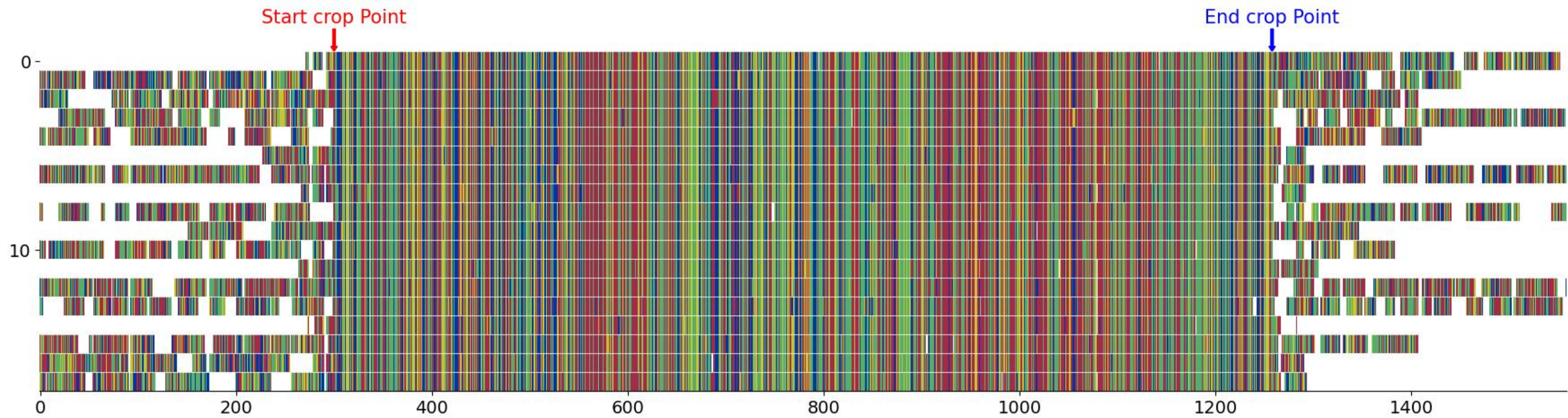


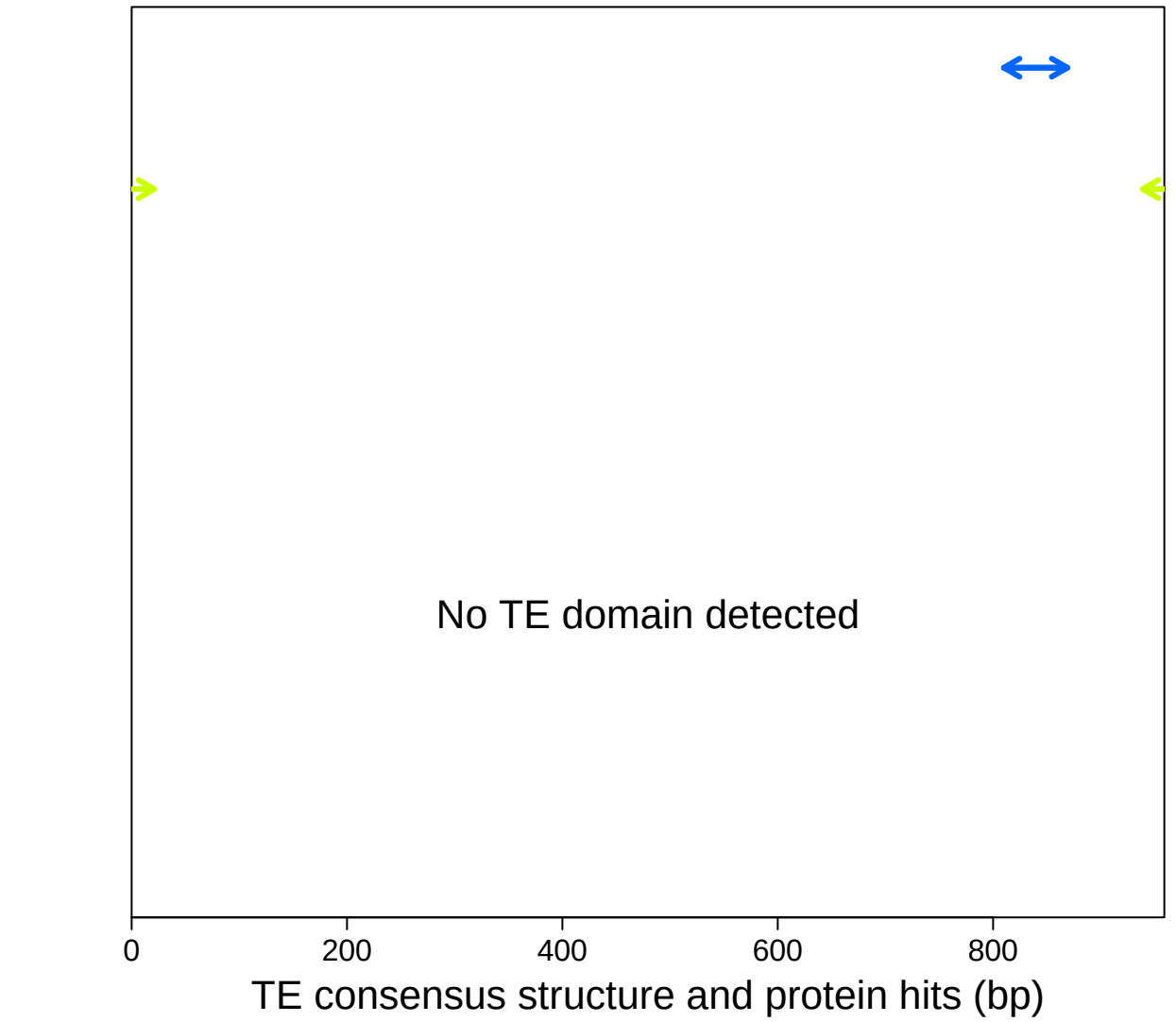
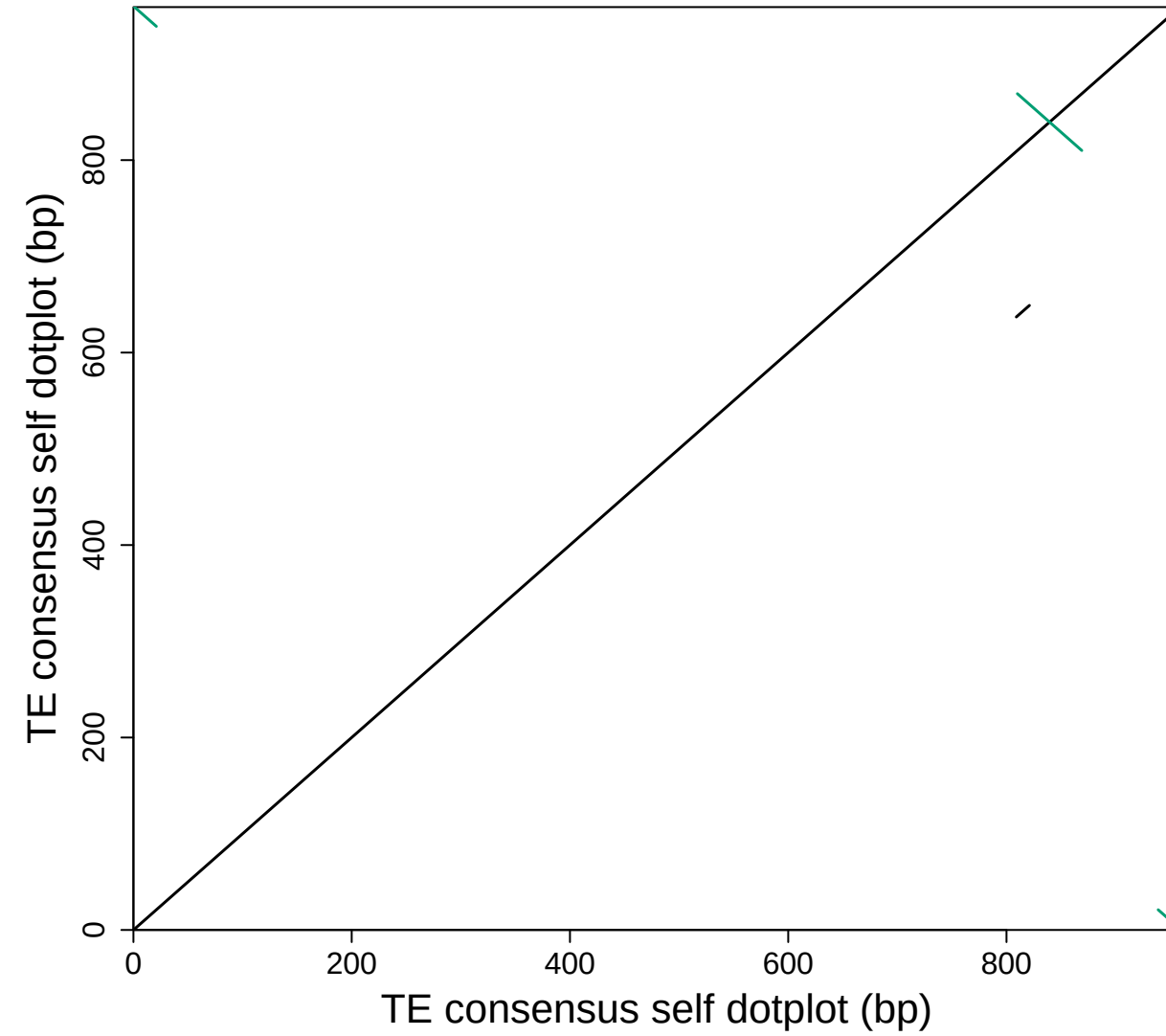
