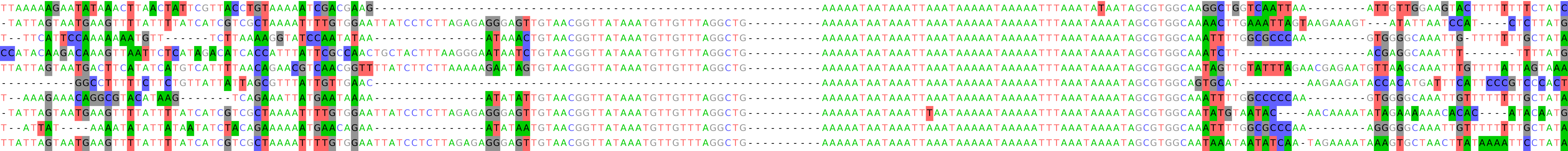


Start Crop Point

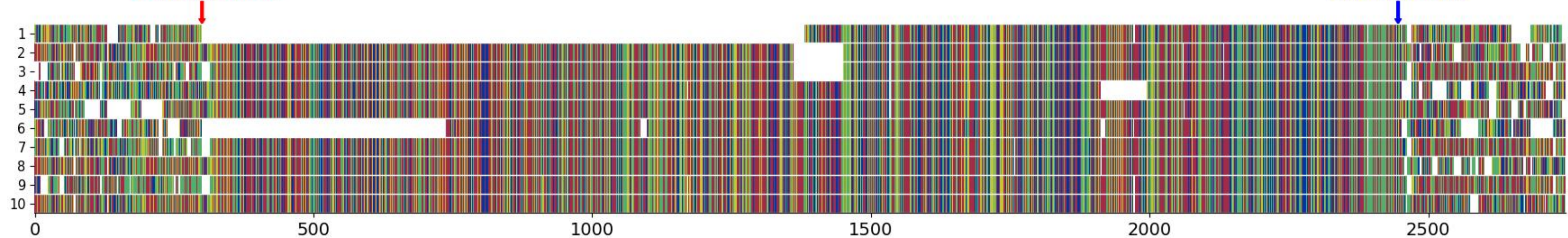
