

Start crop Point

End crop Point

MSA length = 2436



Start crop Point

End crop Point

0

10

0

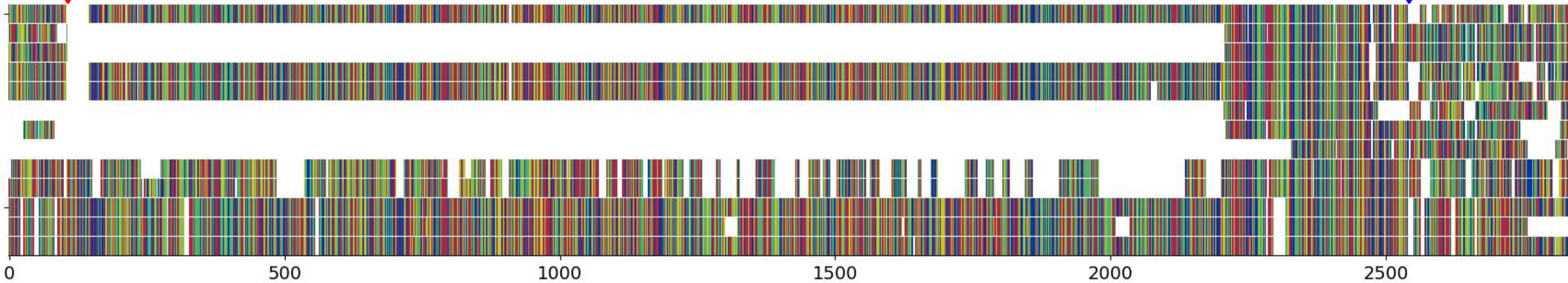
500

1000

1500

