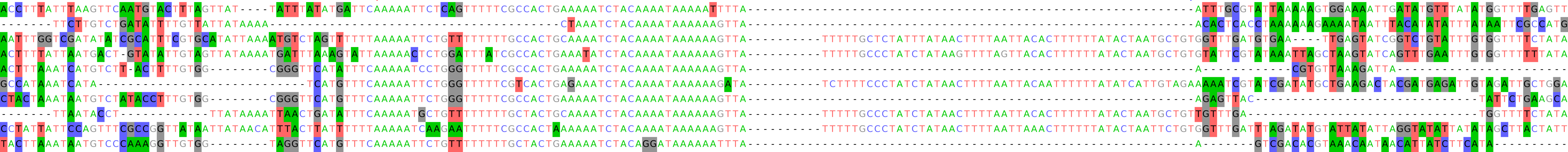


Start Crop Point

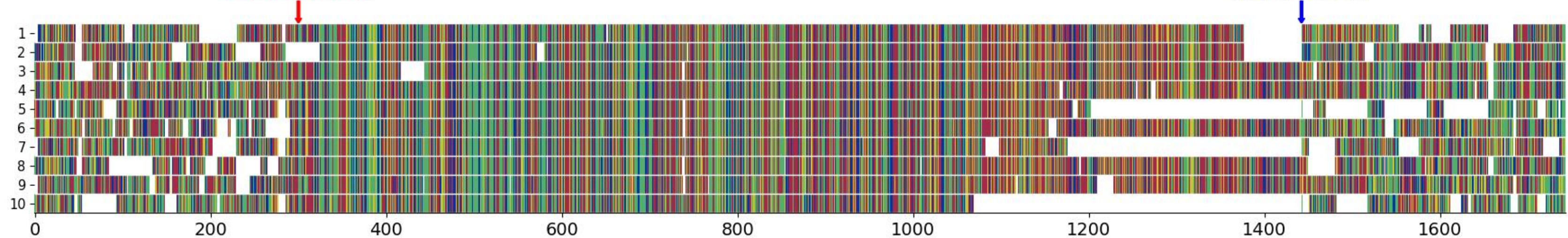
End Crop Point

MSA length = 1143