End Crop Point

MSA length = 2147



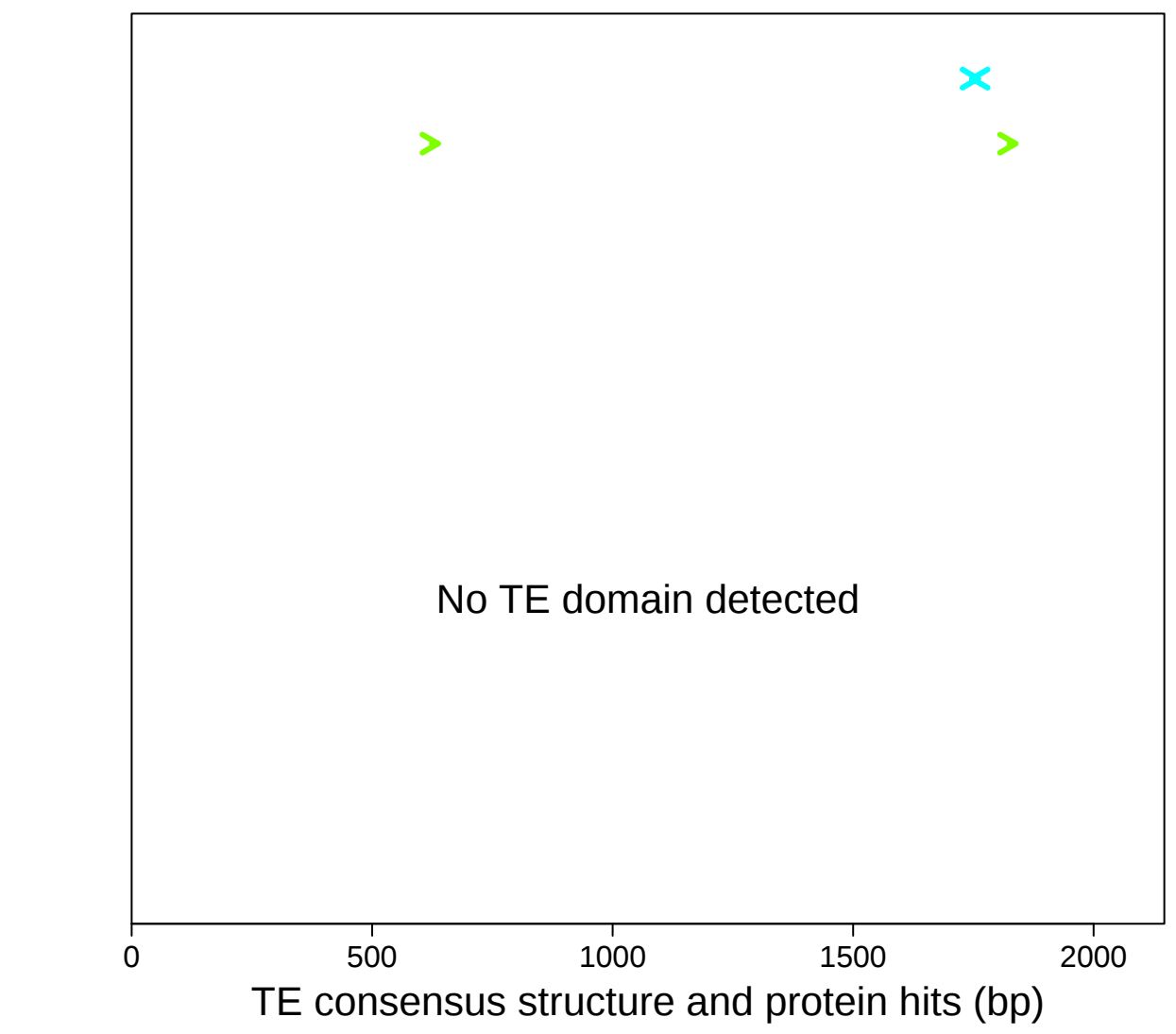
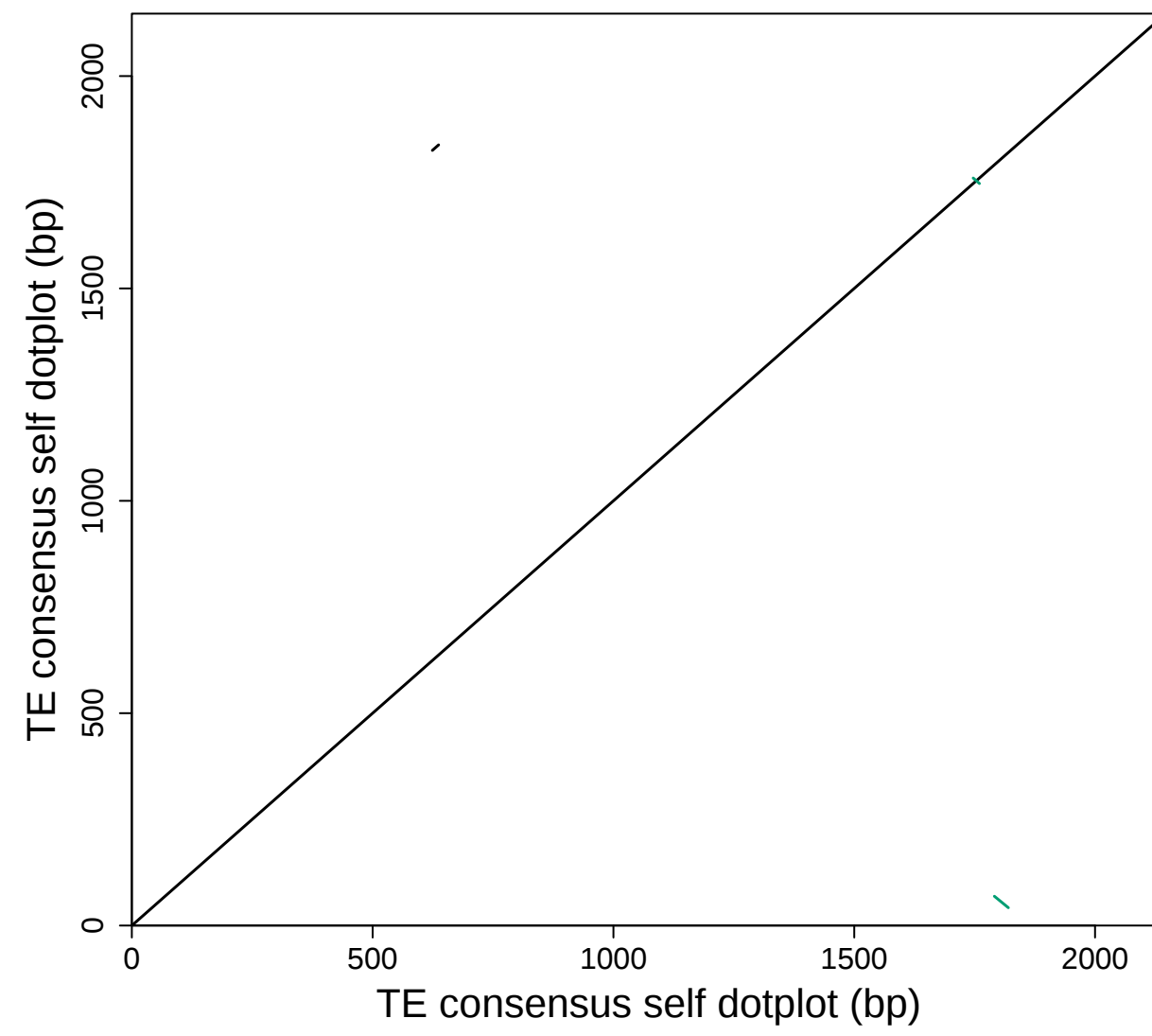
