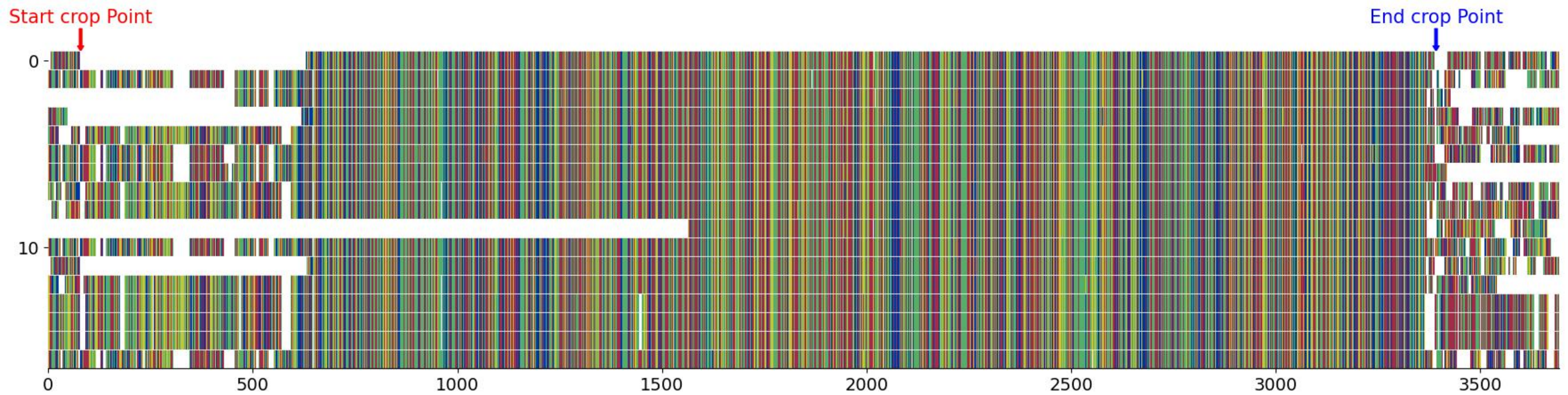
