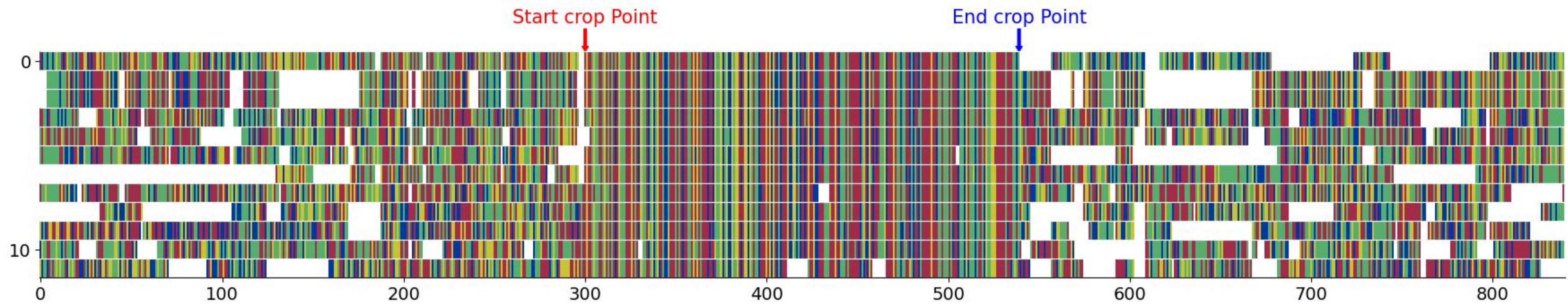


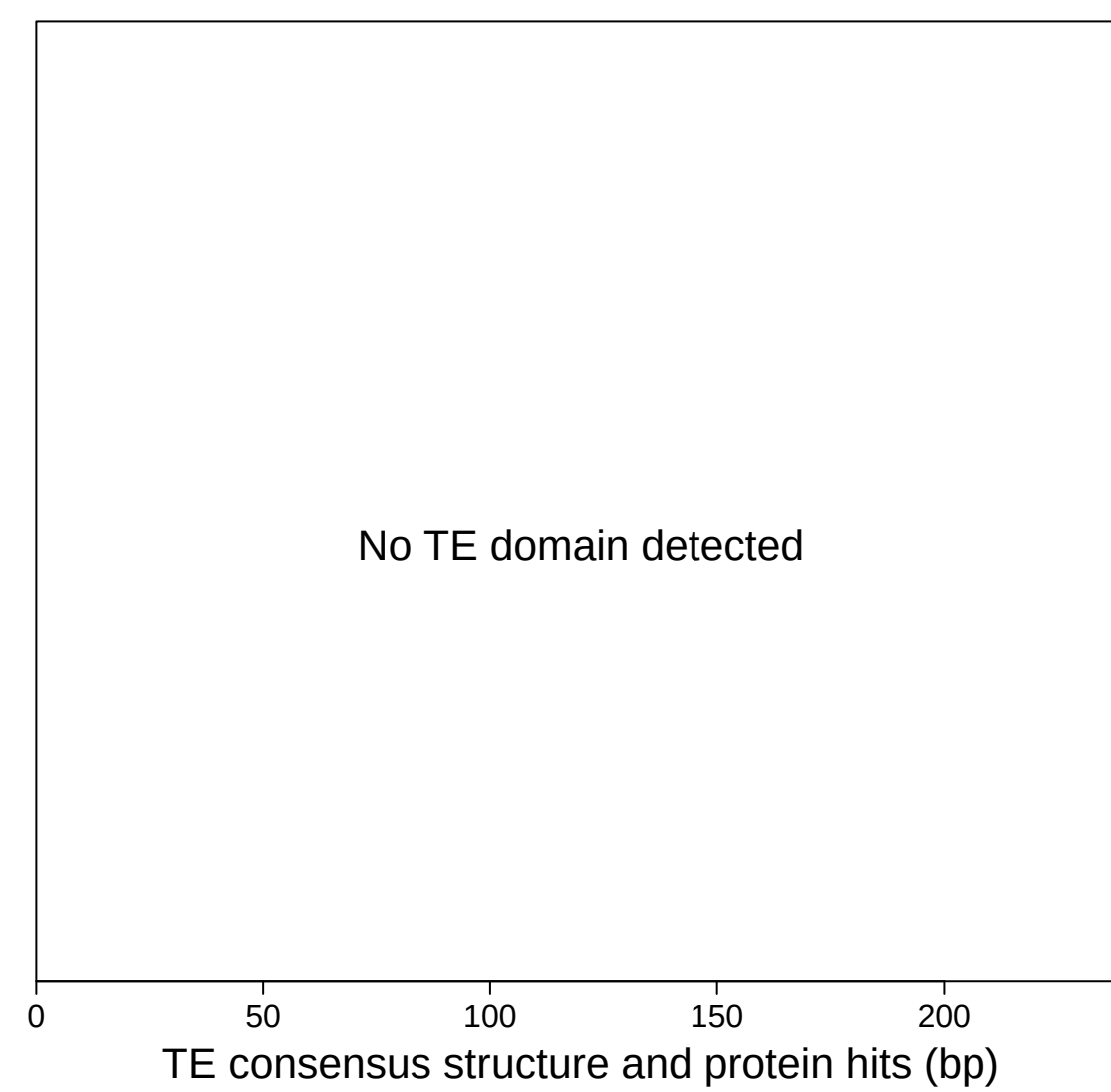
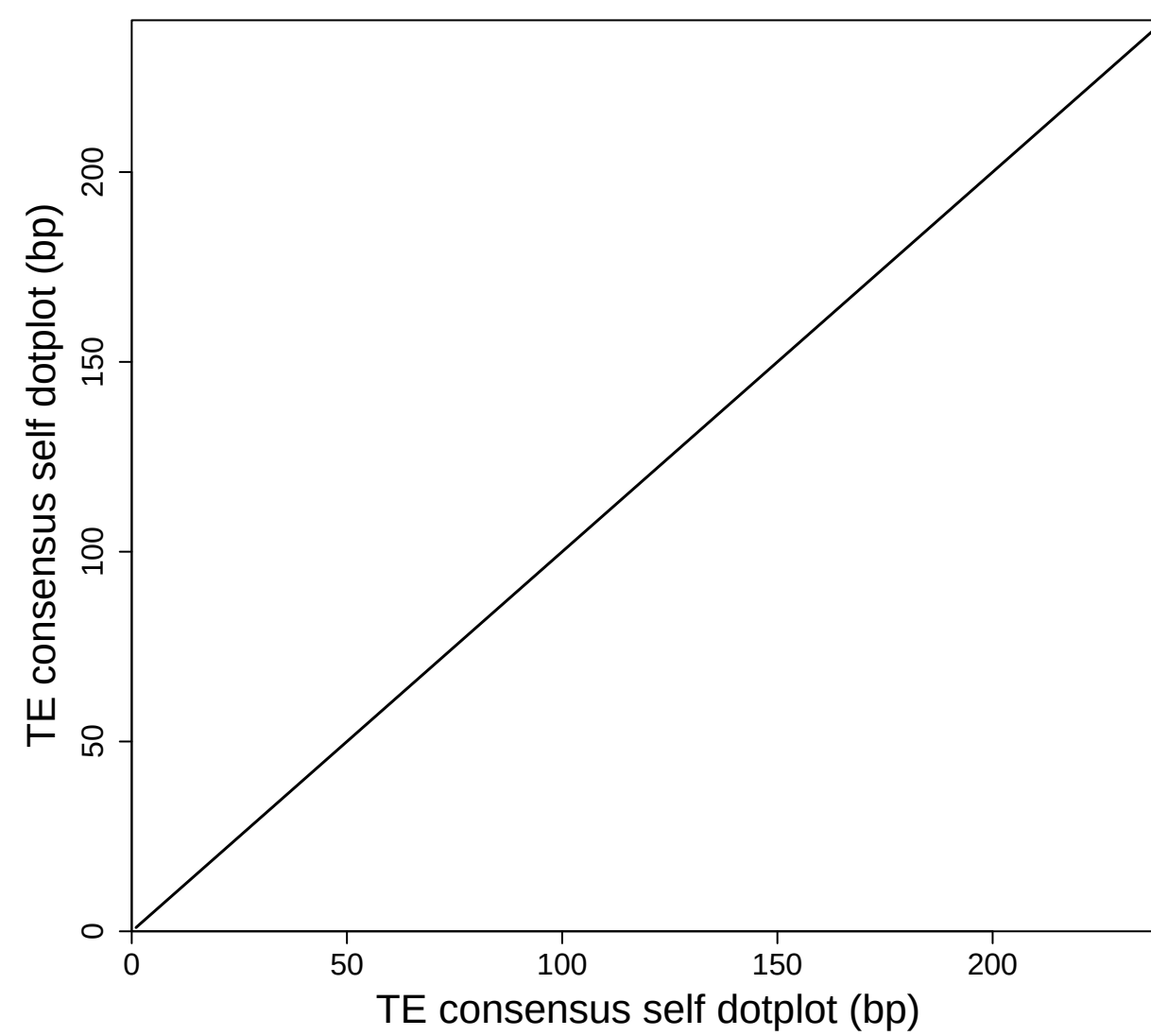