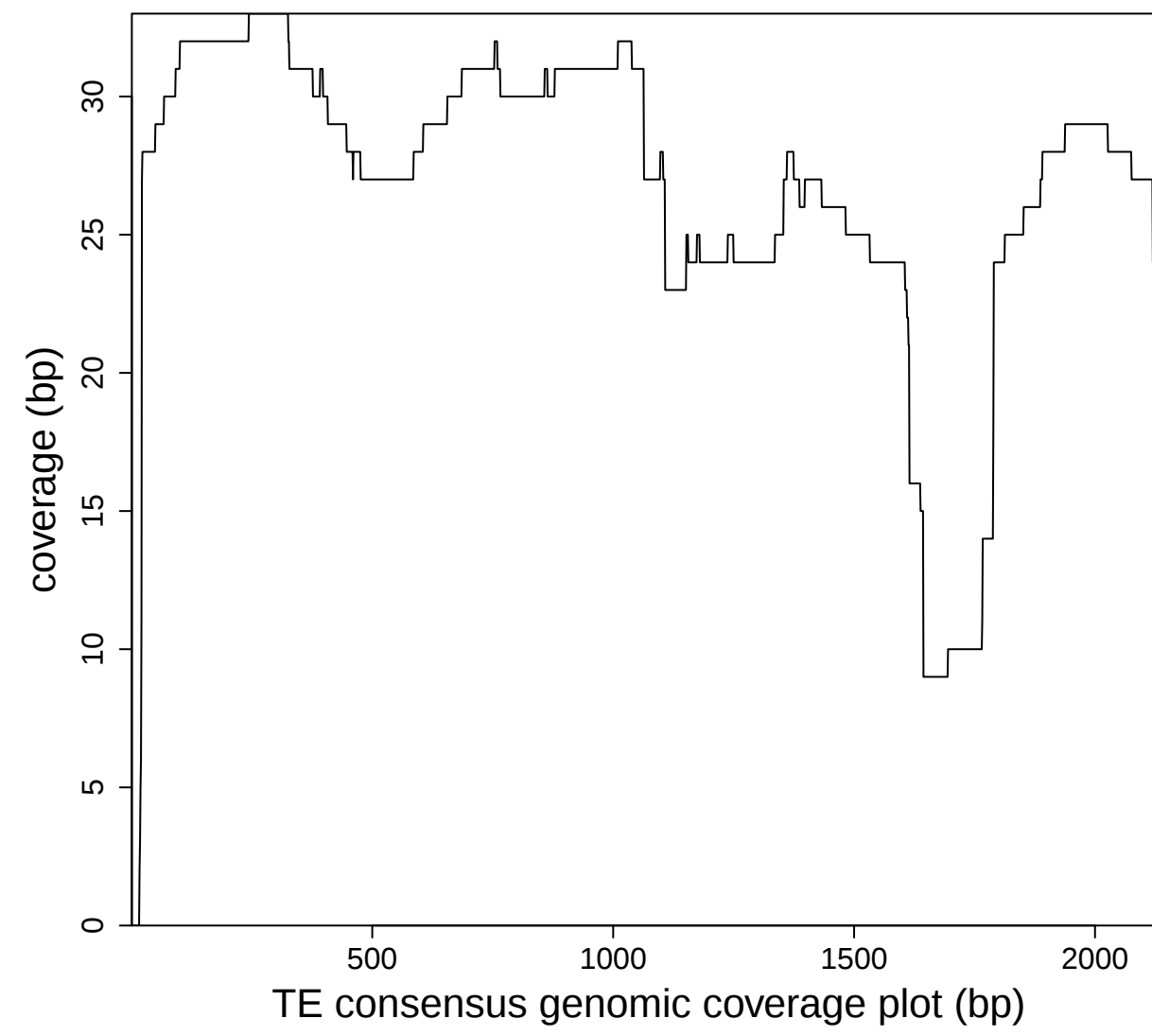
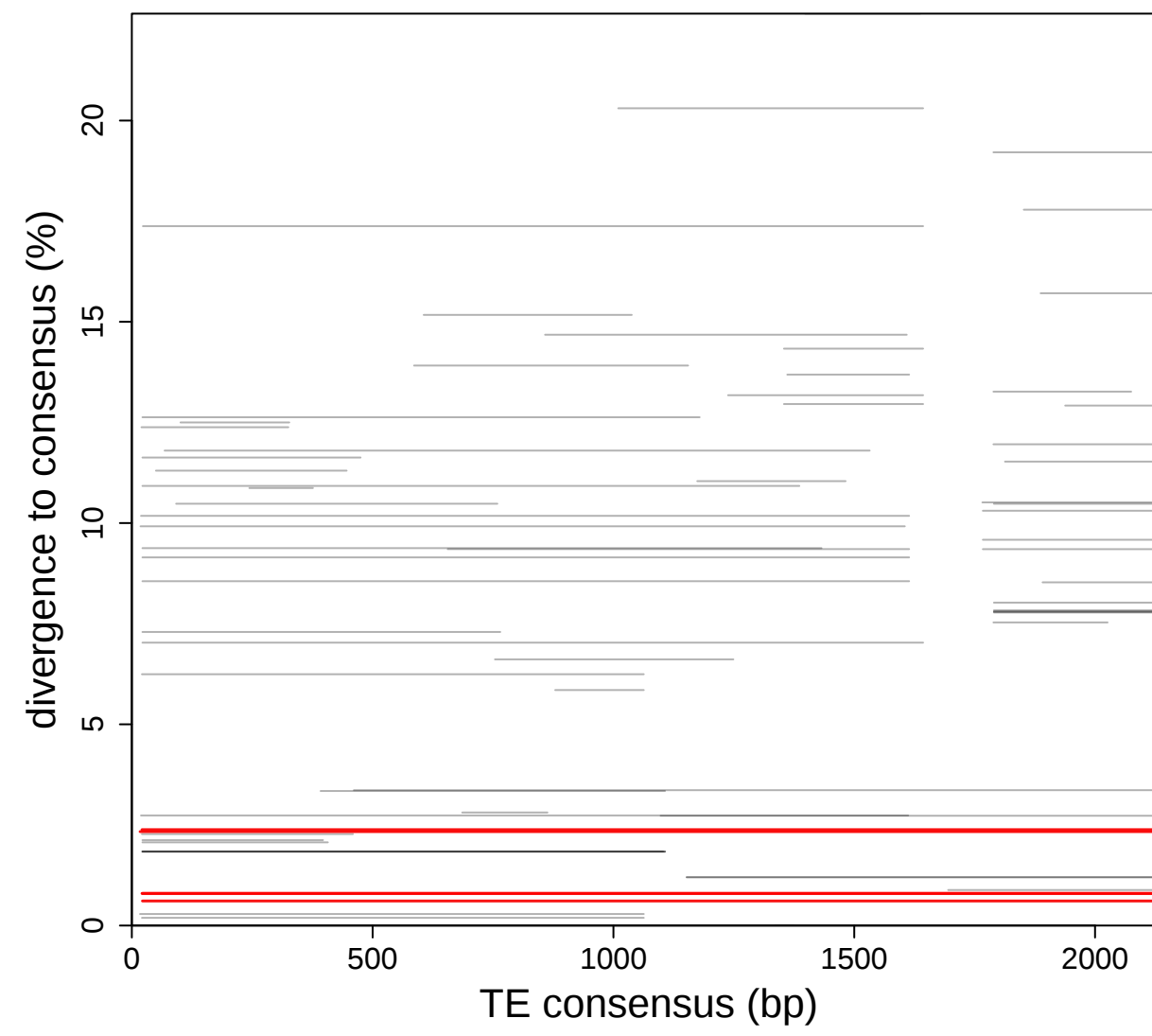
Start crop Point