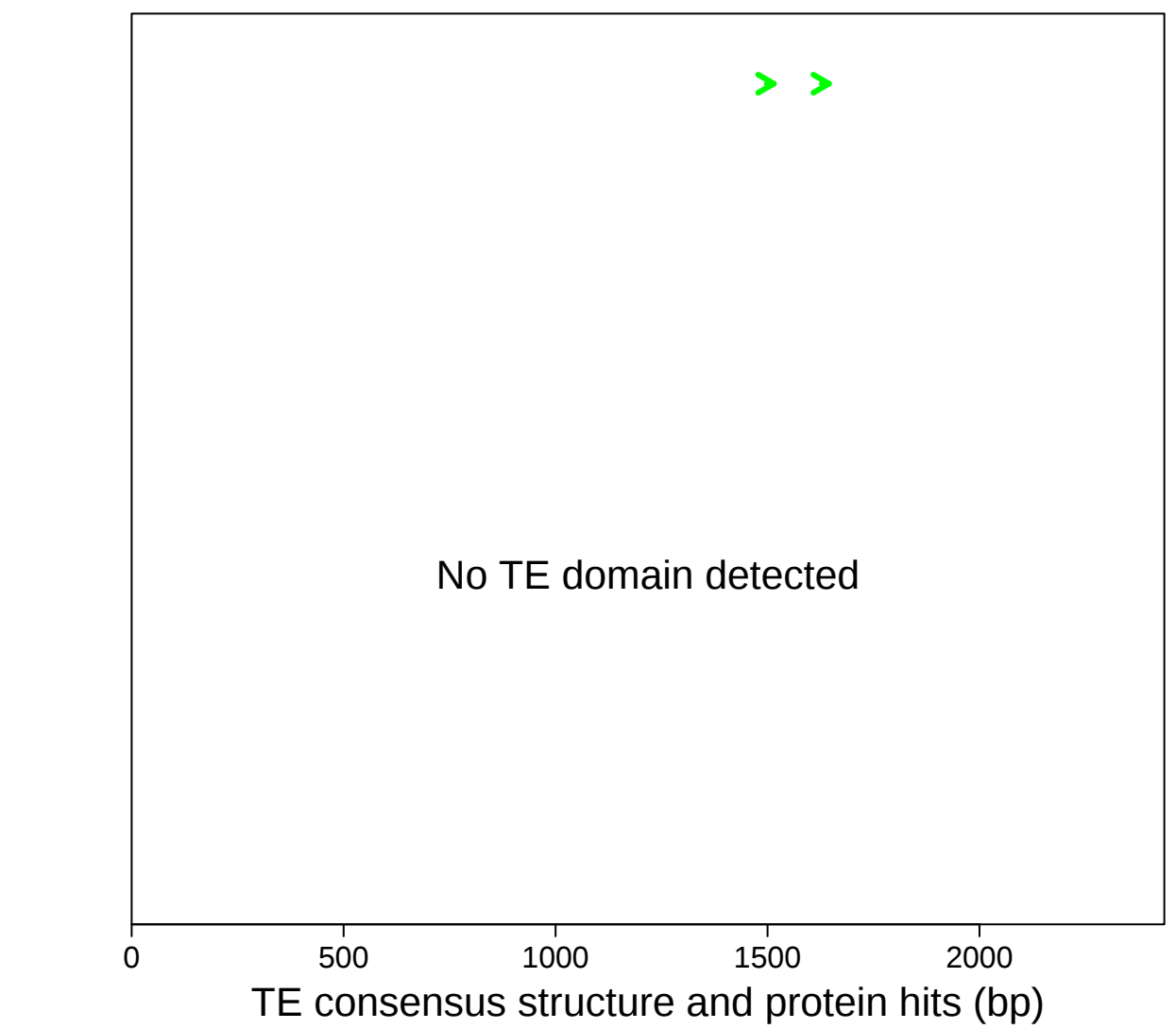
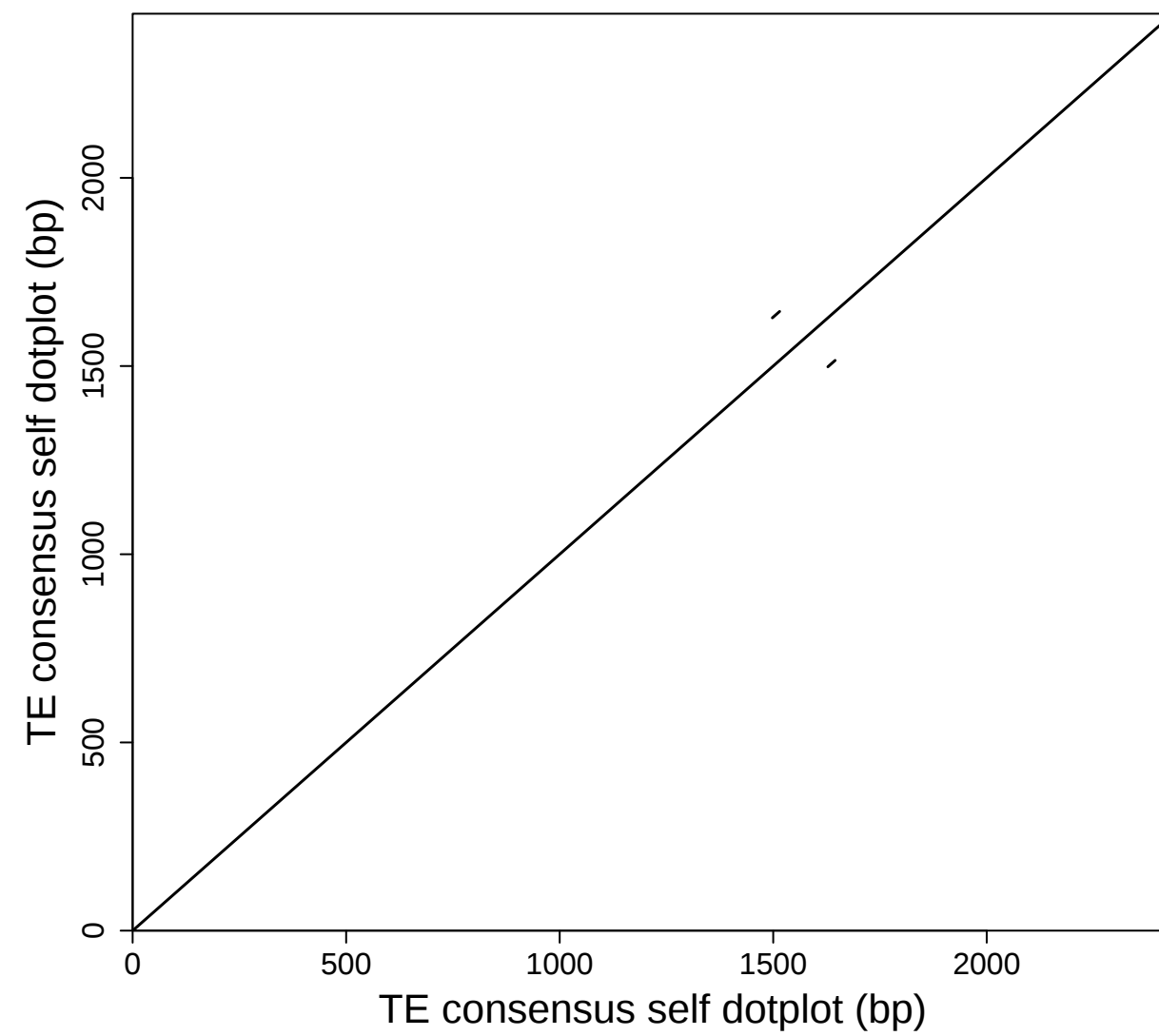
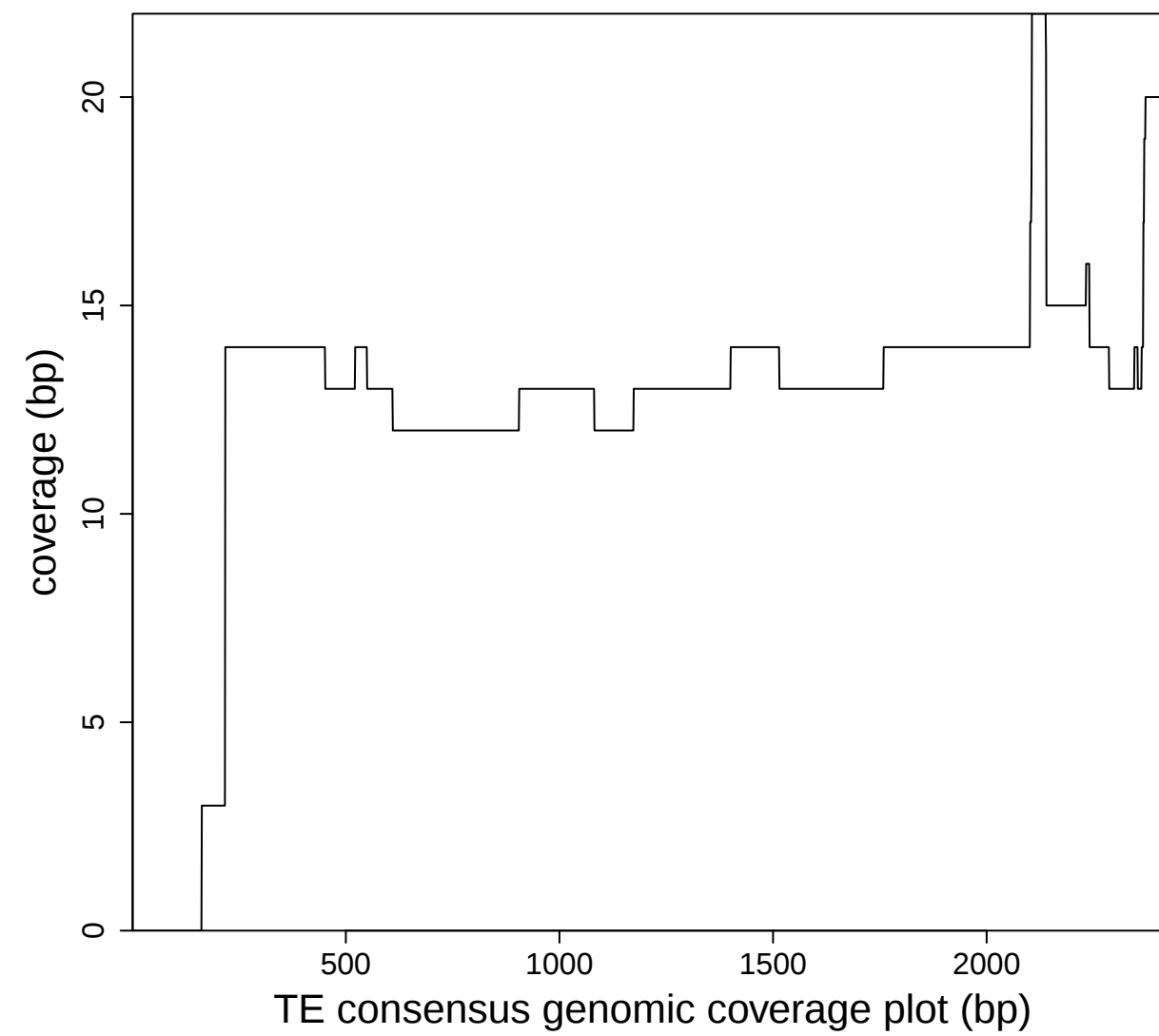
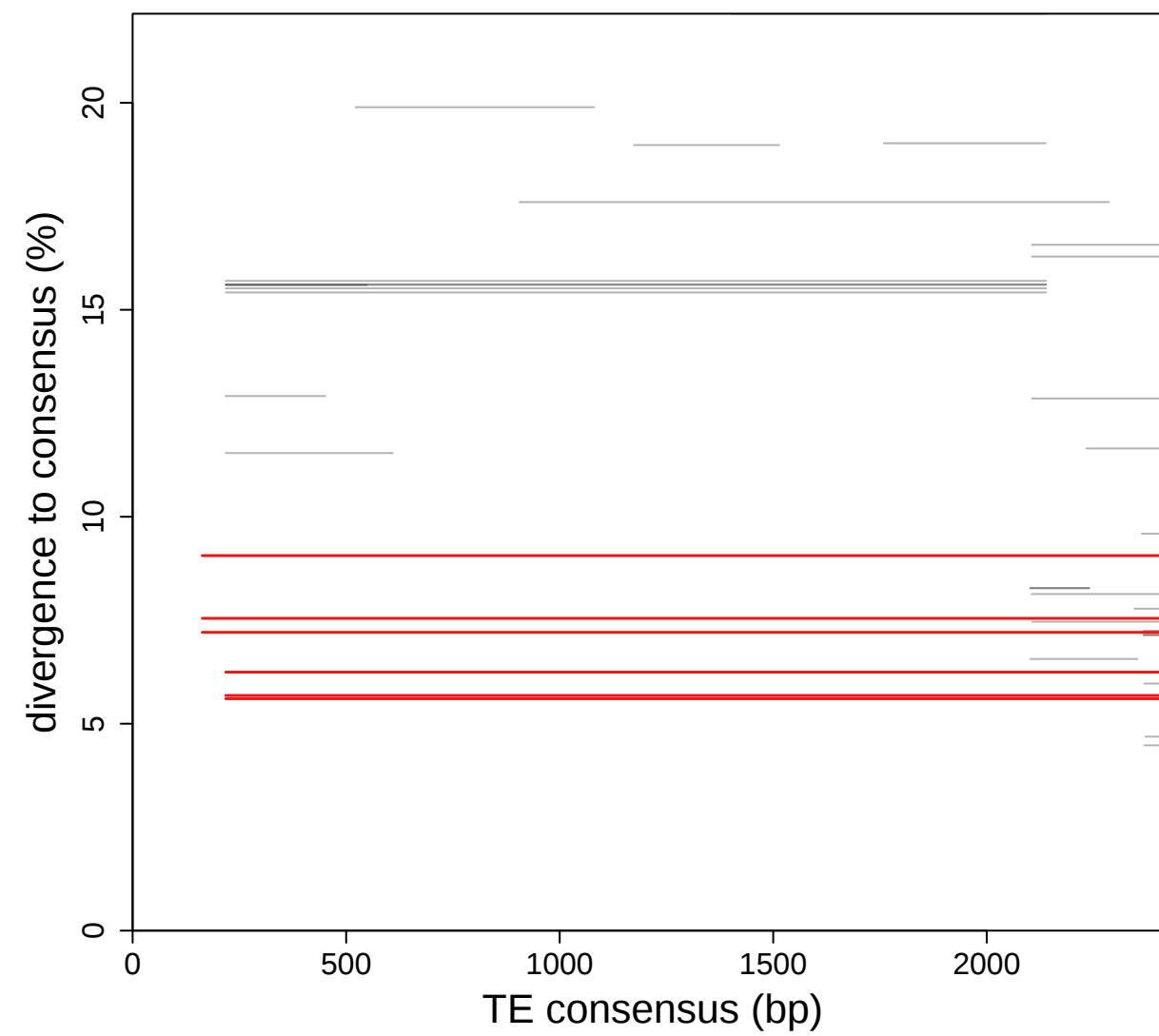
2000