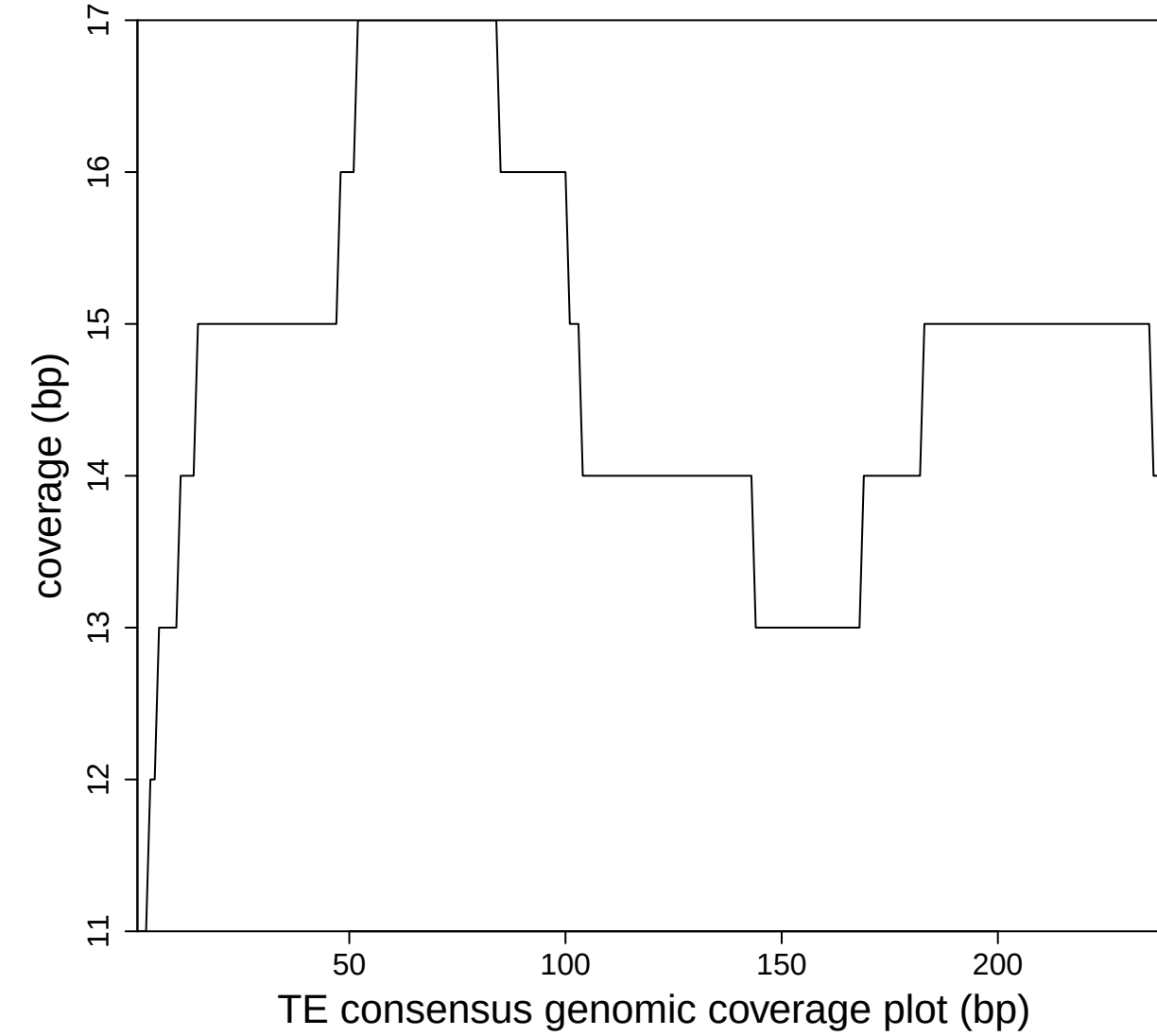
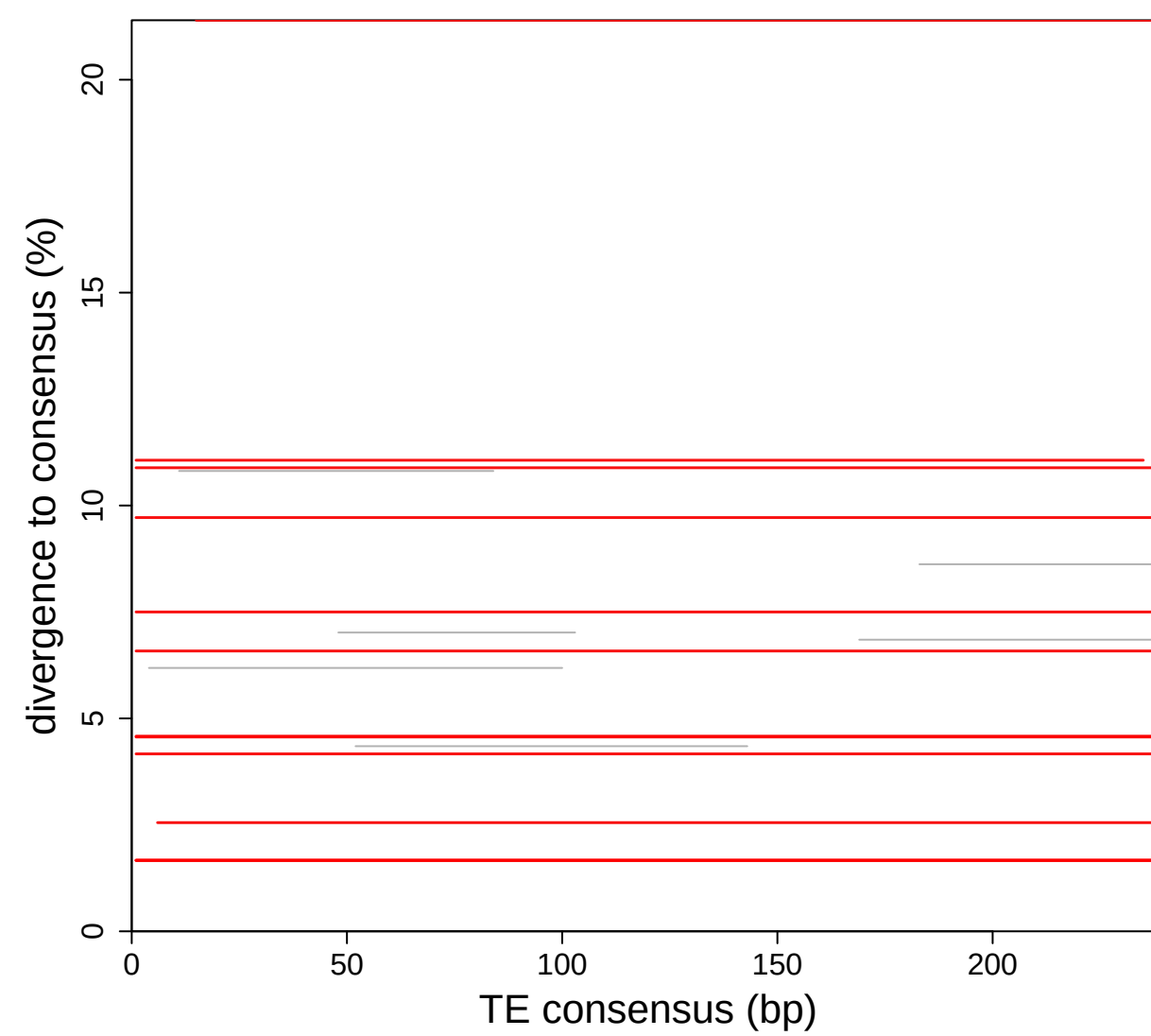
MSA length =

[illegible]



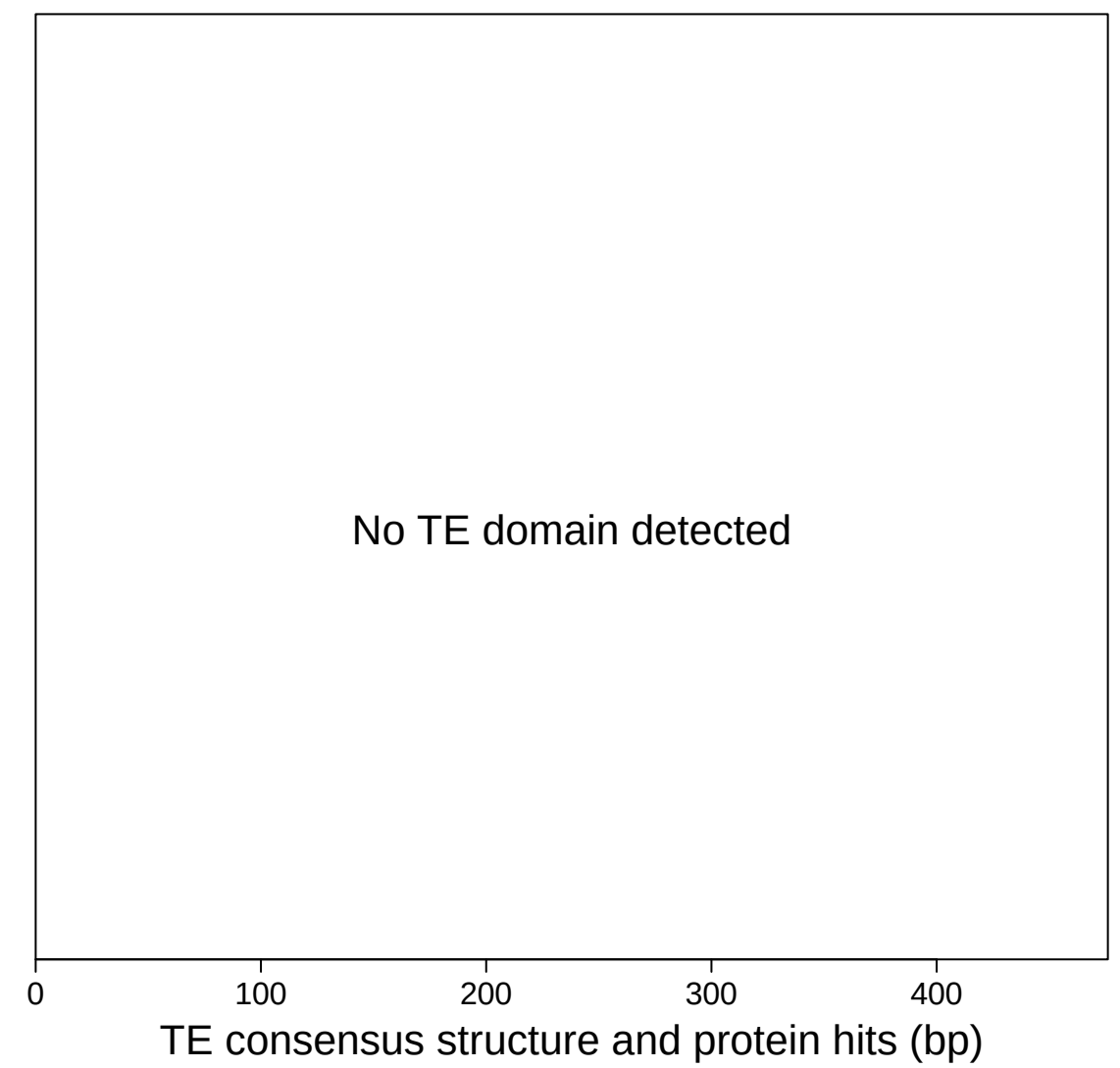