End crop Point



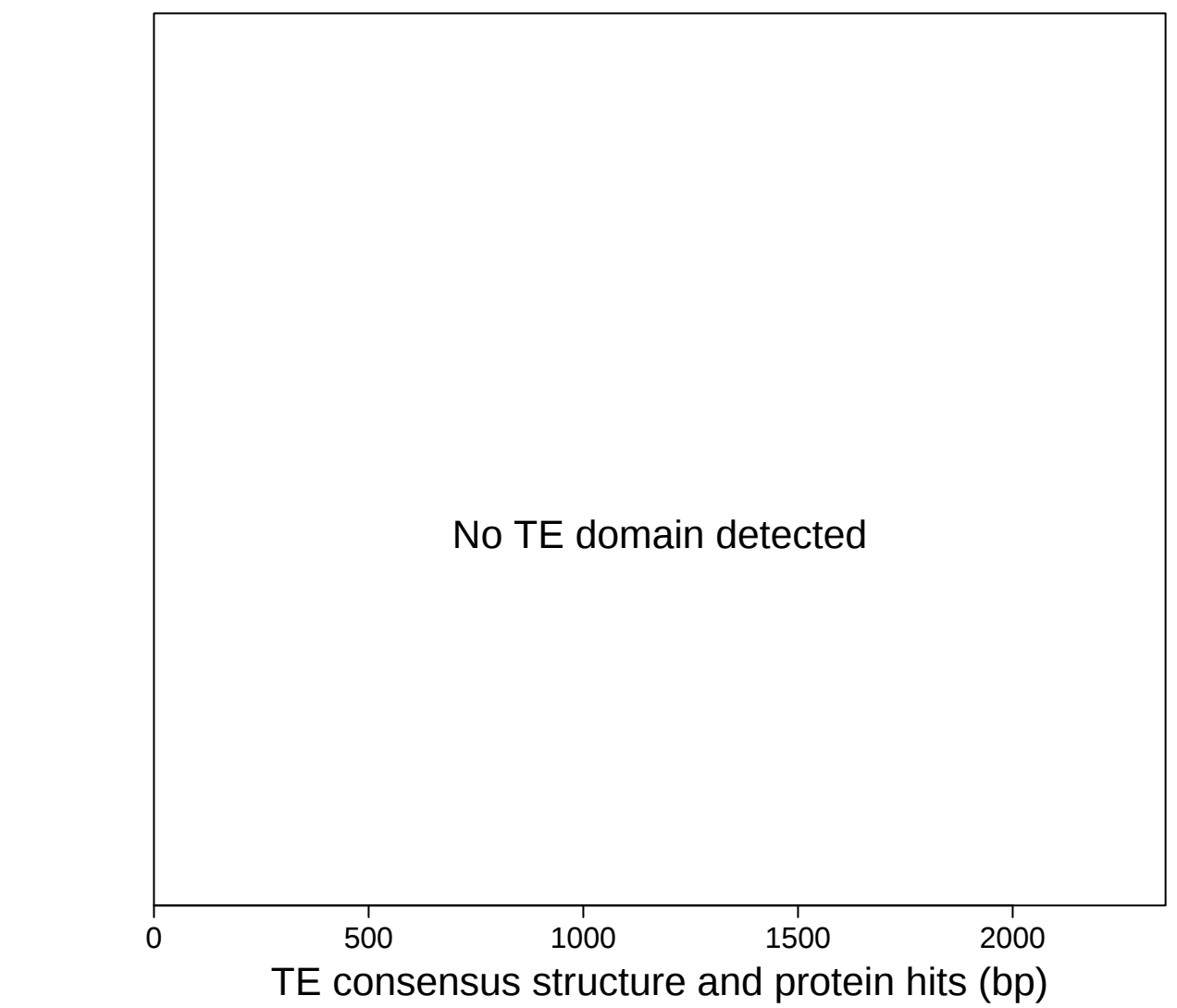
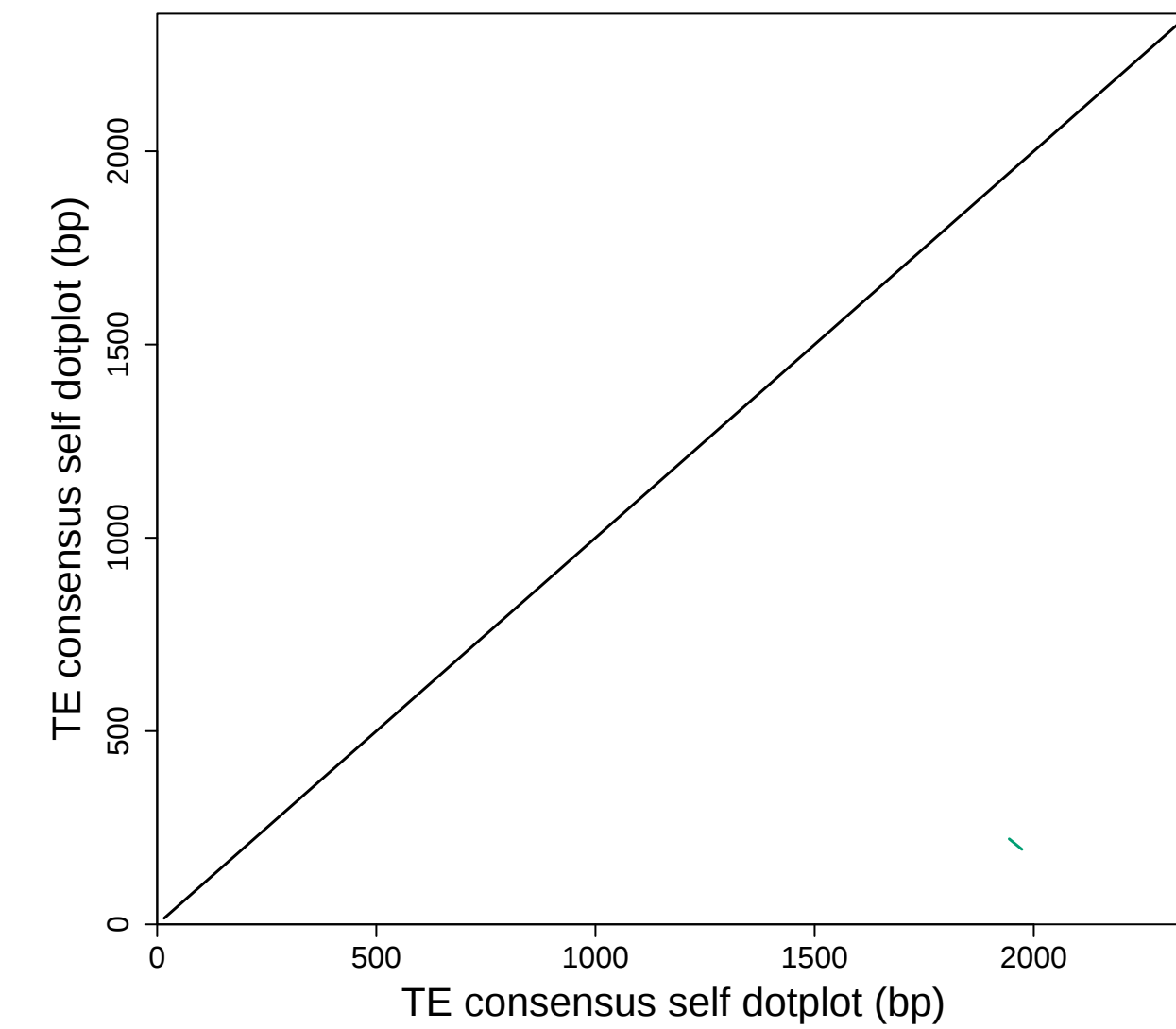
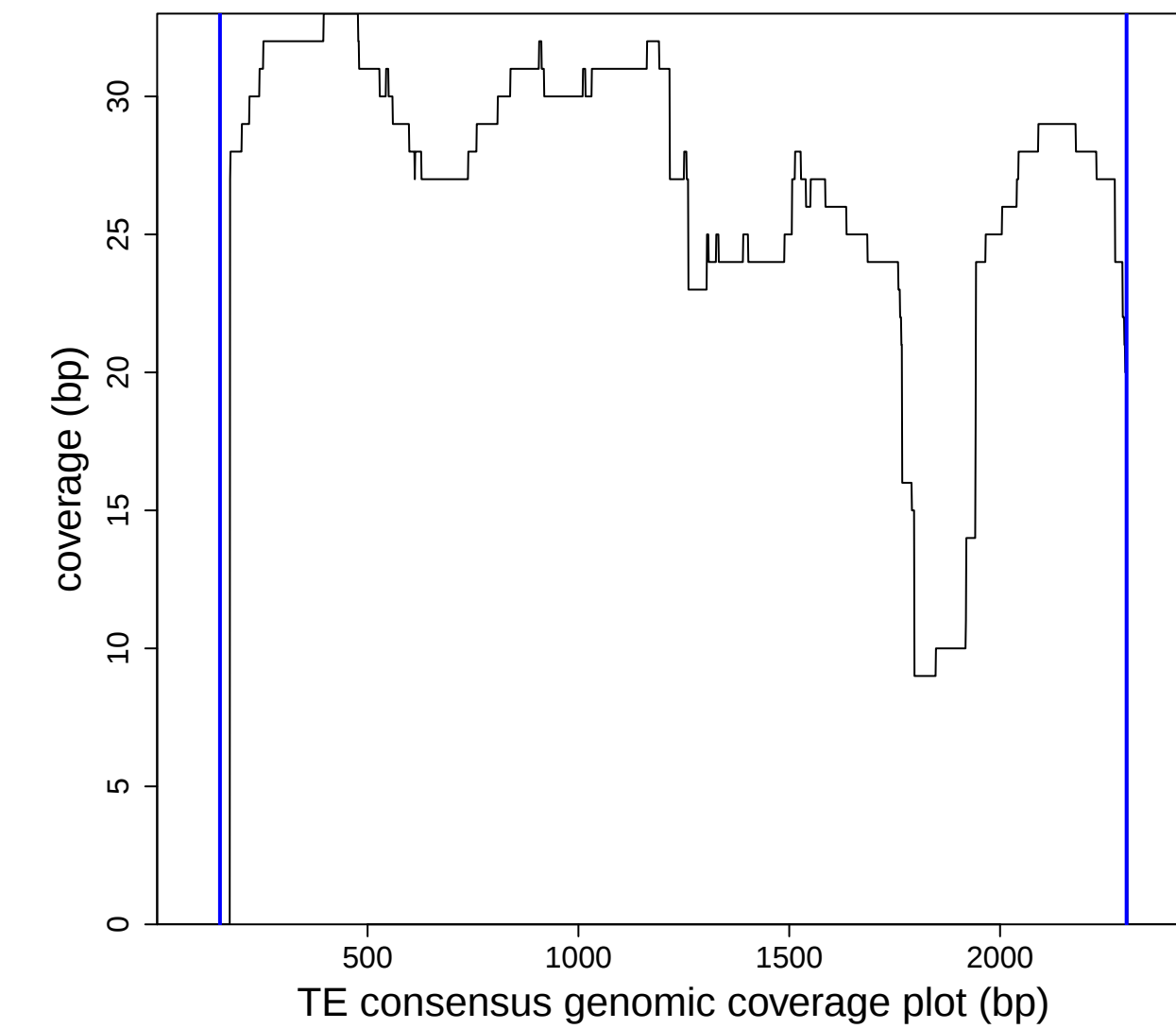