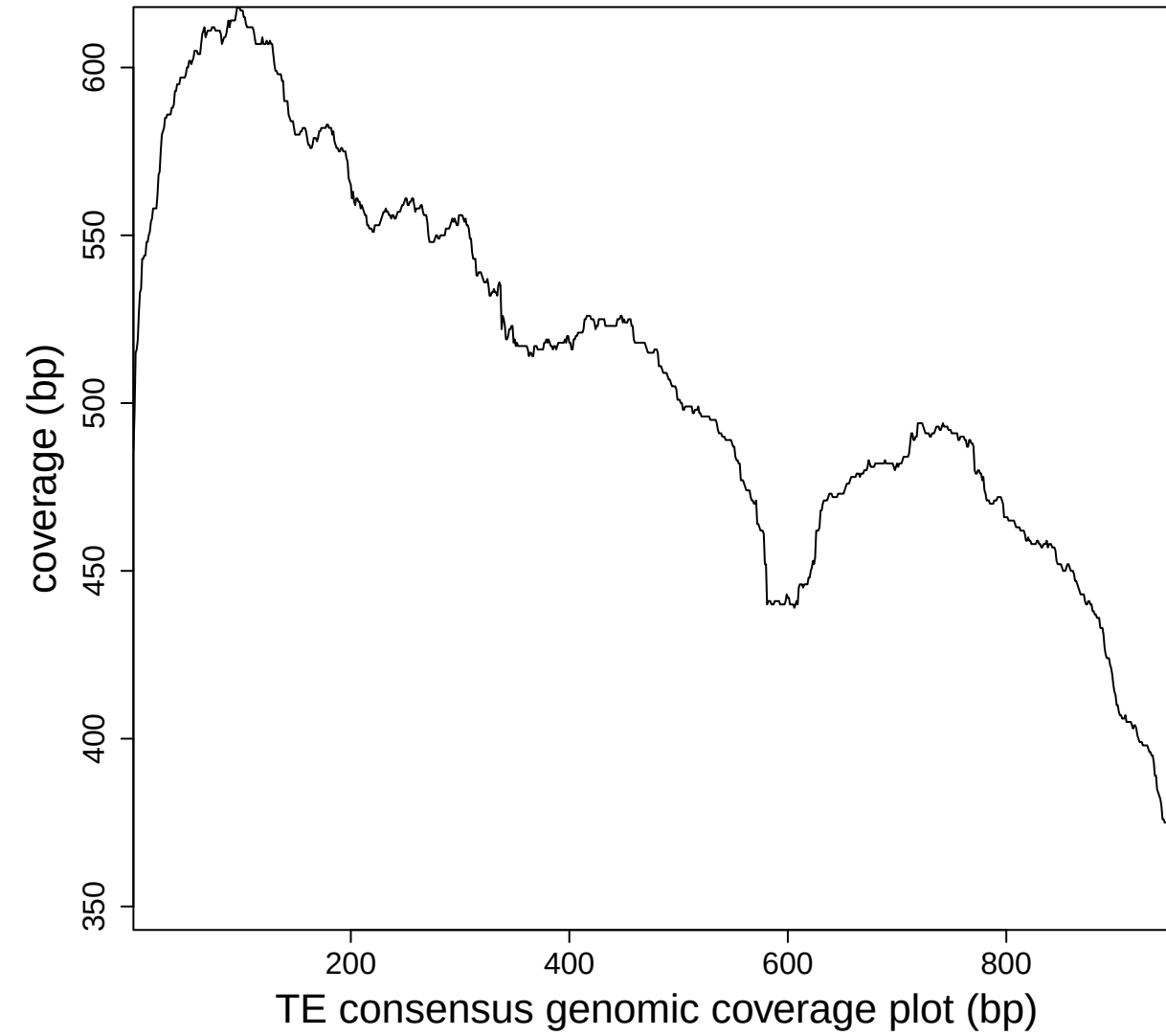
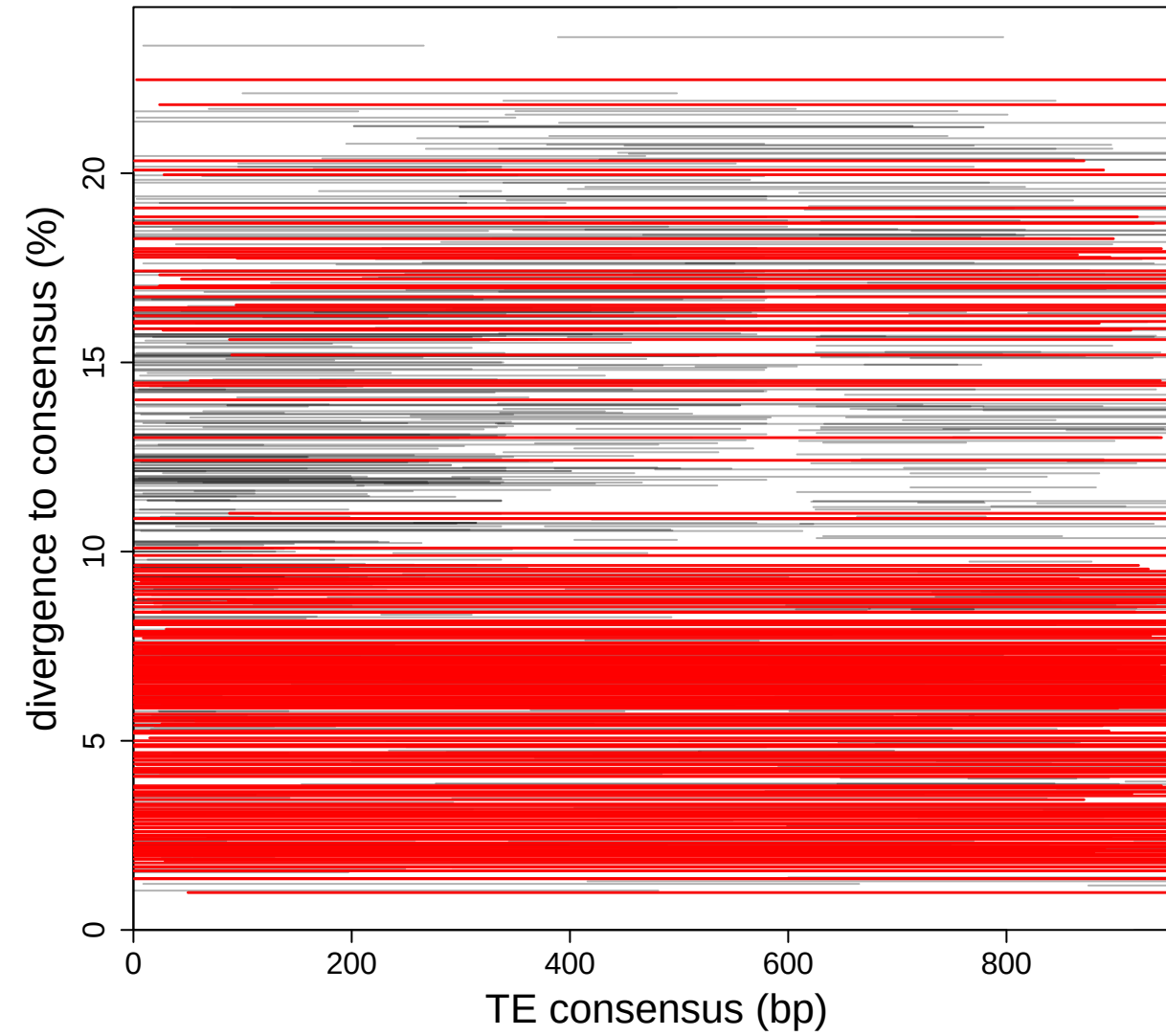
Start crop Point

End crop Point



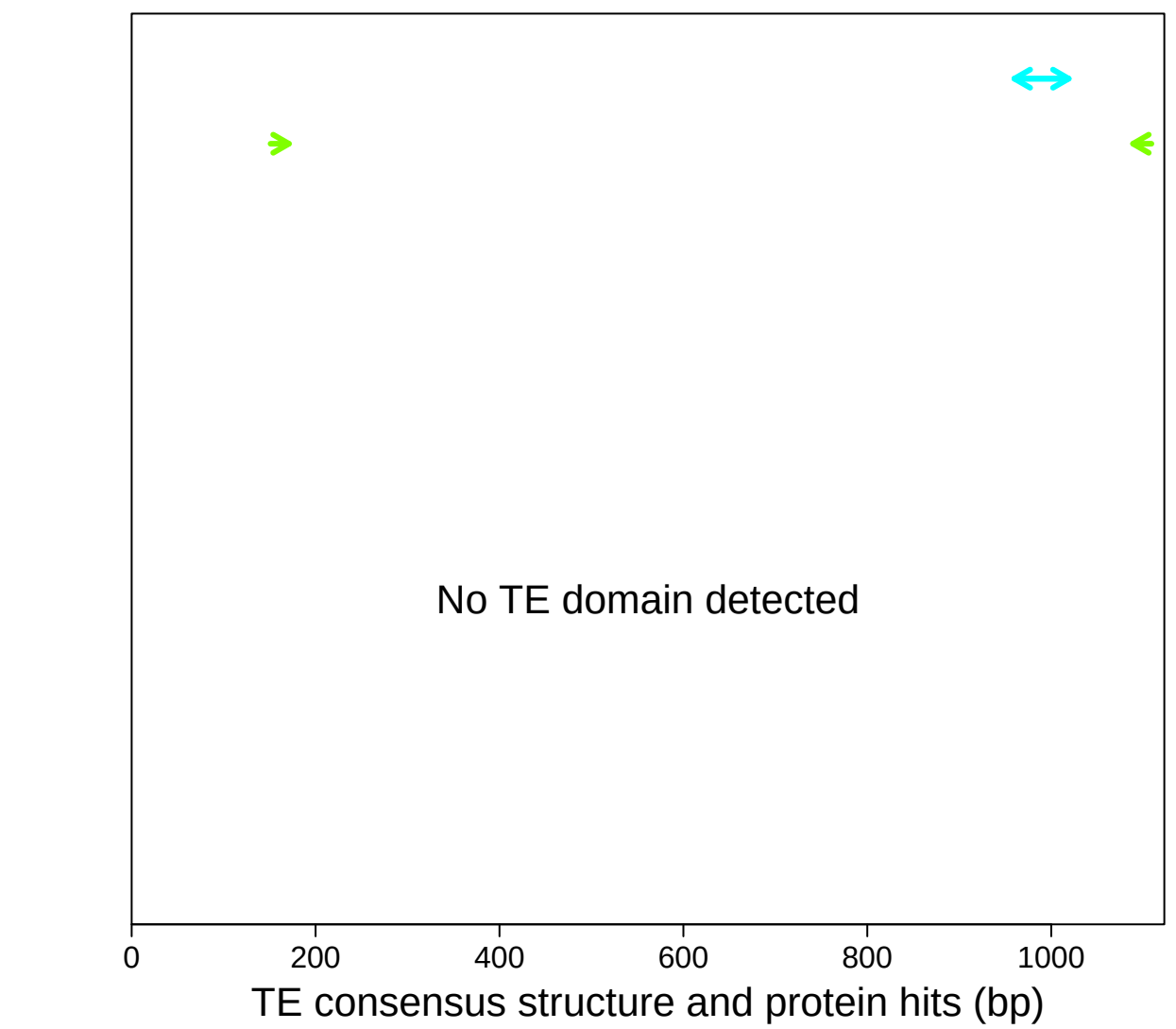
