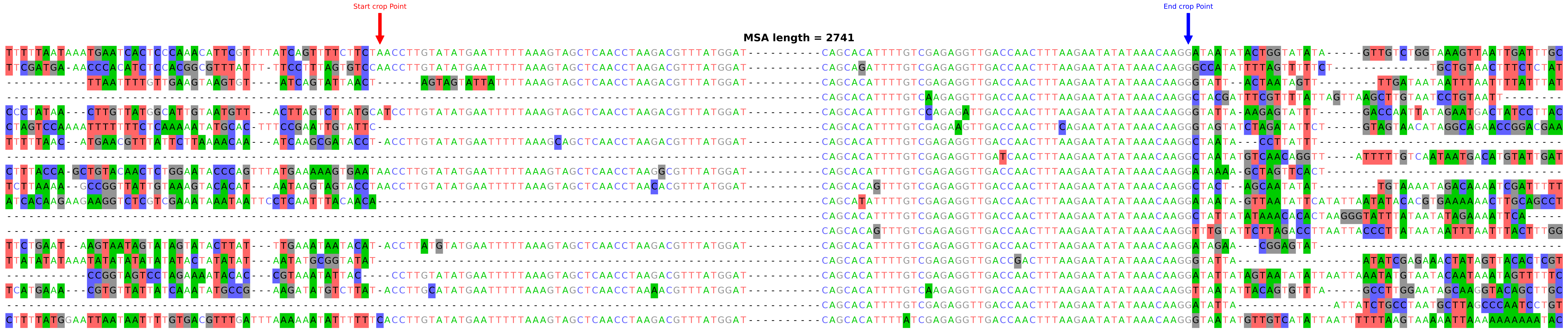
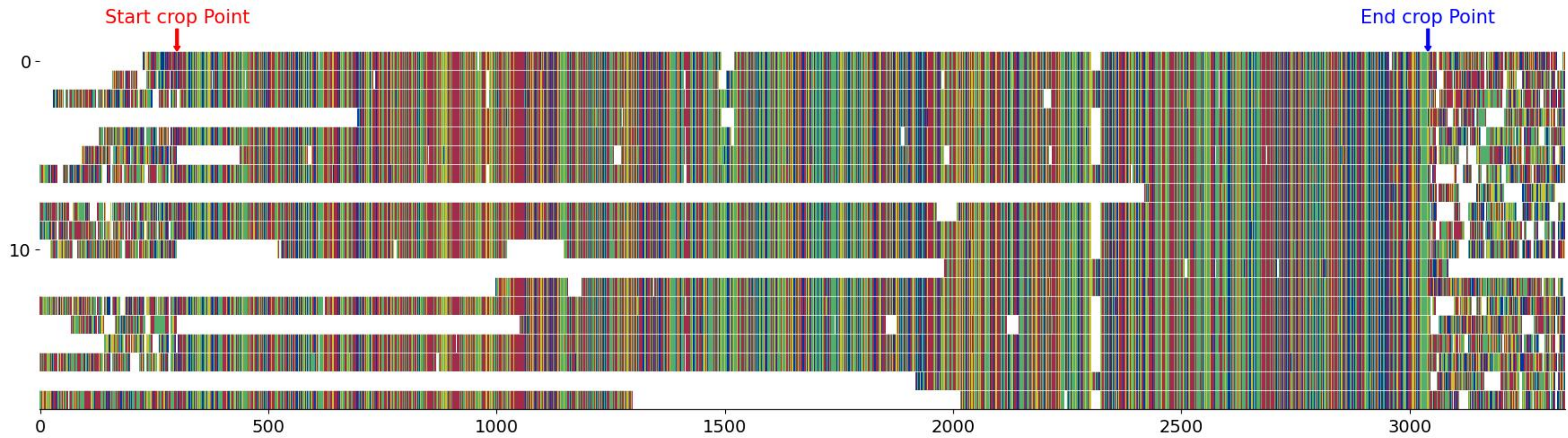




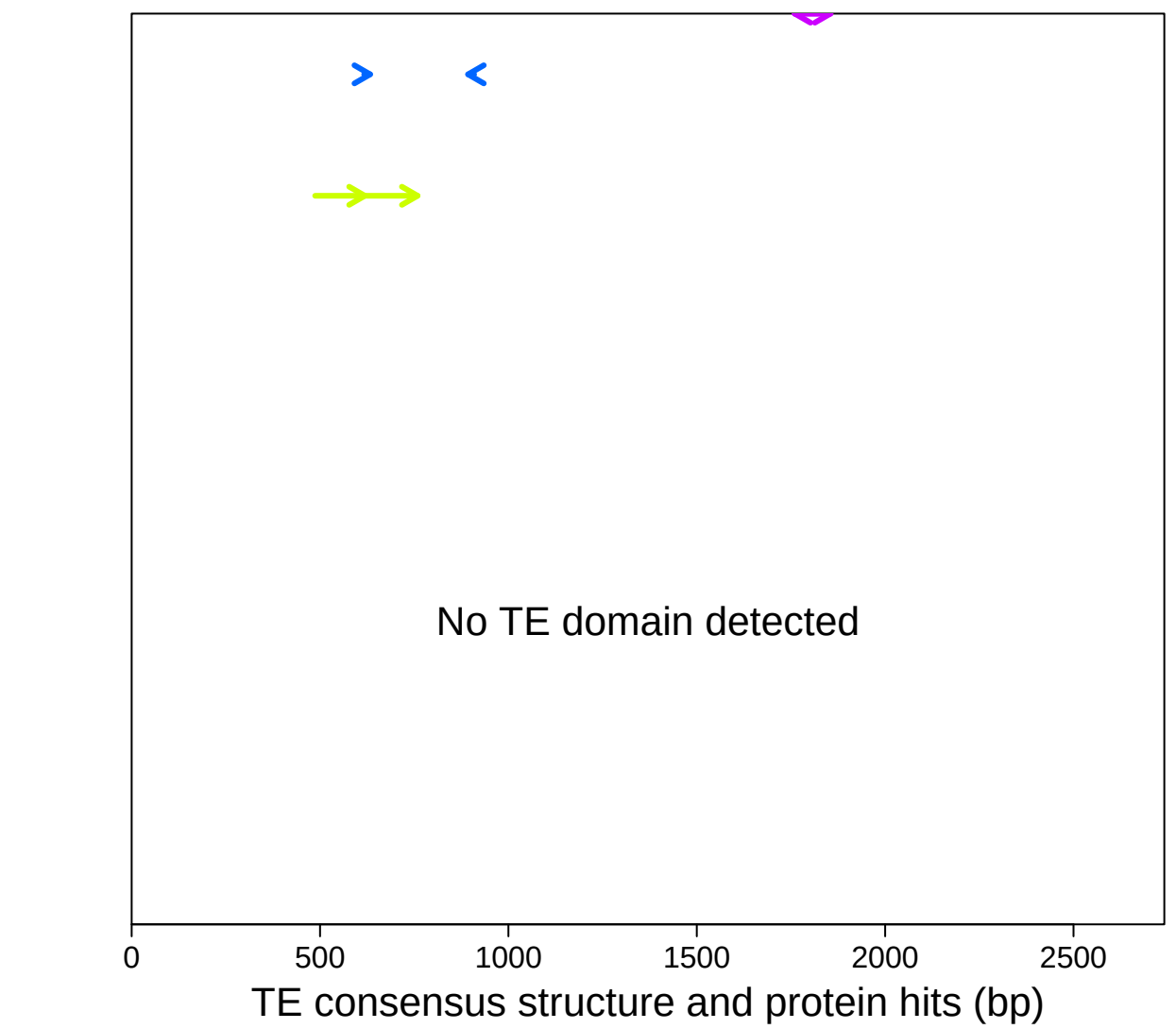
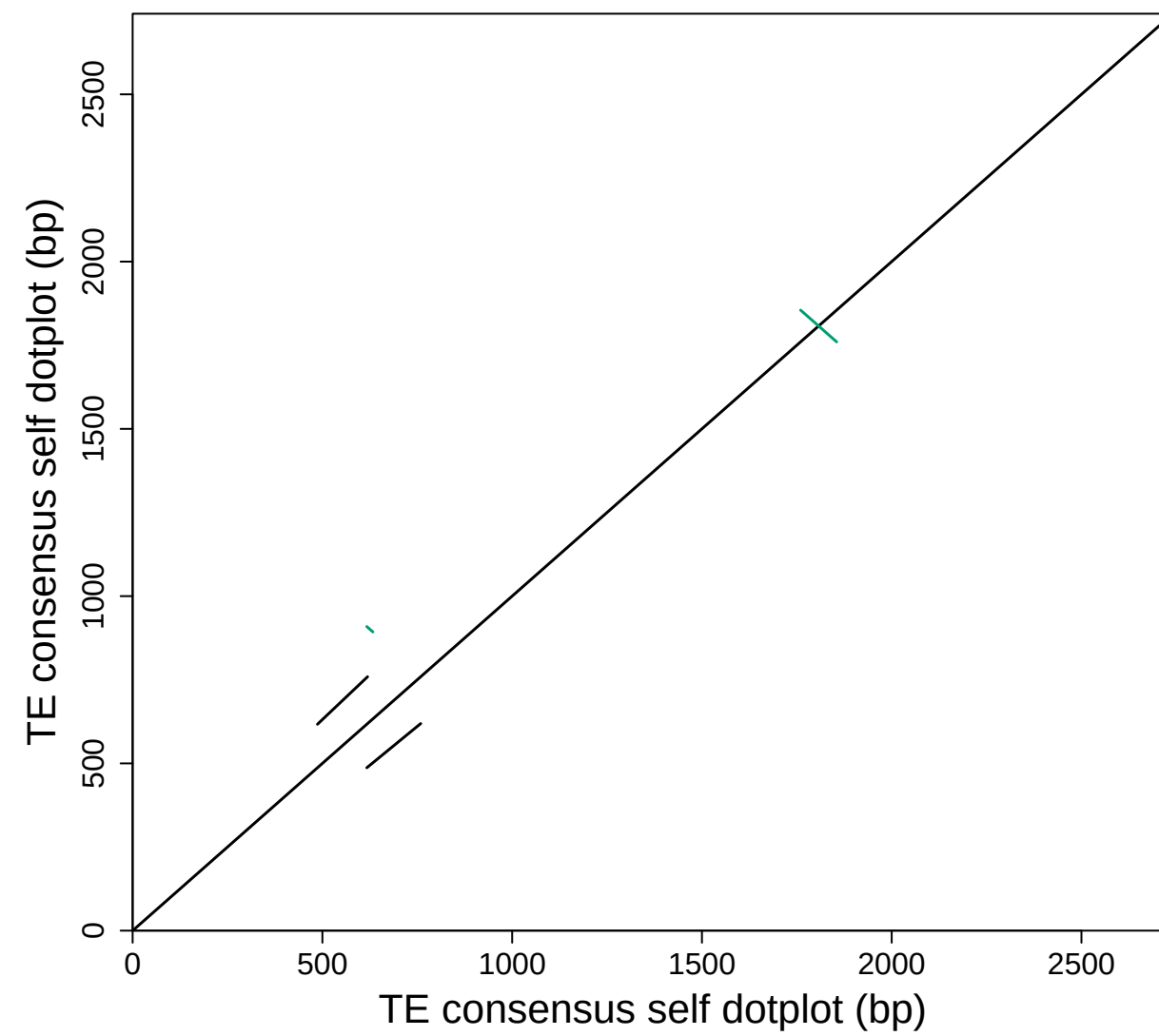