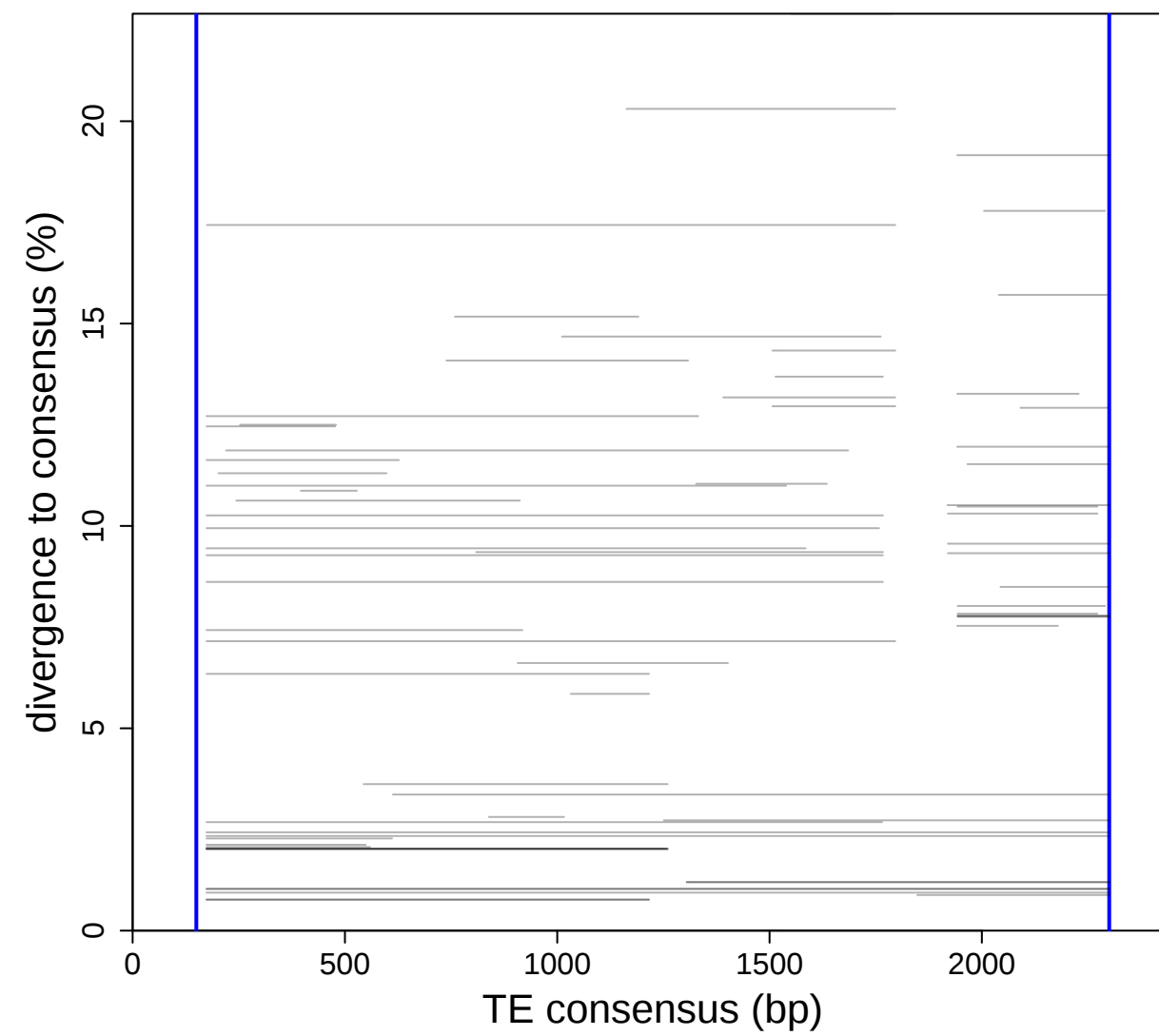
b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 2147bp; fragments: 73; full length: 5 (≥ 1932.3 bp)