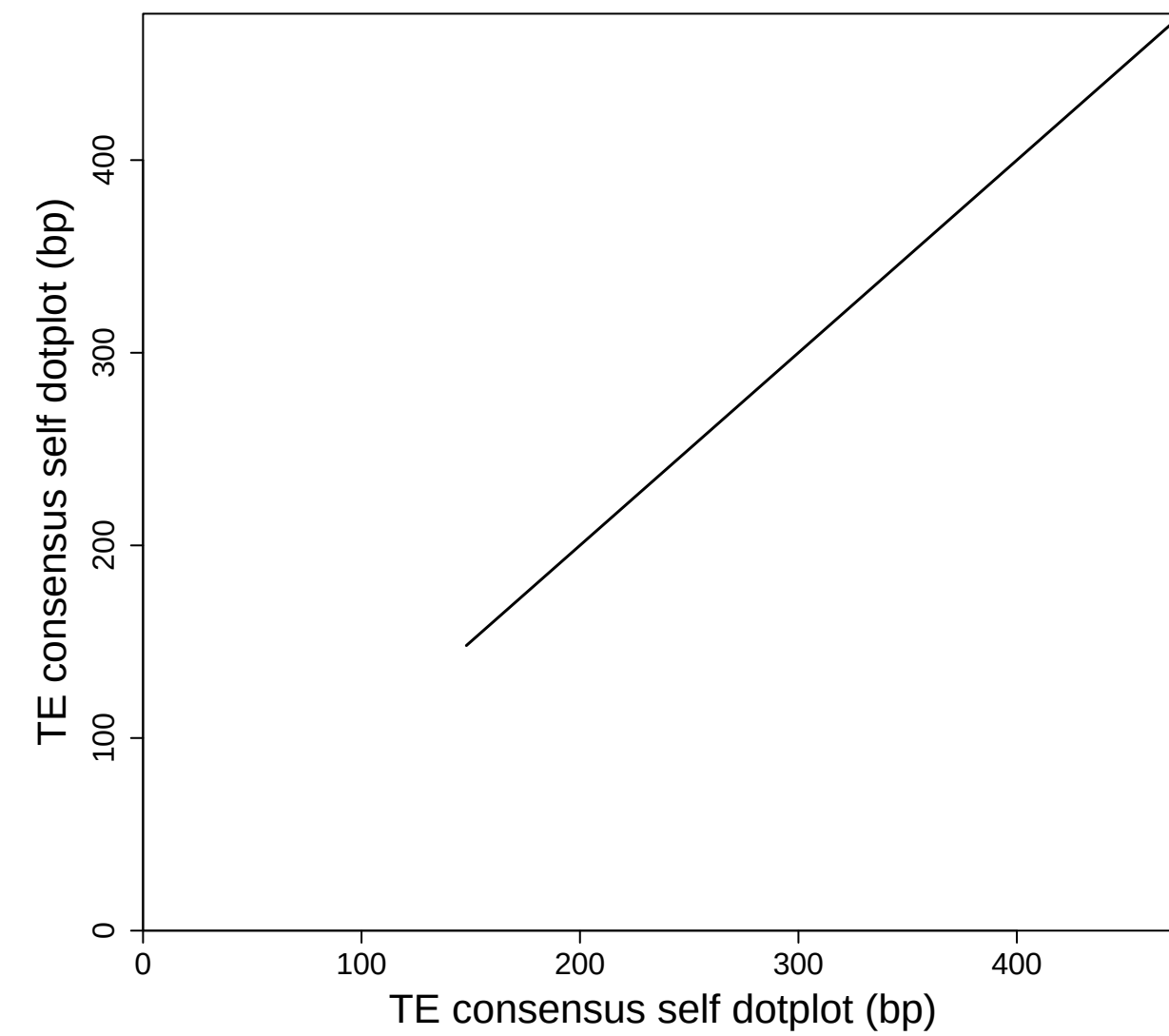
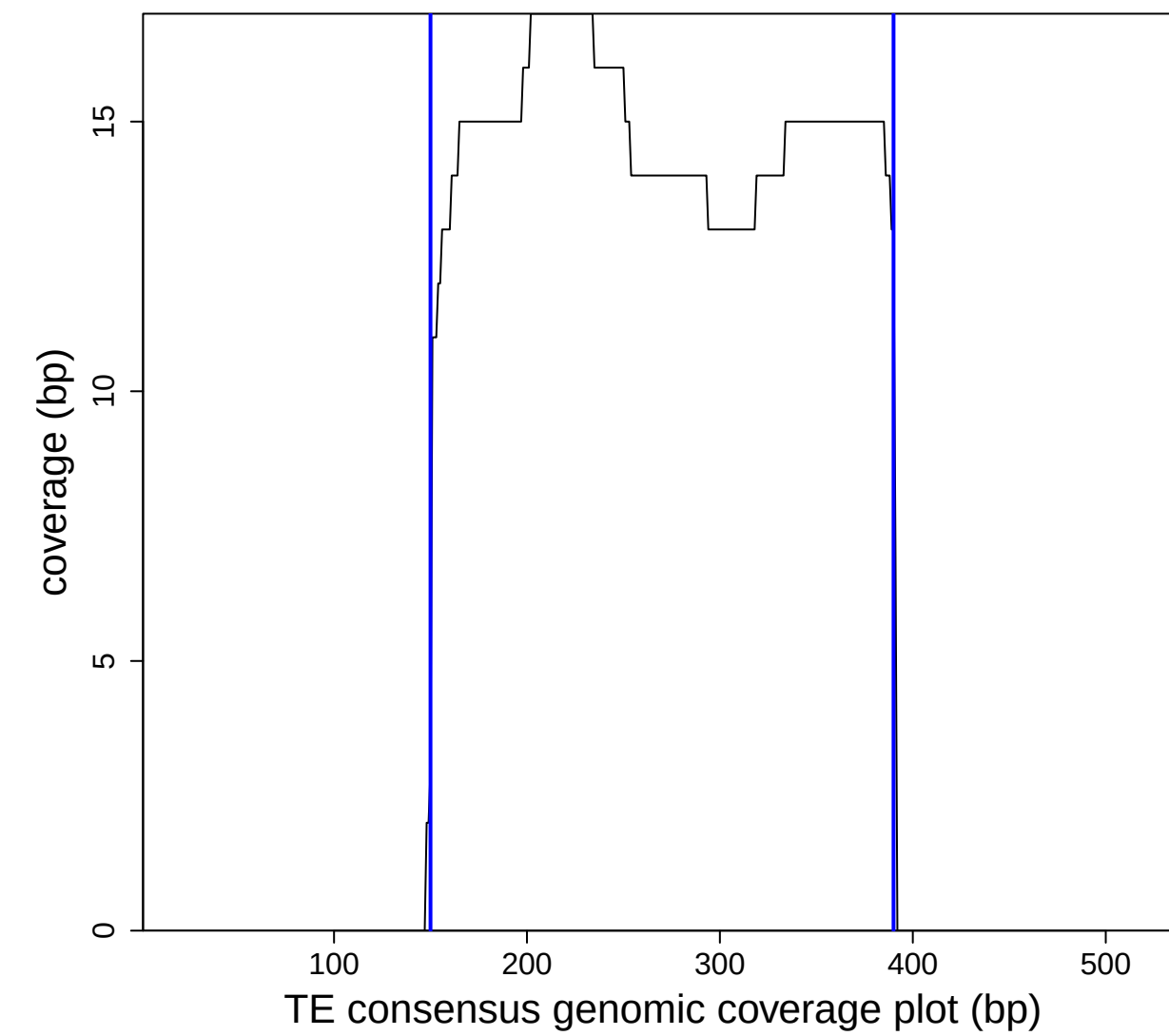
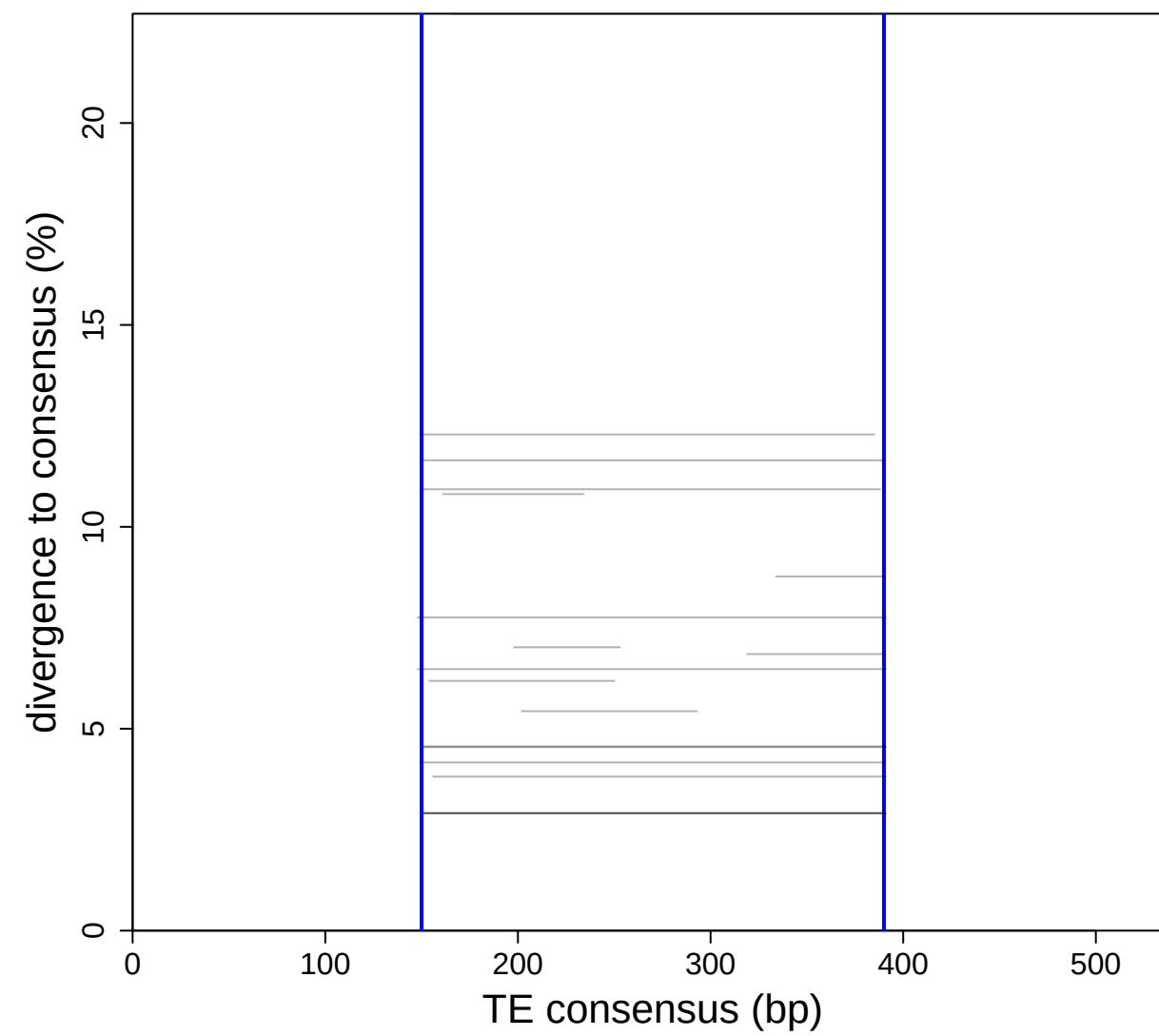
