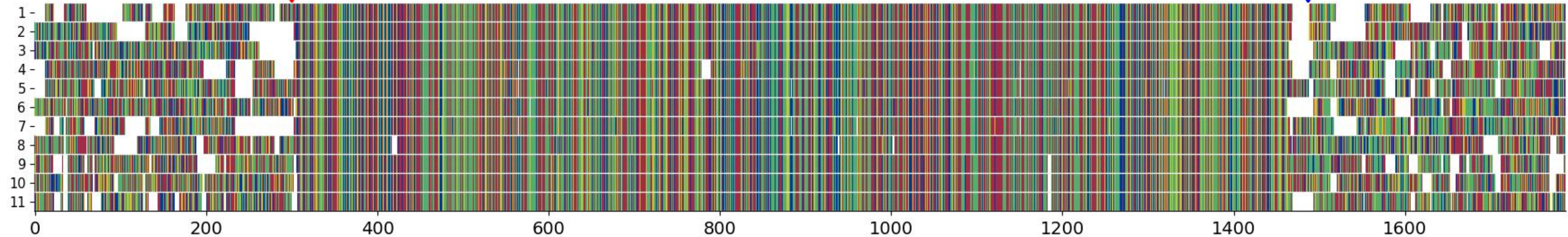


A length

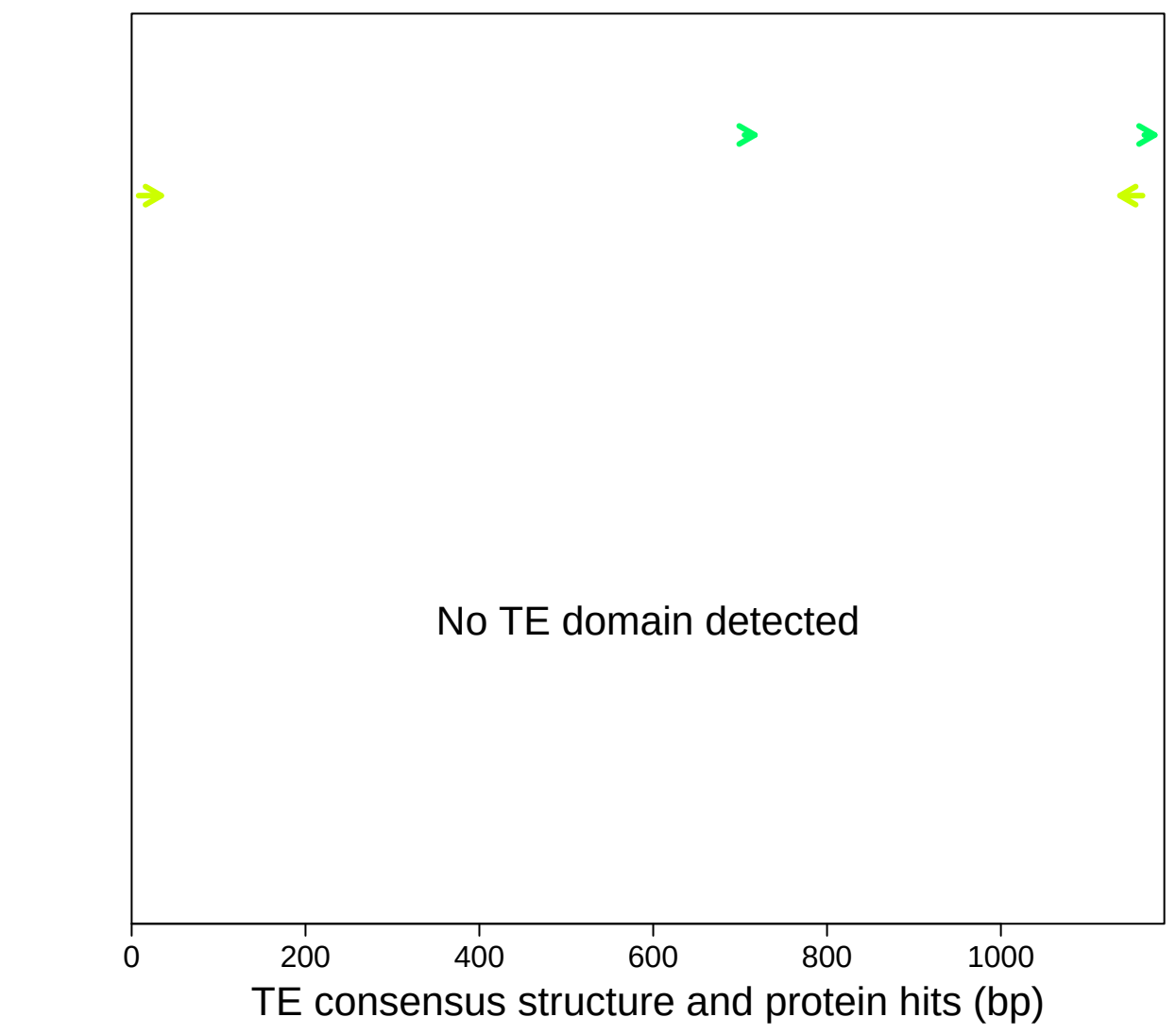
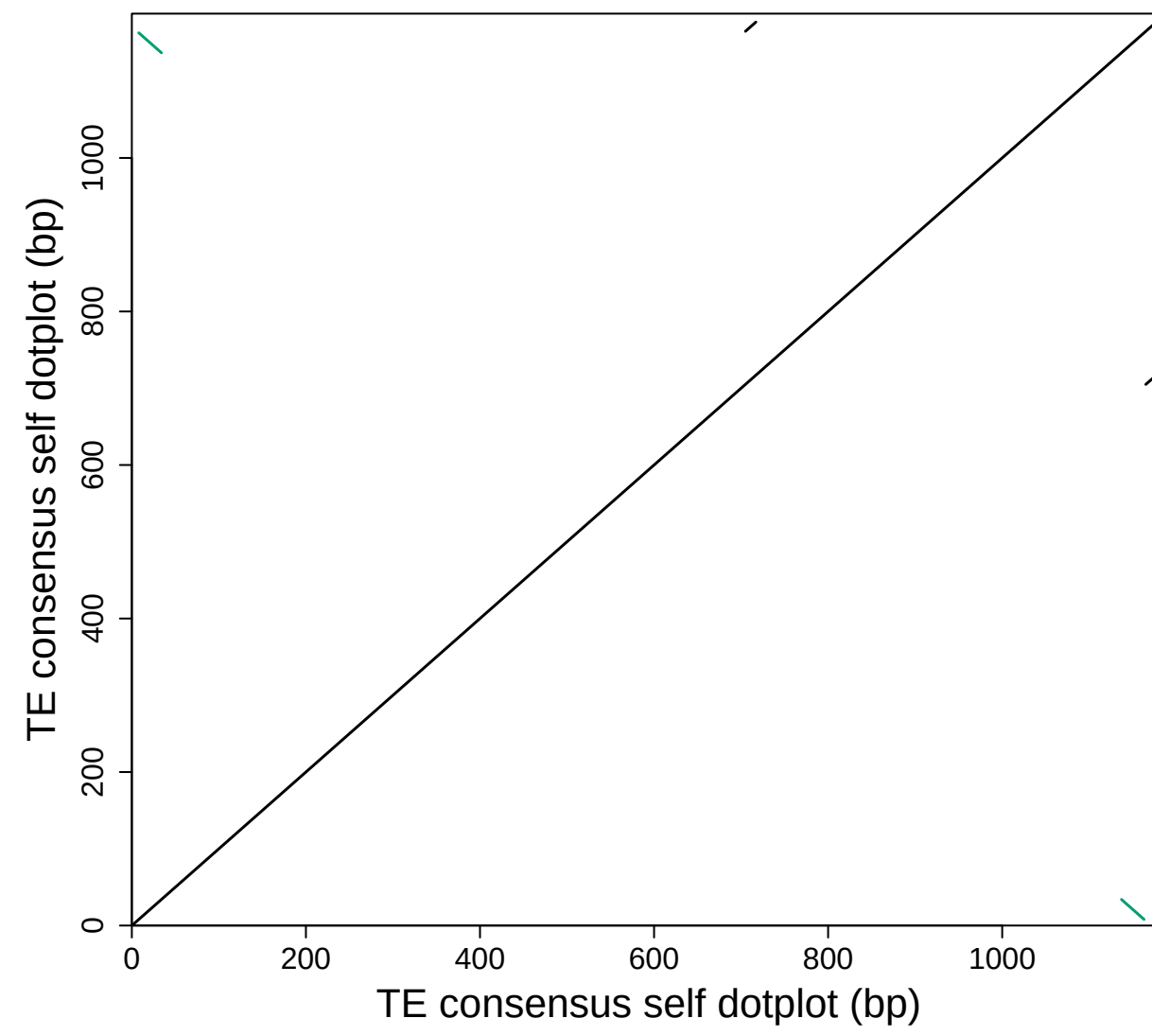