After TEtrimmer 2147 bp

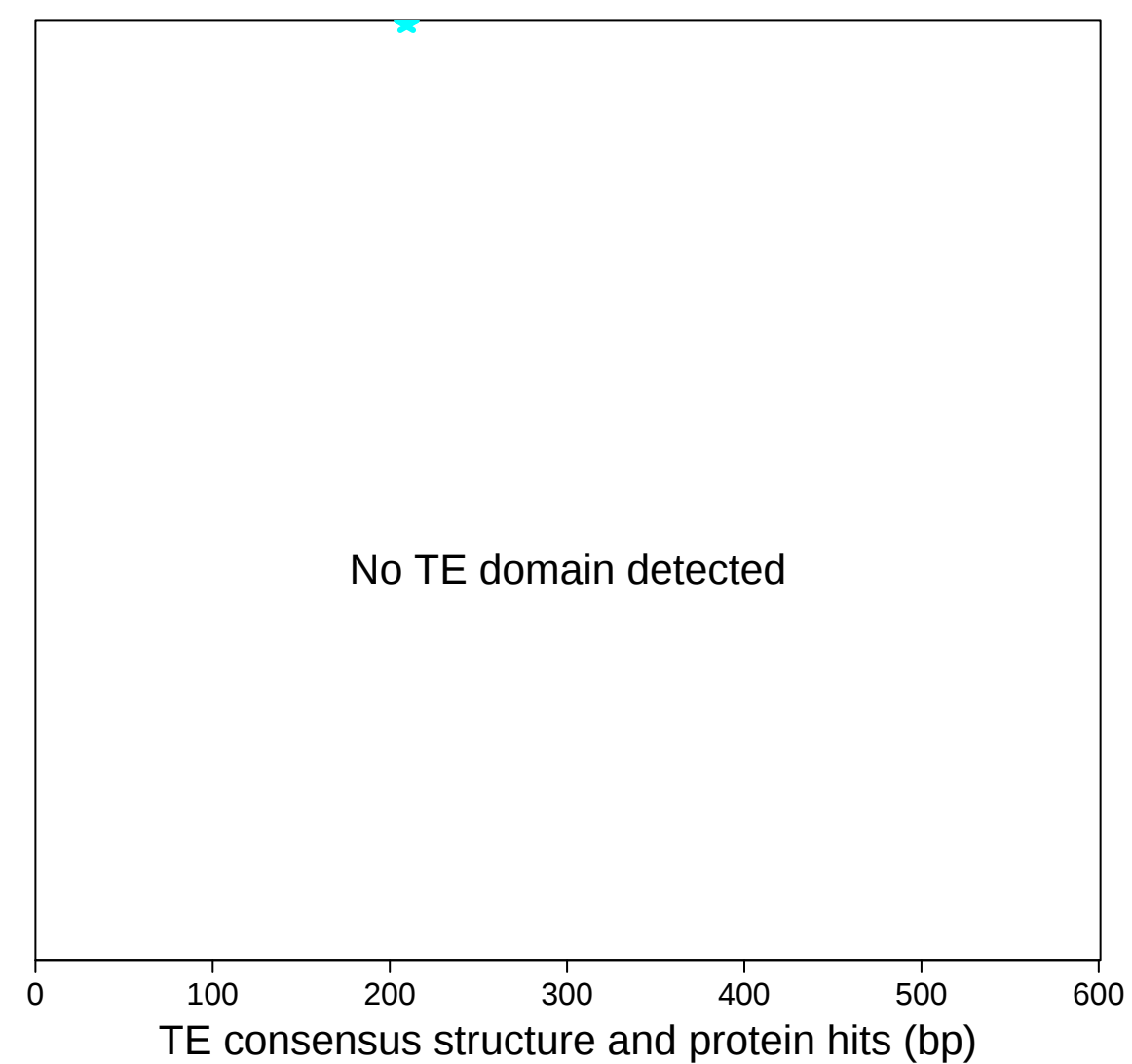
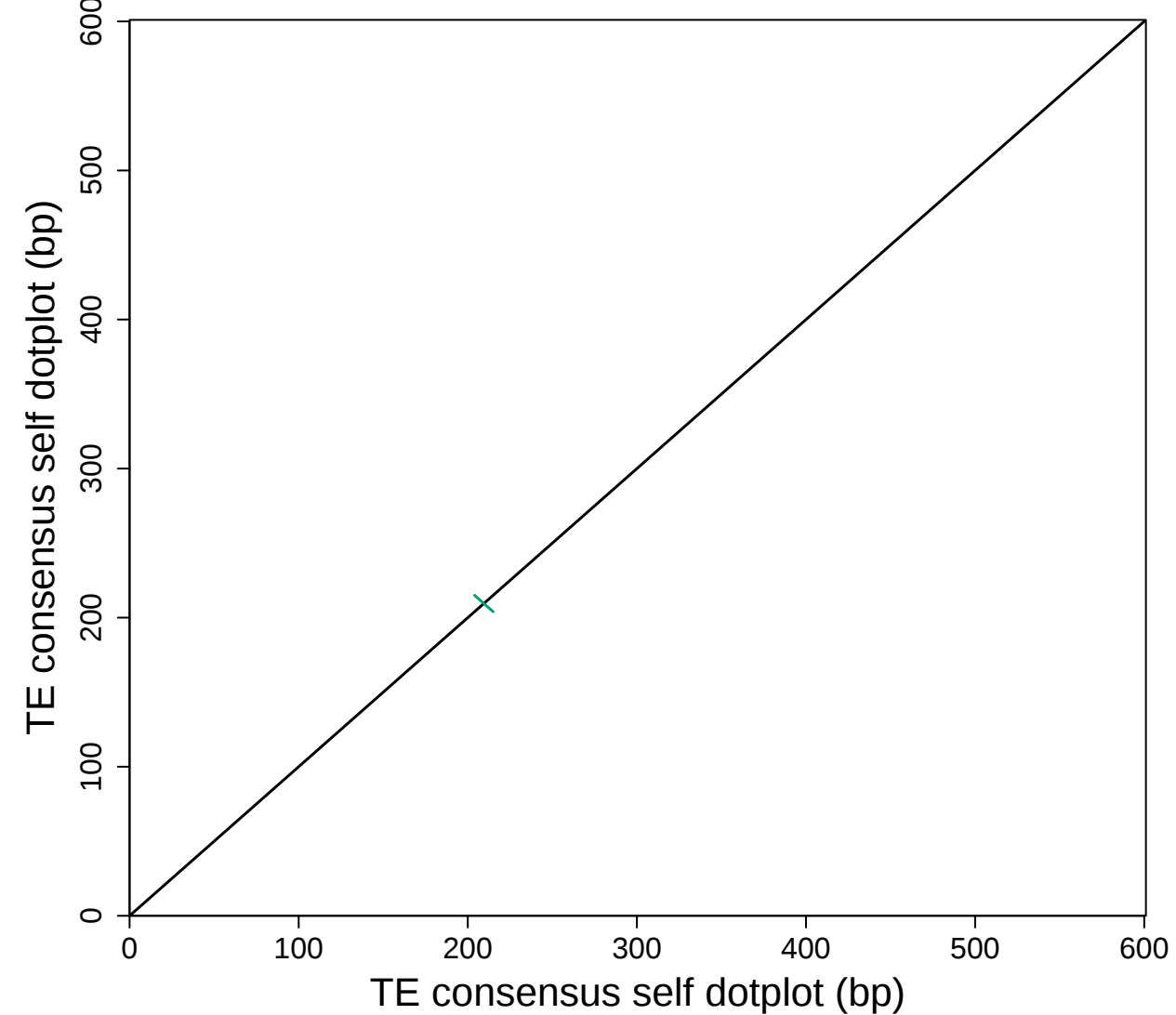