b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 240bp; fragments: 19; full length: 13 (>=216bp)

After TEtrimmer 240 bp



%.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 540bp; fragments: 19; full length: 0 (>=486bp)

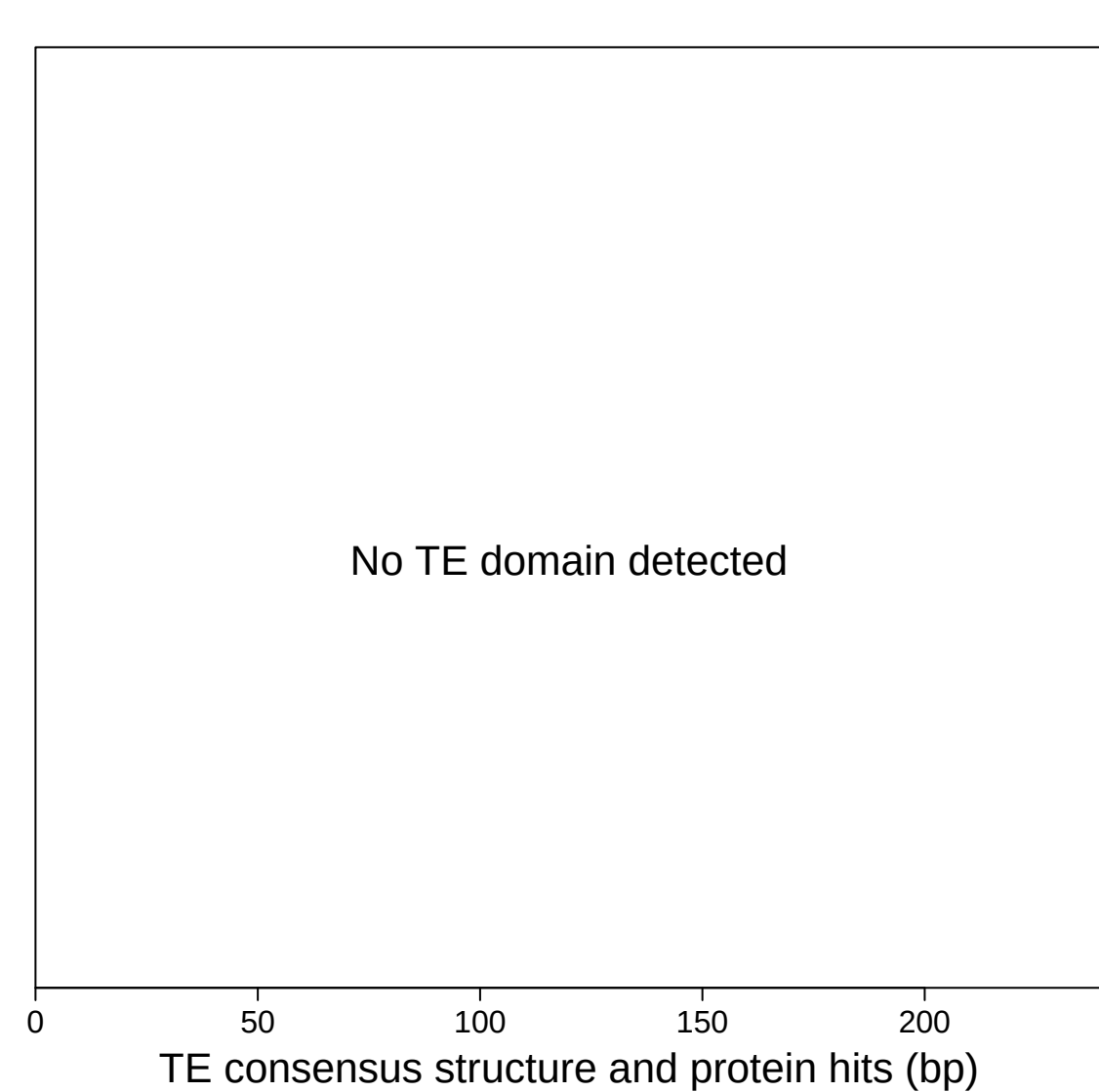