2500



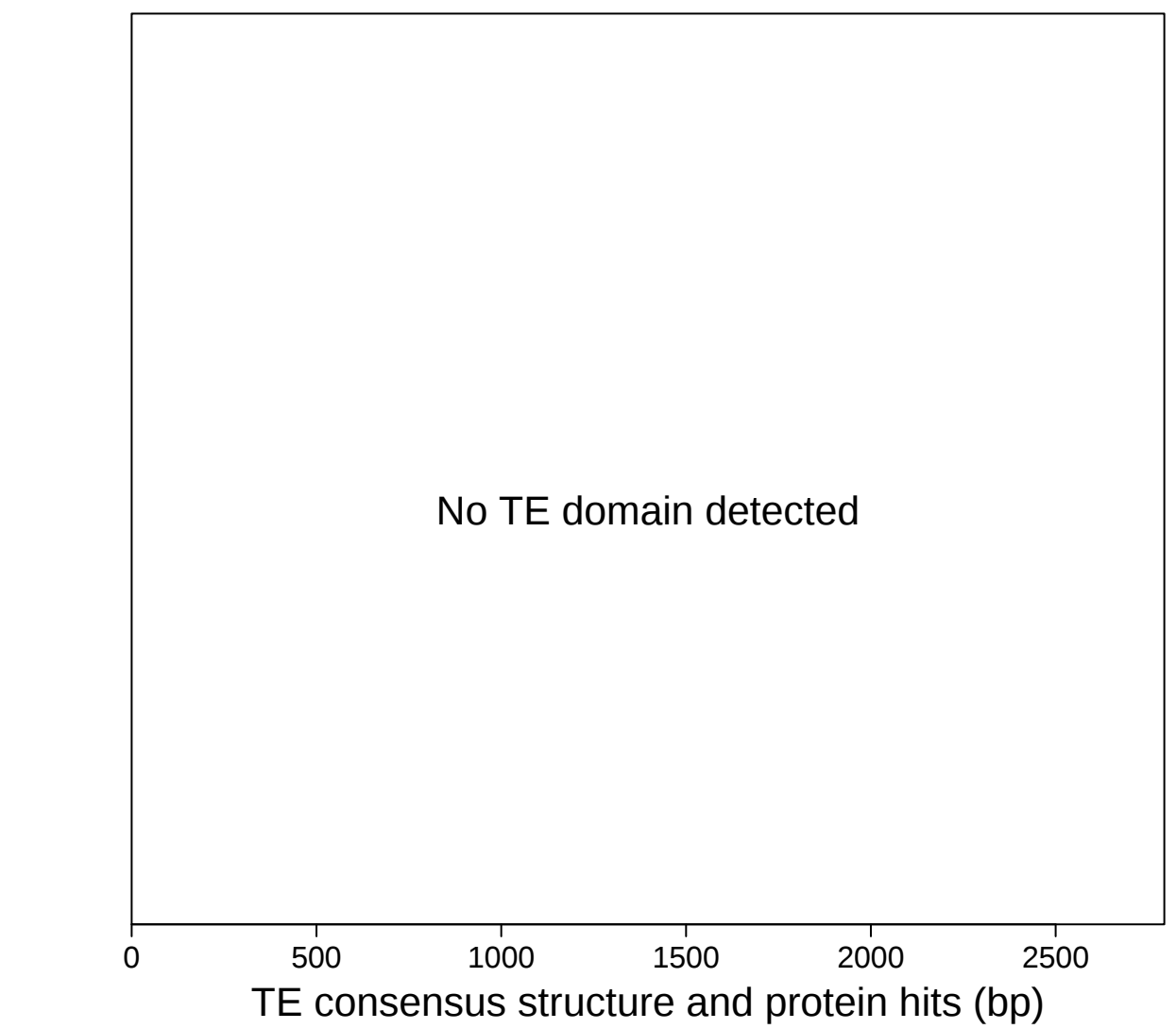
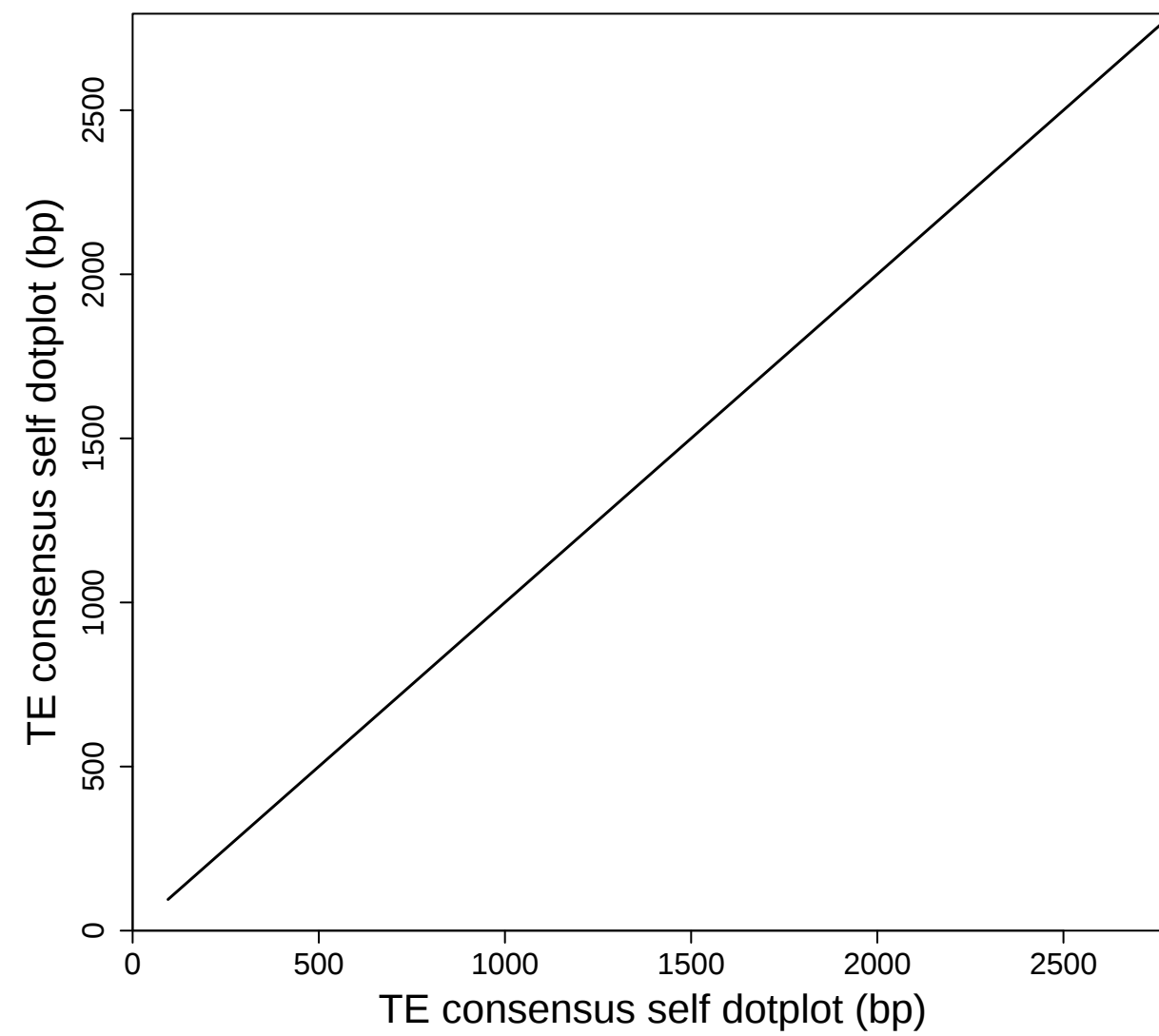
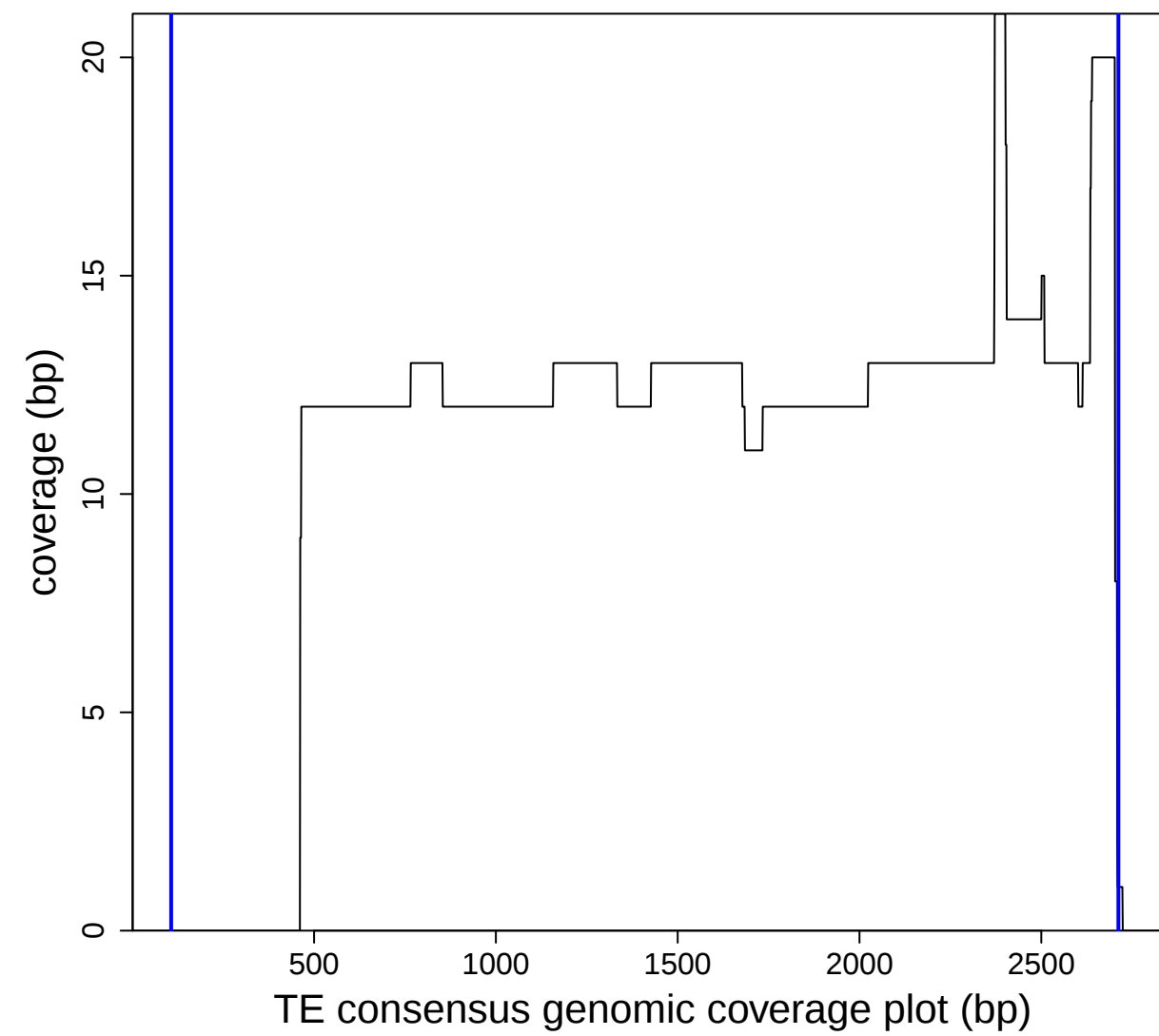