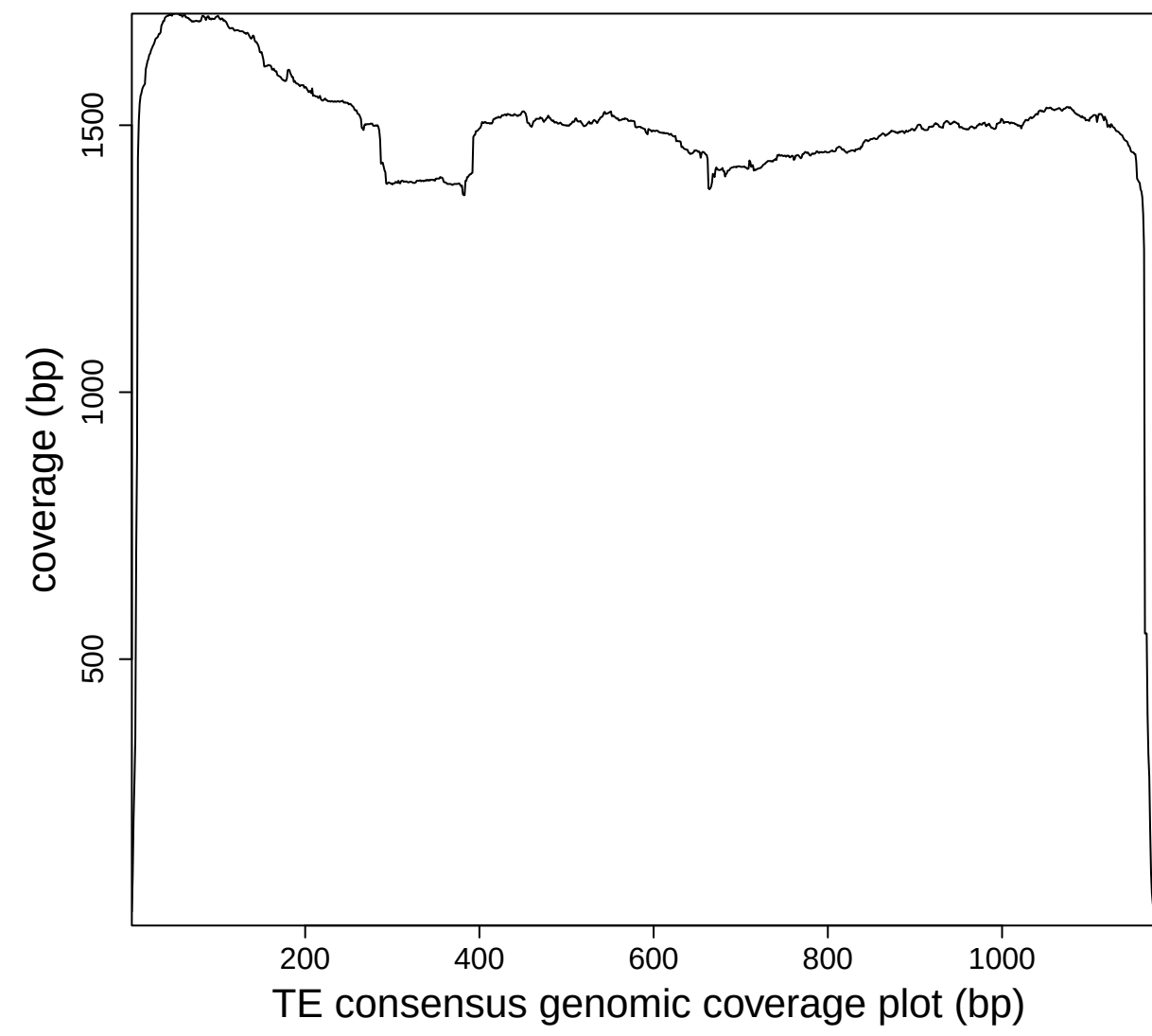
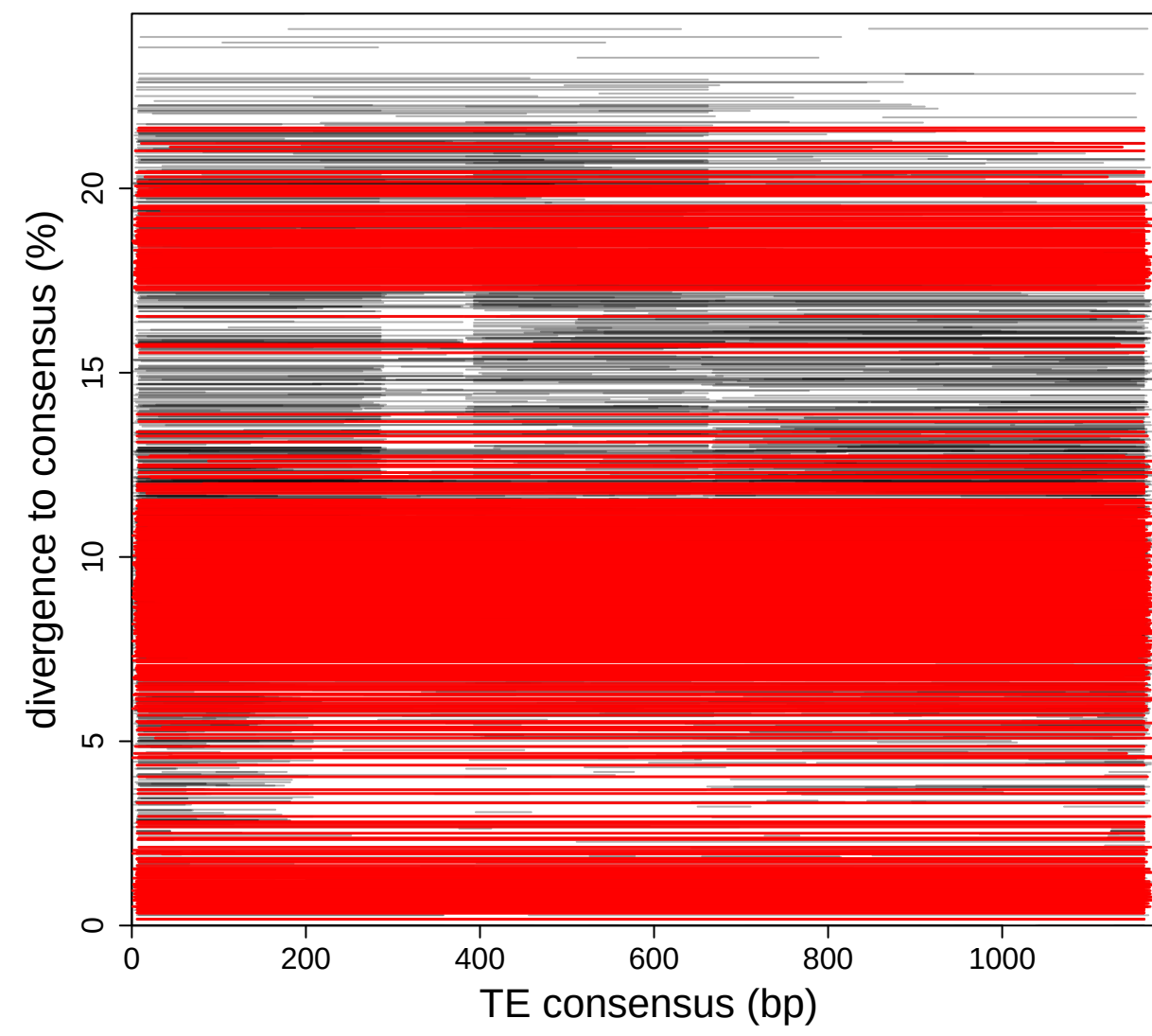
Start crop Point

End crop Point



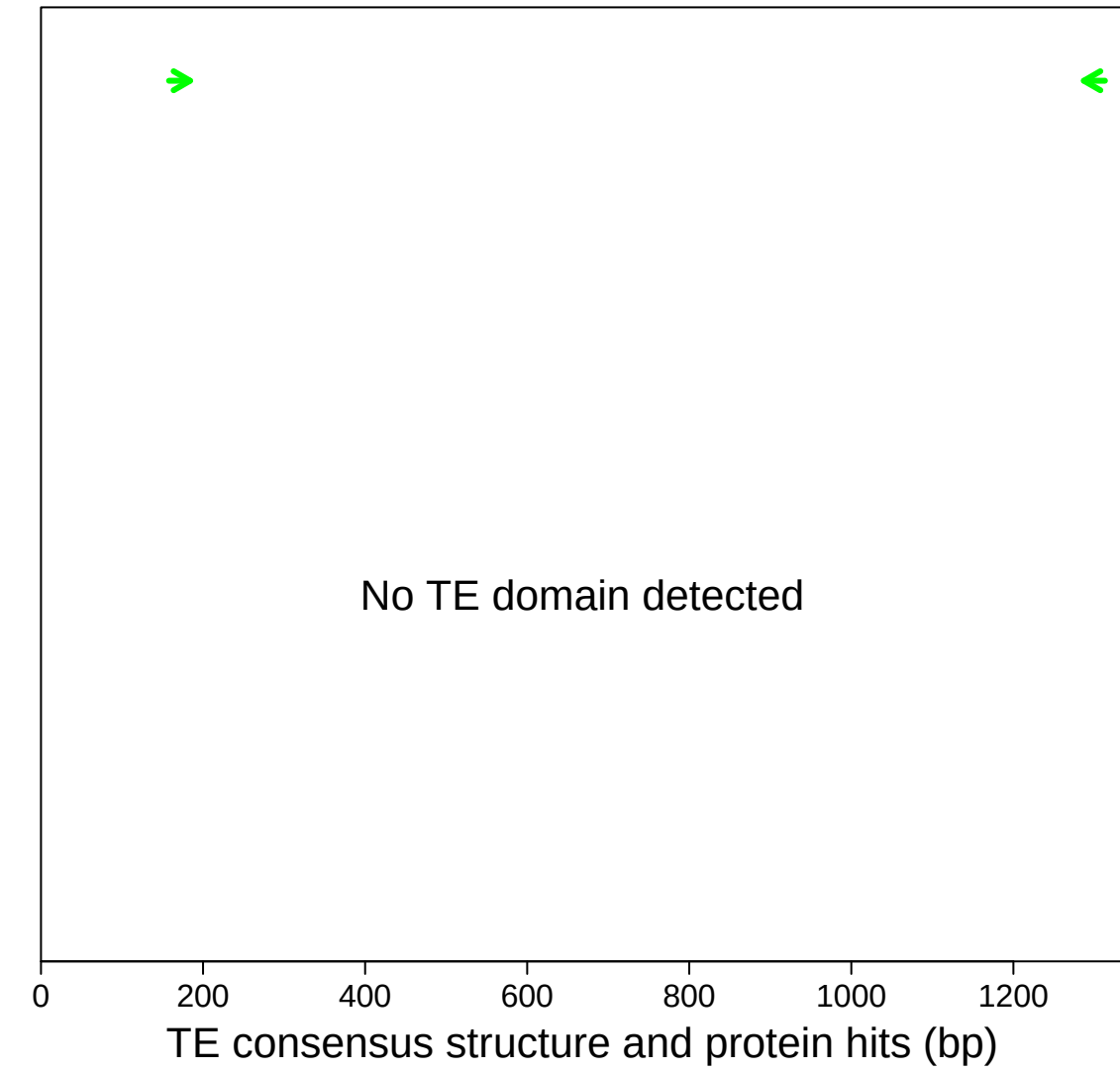