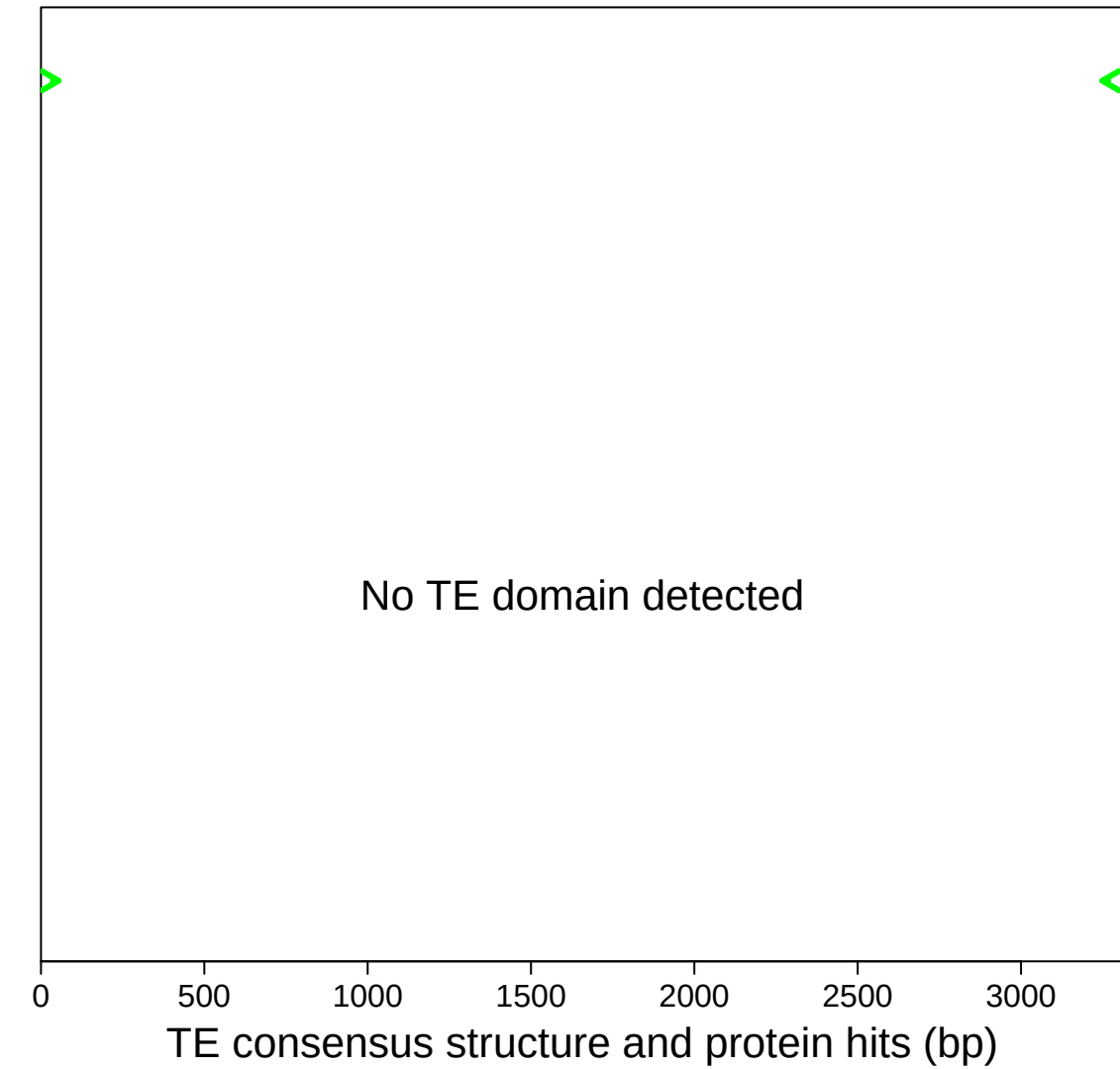