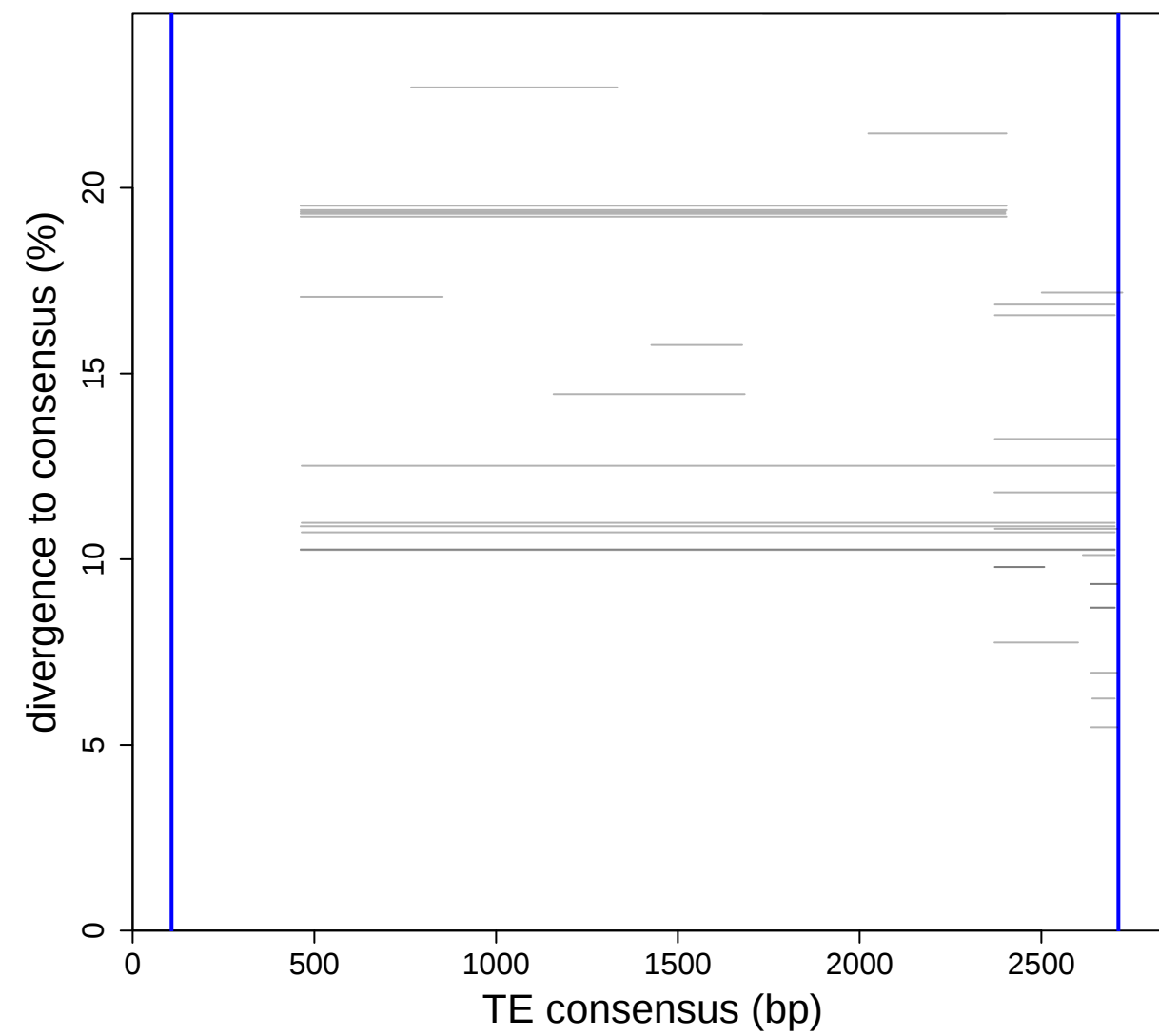
b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 2436bp; fragments: 36; full length: 6 (≥ 2192.4 bp)