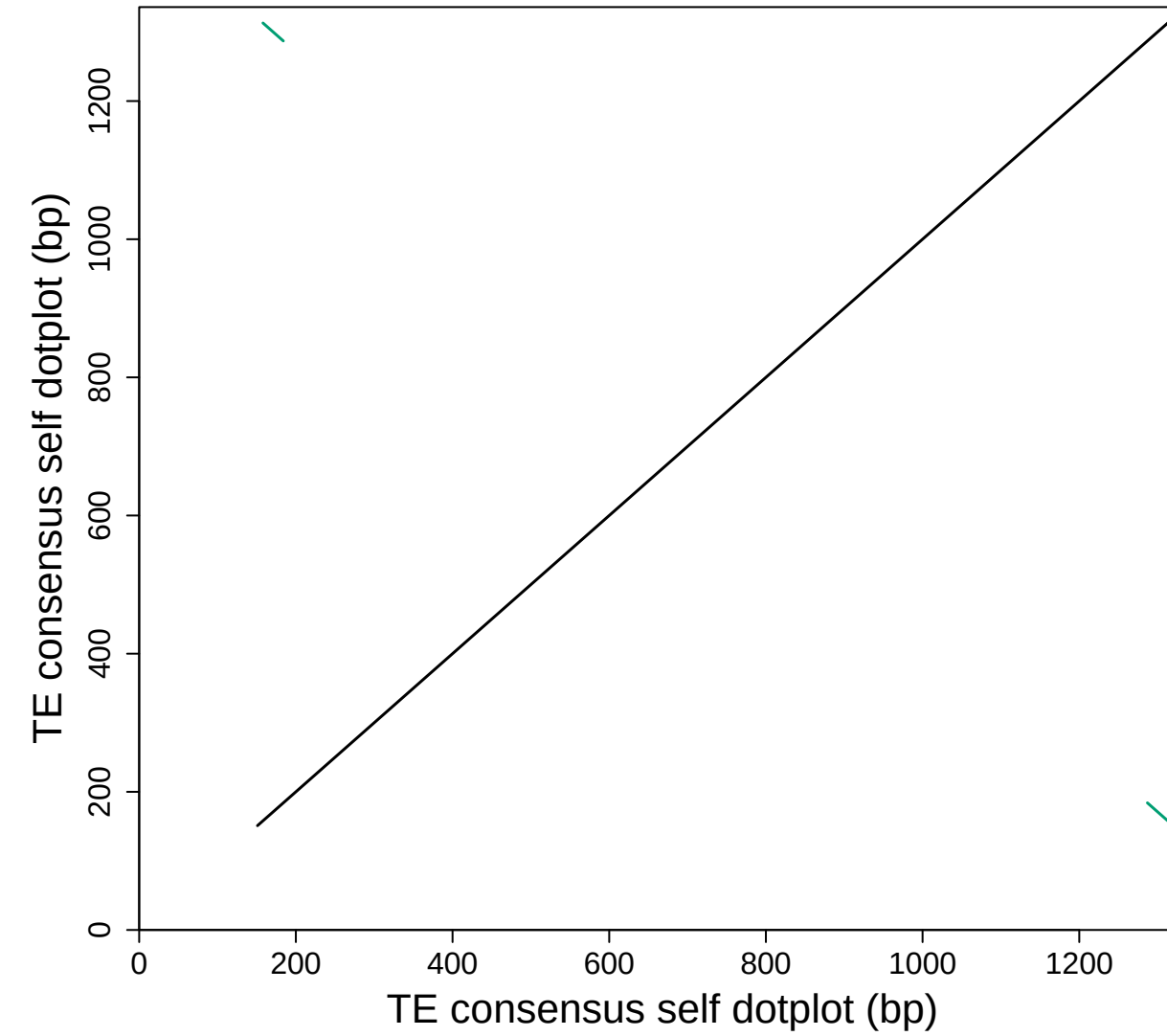
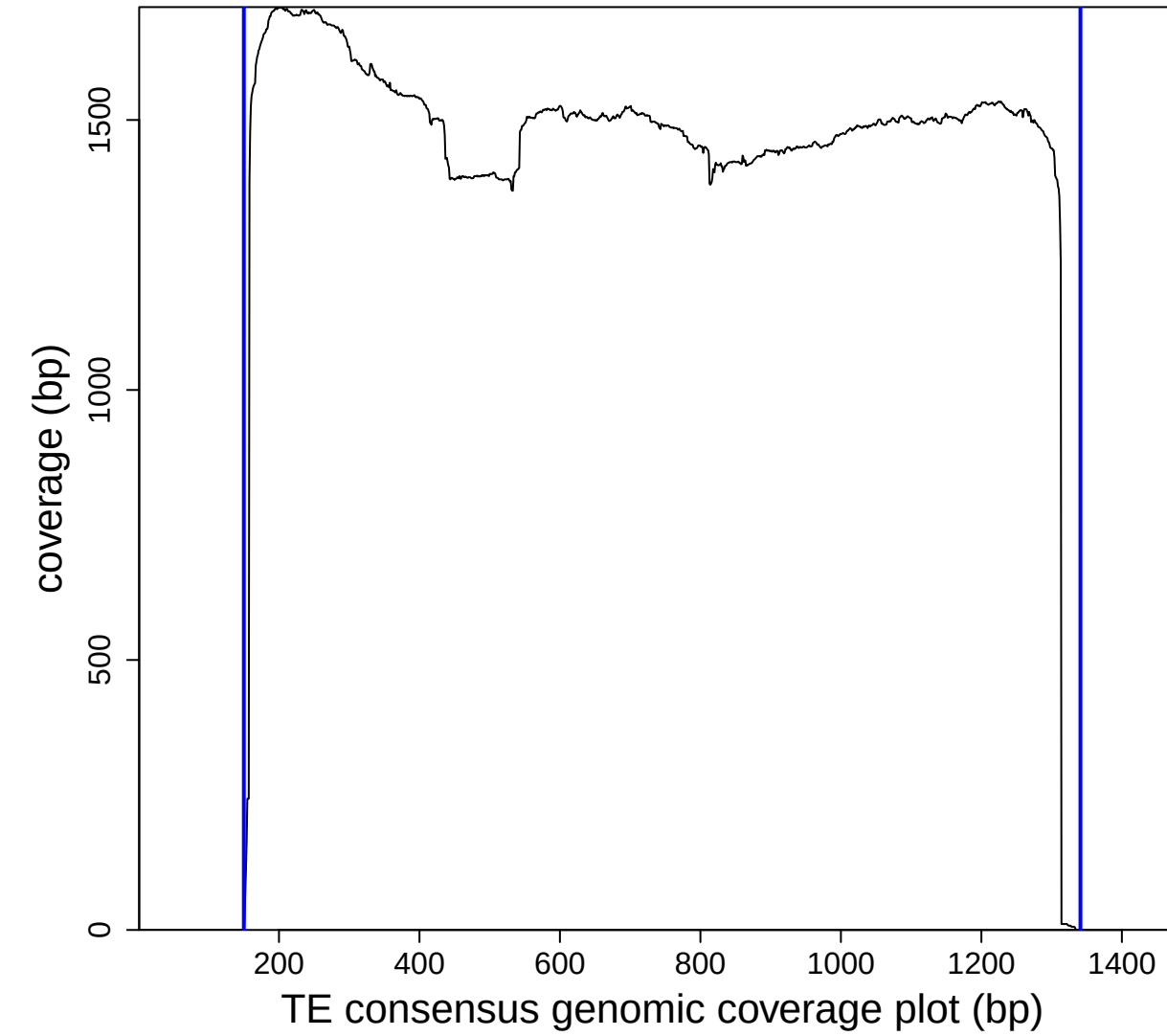
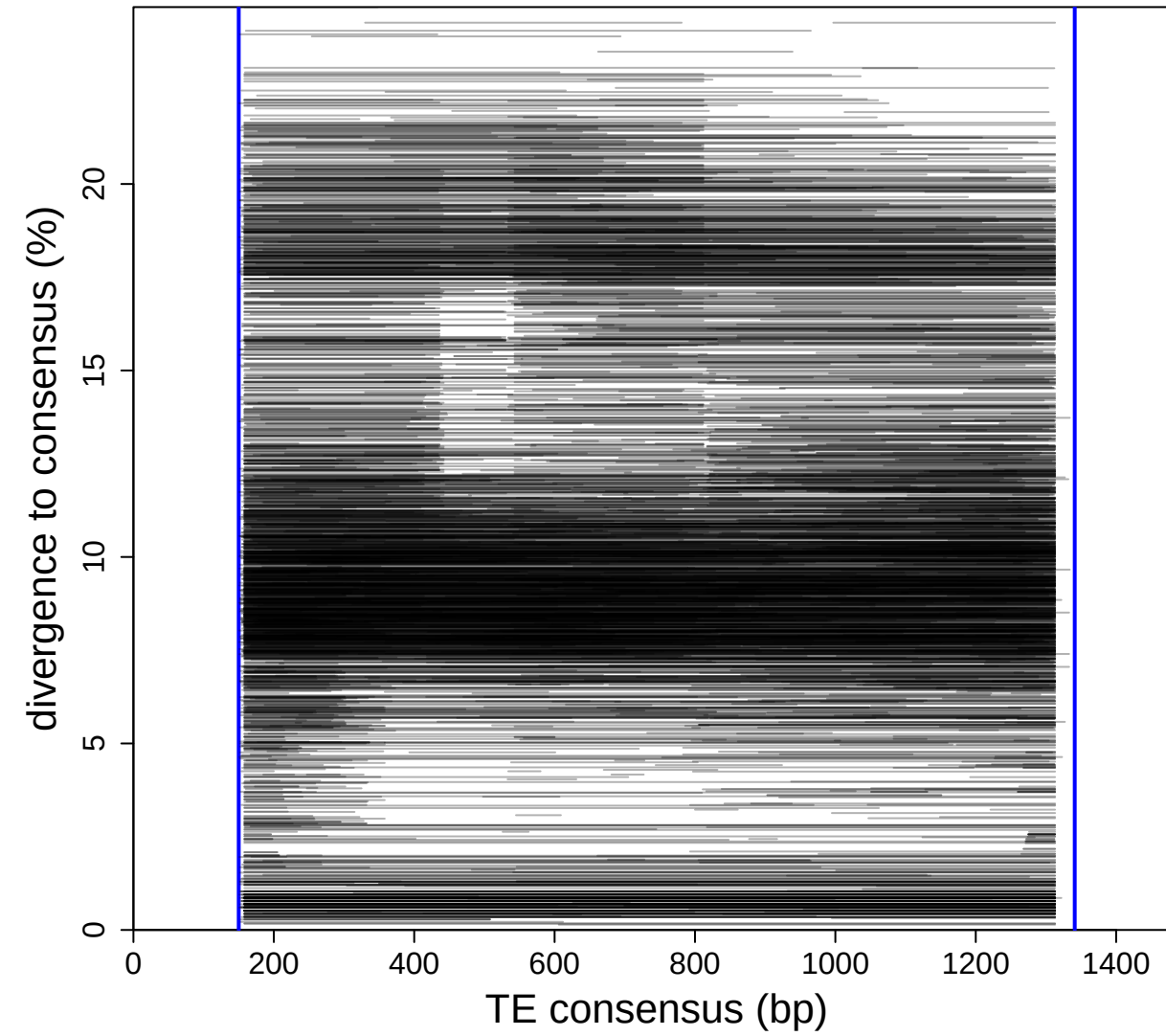