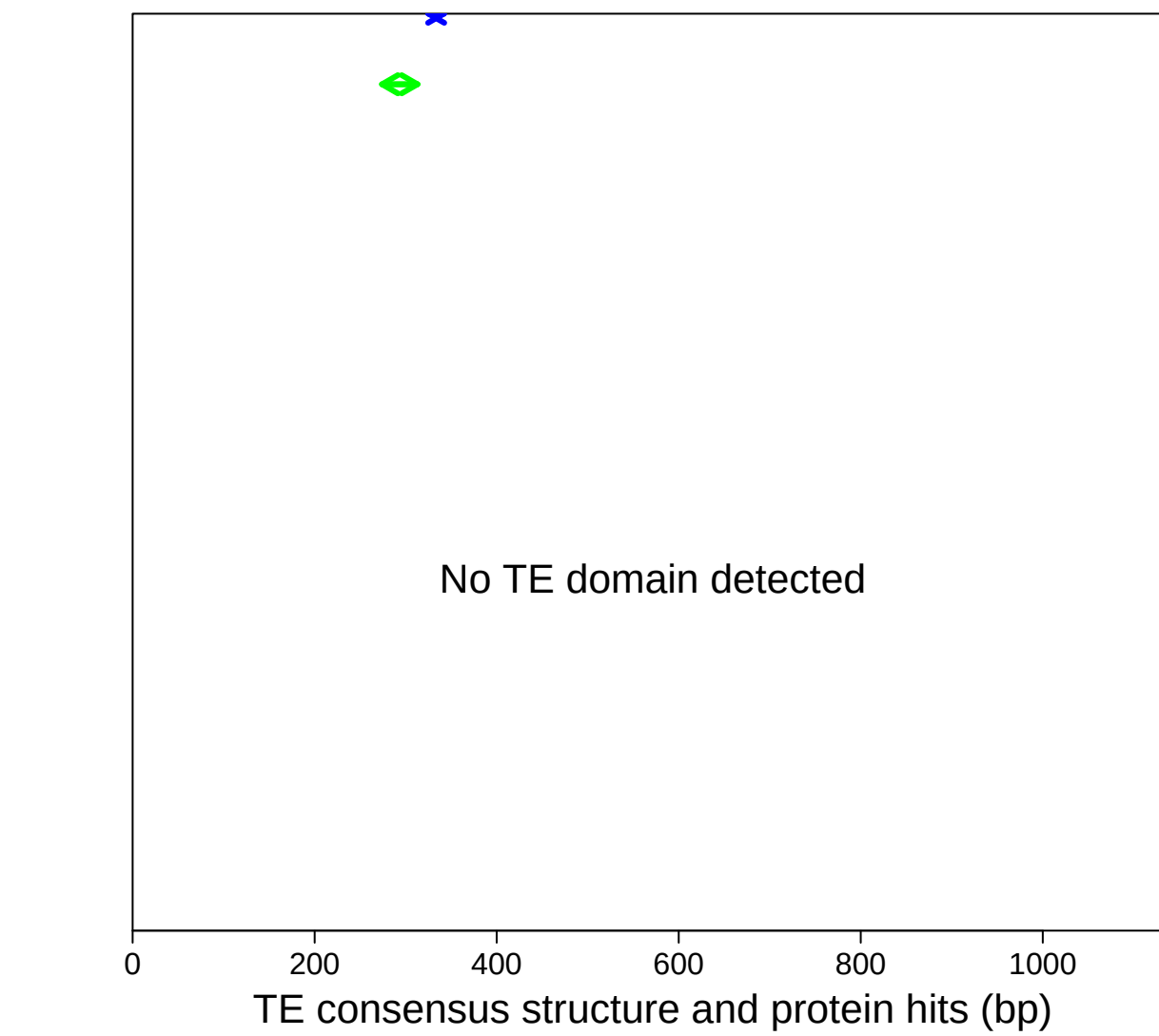
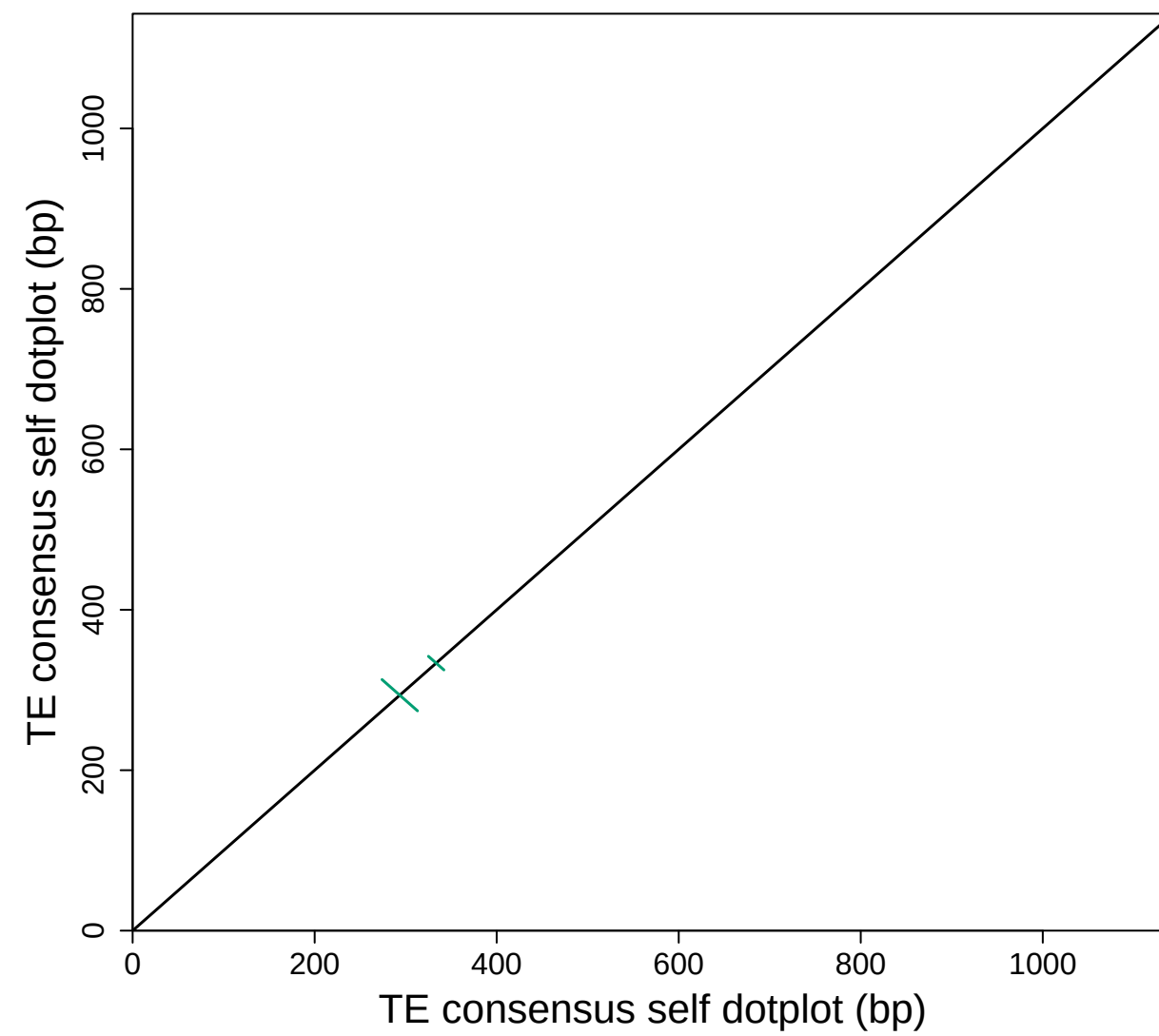
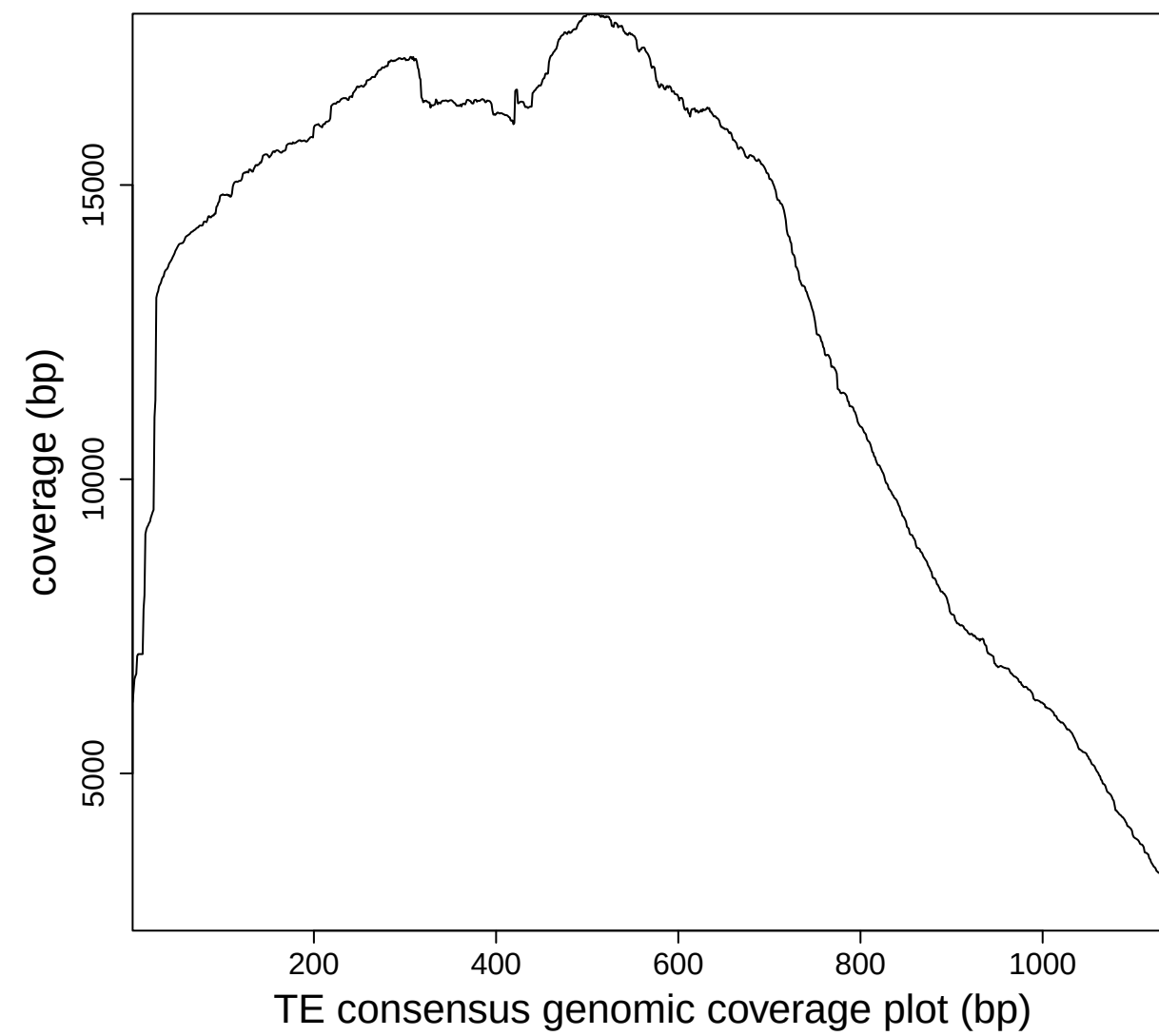
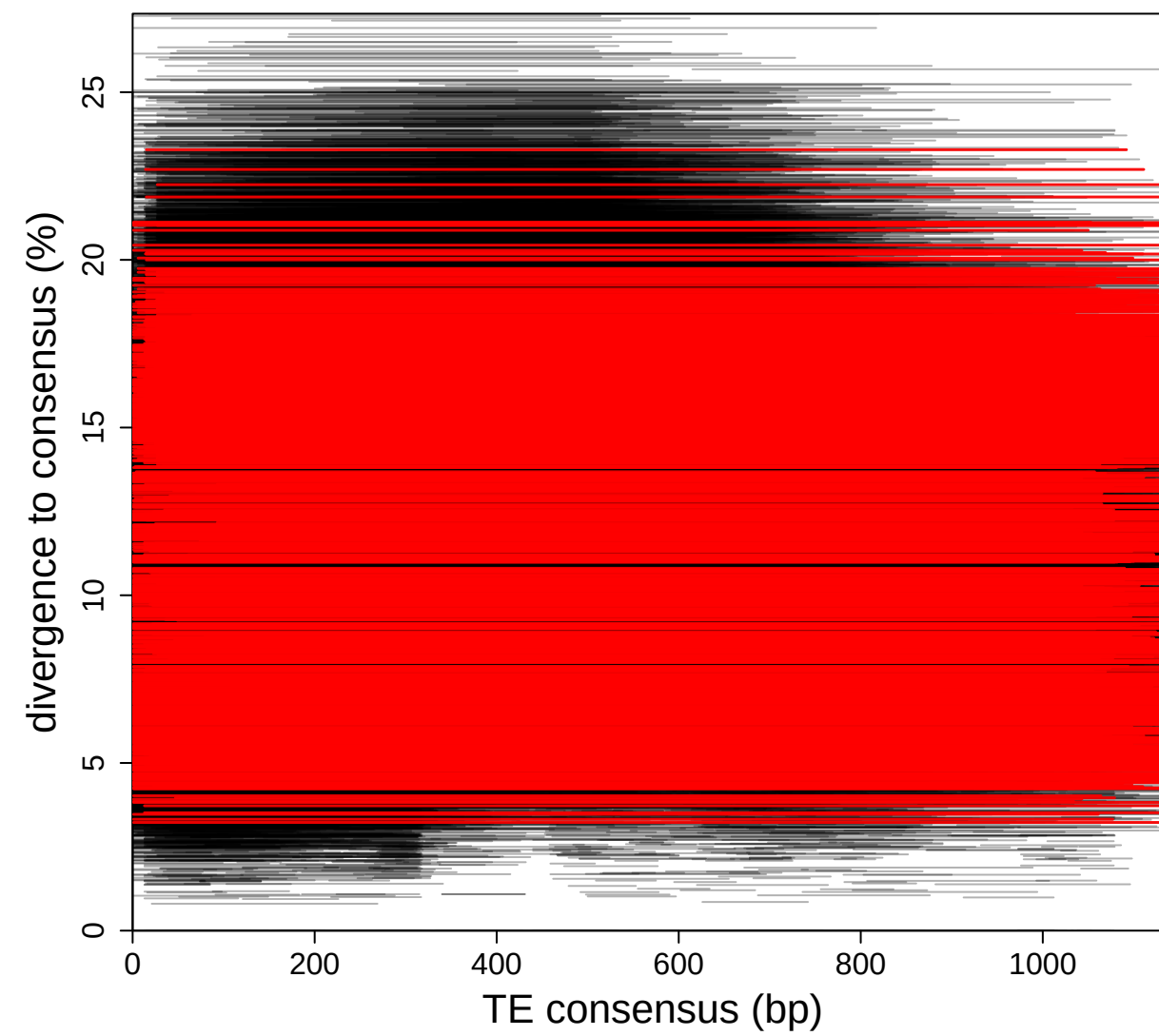
