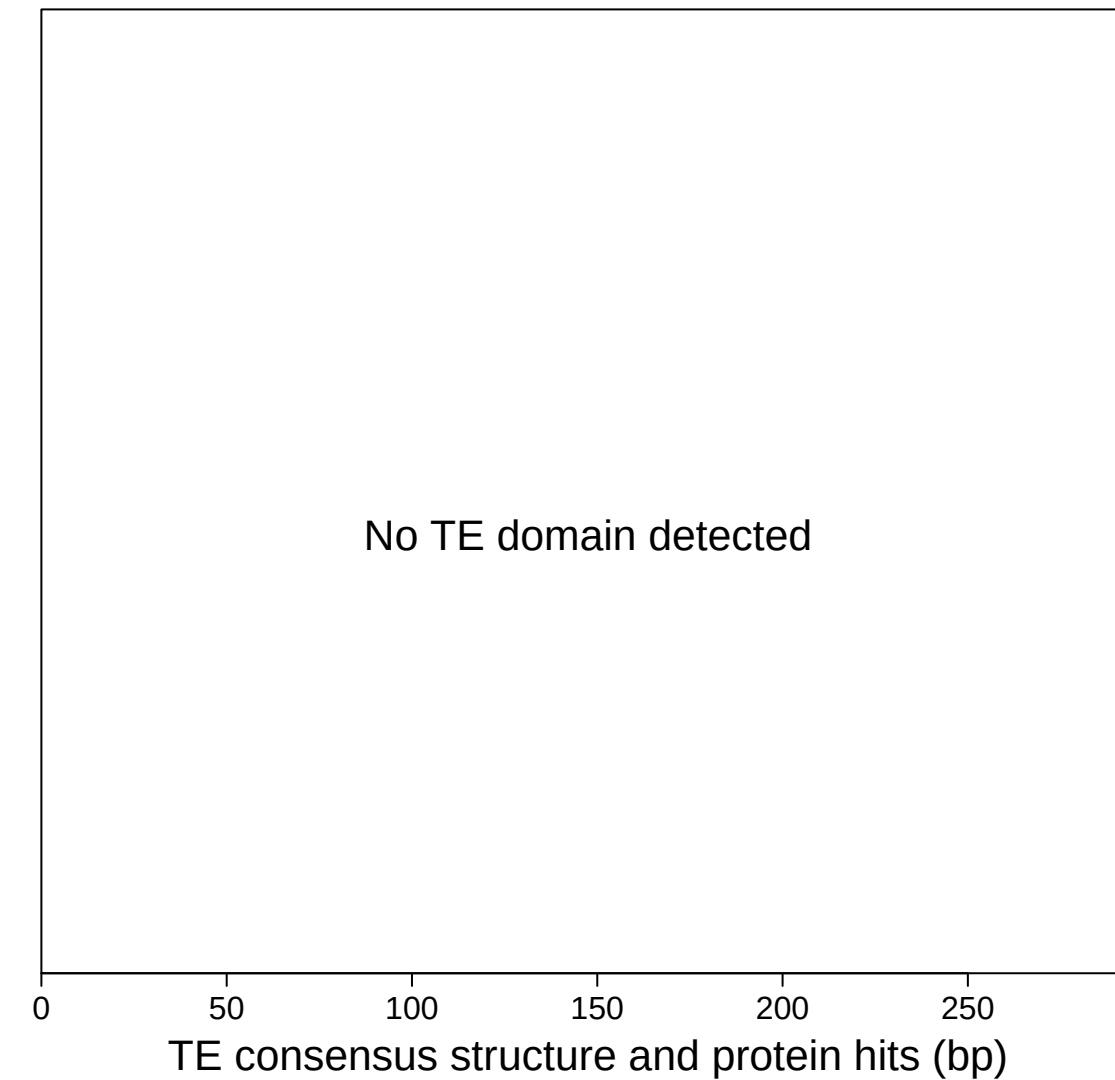
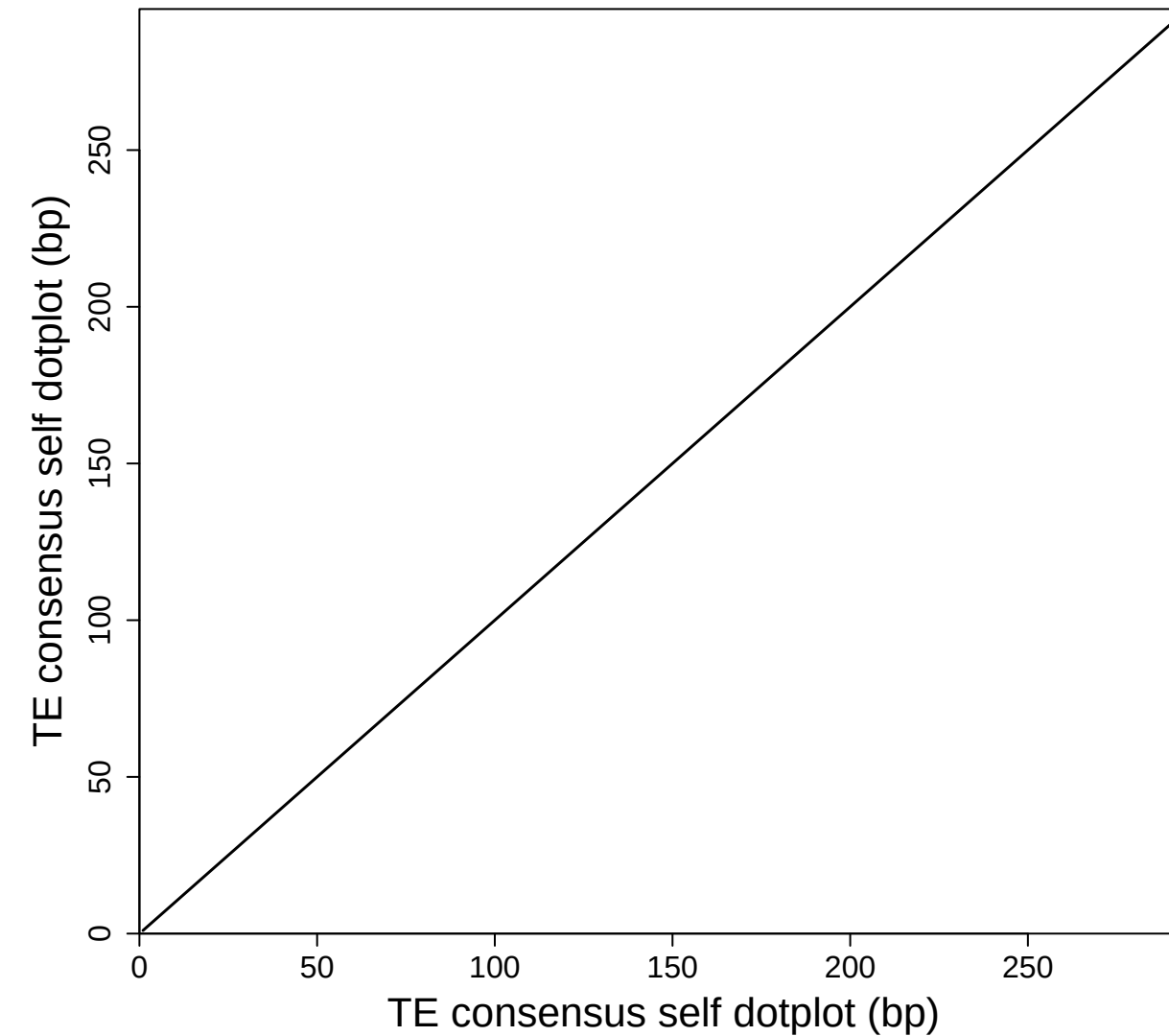
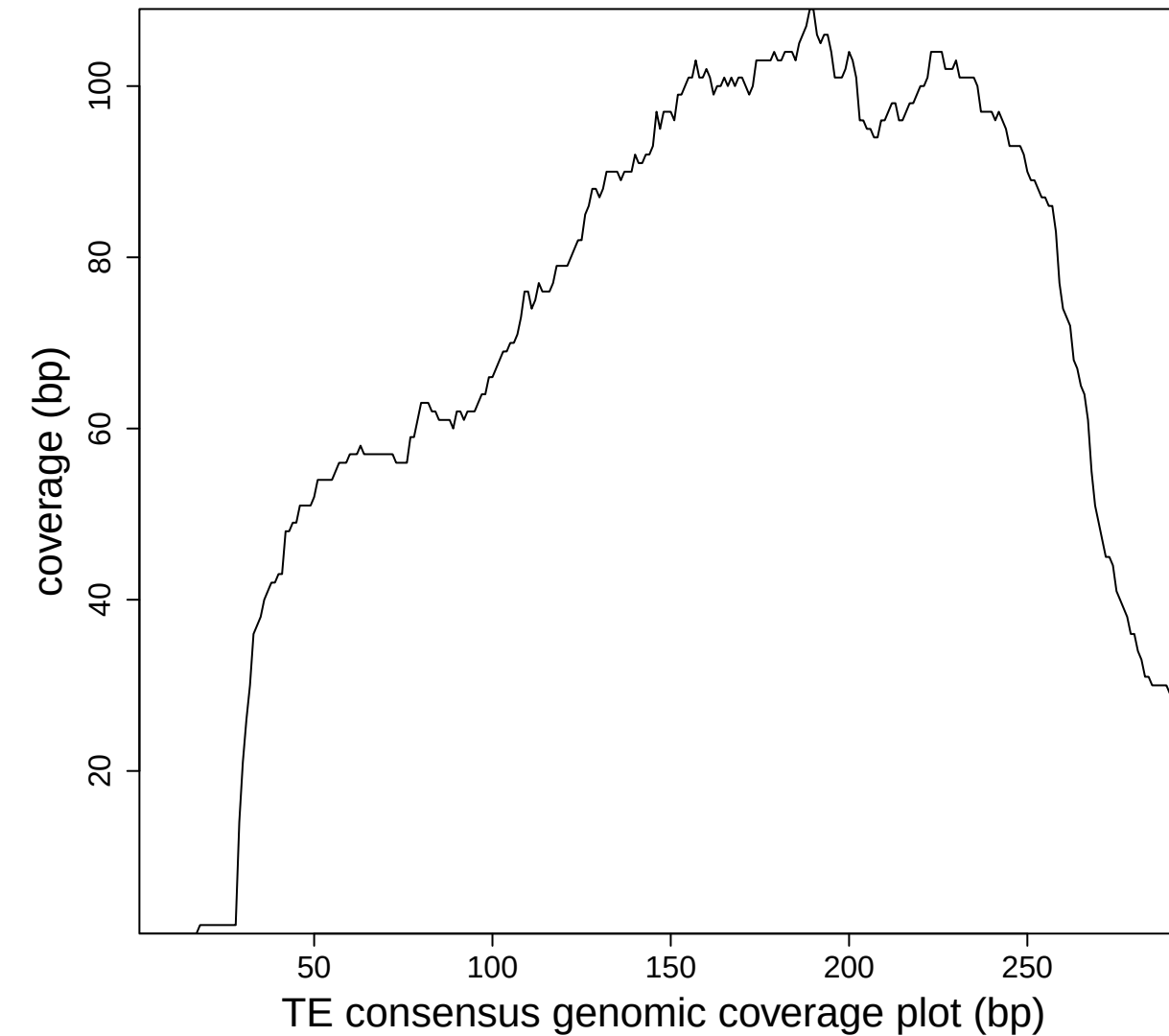