After TEtrimmer 2436 bp

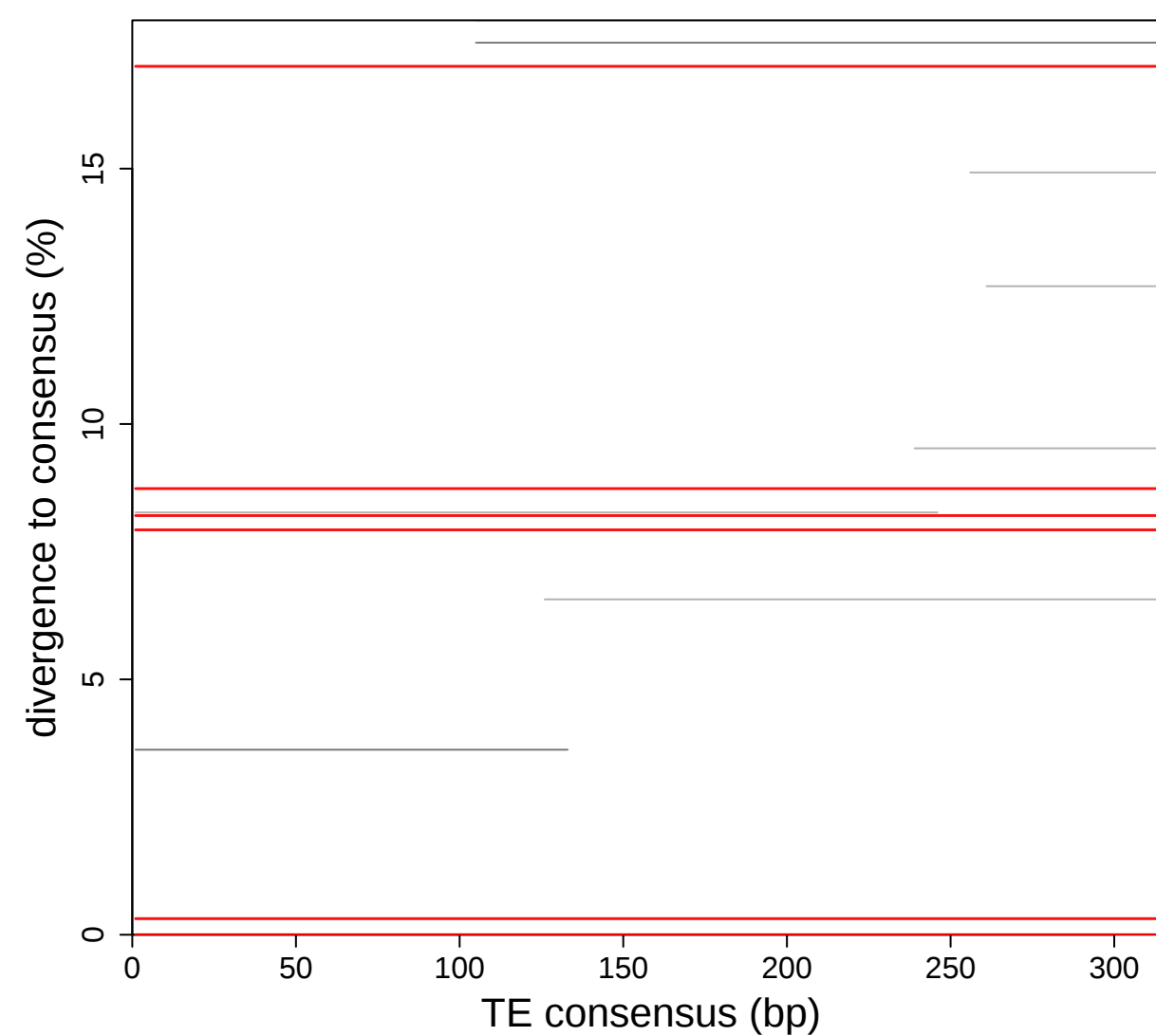


.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 2862bp; fragments: 34; full length: 0 (>=2575.8bp)