Start crop Point

End crop Point



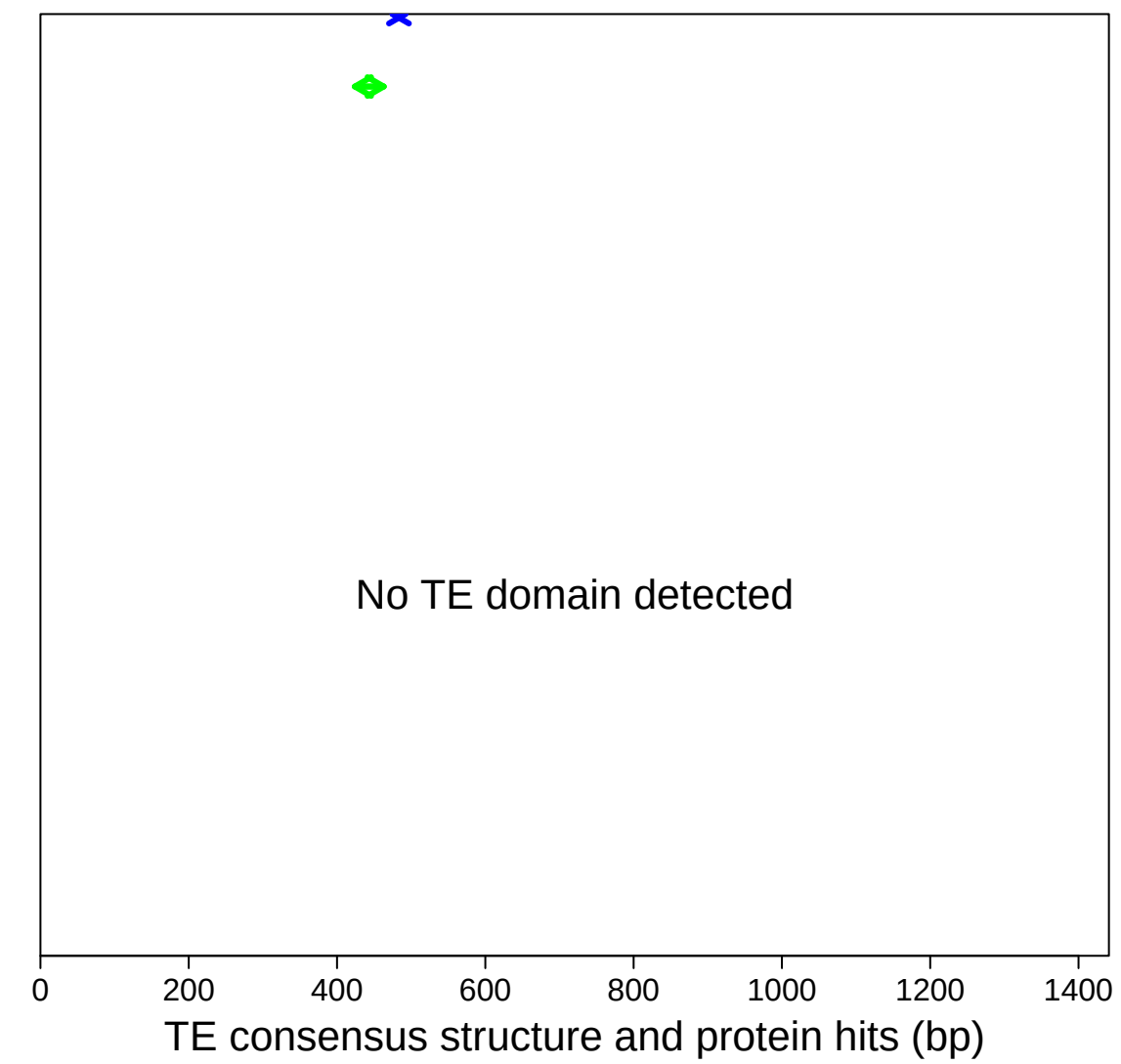
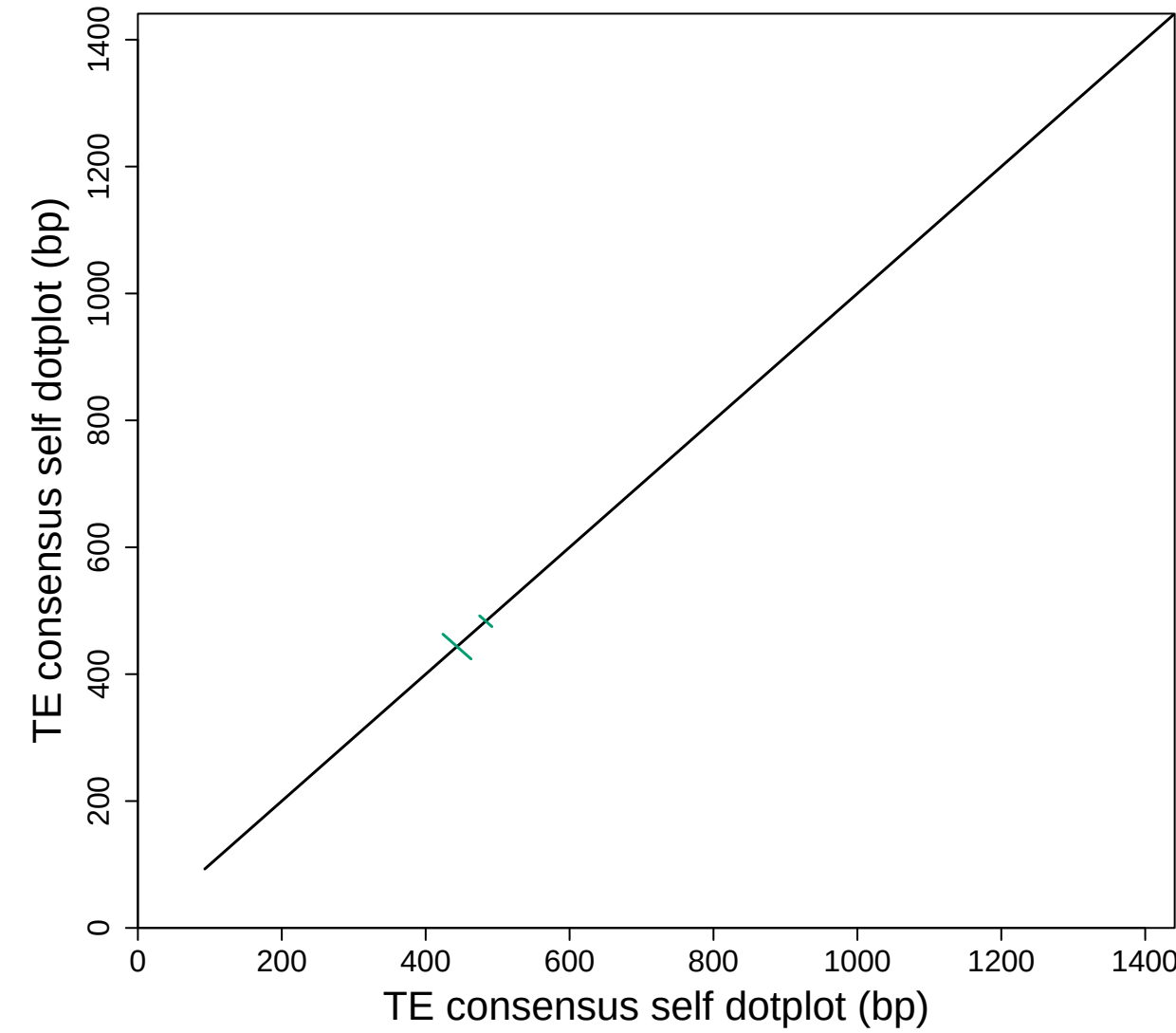
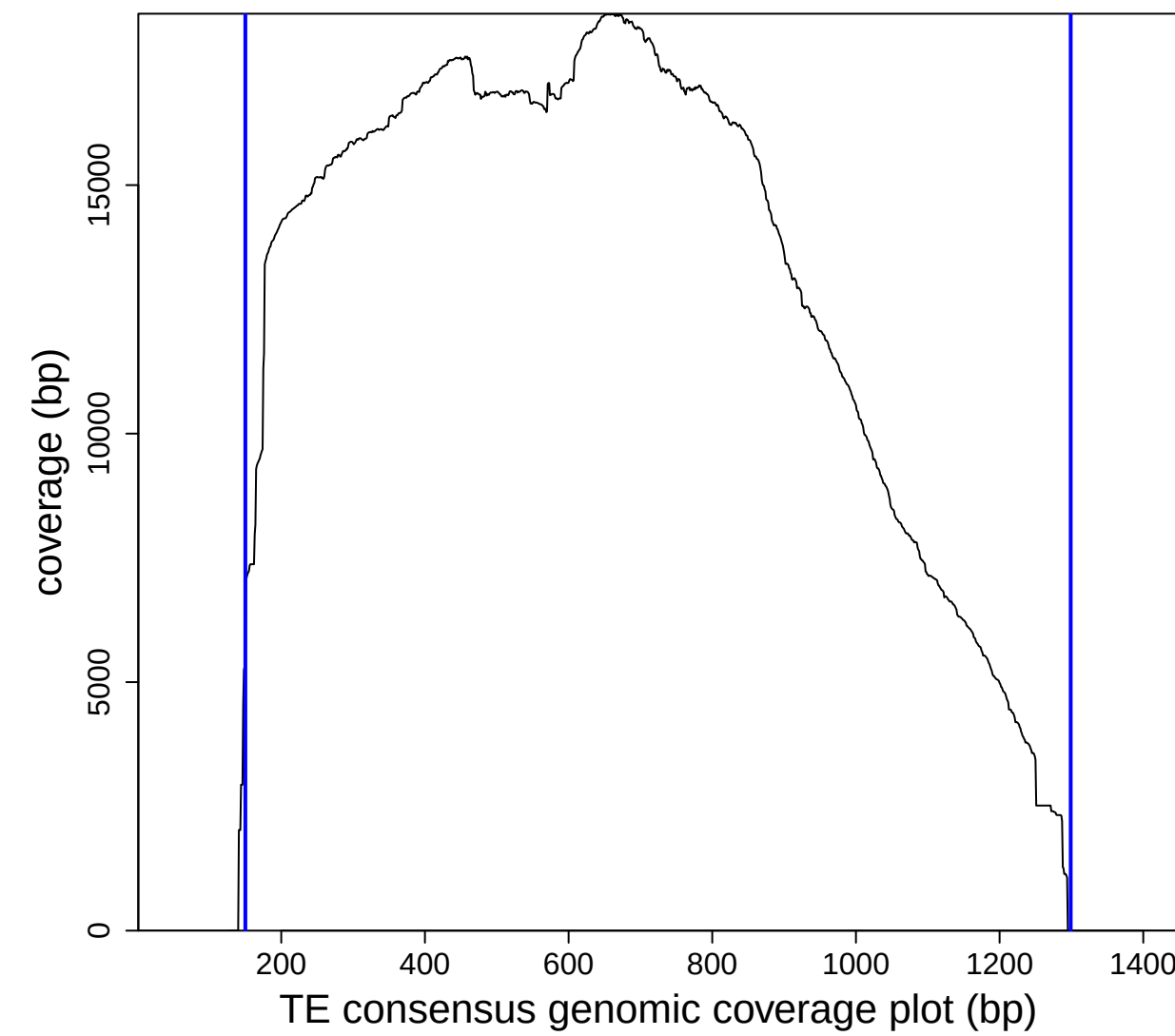