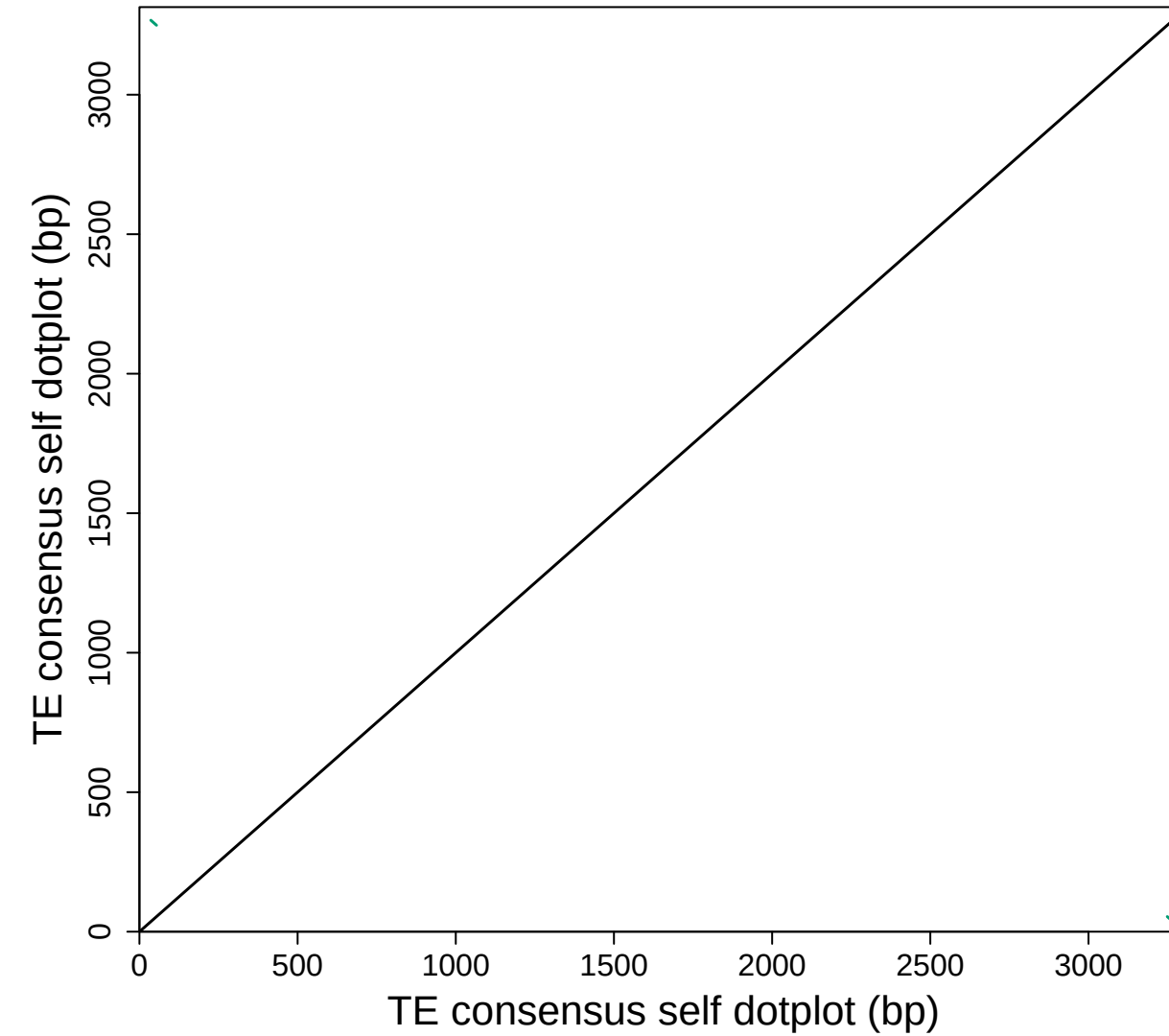