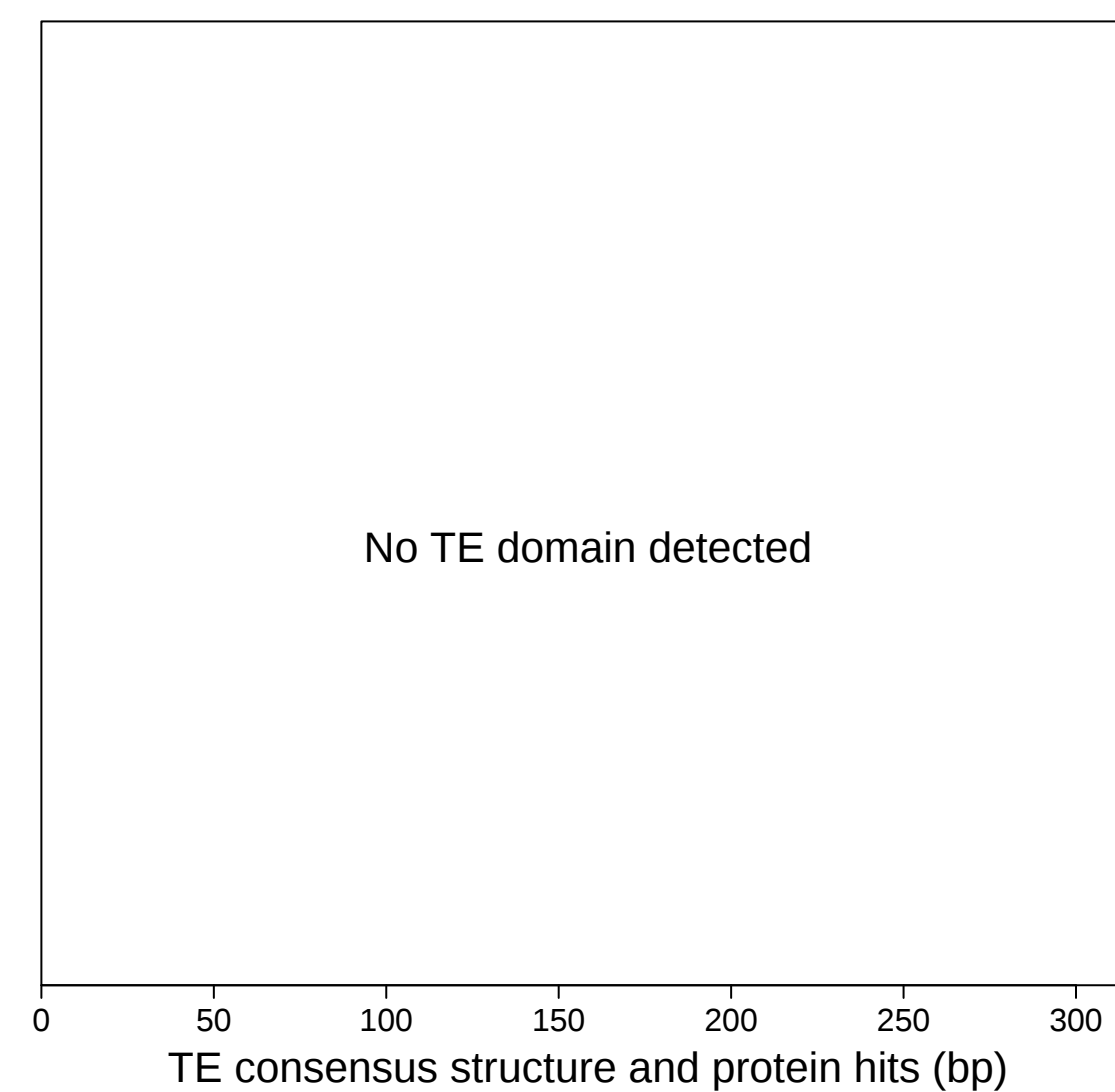
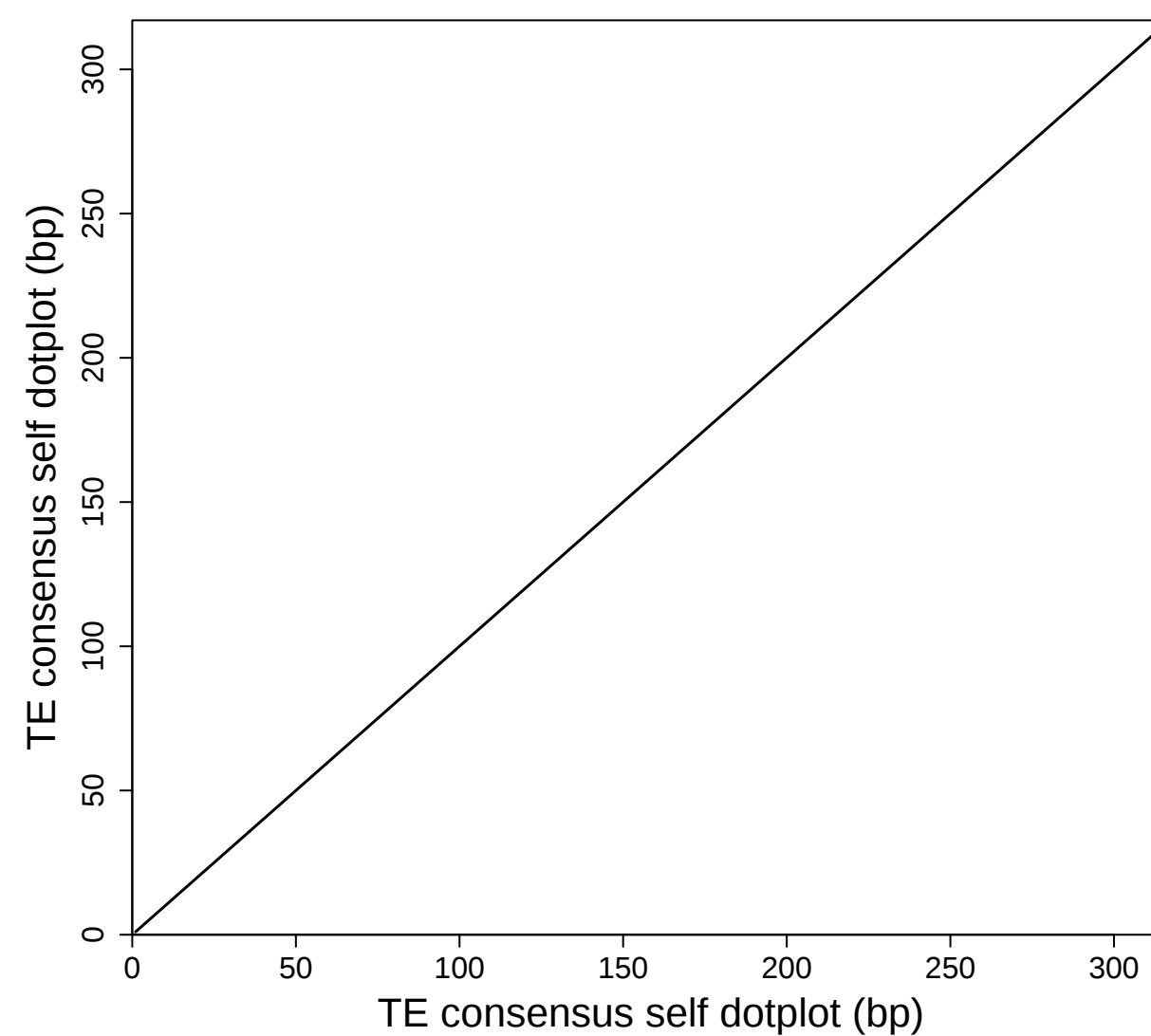