b.bed uf.bed g 4.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 1188bp; fragments: 3064; full length: 701 (≥ 1069.2 bp)

After TEtrimmer 1188 bp



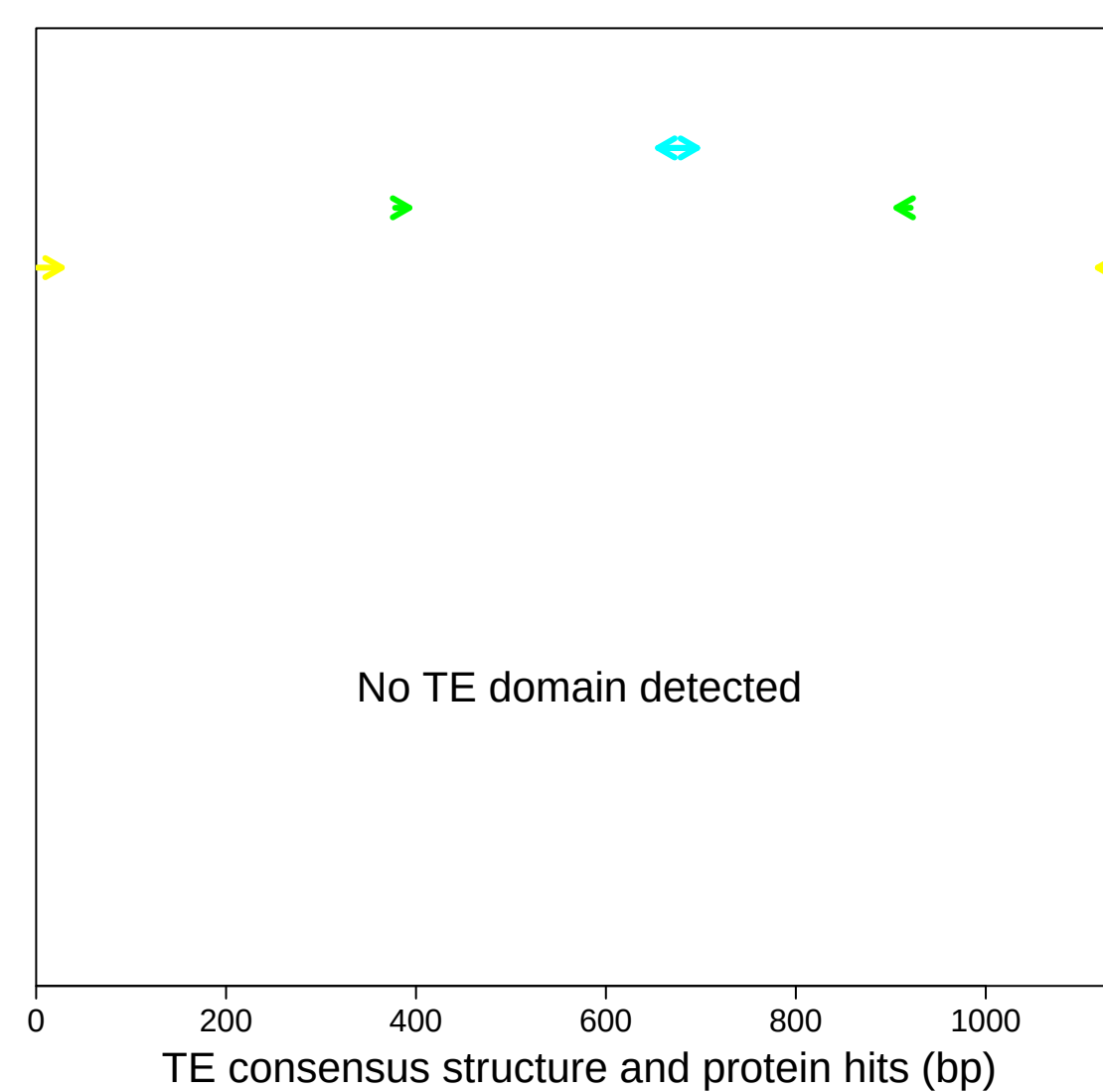
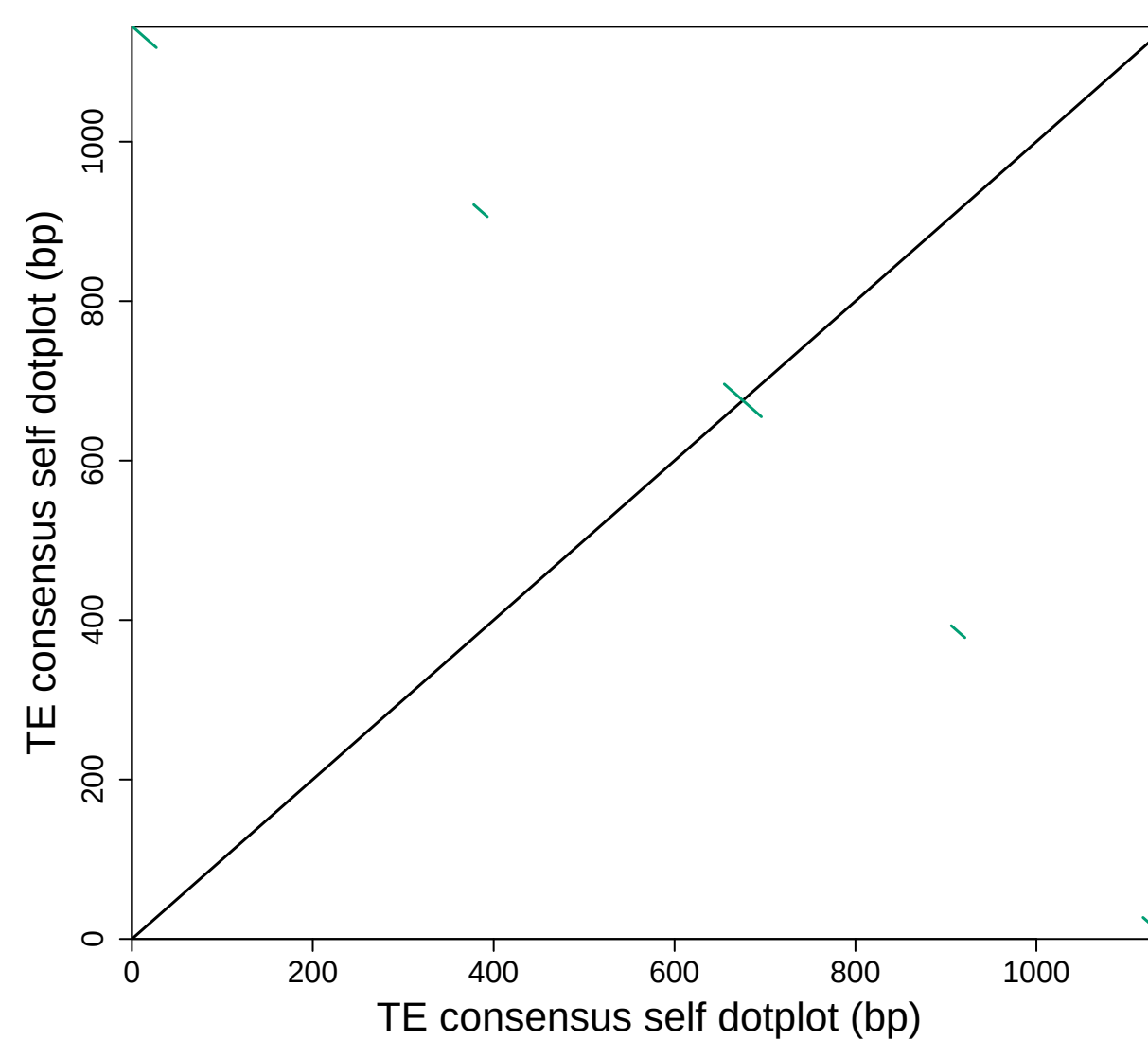
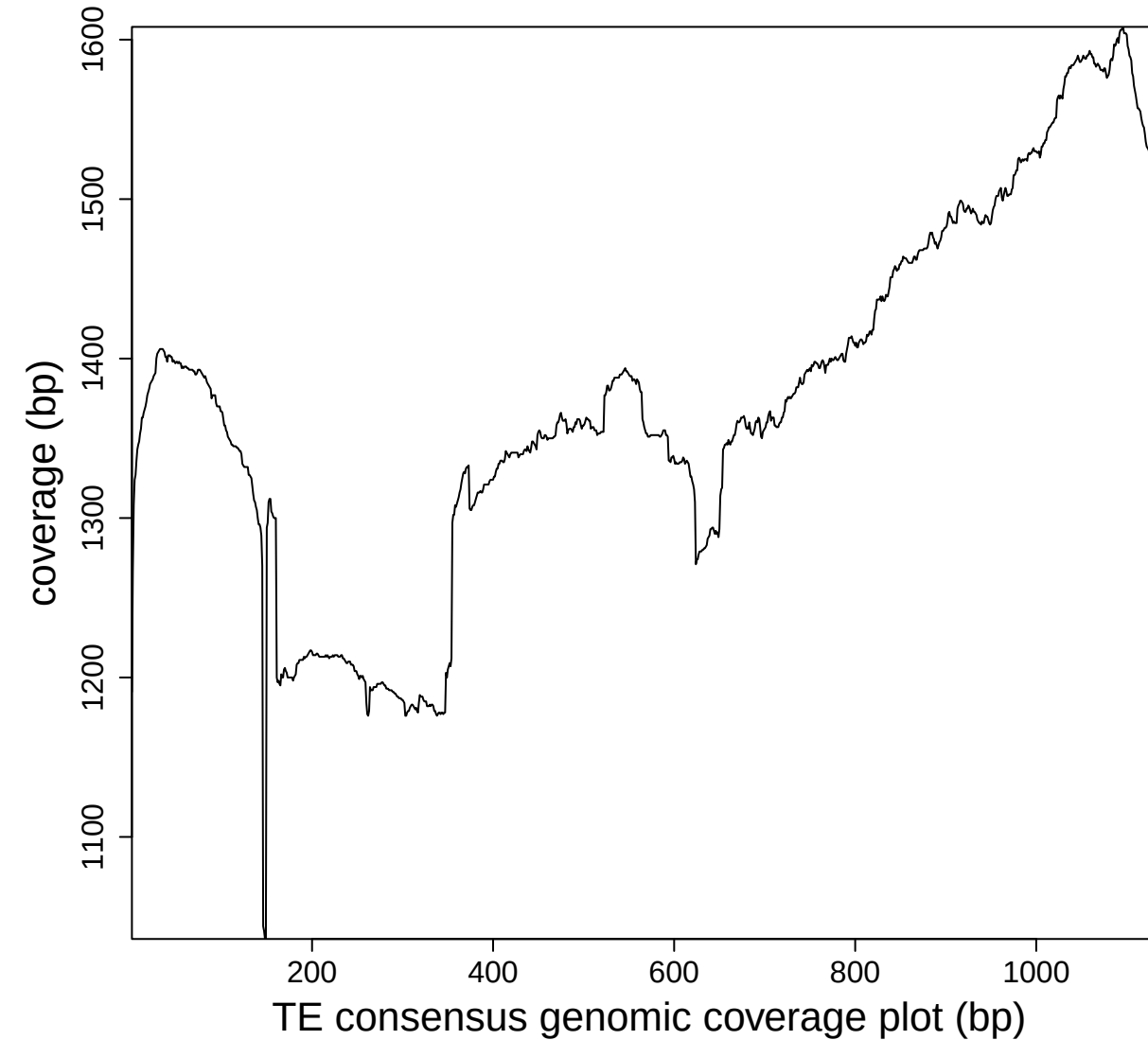
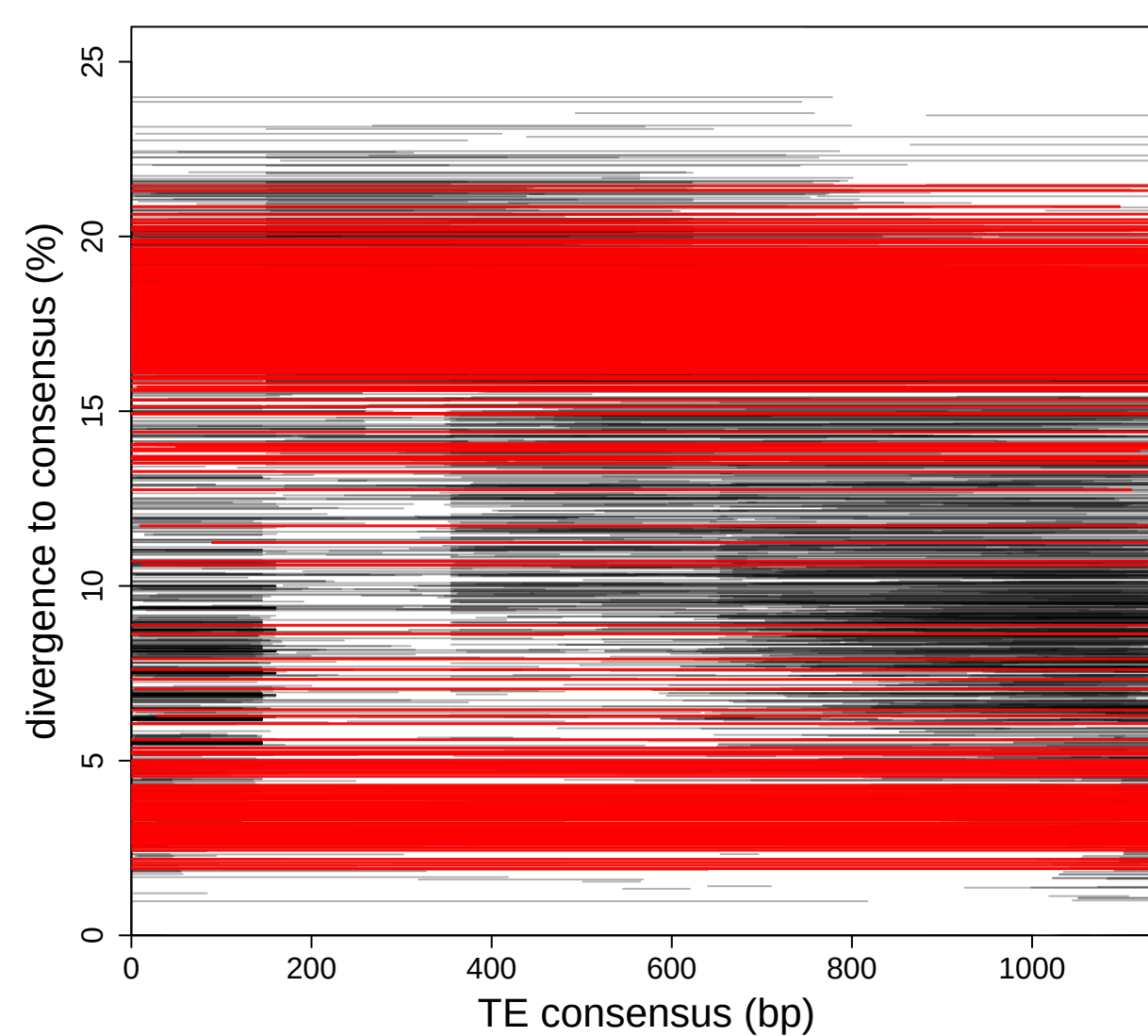