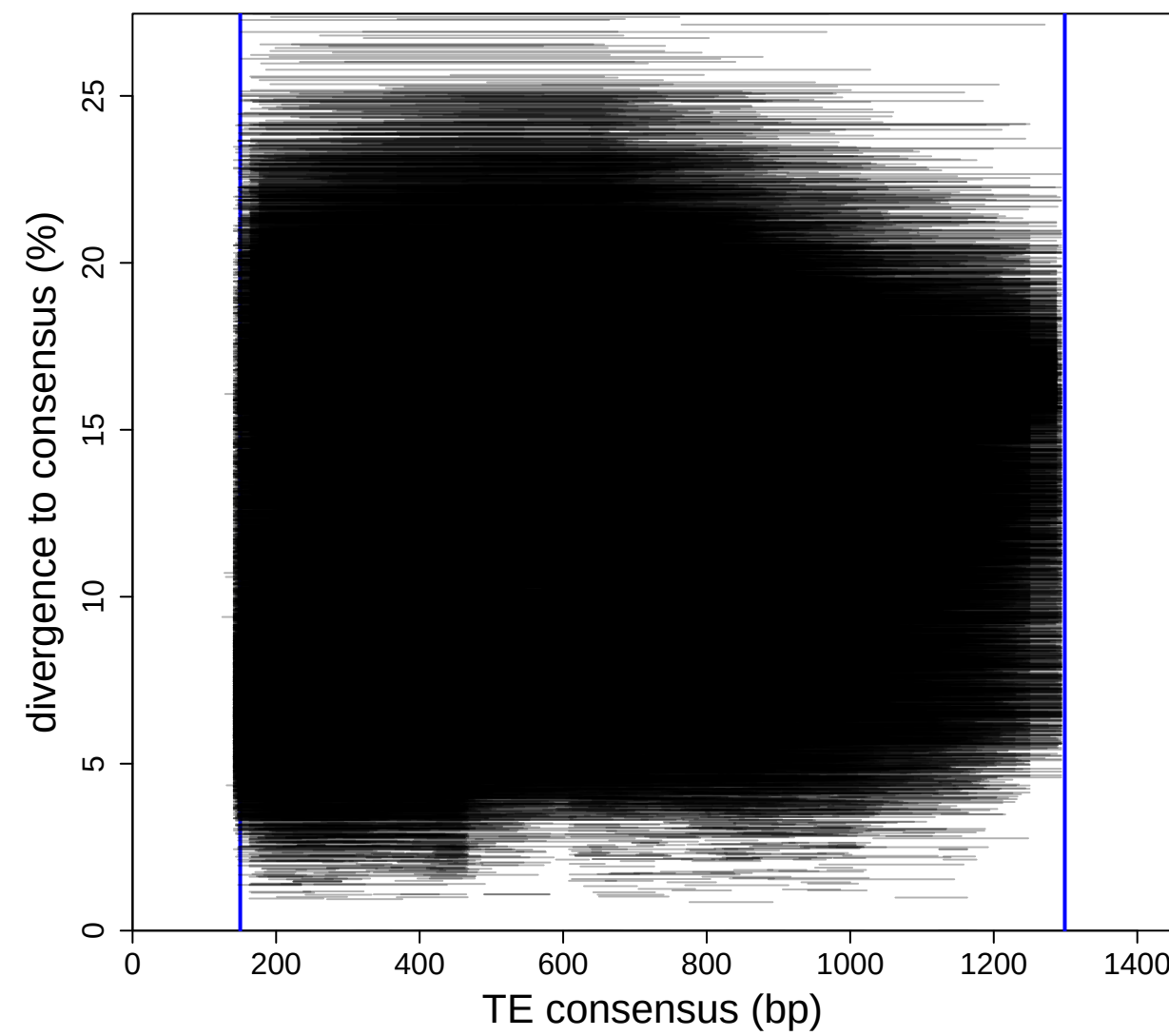
ed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 1143bp; fragments: 30230; full length: 2061 (>=1028.7bp)