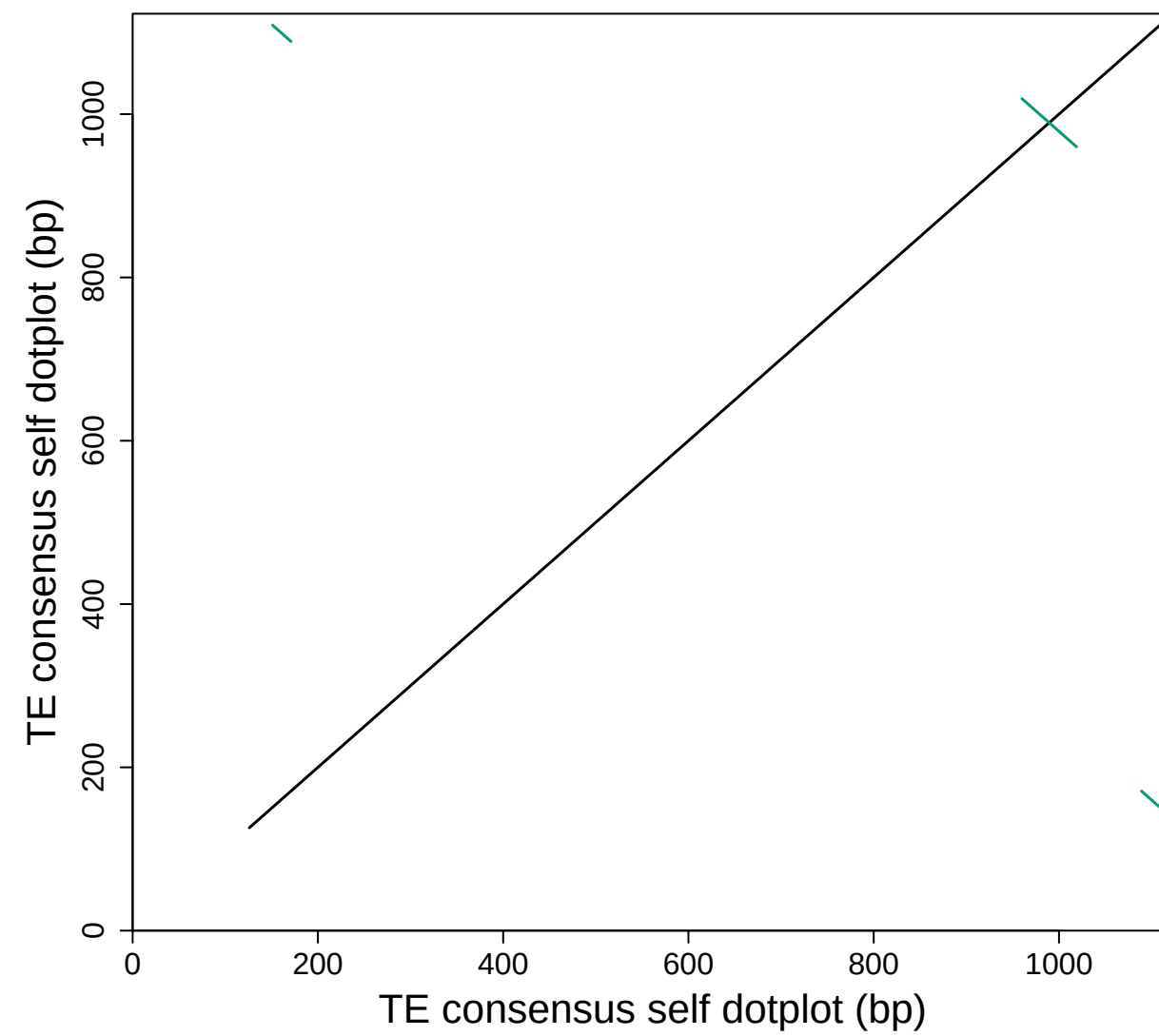
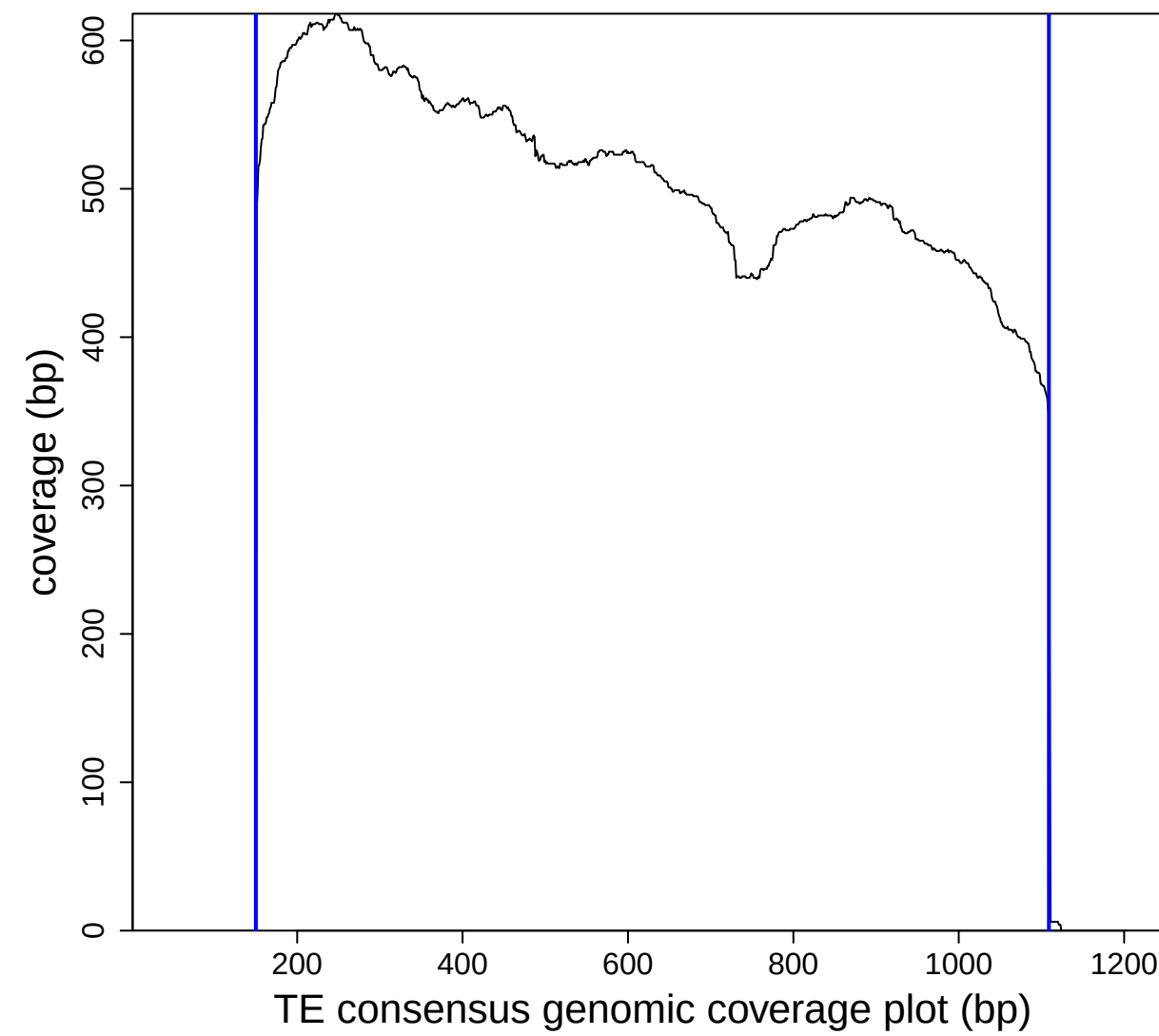
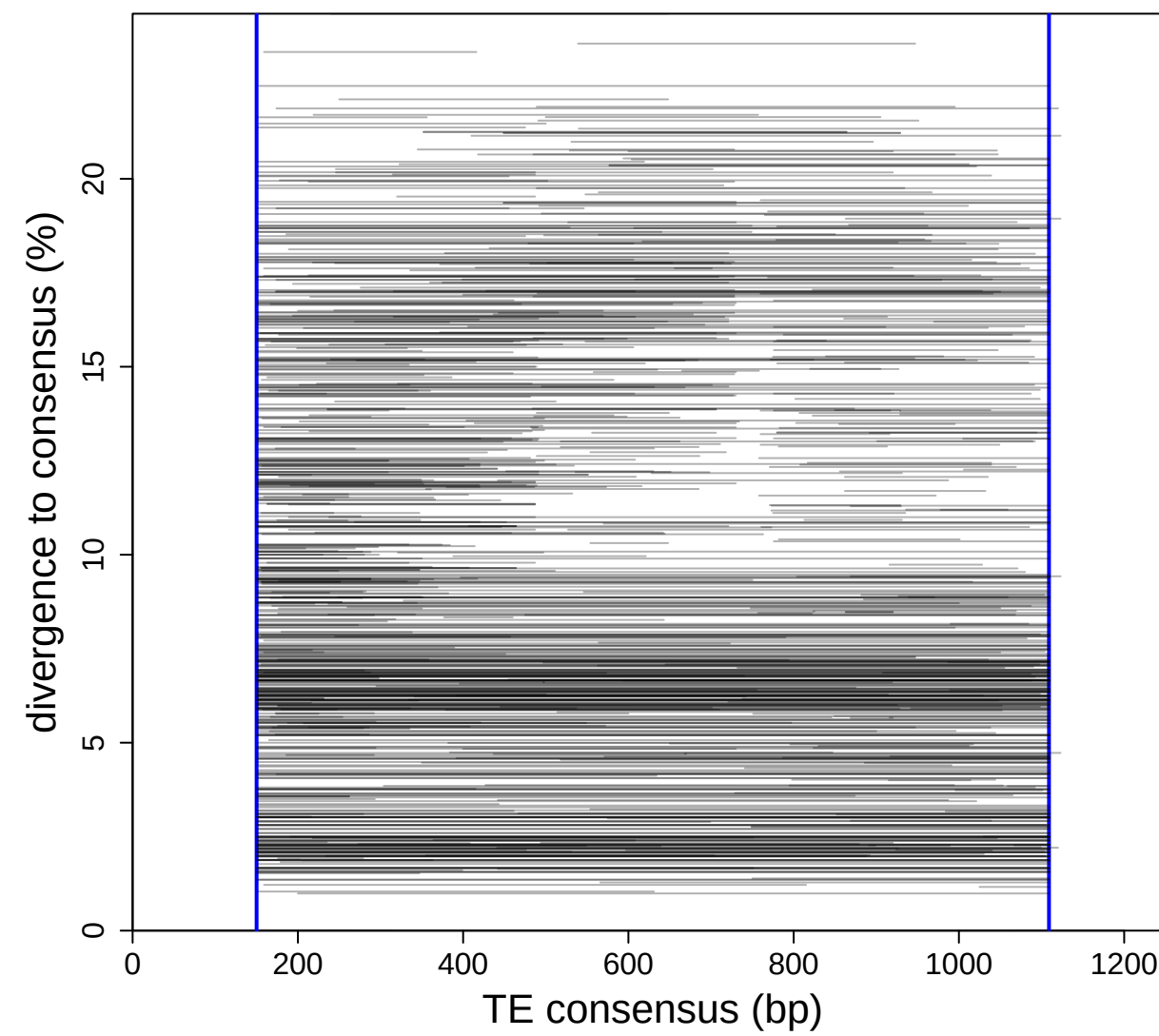