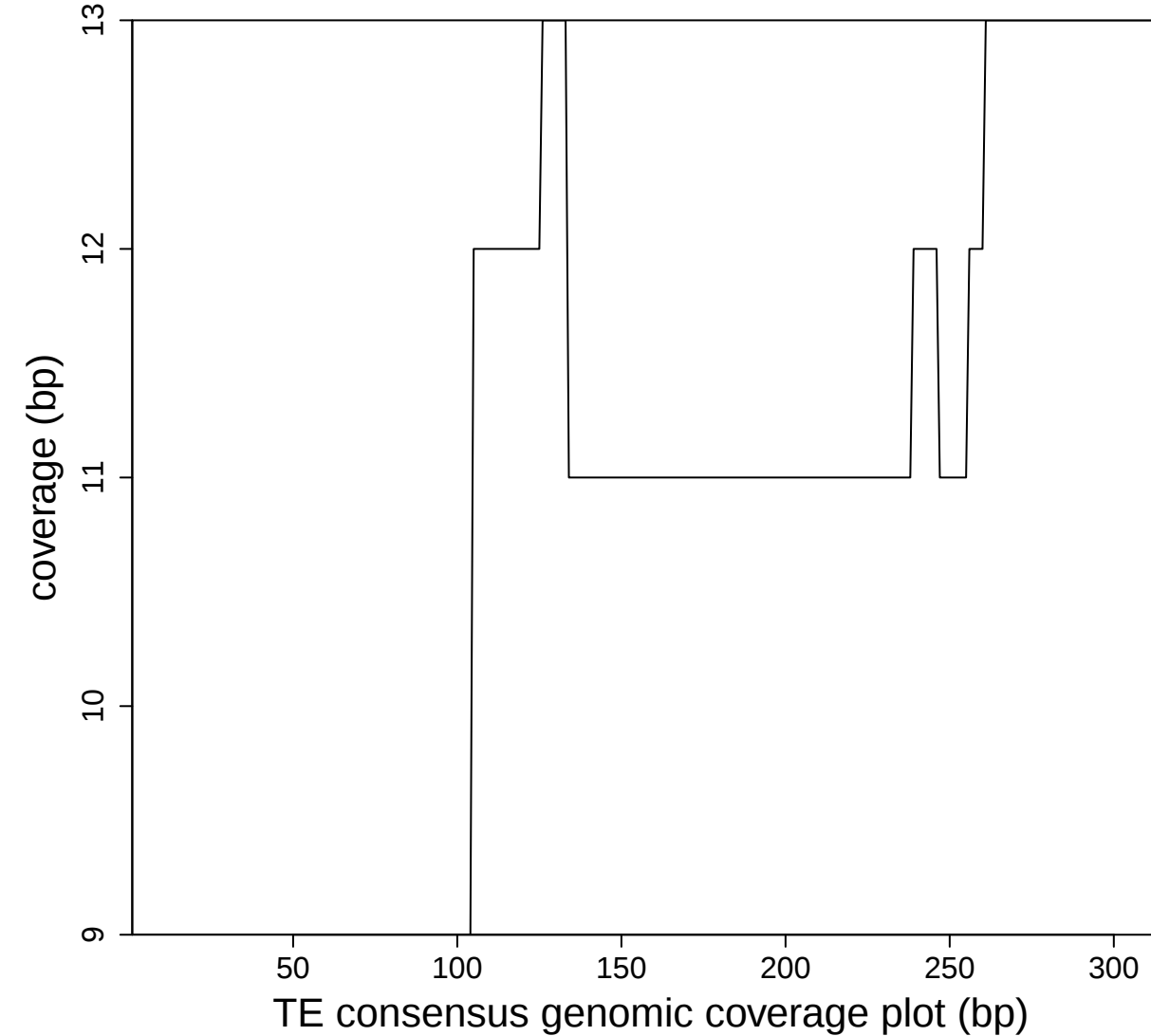
After TEtrimmer Extended plot Blue lines are boundaries