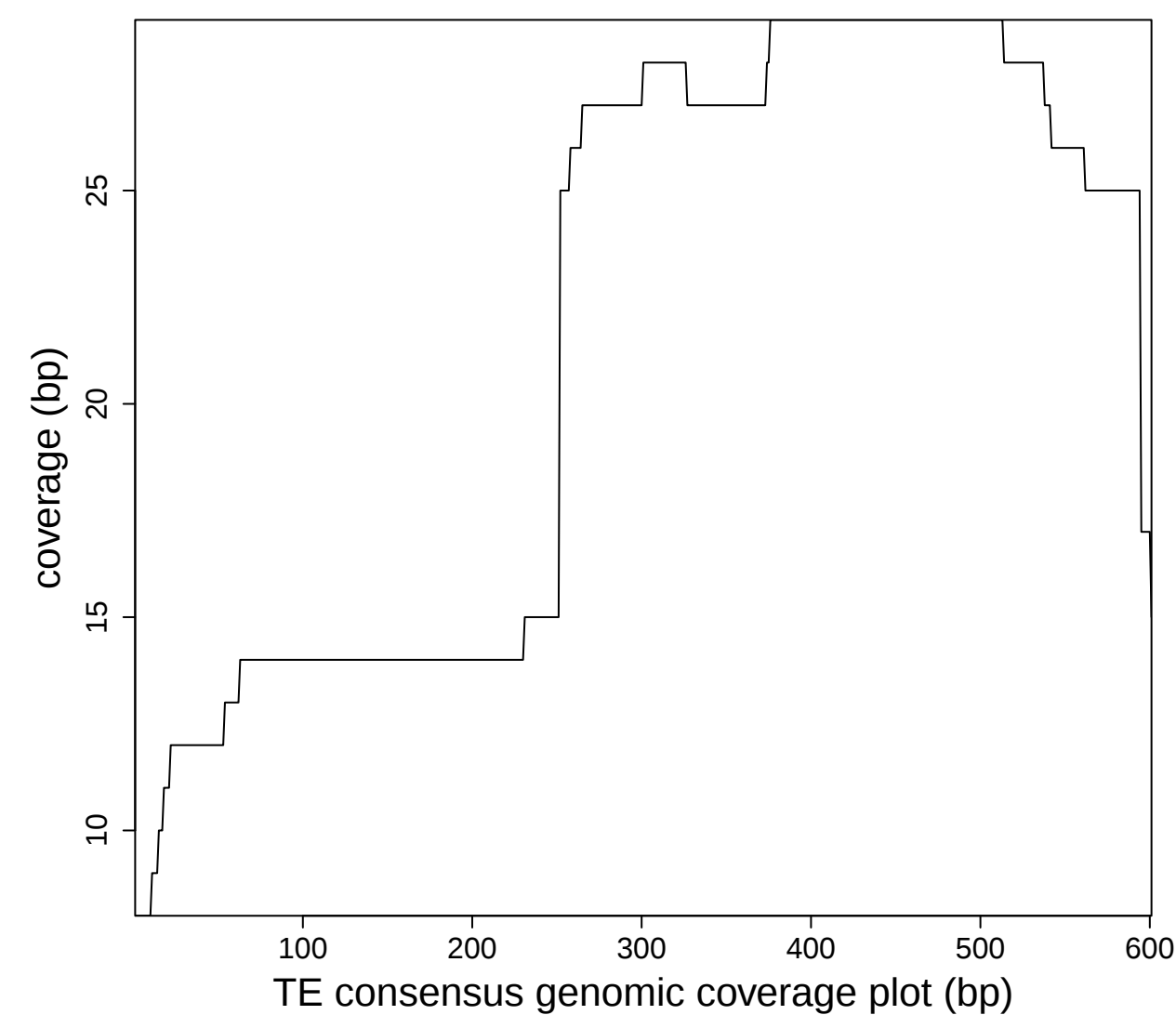
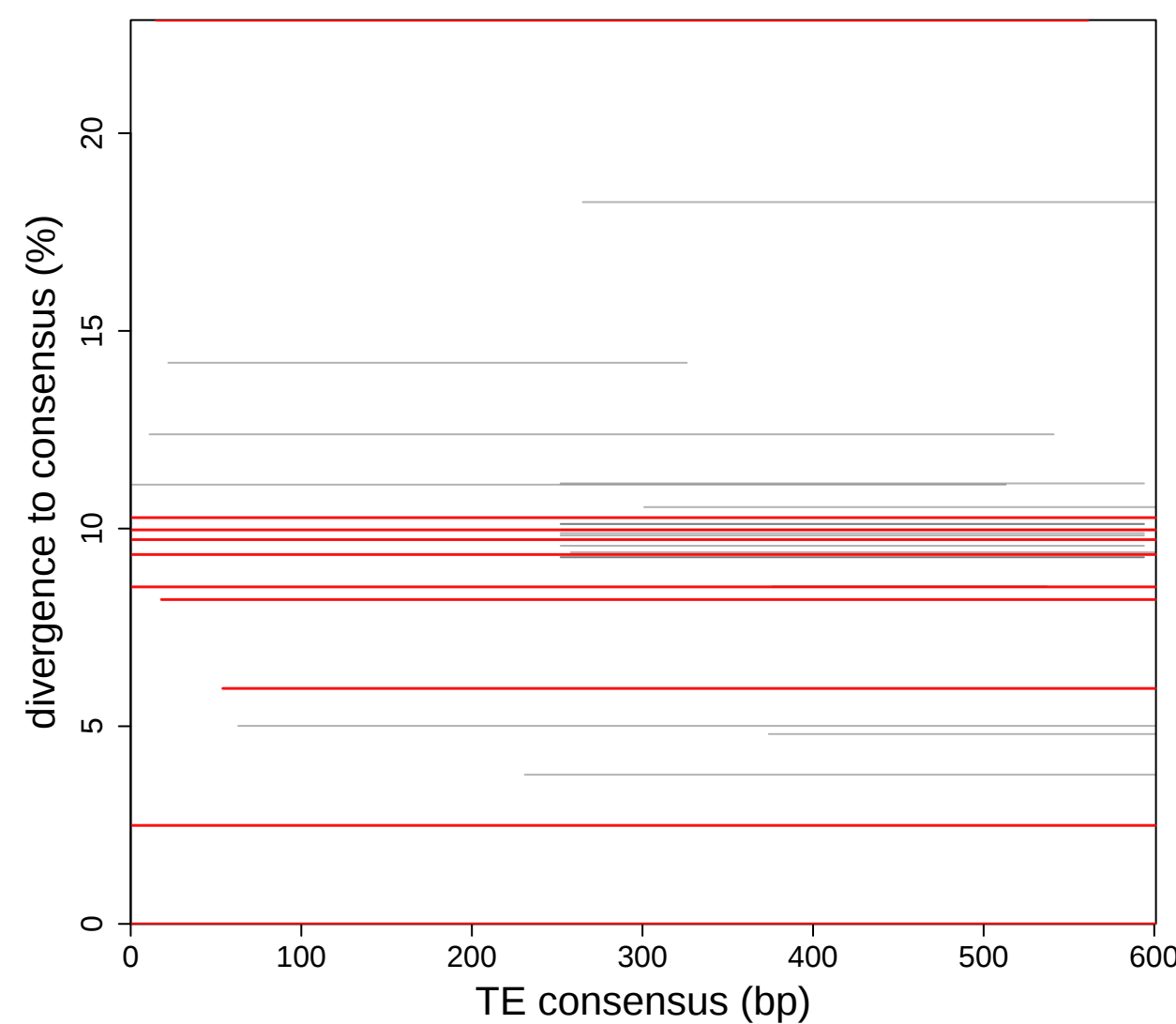