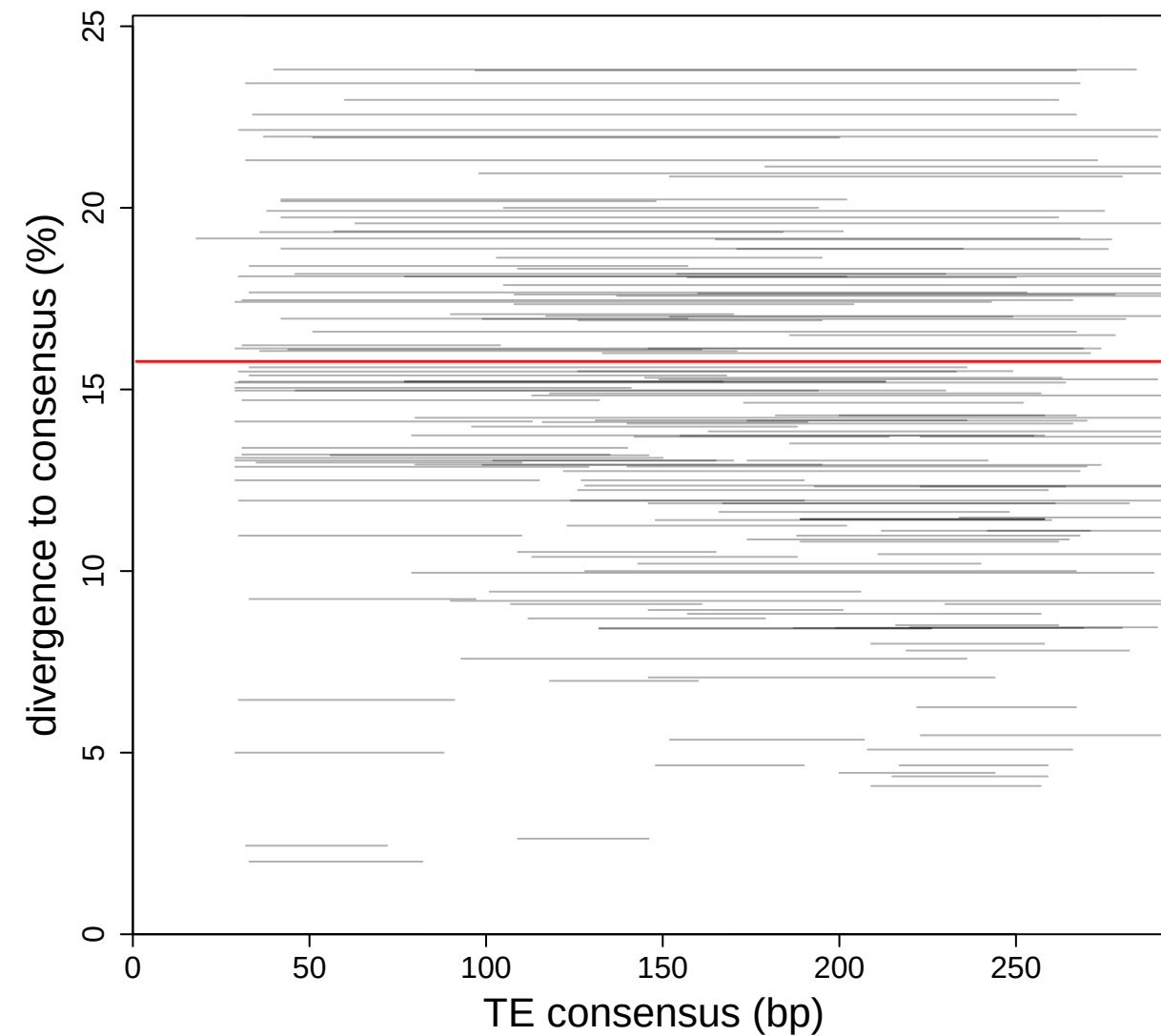
# After TEtrimmer Extended plot Blue lines are boundaries