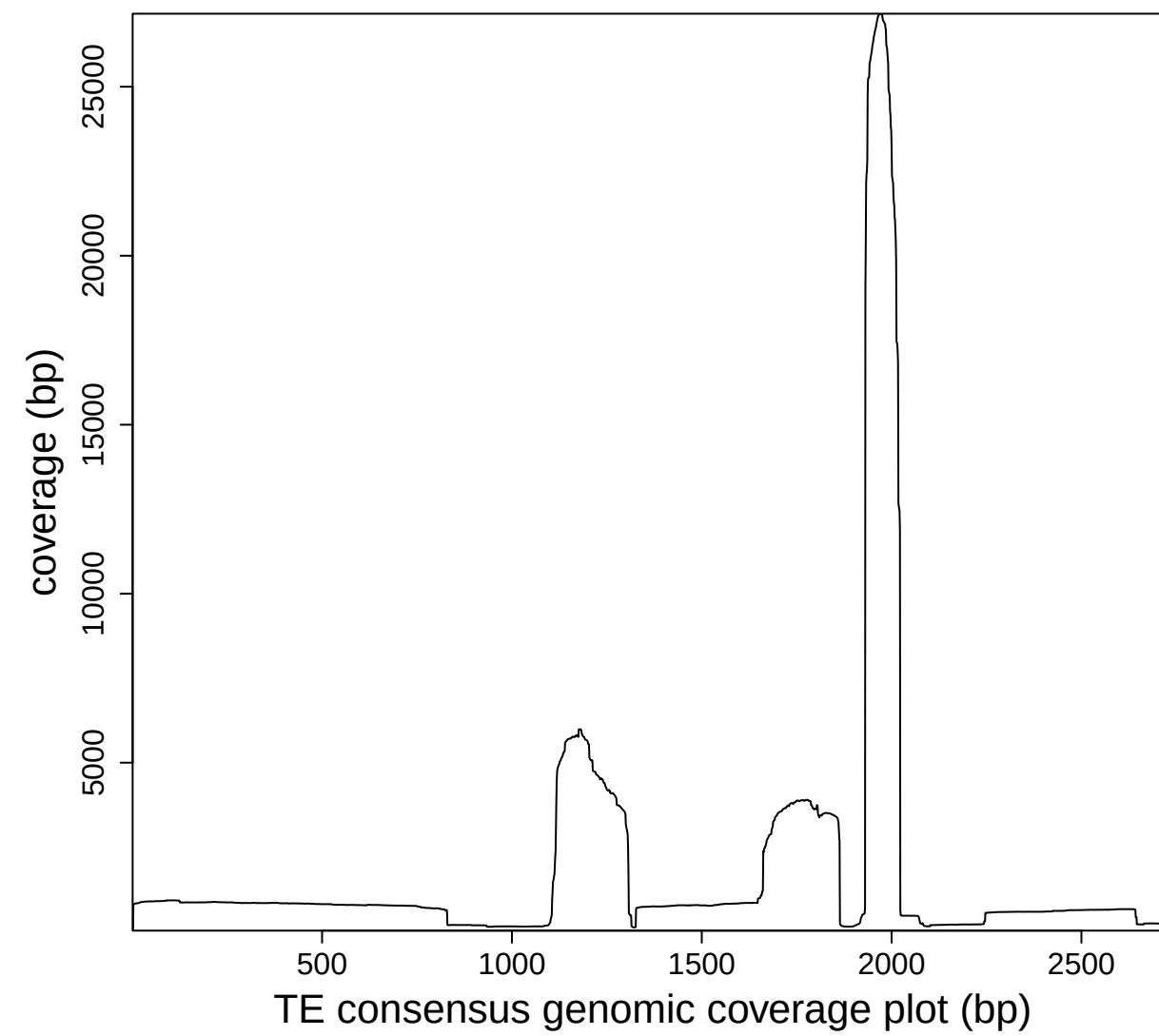
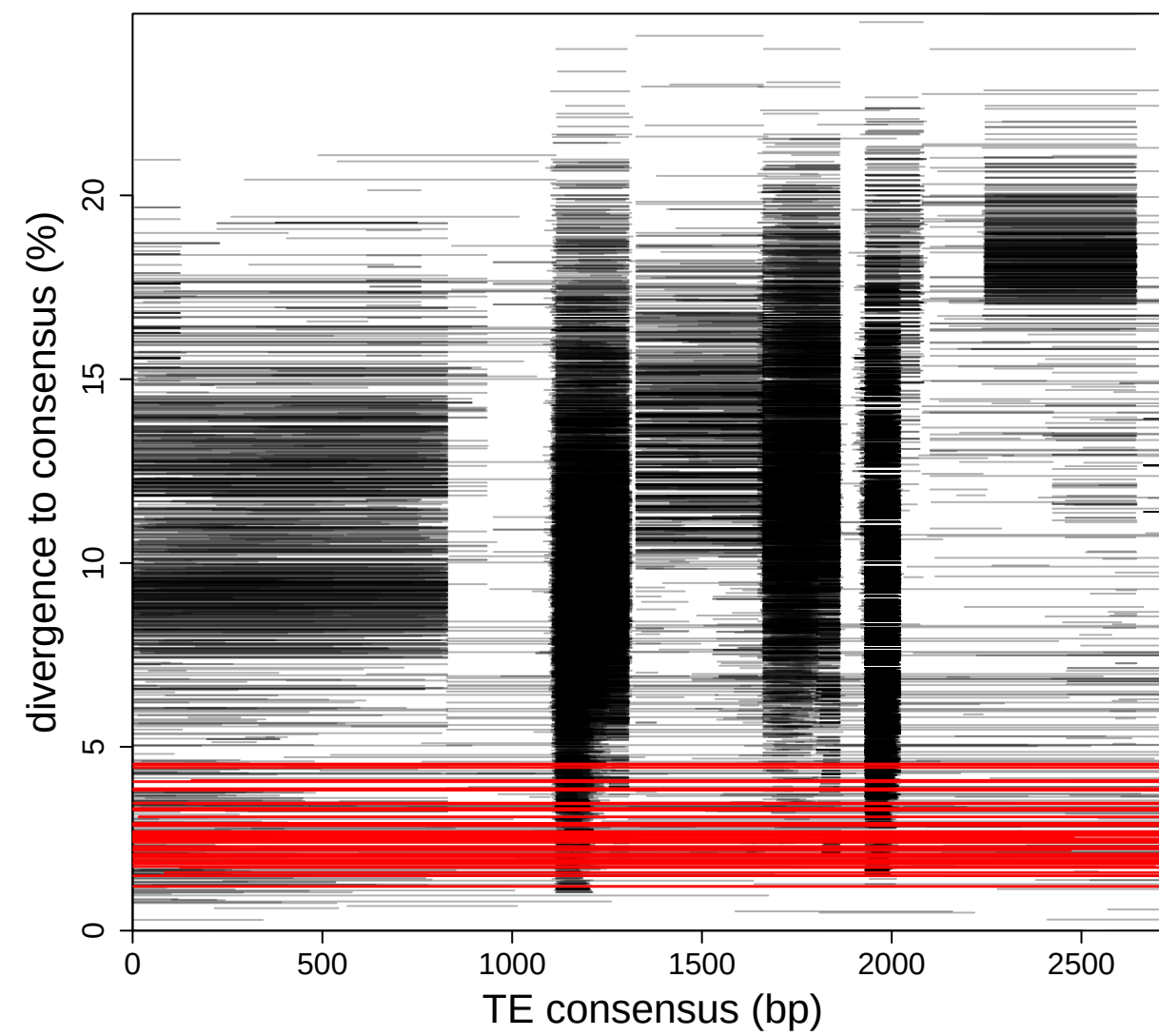
Start crop Point

End crop Point