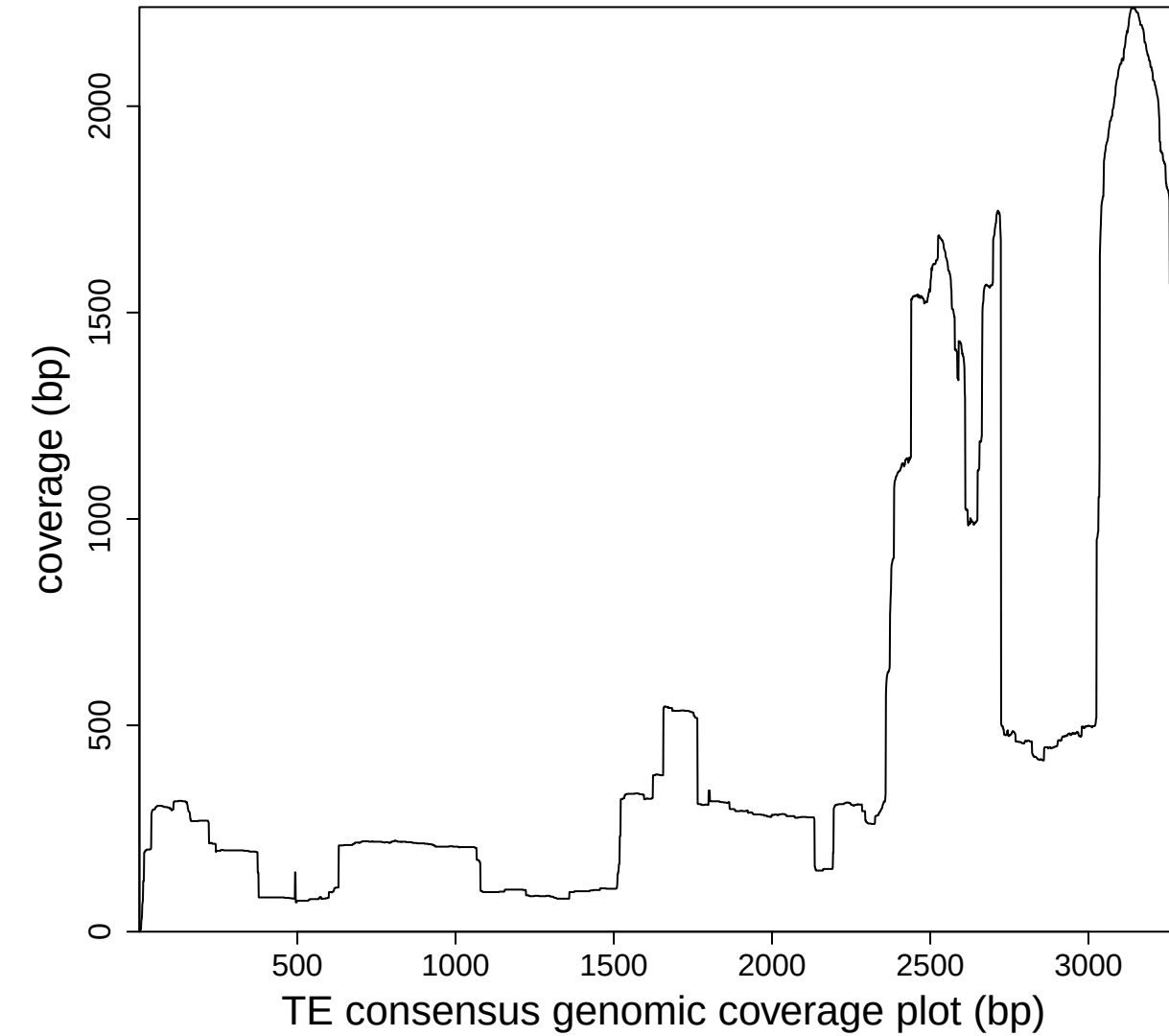
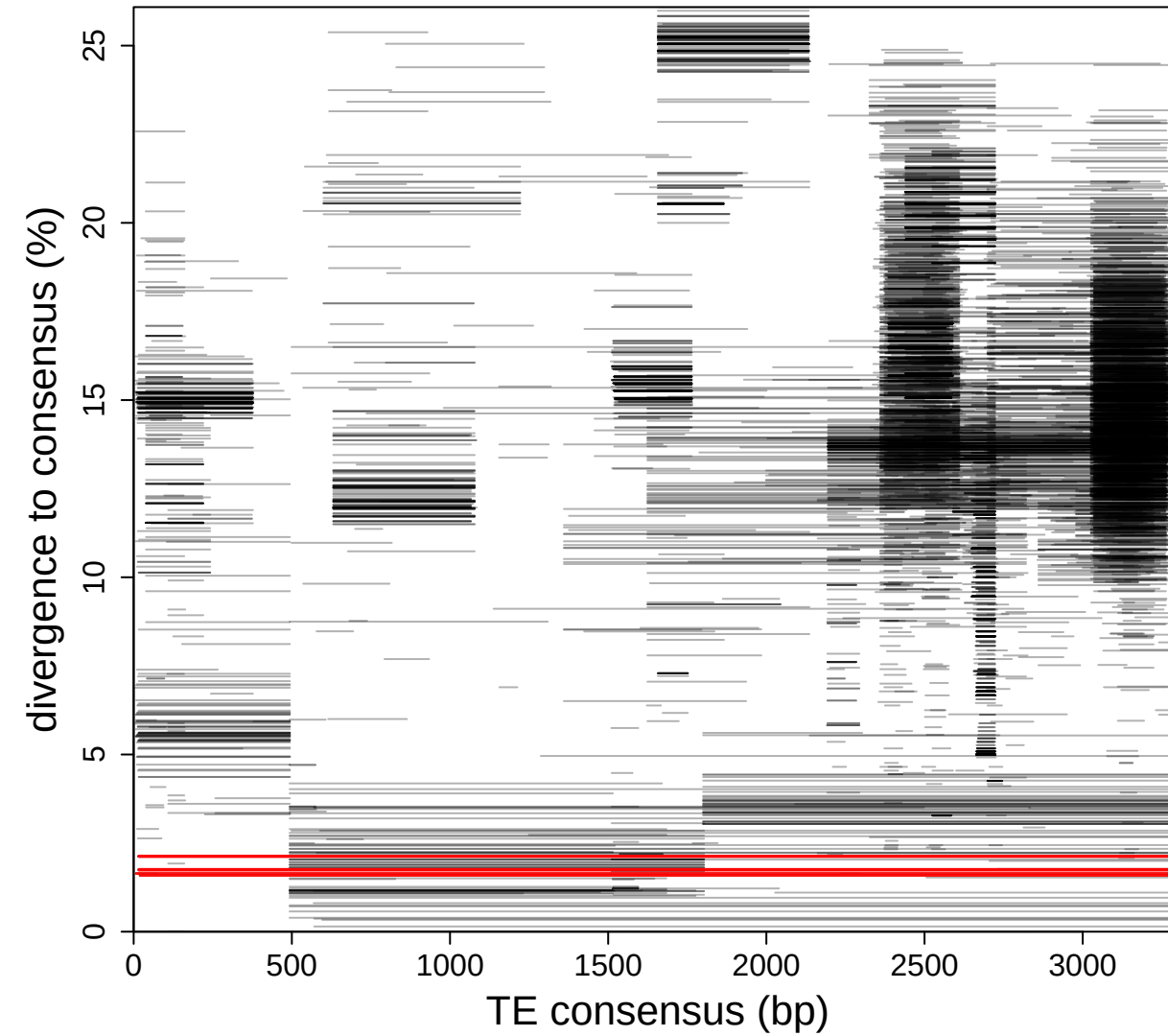