b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 959bp; fragments: 890; full length: 280 (≥ 863.1 bp)

After TEtrimmer 959 bp



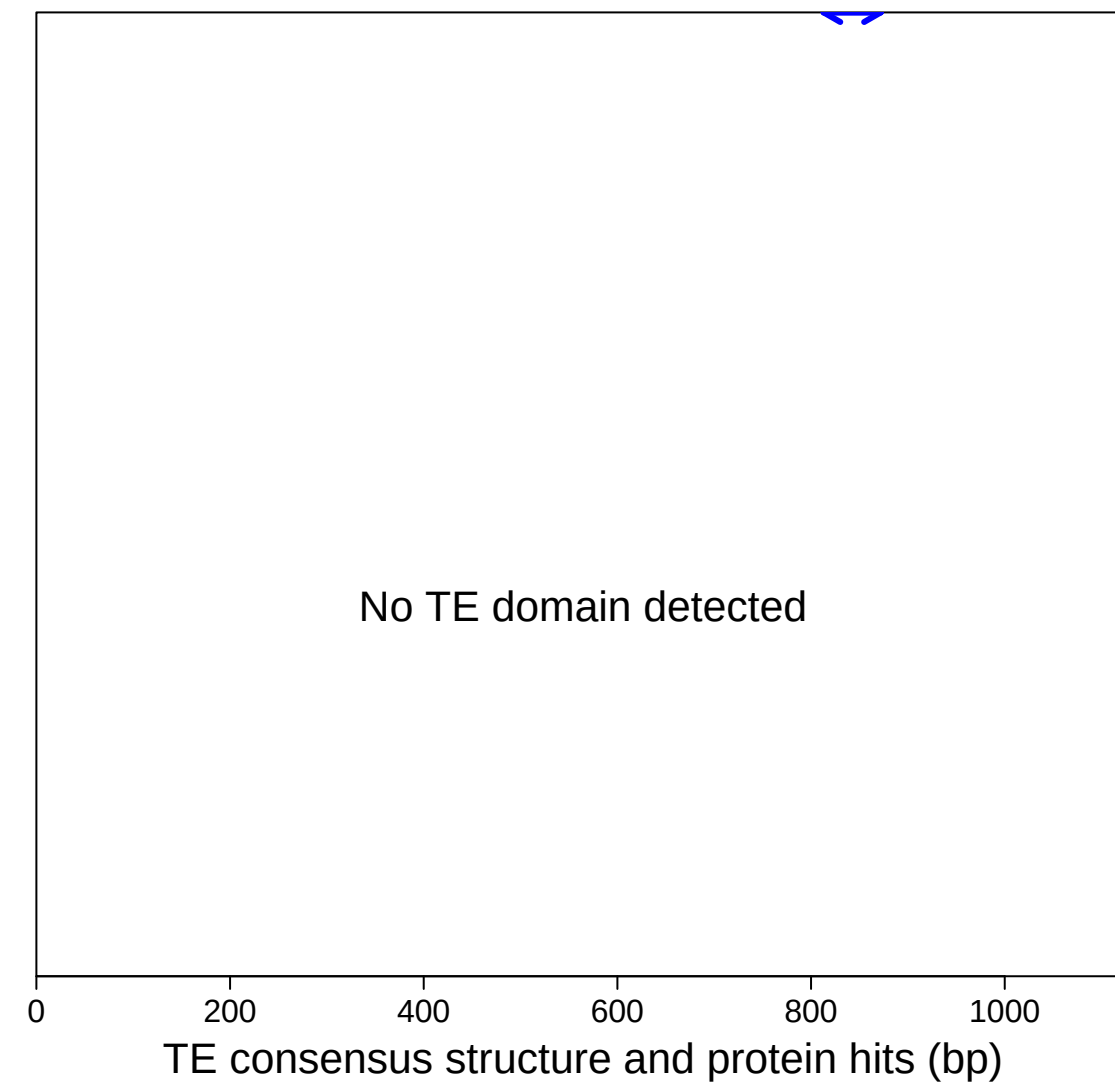
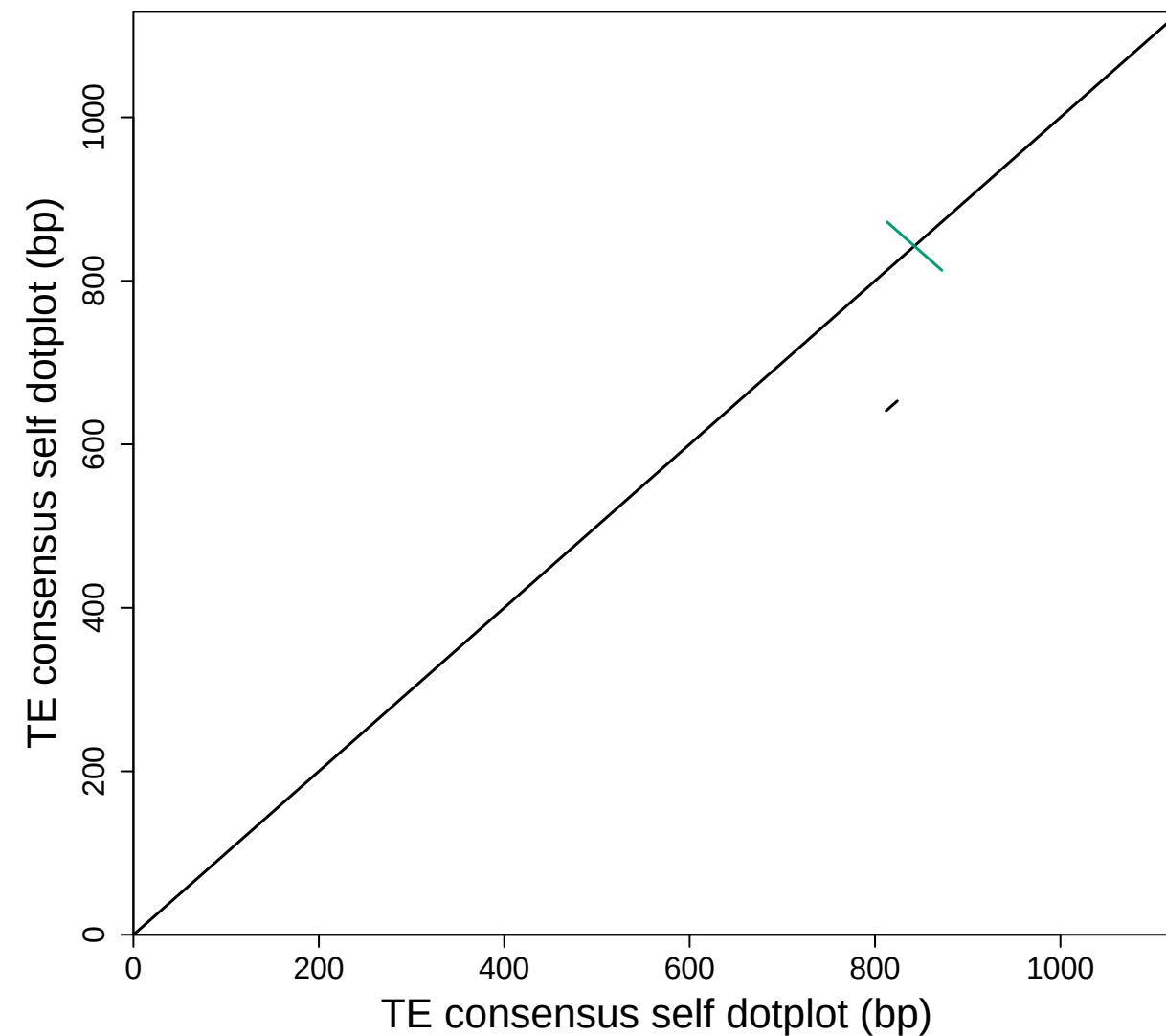
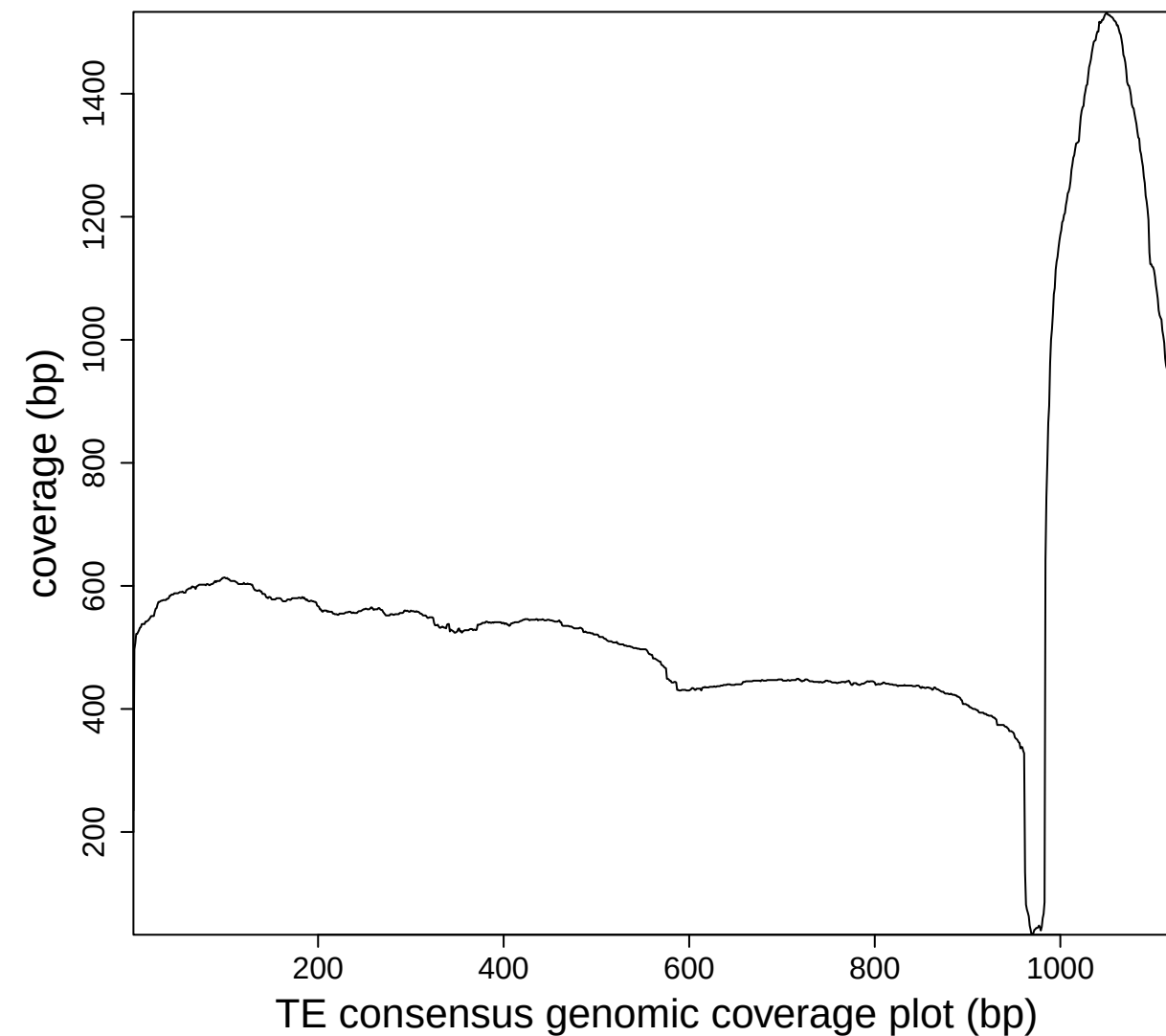
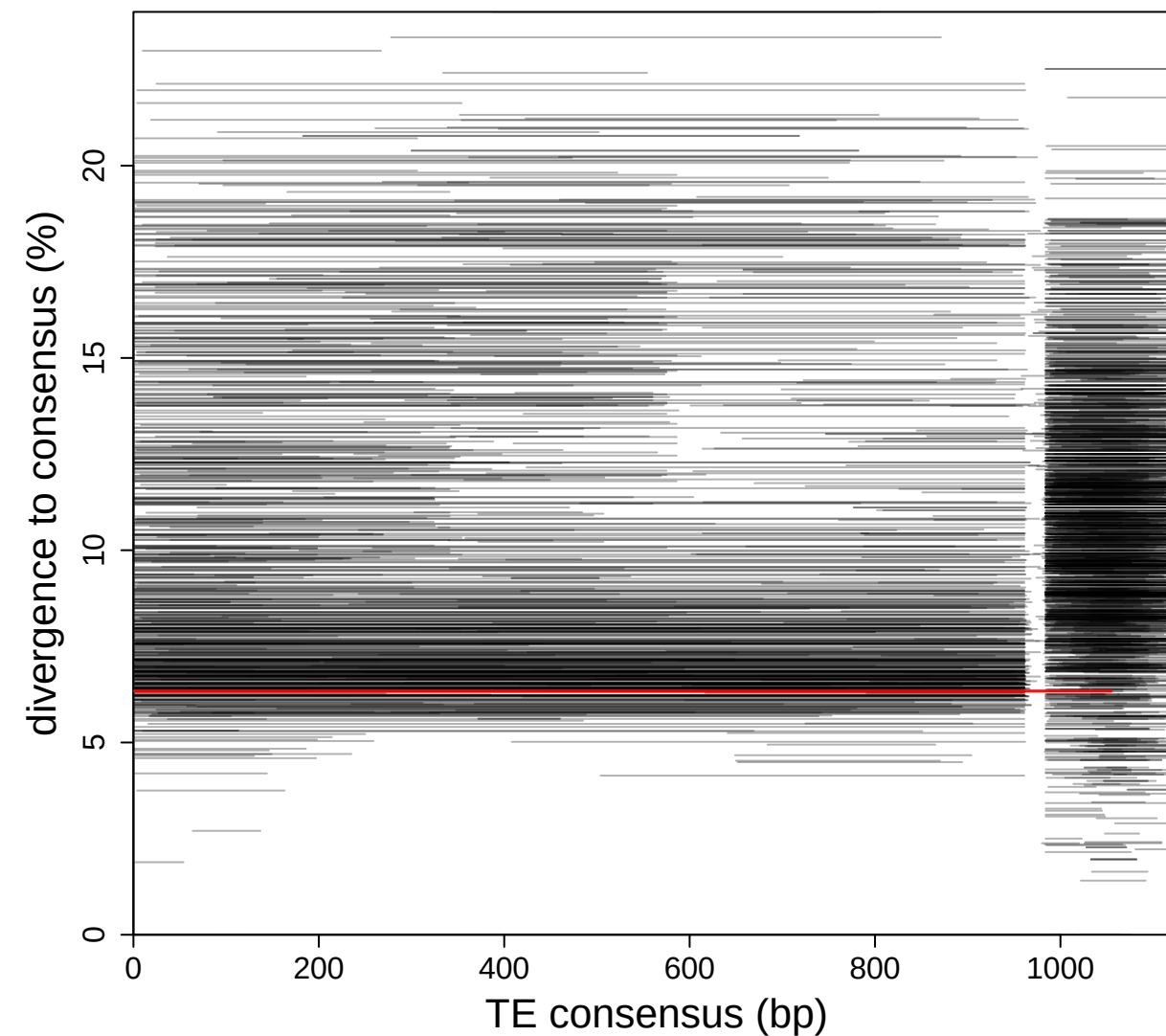