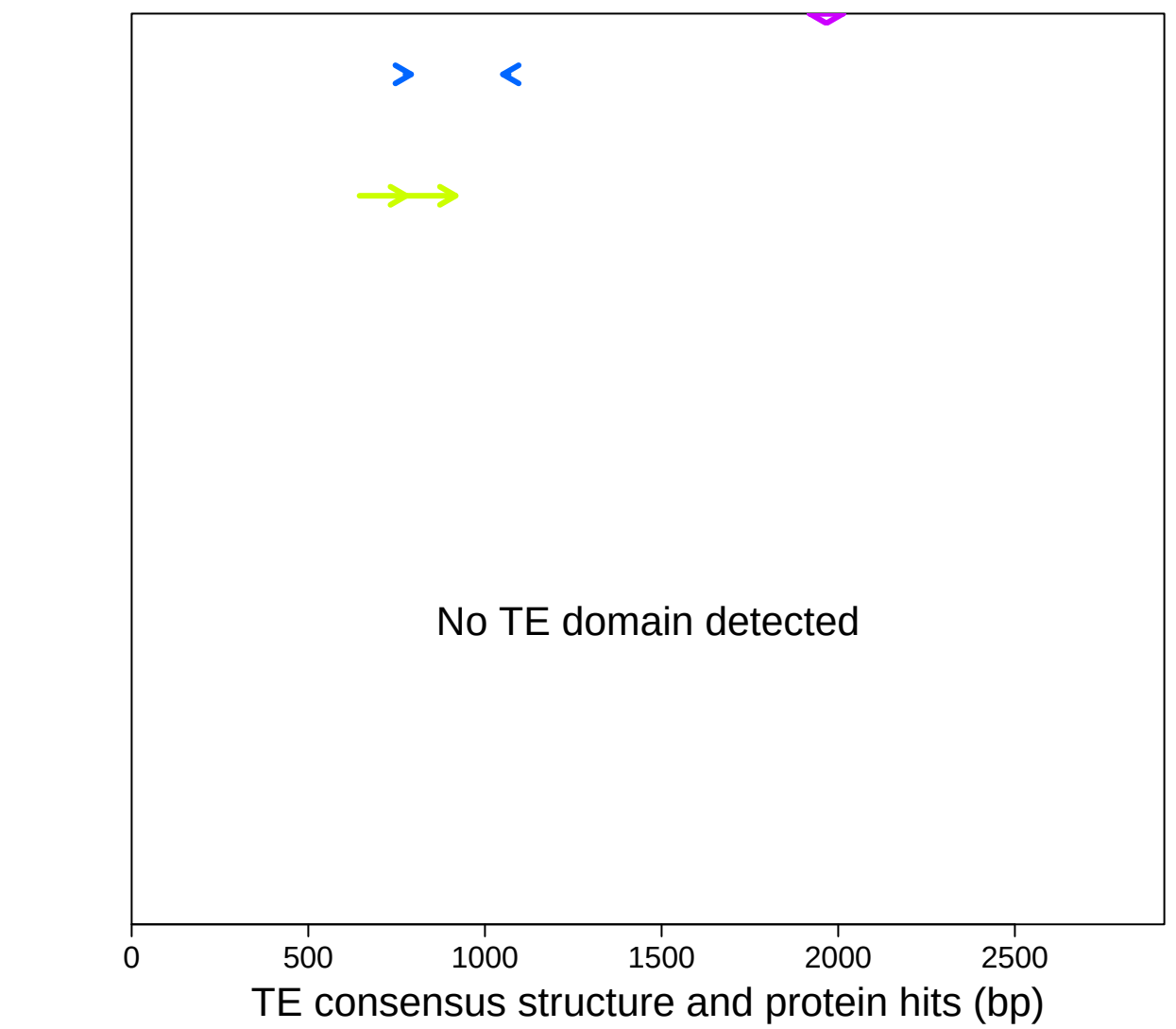
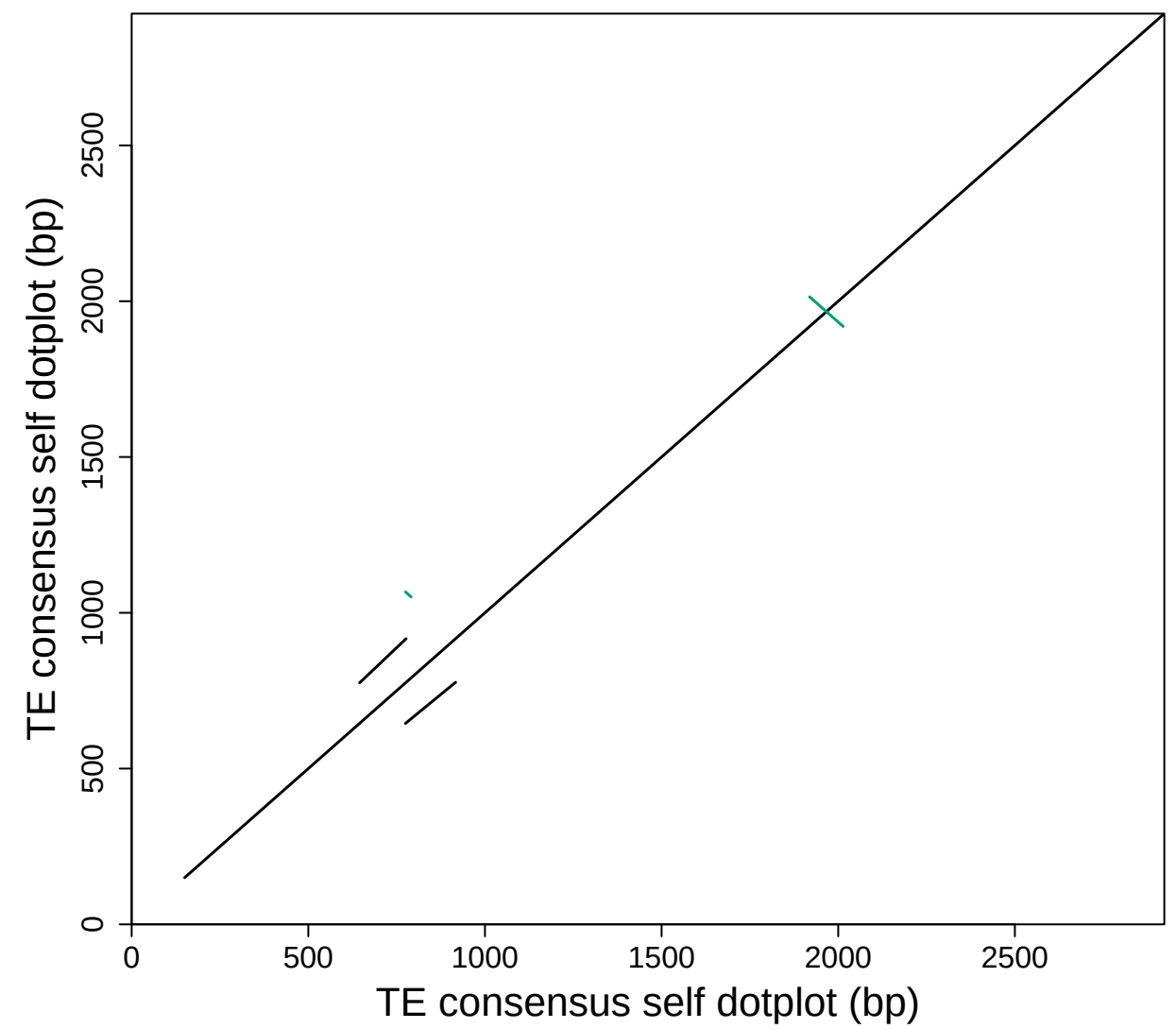
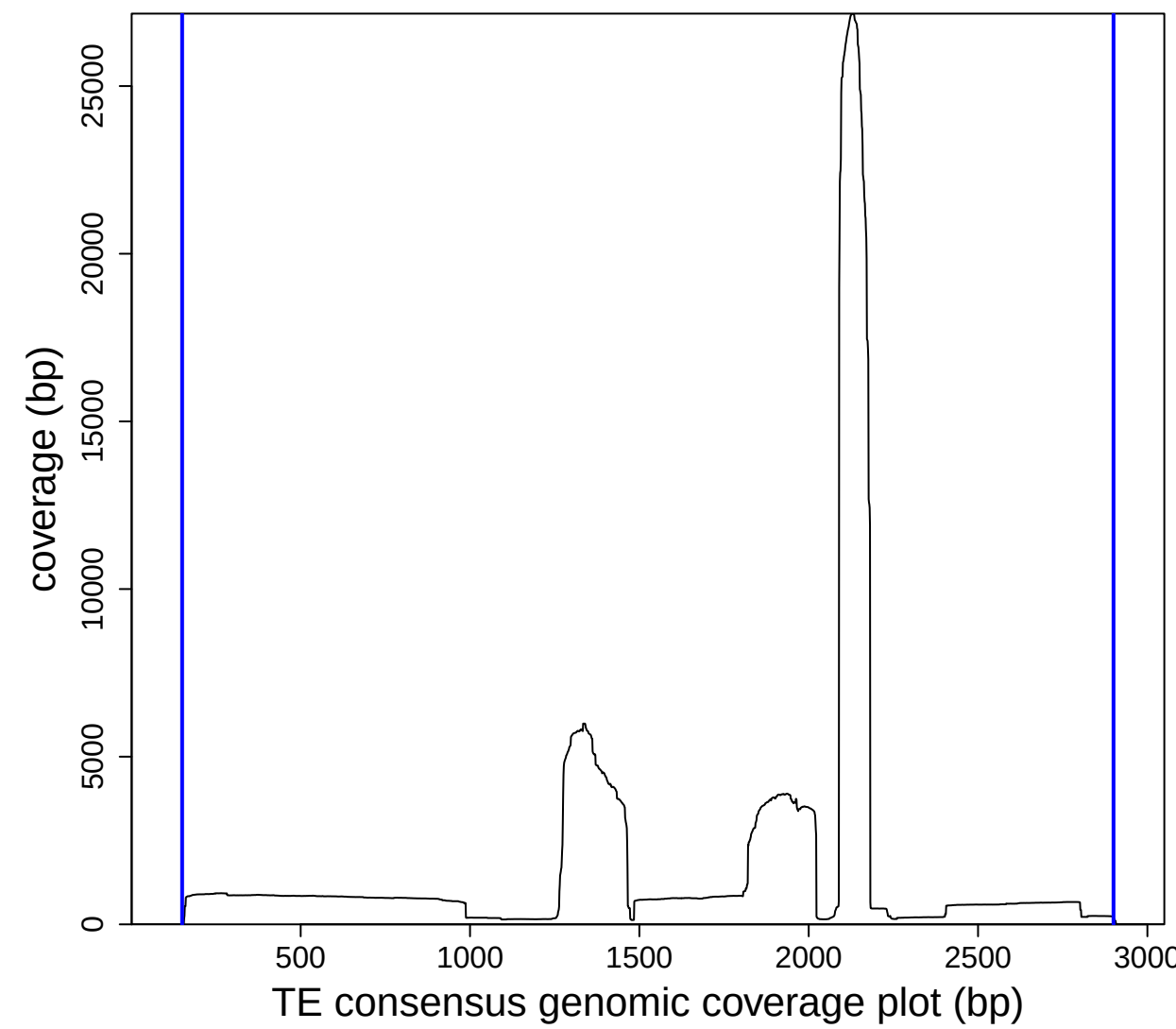
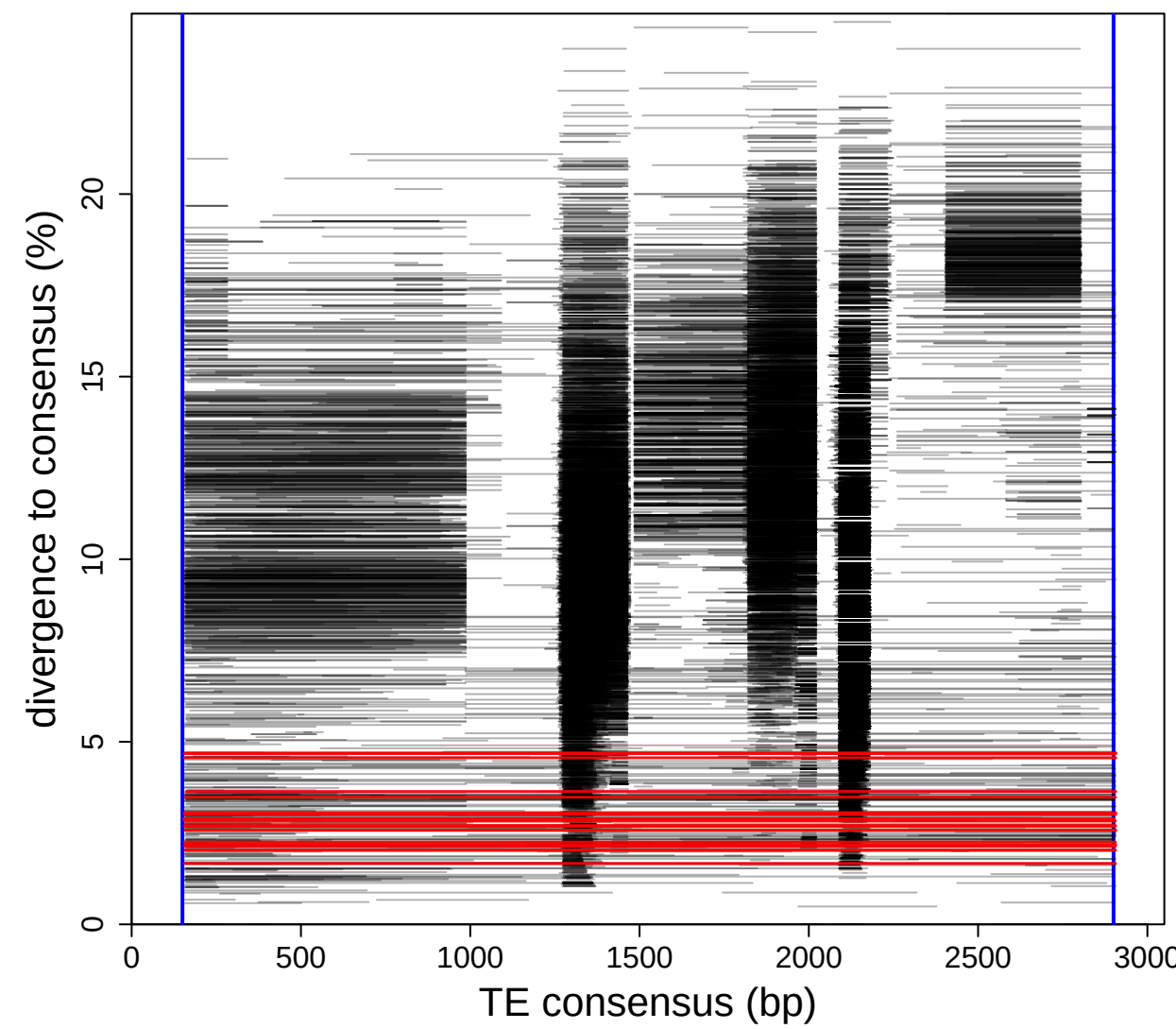