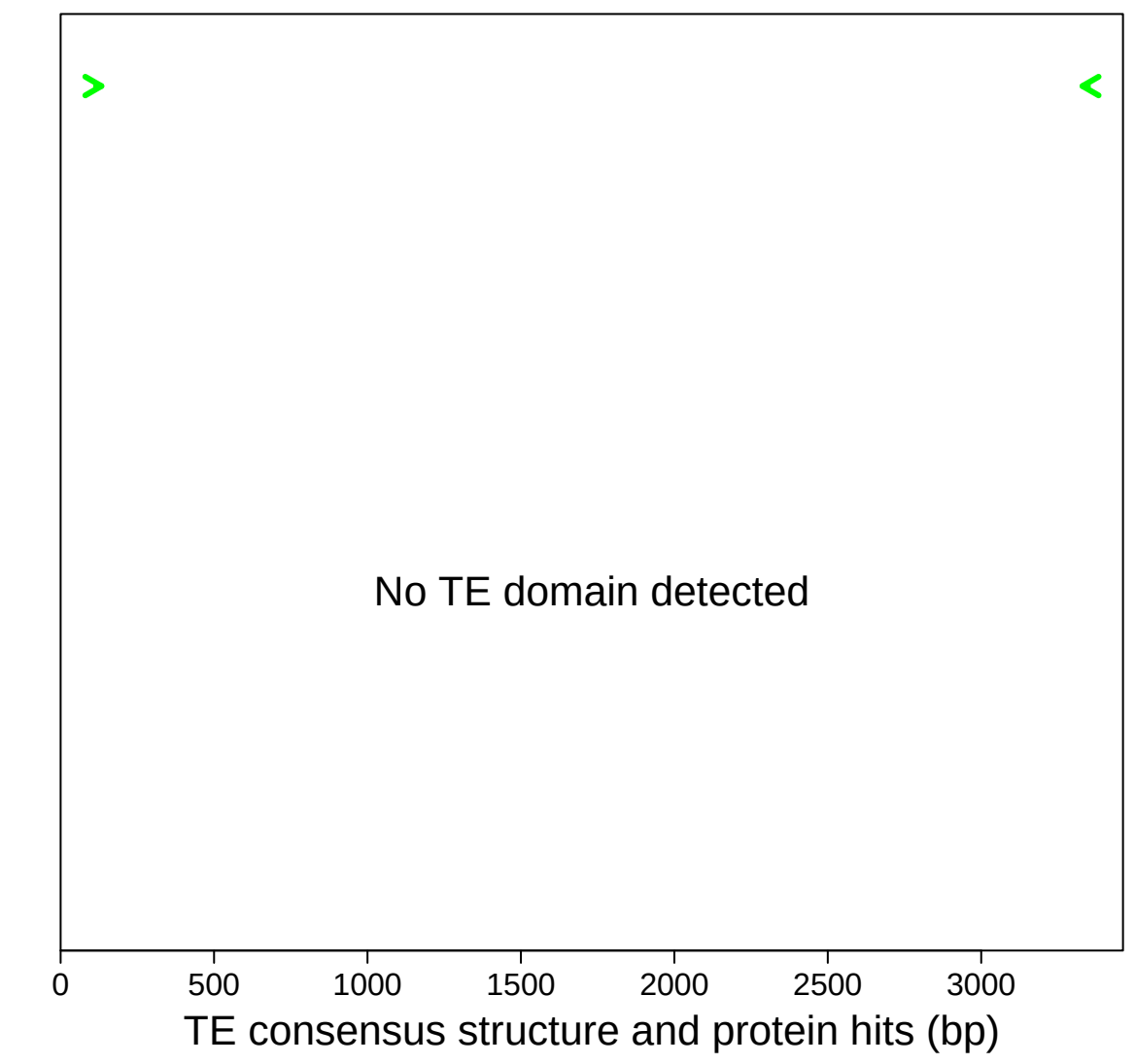