After TEtrimmer 1143 bp



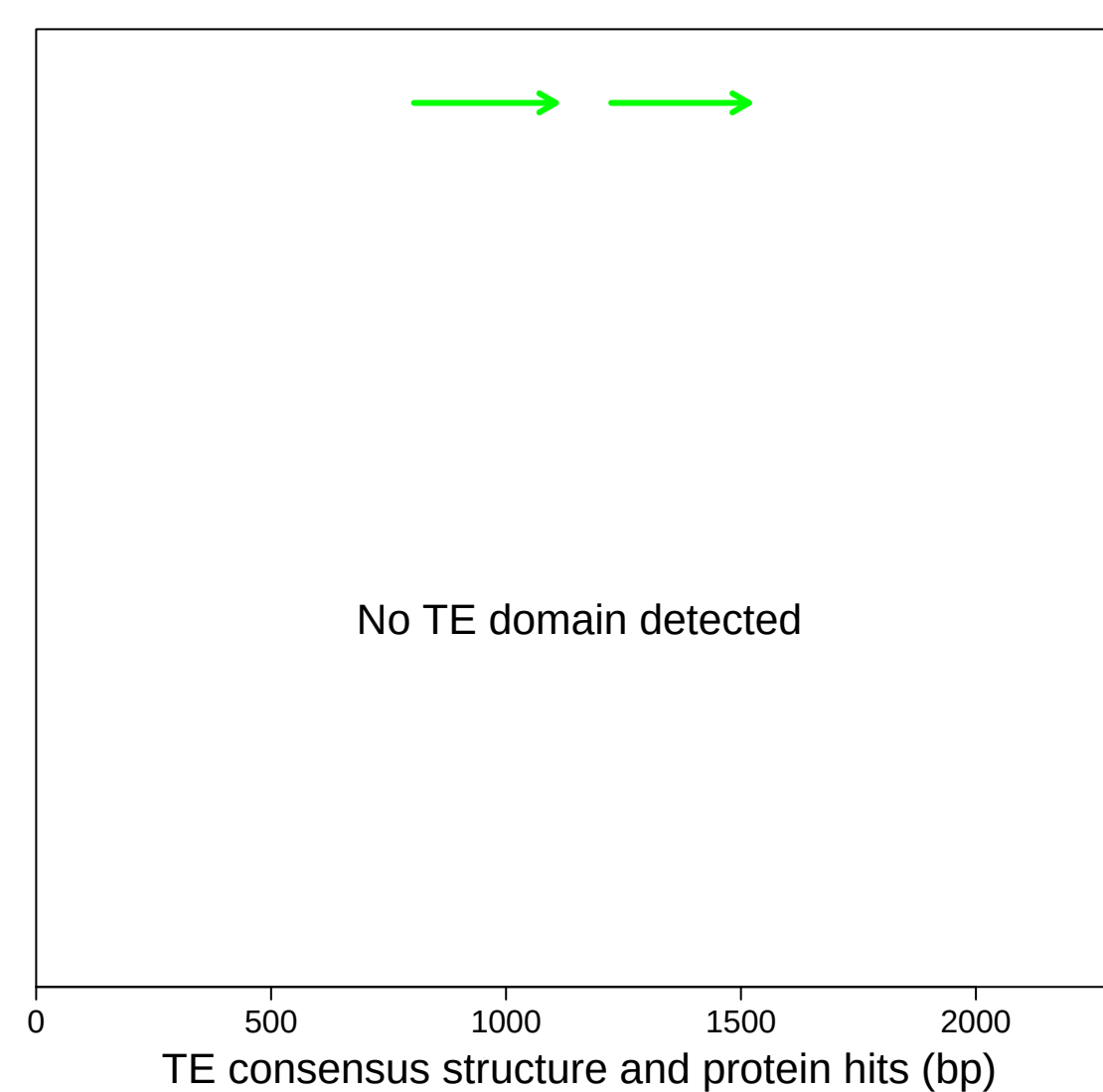
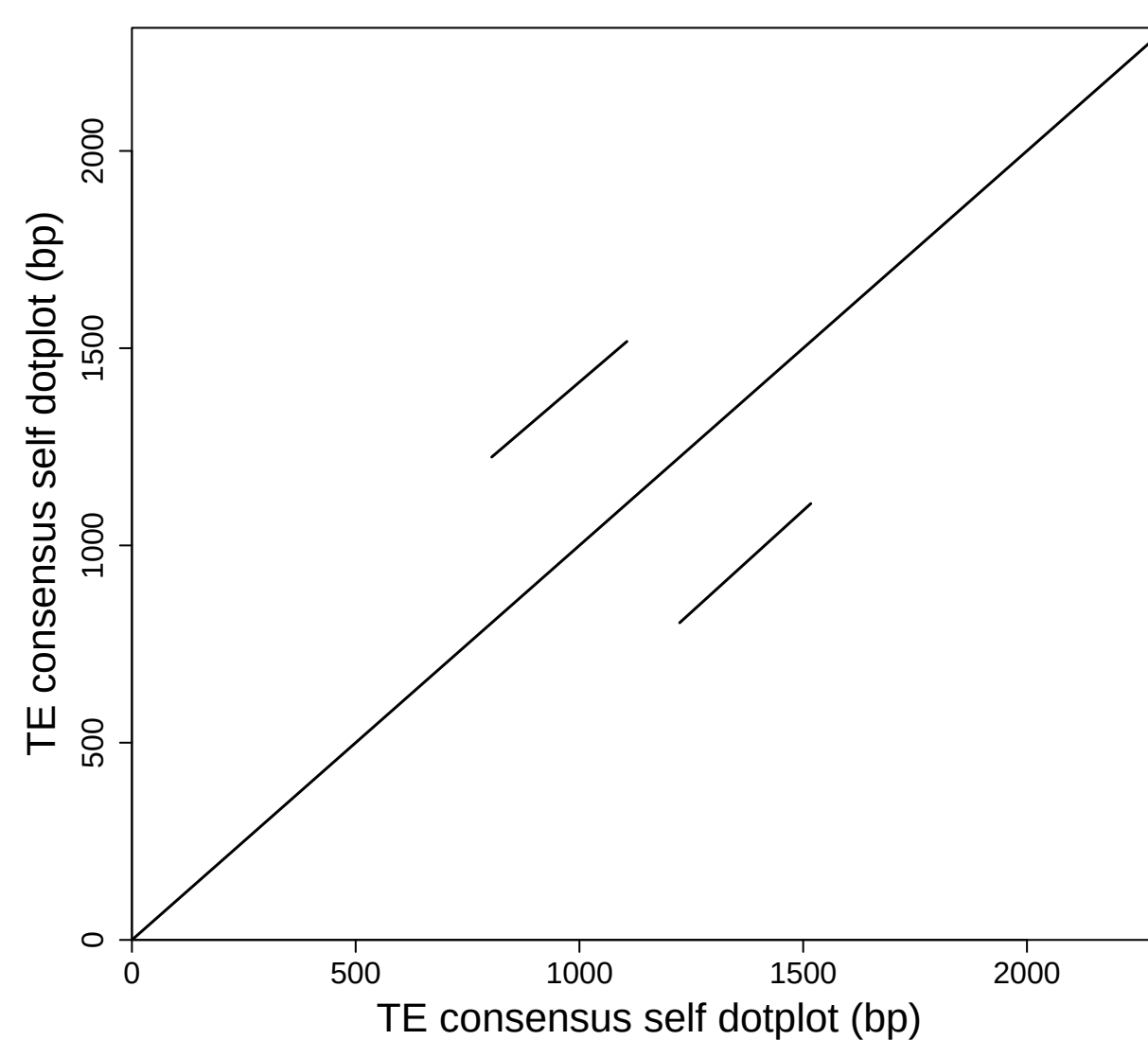
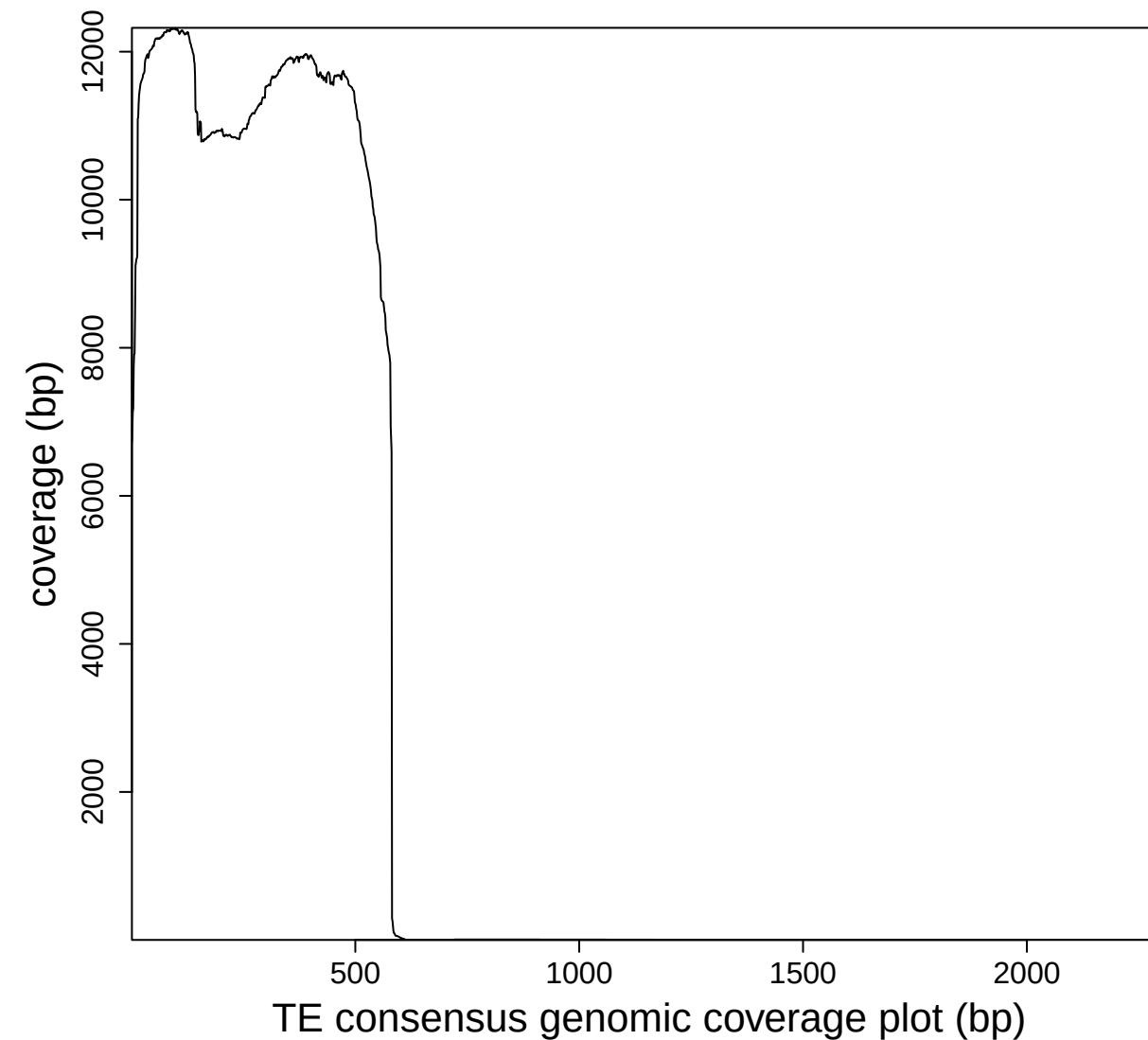