l.b.bed uf.bed g_2.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 2741bp; fragments: 41008; full length: 45 (>=2466.9bp)

After TEtrimmer 2741 bp



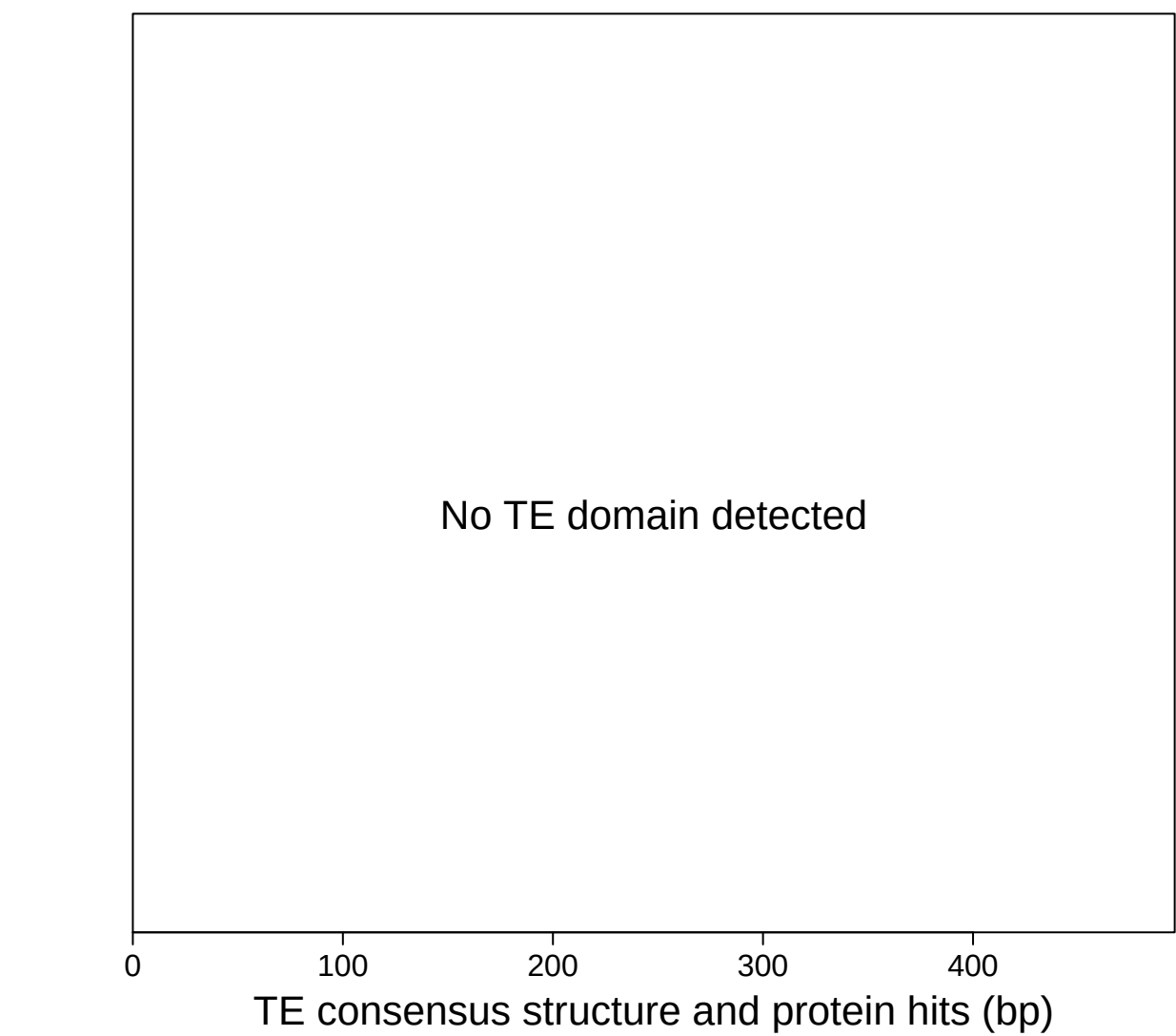
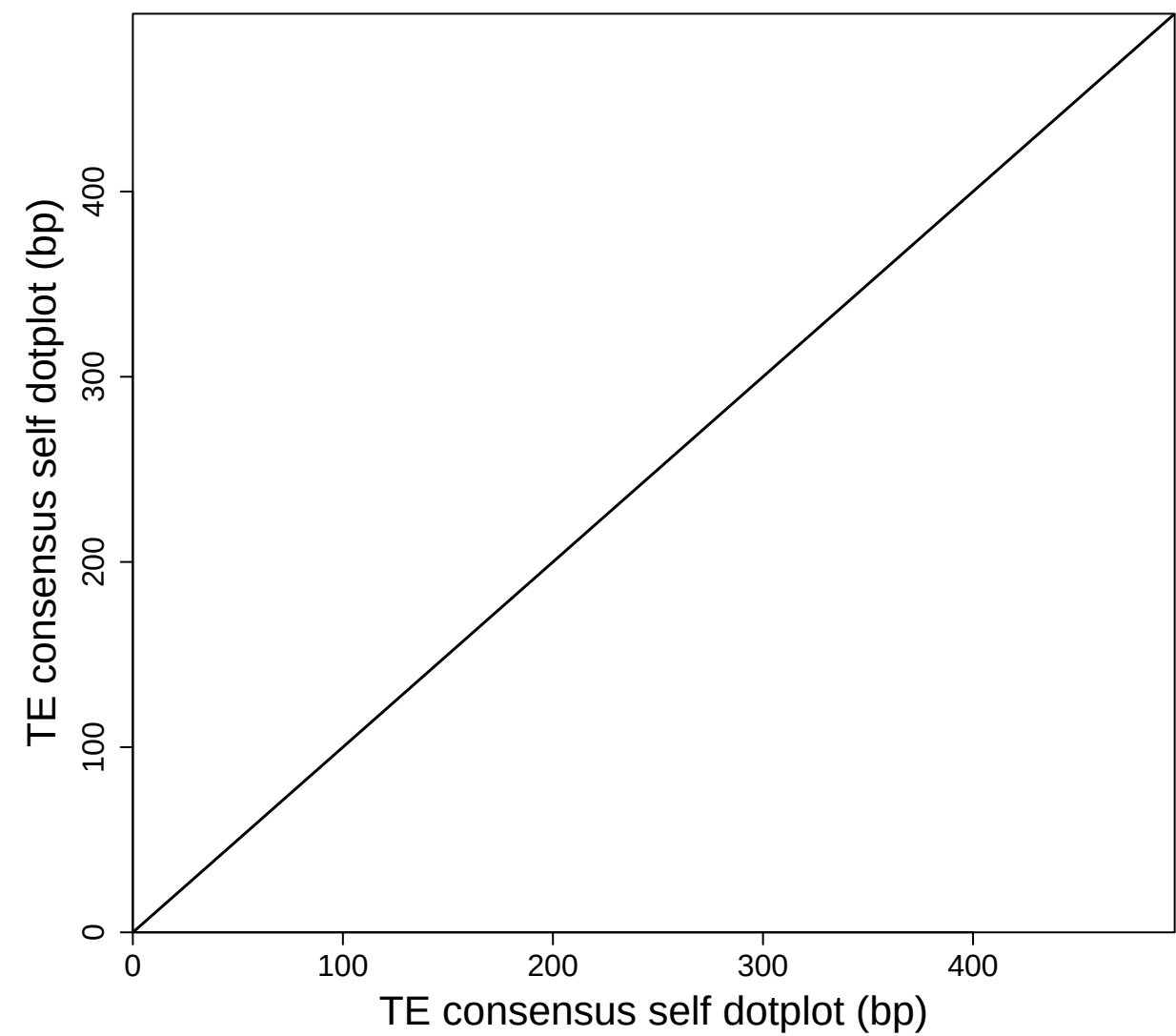
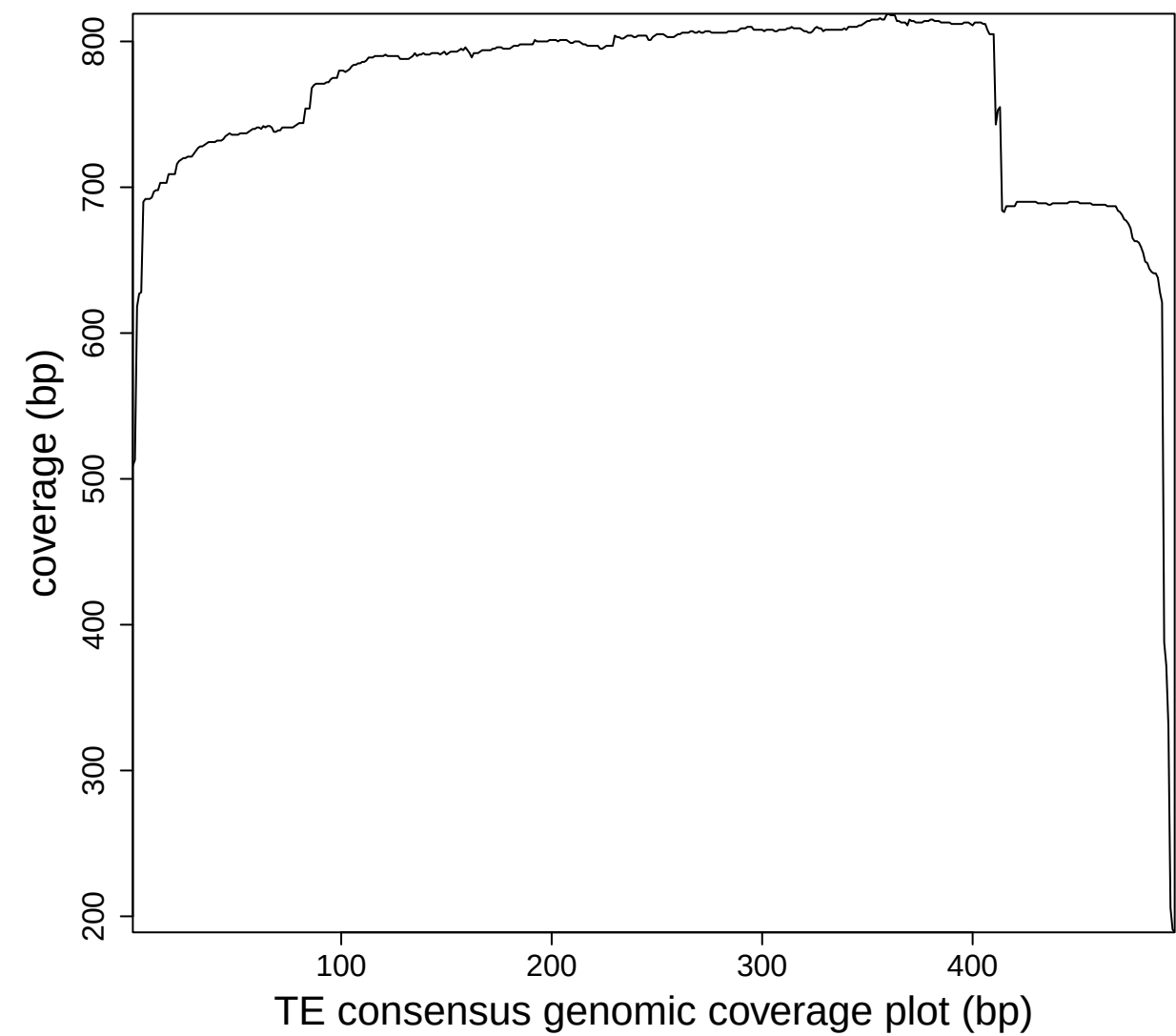