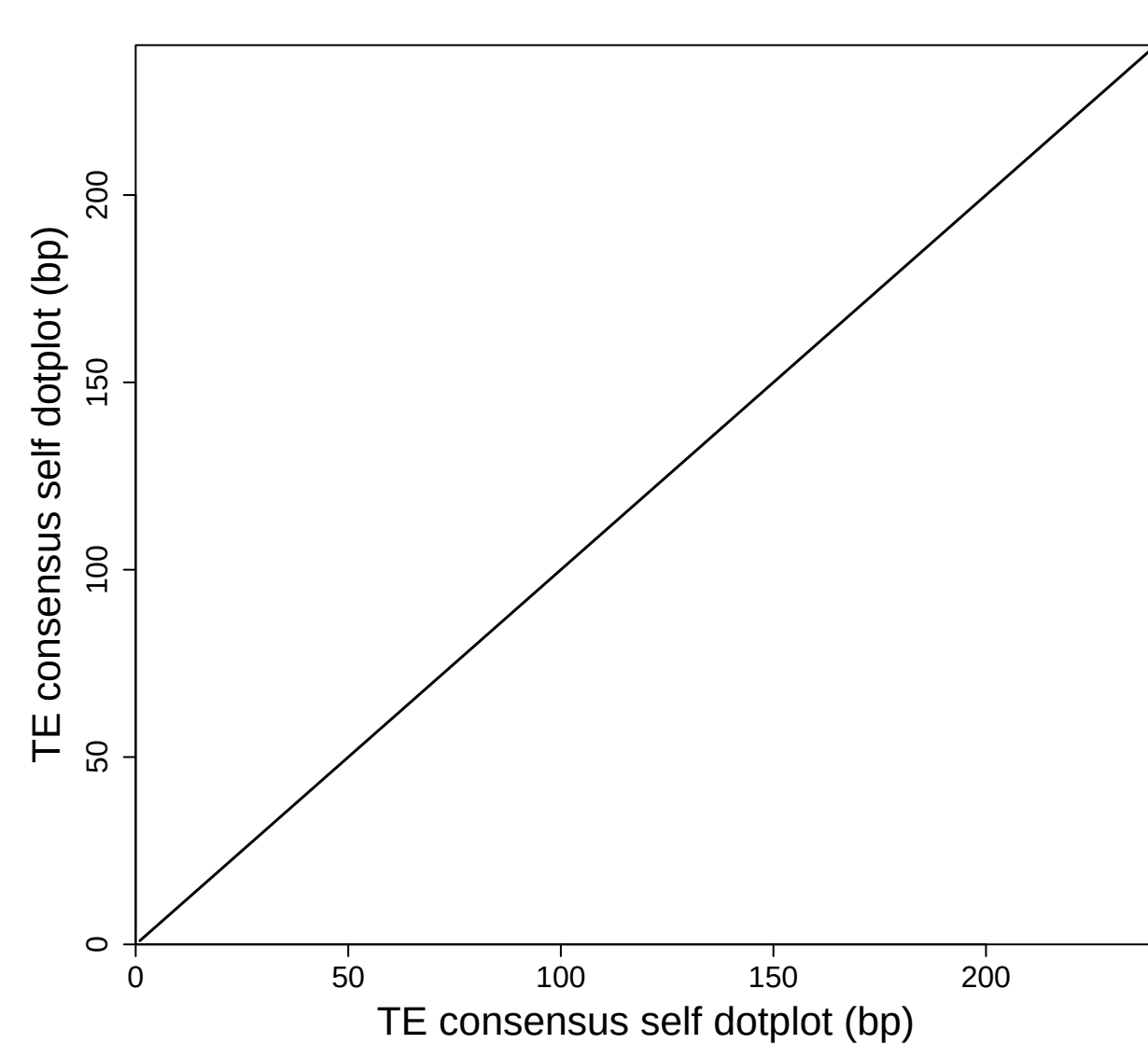
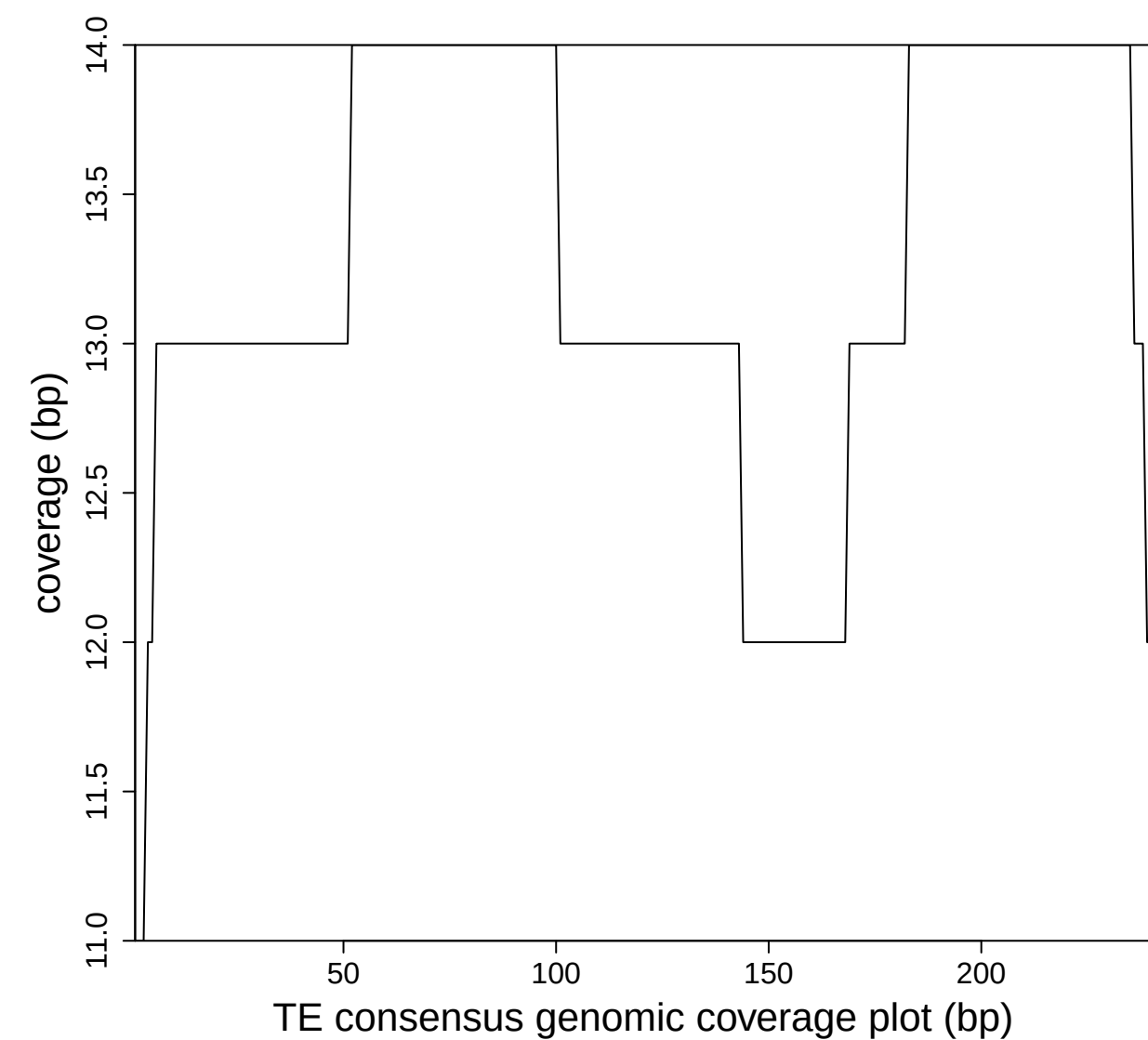
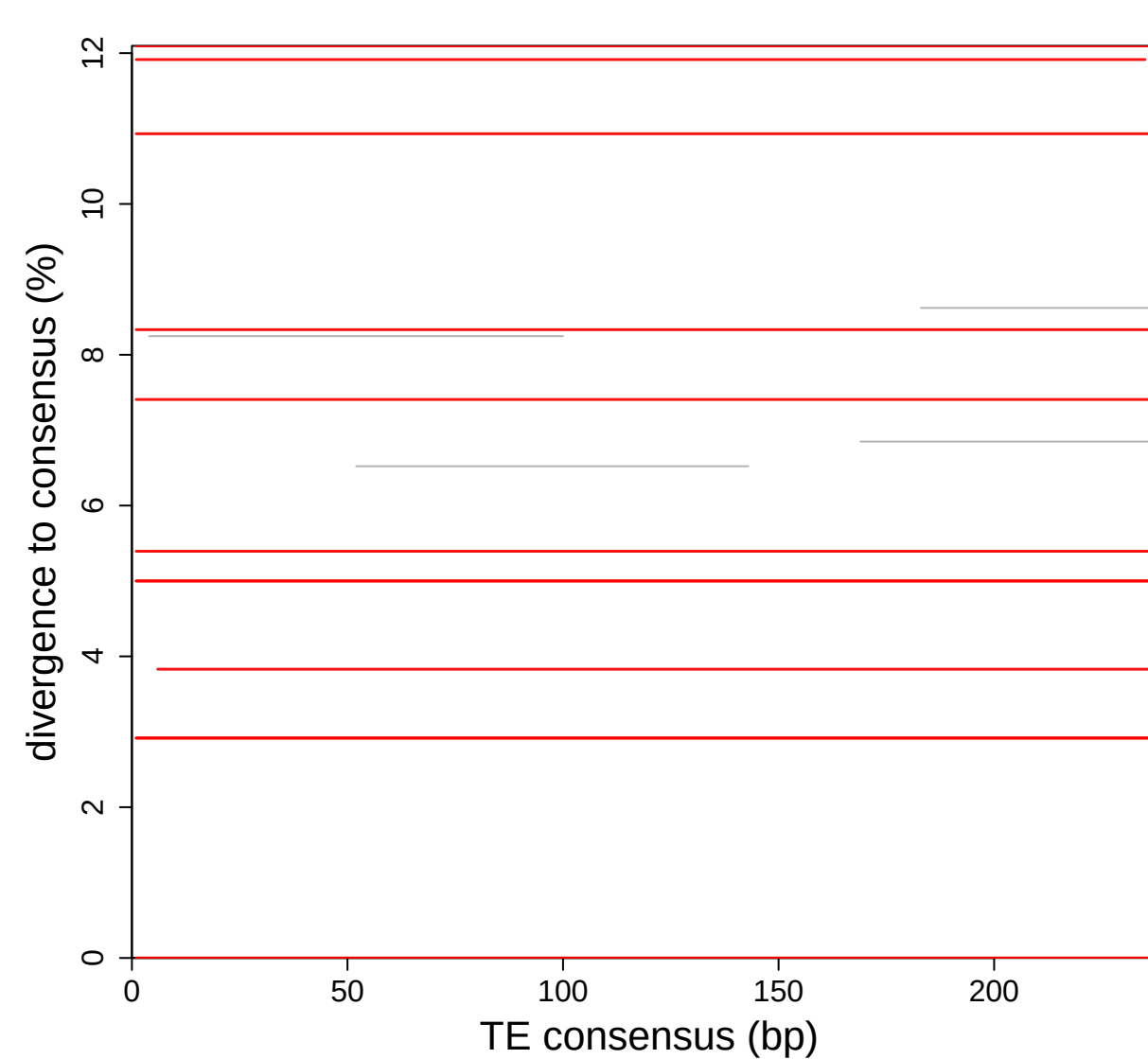
After TEtrimmer Extended