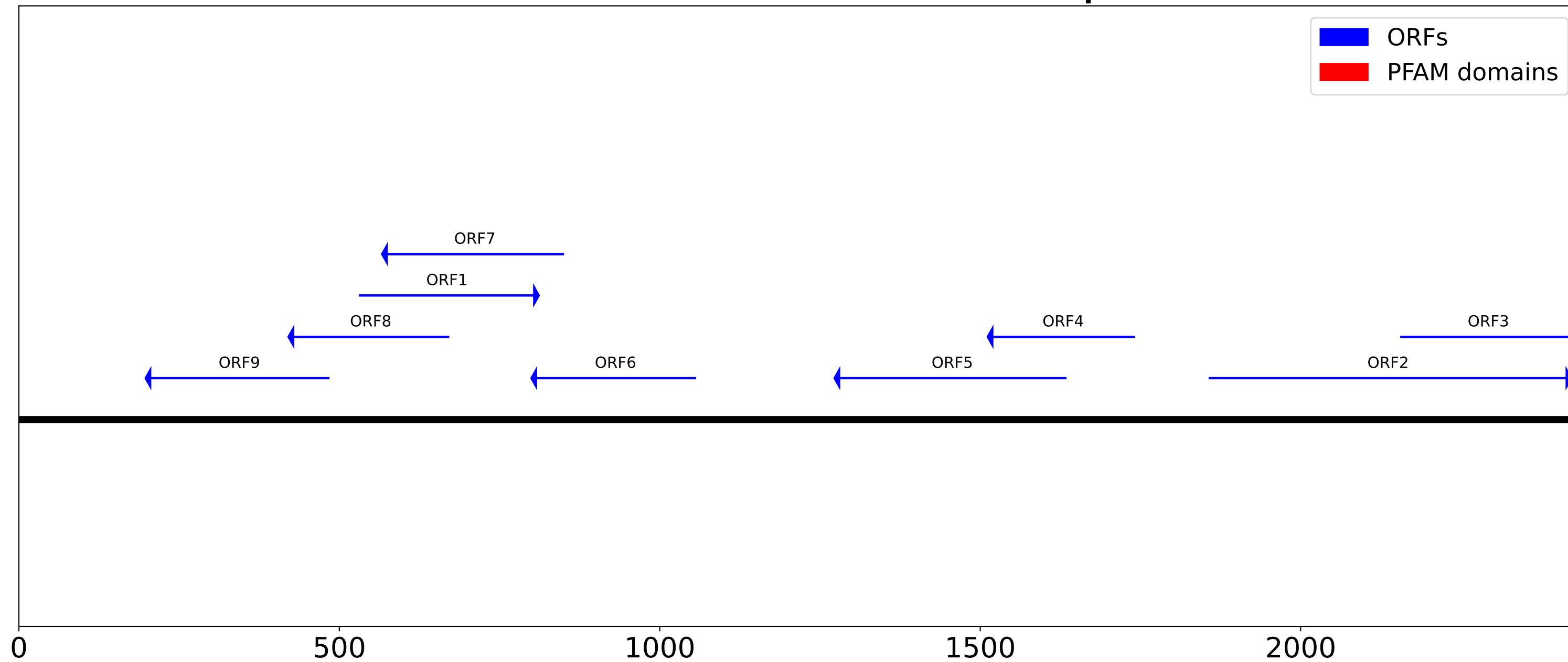
TE: ltr_1_family_254
size: 317bp; fragments: 16; full length: 6 (≥ 285.3 bp)



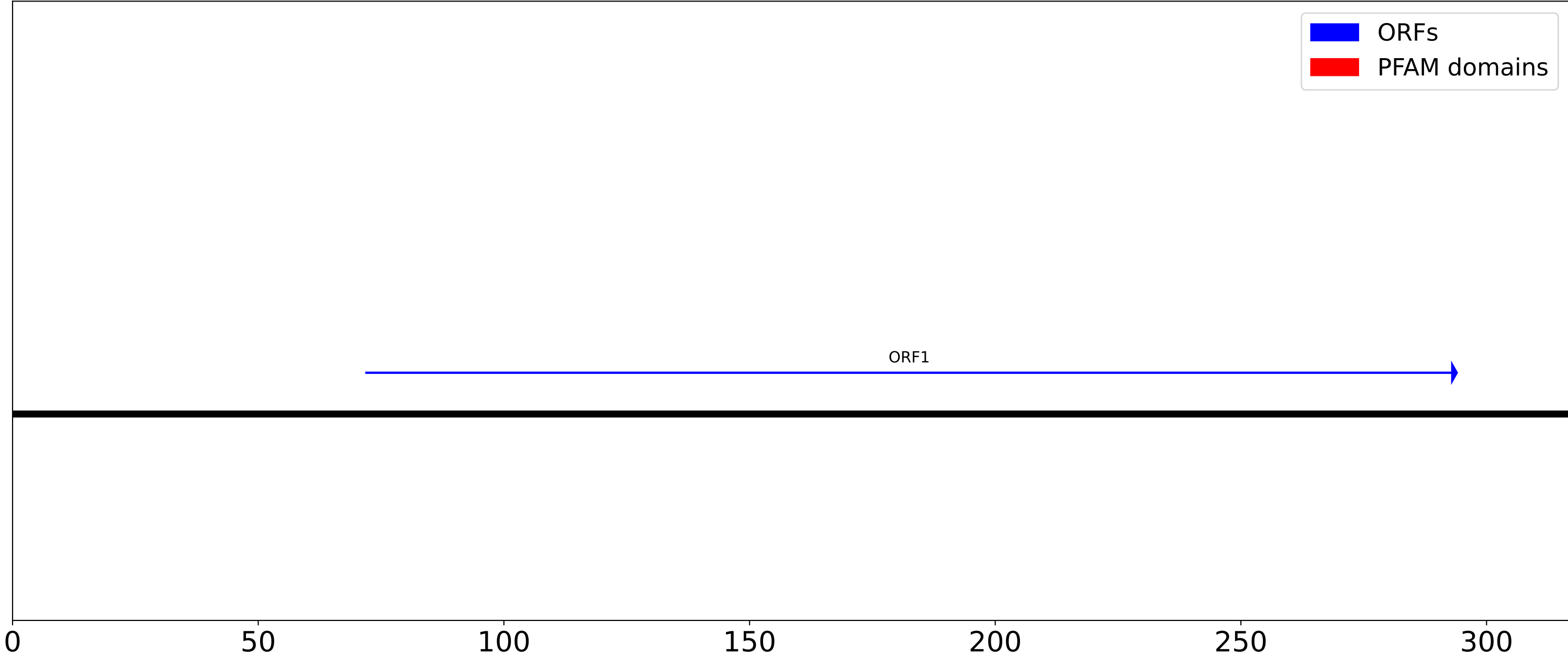
Before TEtrimmer 317 bp



After TEtrimmer ORF and PFAM domain plot



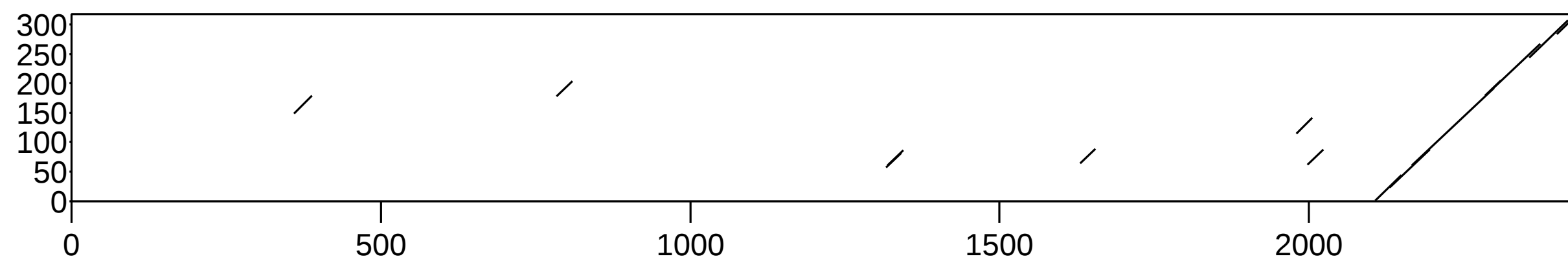
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

sensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)