2.fasta.b.bed_uf.bed_g_4.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 1491bp; fragments: 3059; full length: 0 (>=1341.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

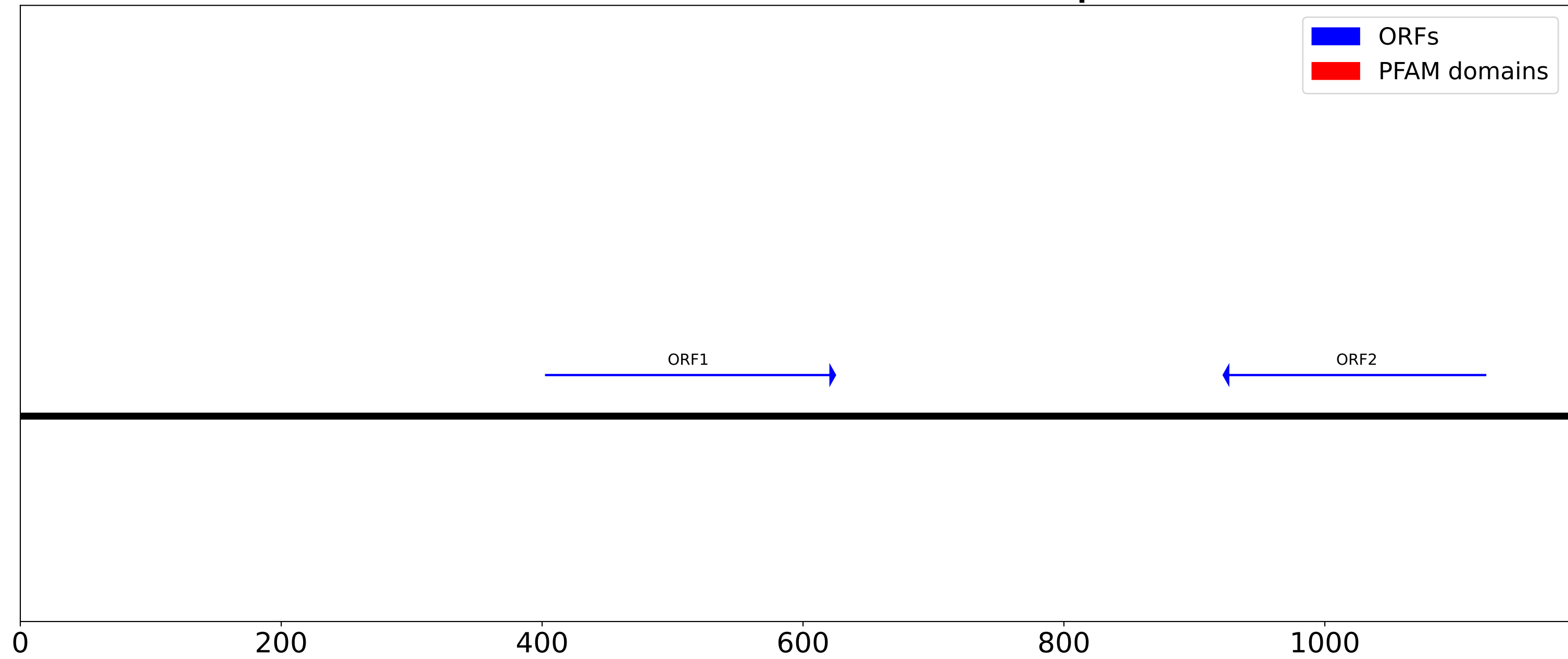


TE: rnd_1_family_92
size: 1144bp; fragments: 3085; full