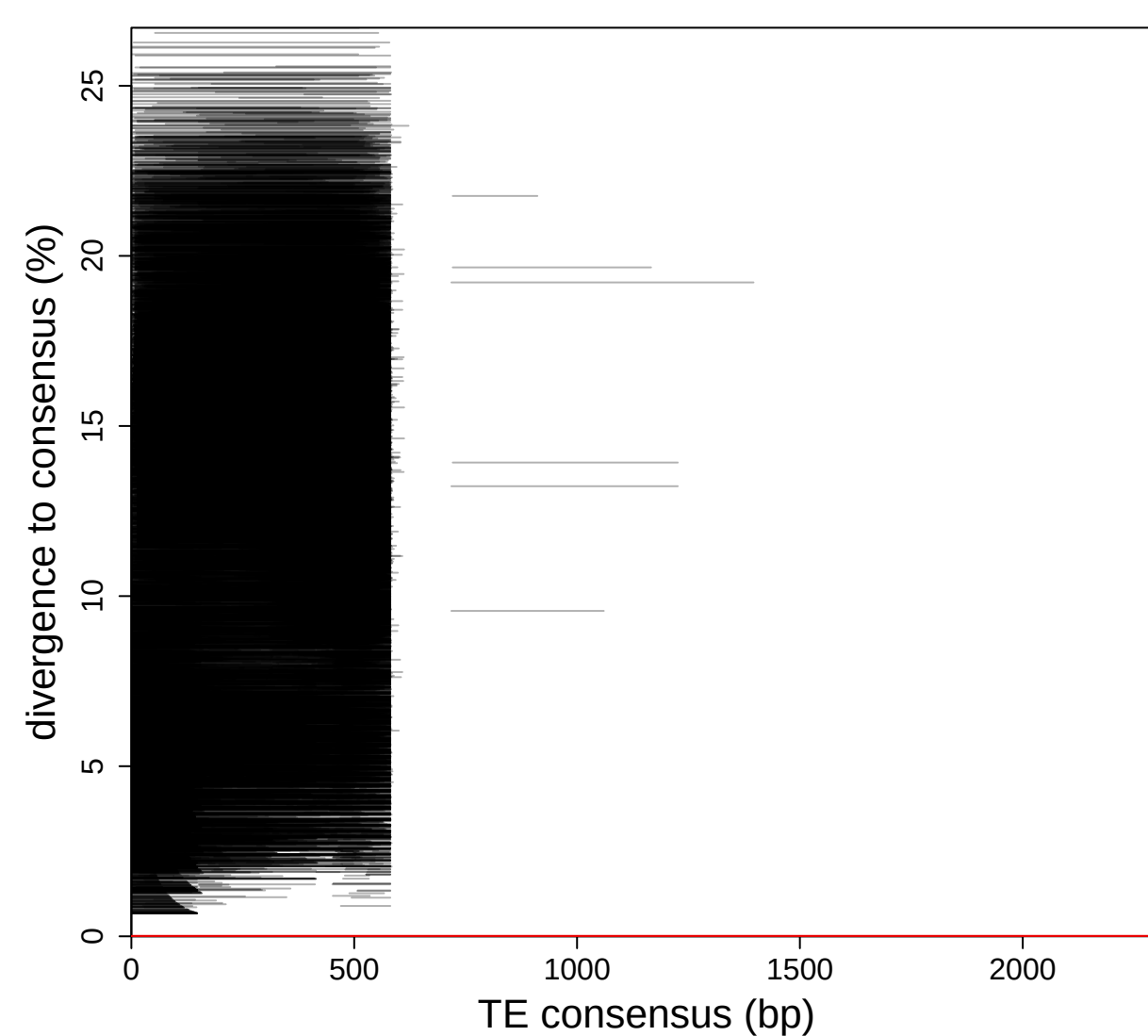
3.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 1449bp; fragments: 31031; full length: 0 (>=1304.