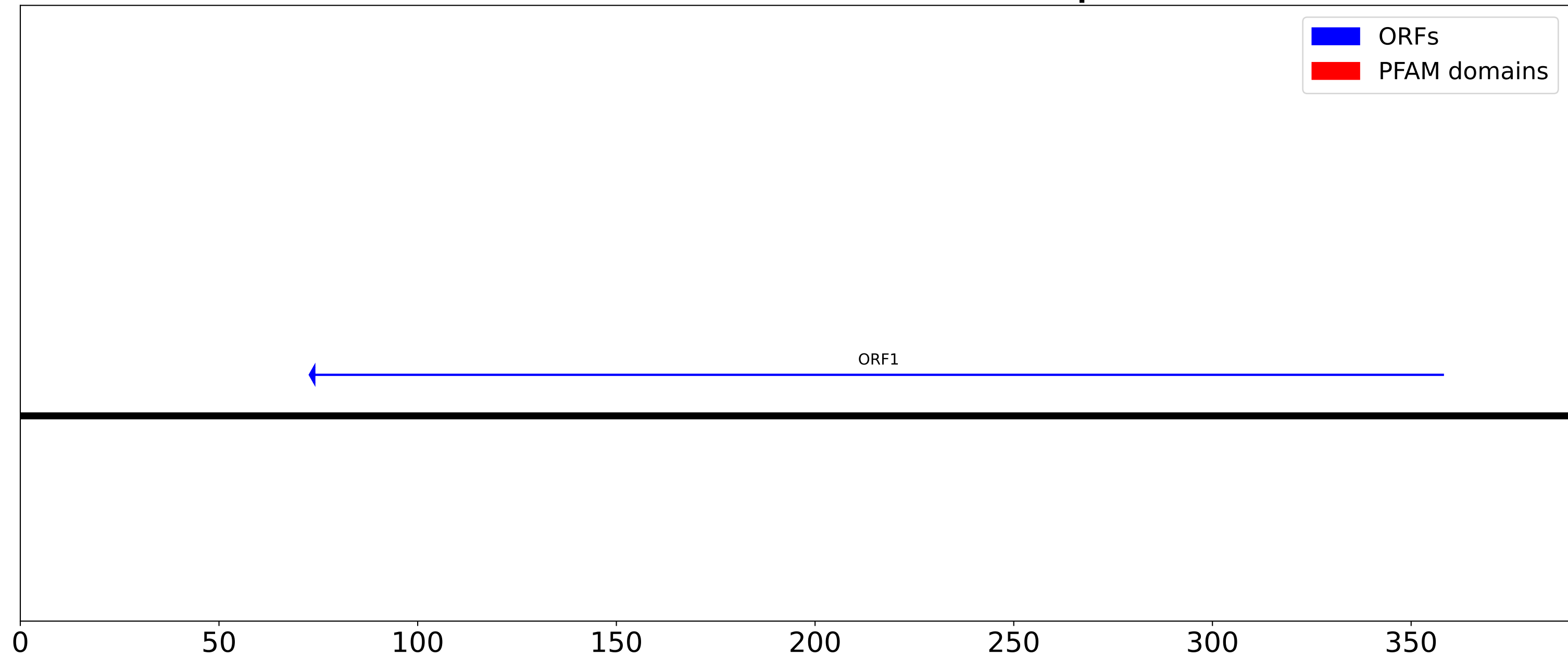
## Before TEtrimmer 295 bp

TE: rnd\_5\_family\_721

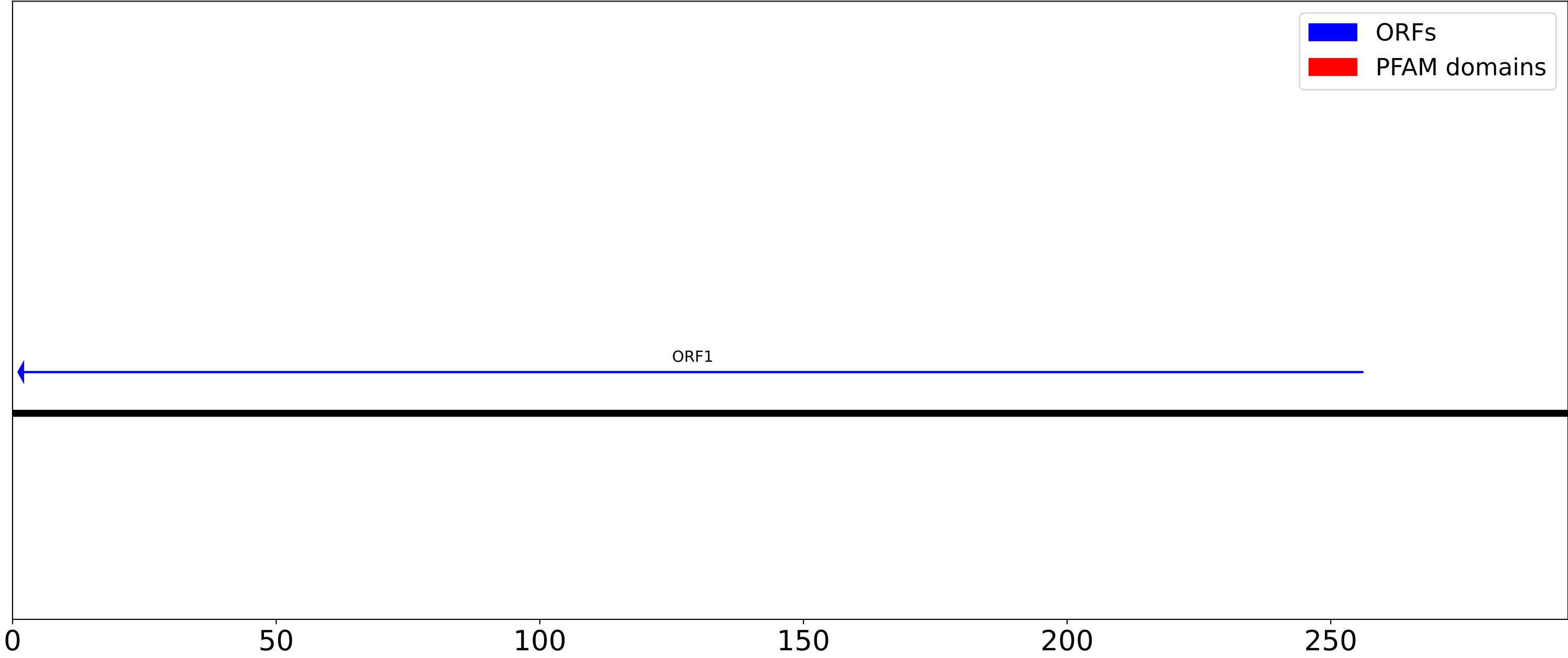
size: 295bp; fragments: 167; full length: 1 ( $\geq 265.5$ bp)



## After TEtrimmer ORF and PFAM domain plot



**Before TEtrimmer ORF and PFAM domain plot**





# Dotplot

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