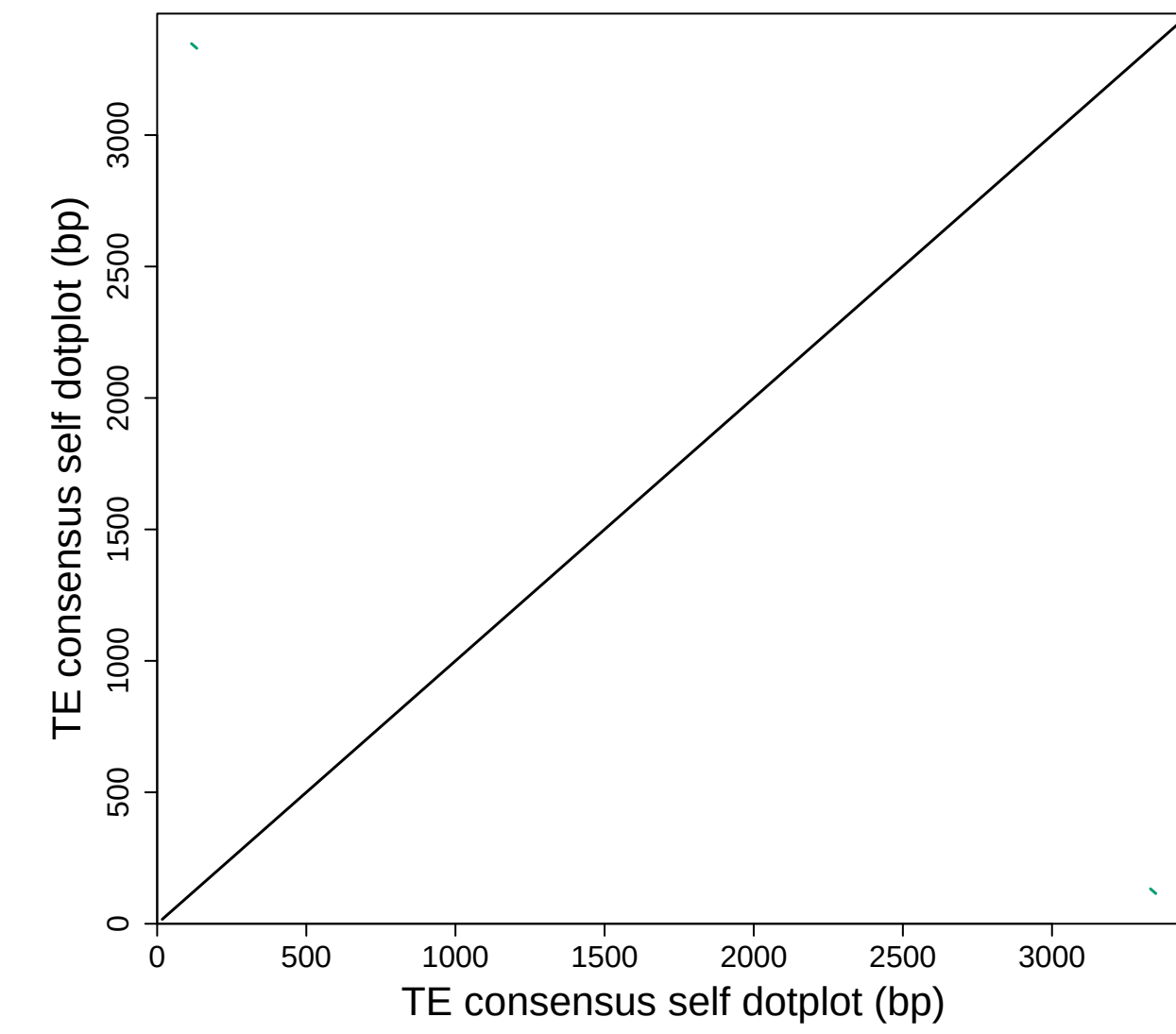