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

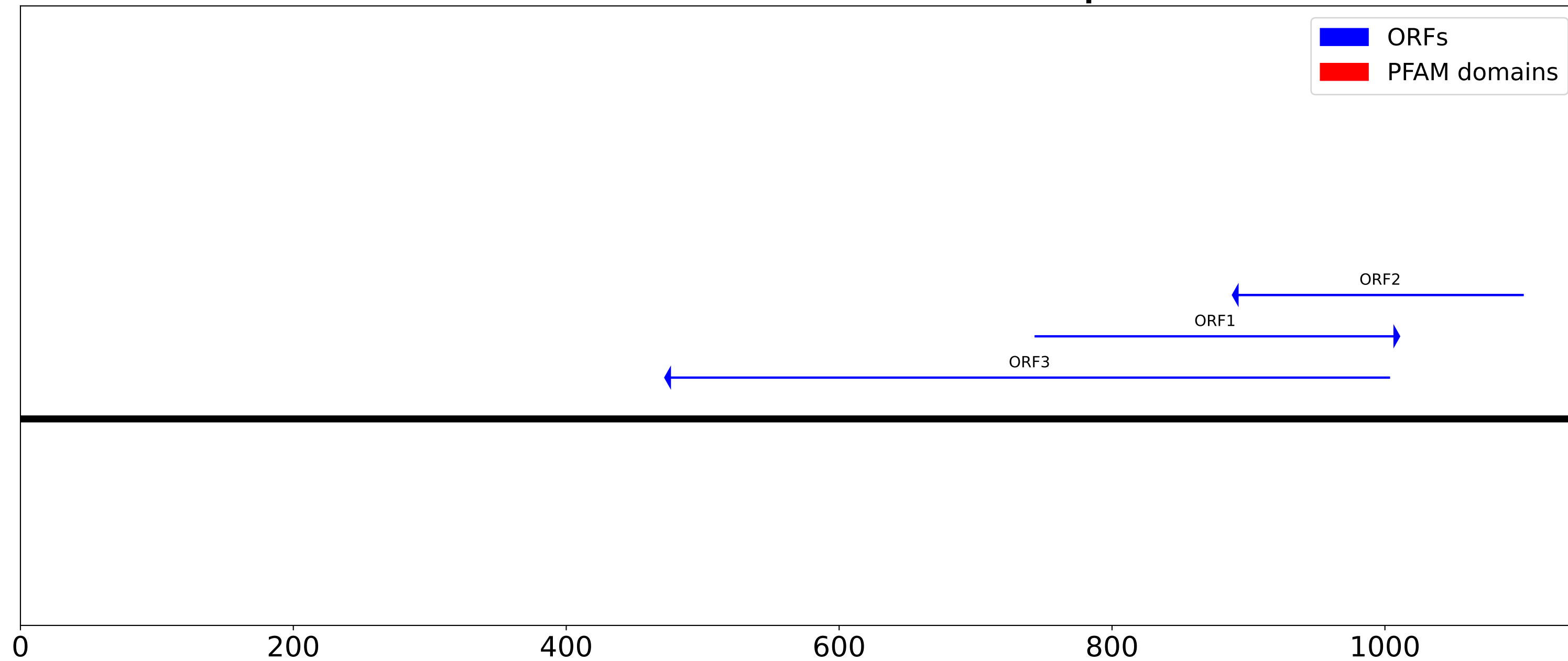


TE: ltr_1_family_433
size: 2312bp; fragments: 