plot Blue lines are boundaries



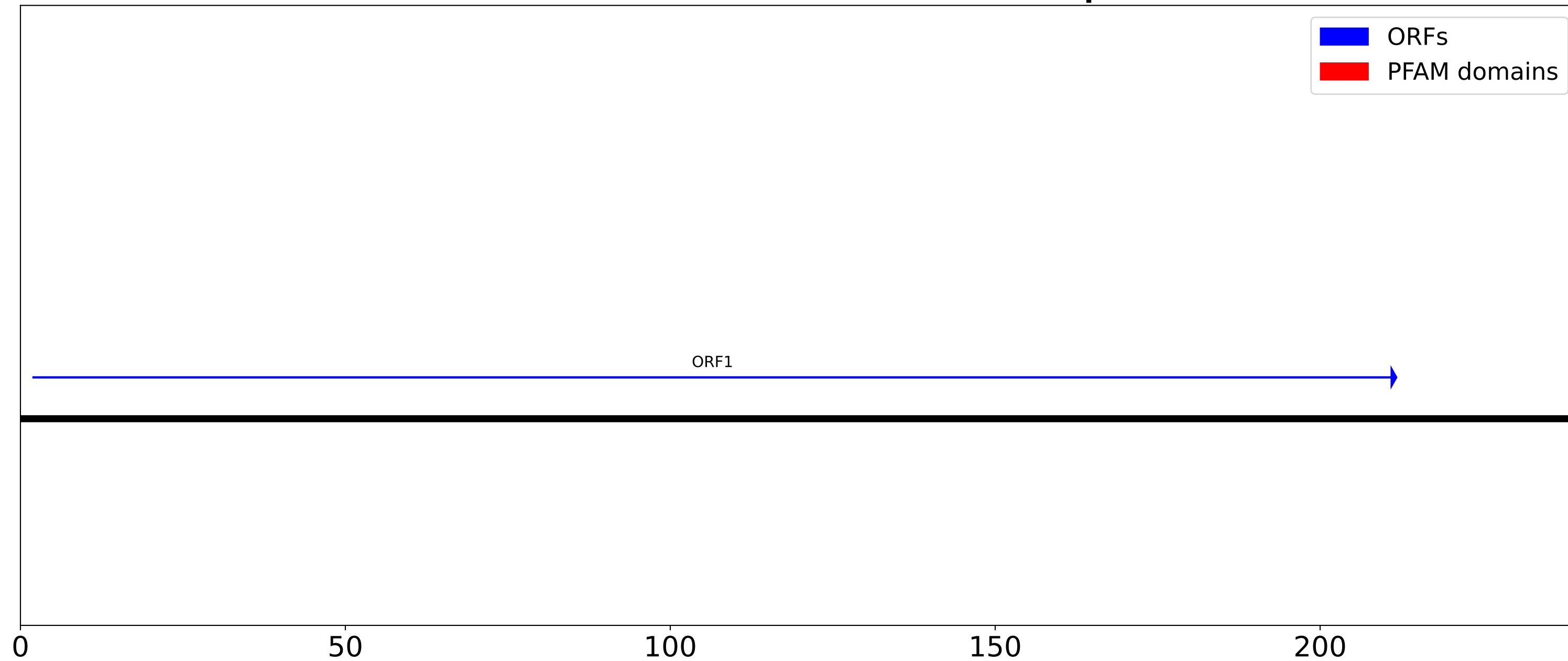
Before TEtrimmer 240 bp

TE: ltr_1_family_317