l.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 1259bp; fragments: 890; full length: 0 (>=1133.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

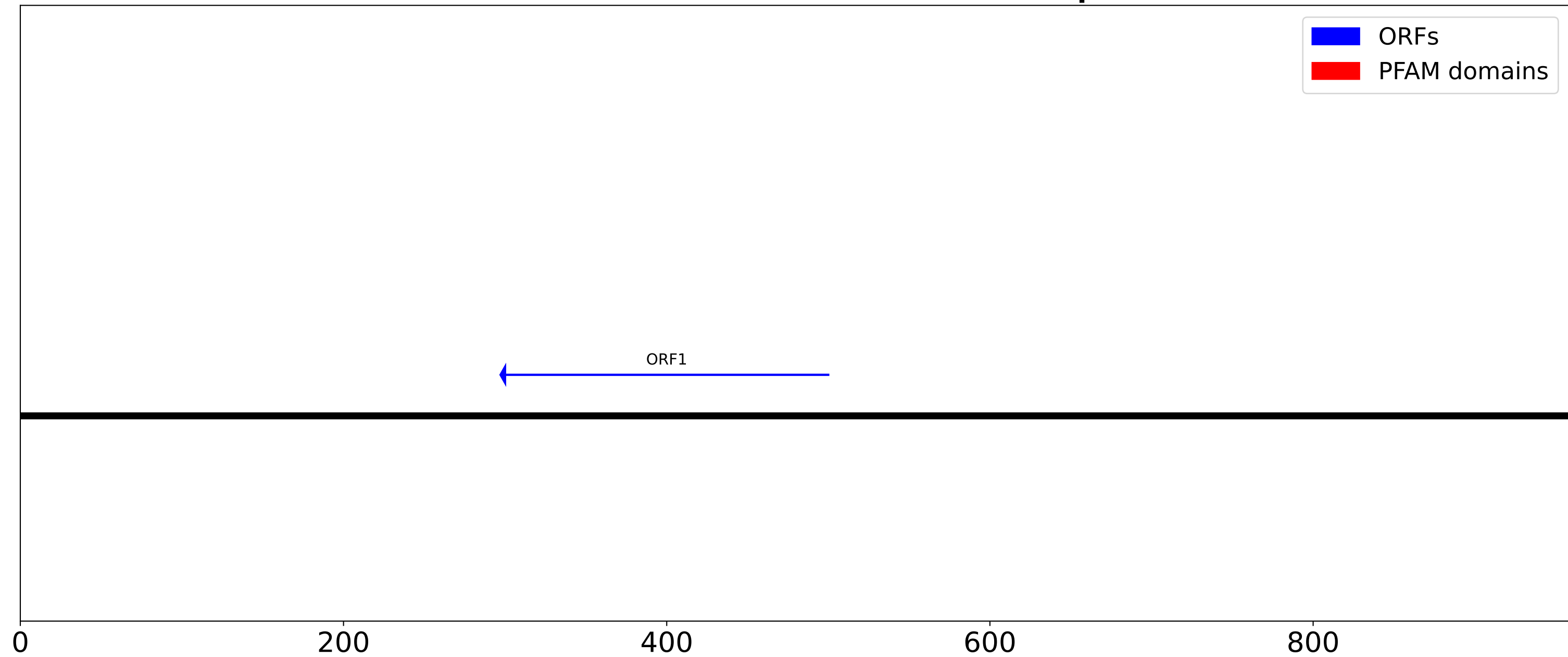


TE: rnd_3 family_24
size: 1129bp; fragments: 2466; full length: 1 (≥ 1016.1 bp)