length: 480 (≥ 1029.6 bp)

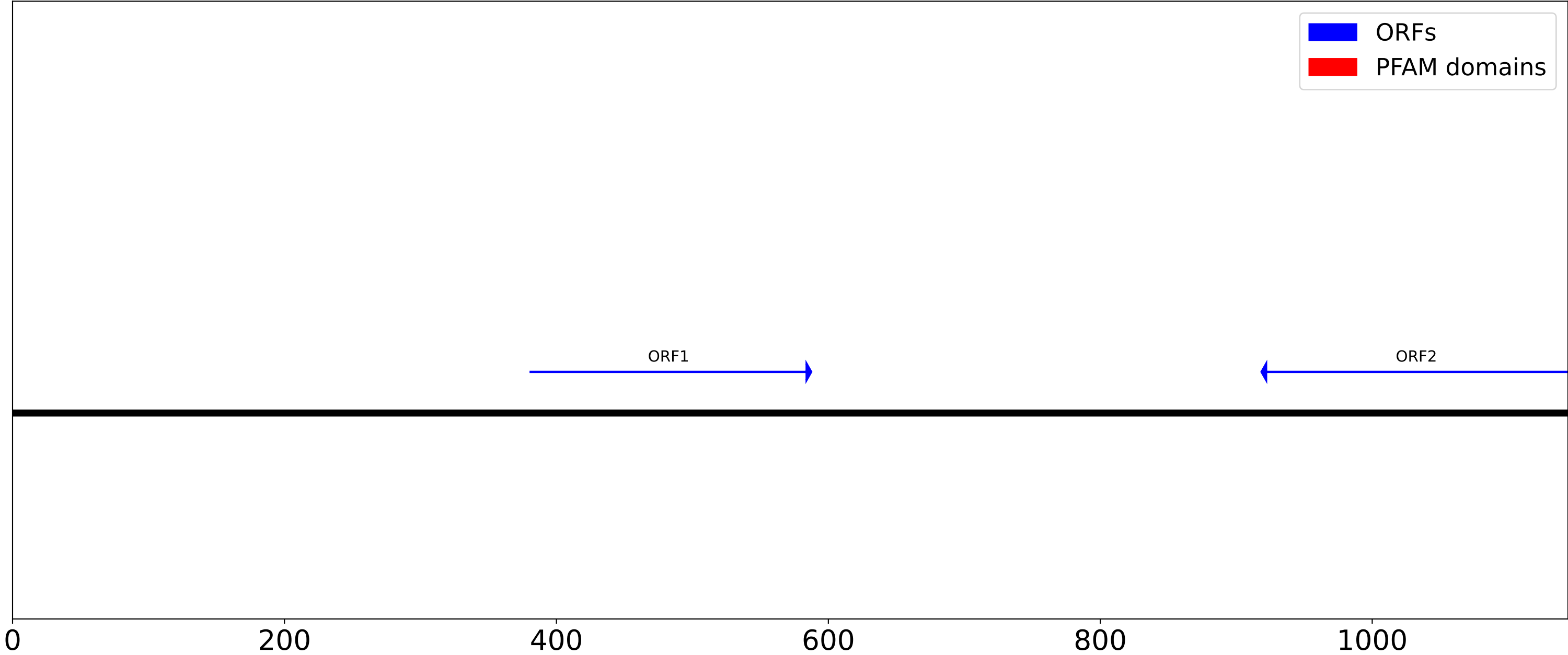
Before TEtrimmer 1144 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