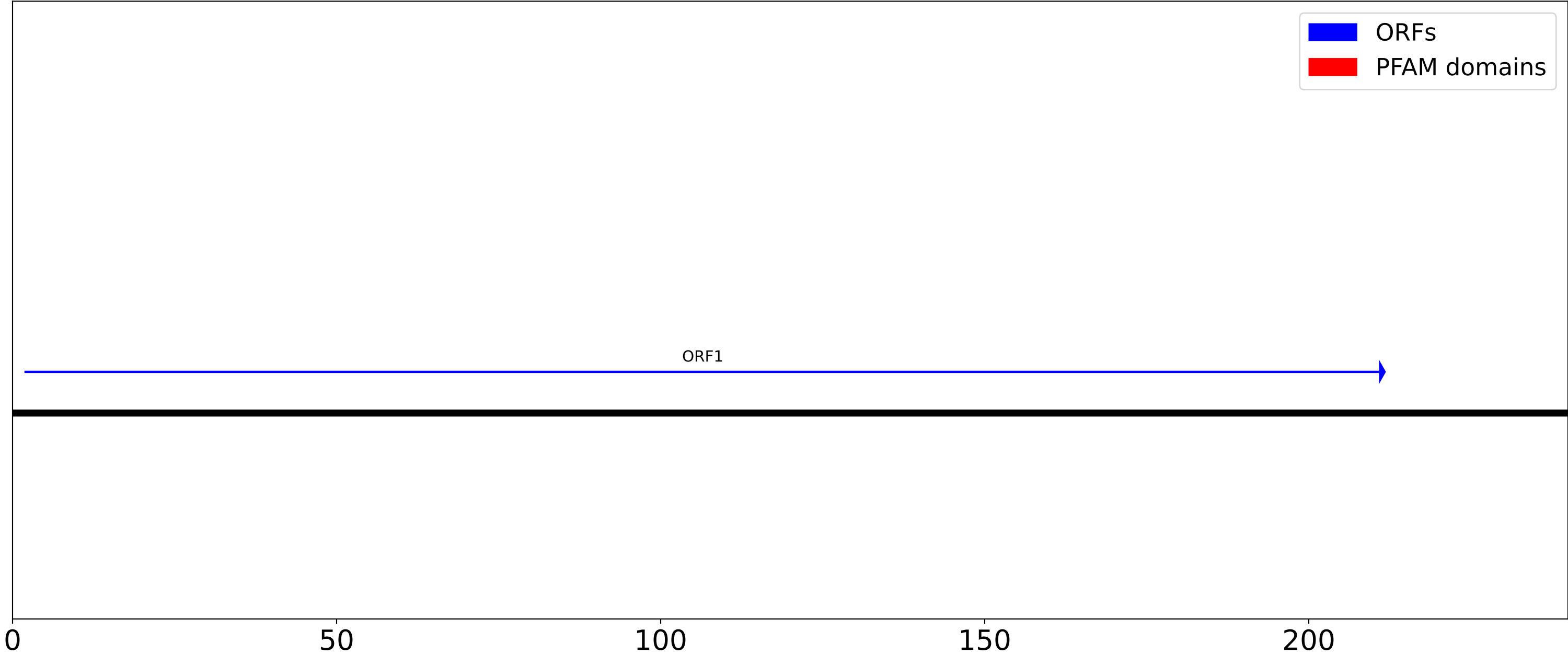
size: 240bp; fragments: 16; full length: 12 (≥ 216 bp)



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