.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 2450bp; fragments: 73; full length: 0 (>=2205bp)

After TEtrimmer Extended plot Blue lines are boundaries

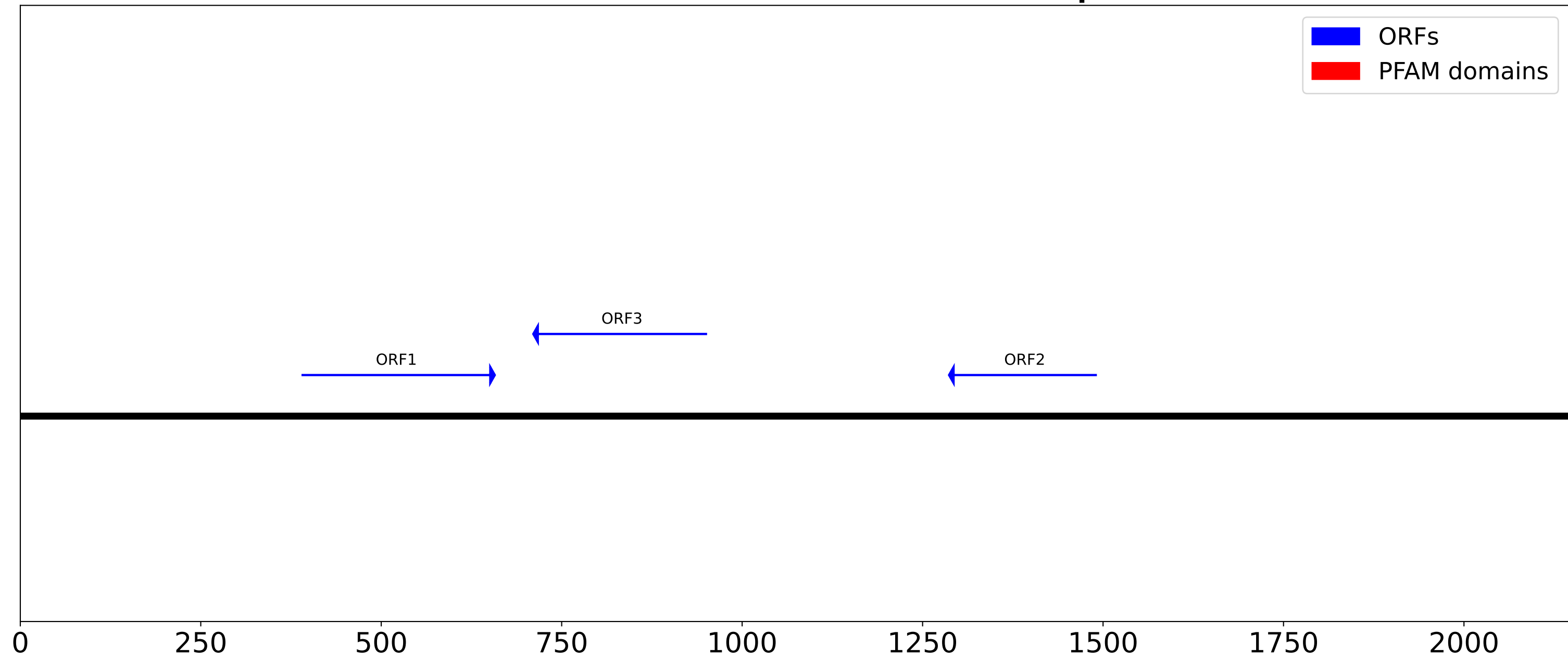


TE: ltr_1_family_300
size: 601bp; fragments: 30; full length: 10 (≥ 540.9 bp)

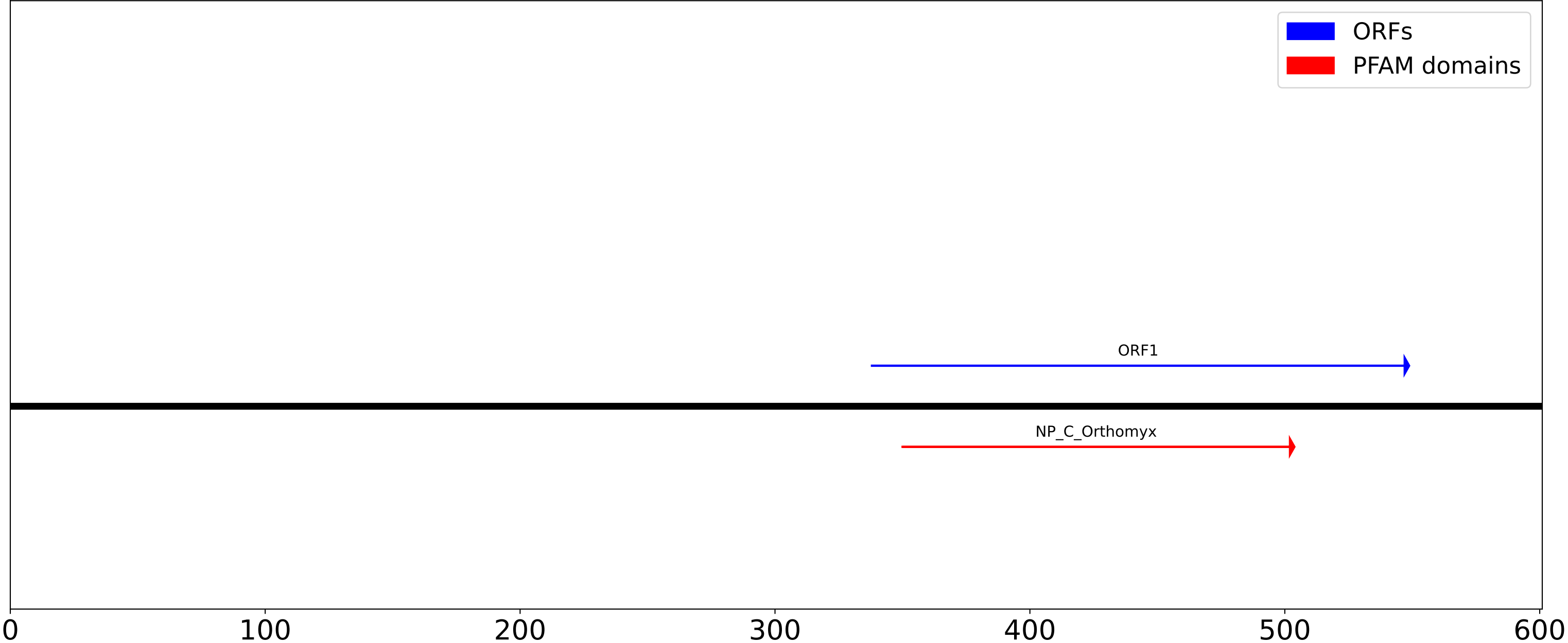


Before TEtrimmer 601 bp

After TEtrimmer ORF and PFAM domain plot



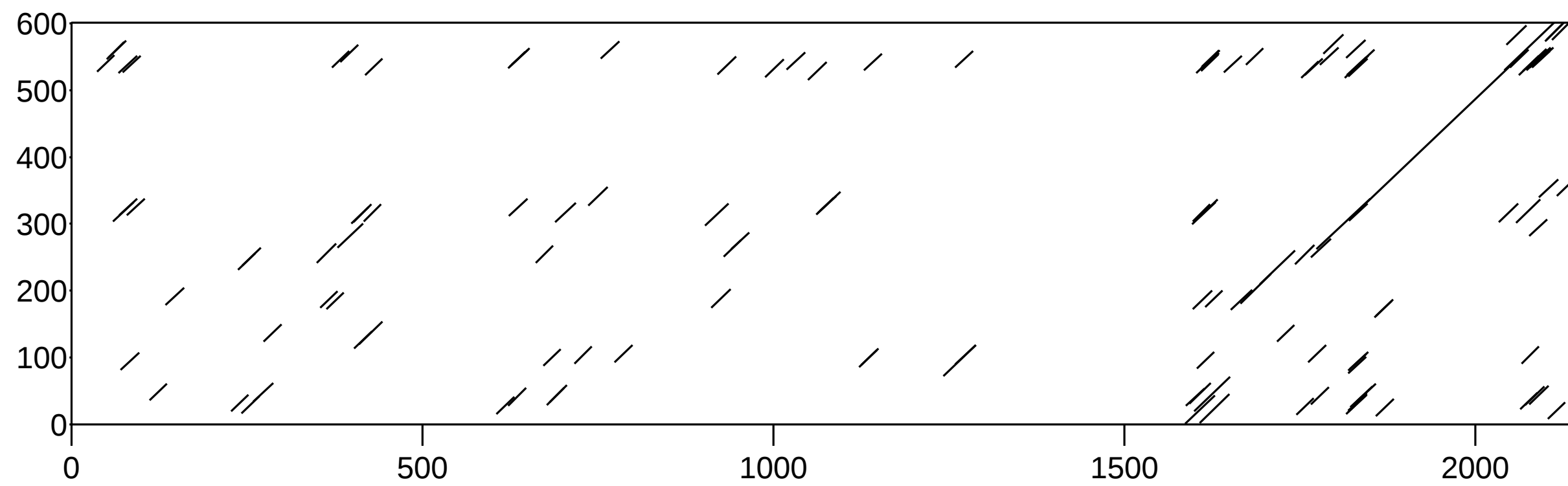
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)