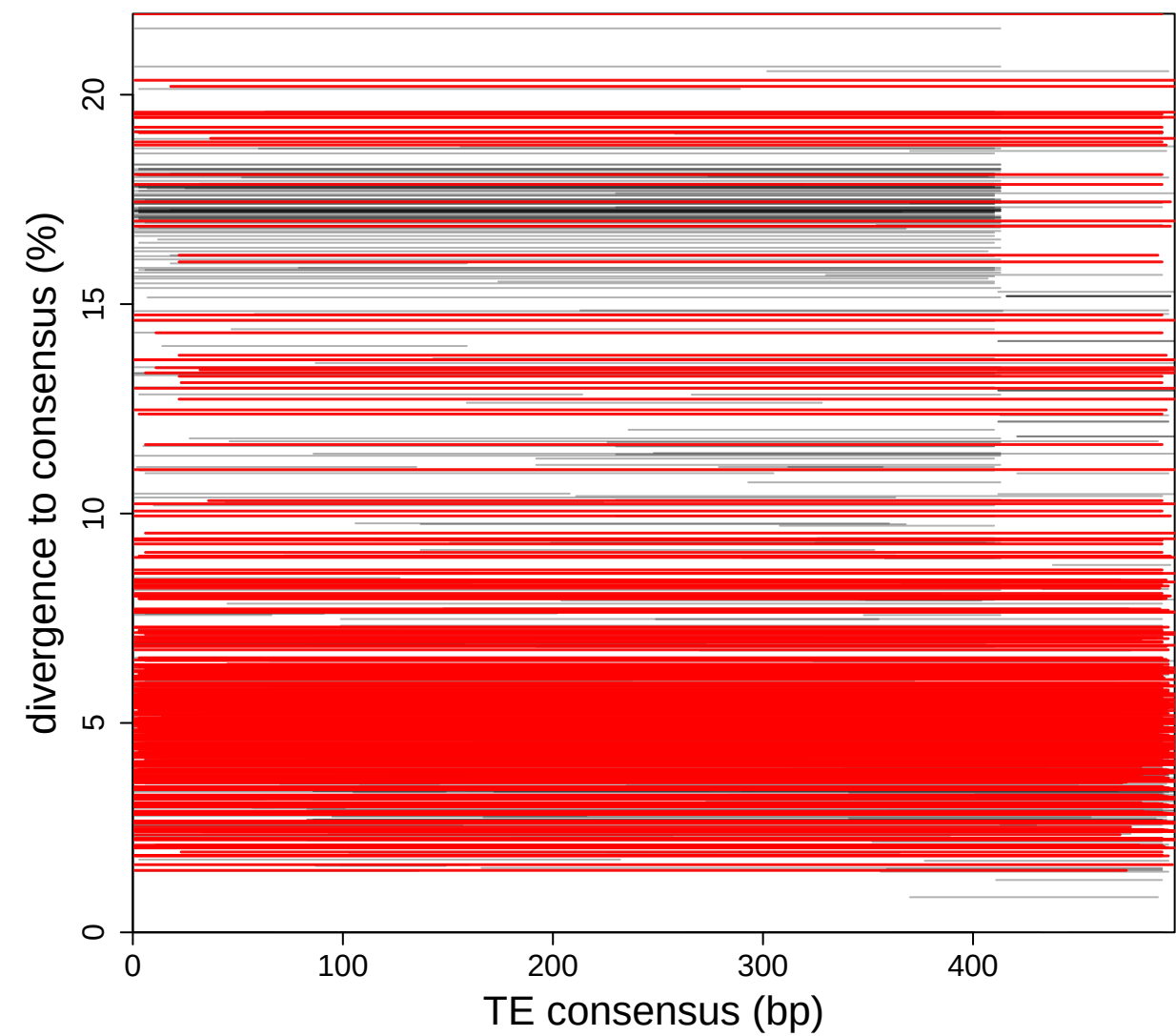
1.fasta.b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 3050bp; fragments: 41003; full length: 19 (>=2745bp)