17485; full length: 1 (≥ 2080.8 bp)

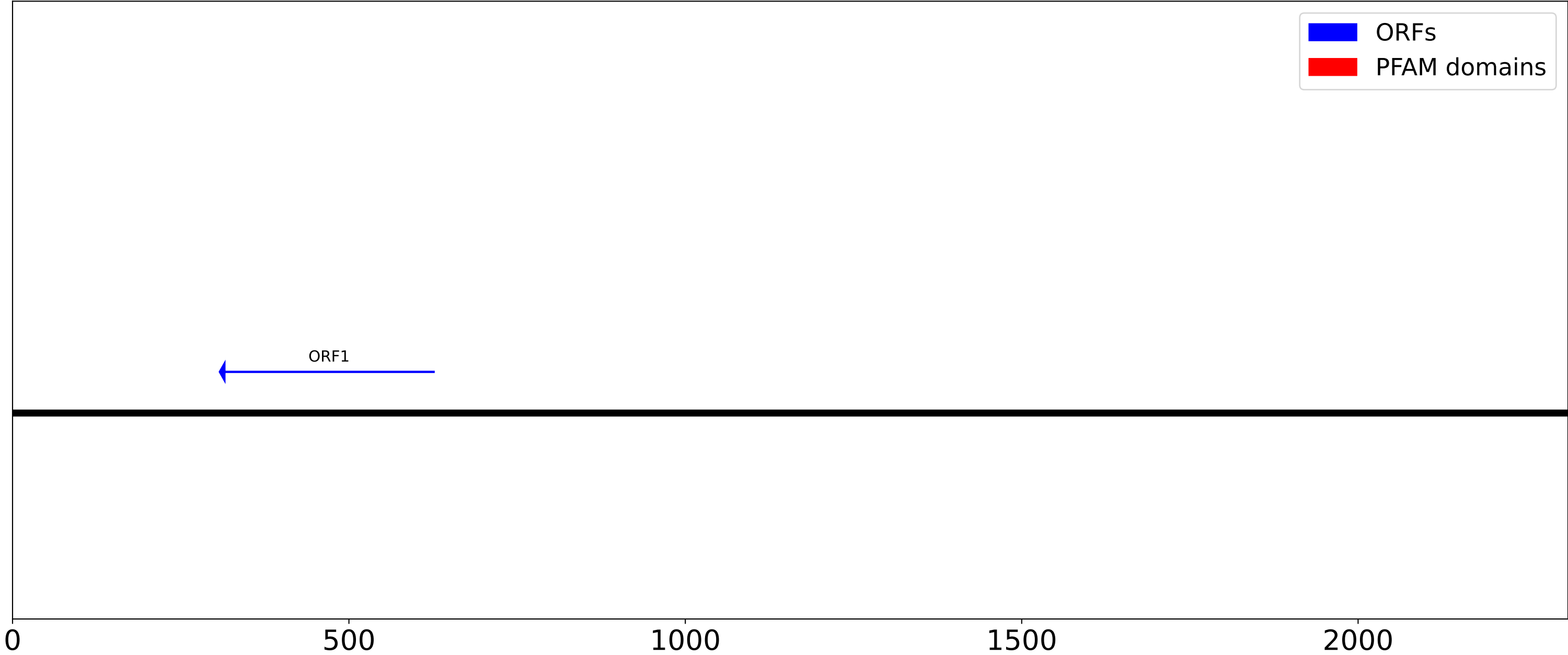
Before TEtrimmer 2312 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