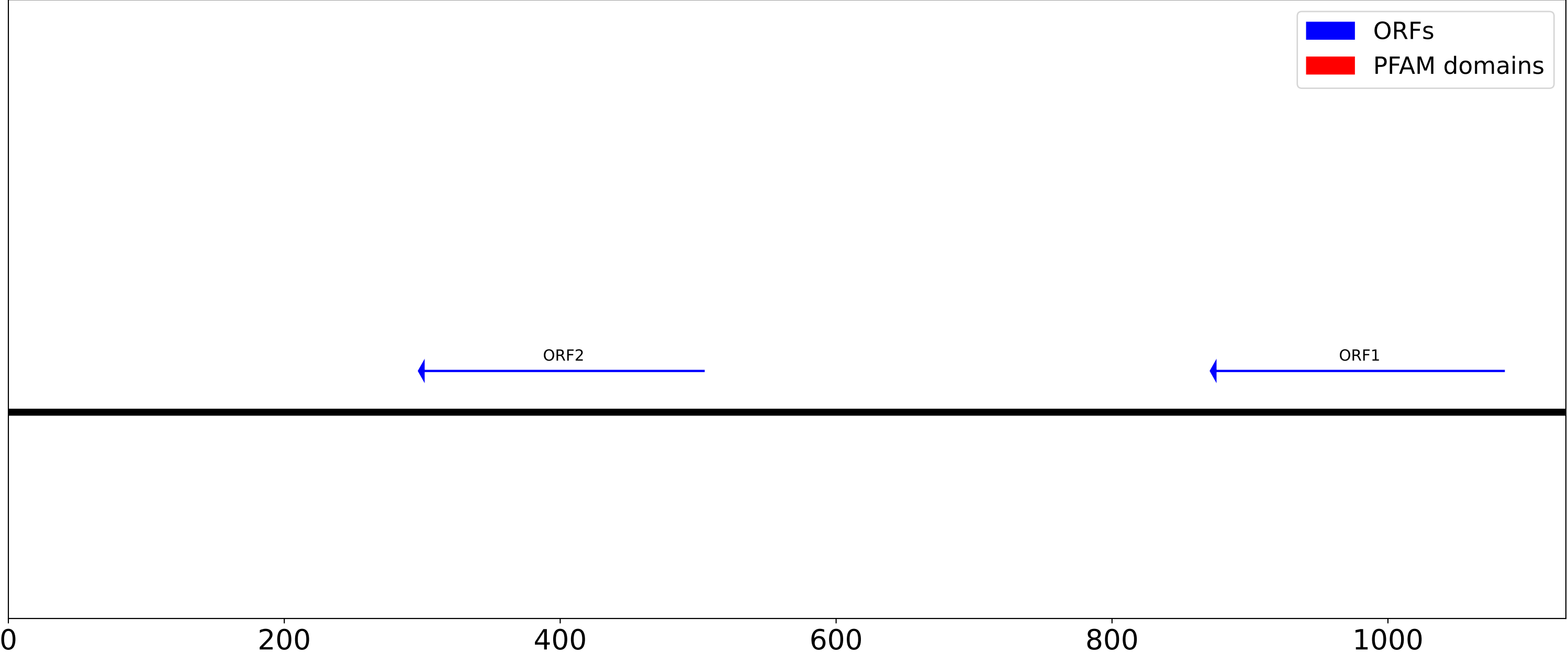
Before TEtrimmer 1129 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