After TEtrimmer Extended plot Blue lines are boundaries

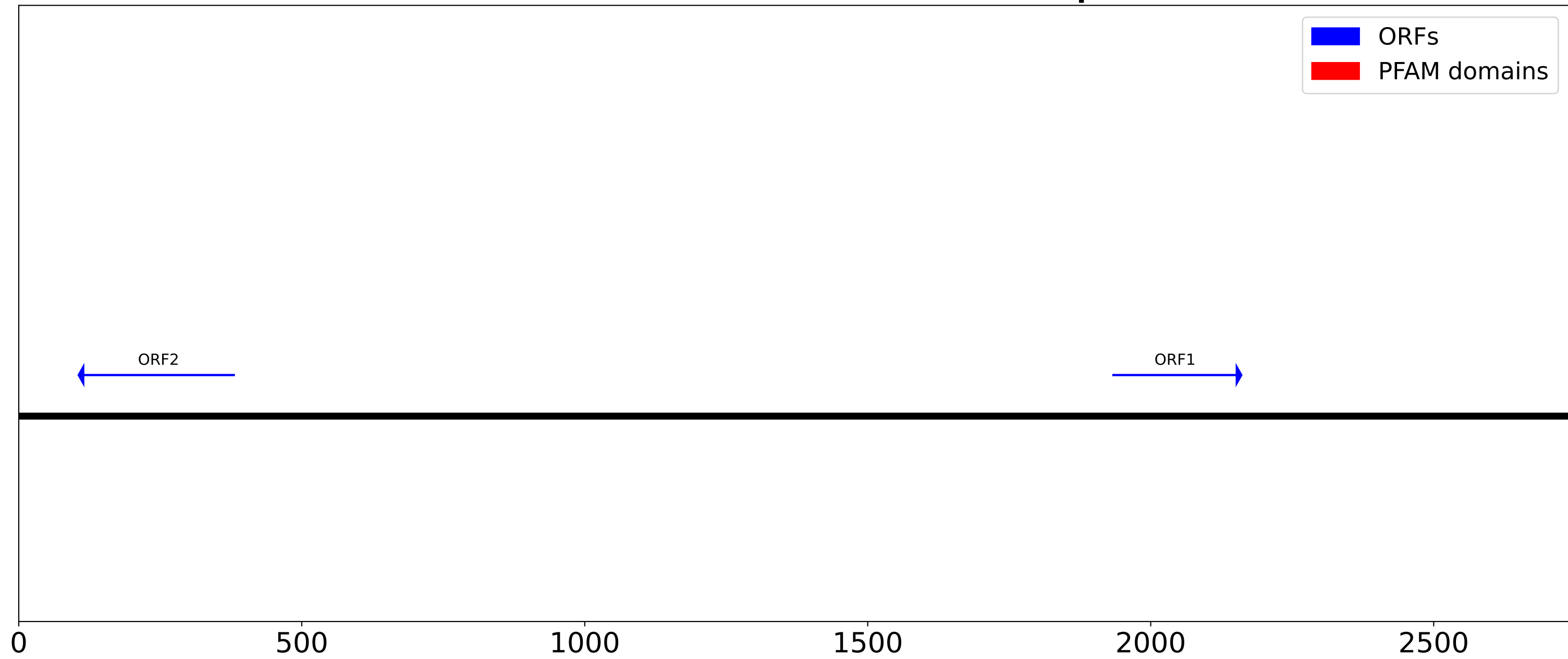


TE: rnd_1_family_471
size: 496bp; fragments: 933; full length: 548 (≥ 446.4 bp)

Before TEtrimmer 496 bp



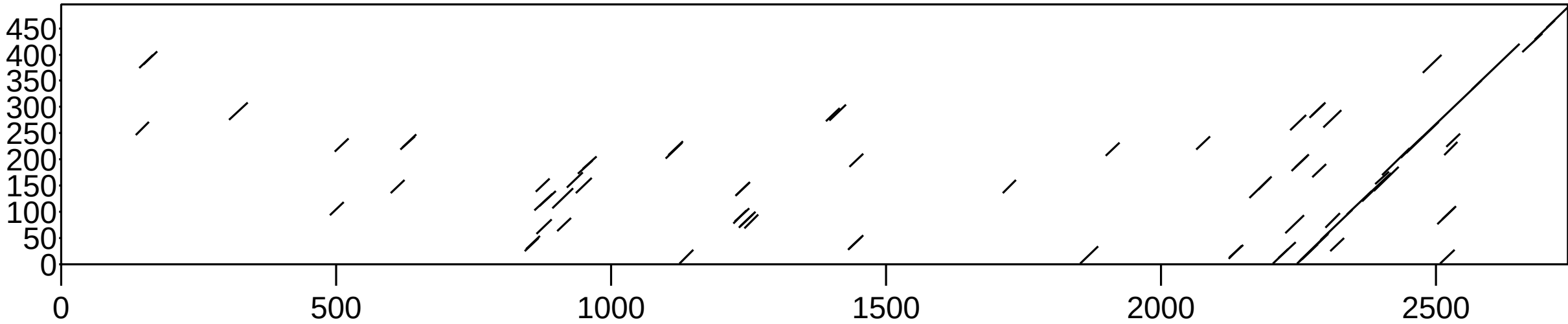
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)