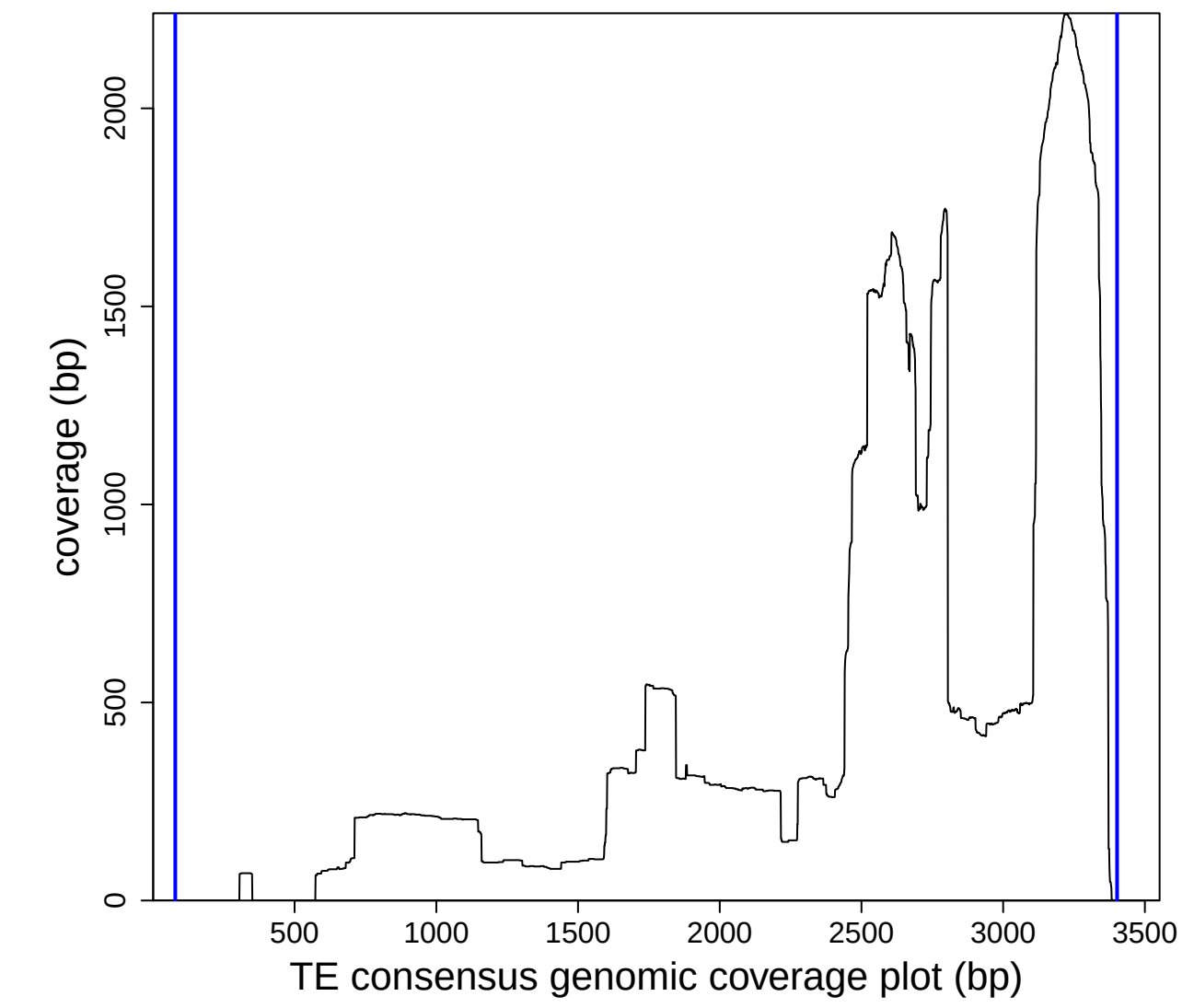
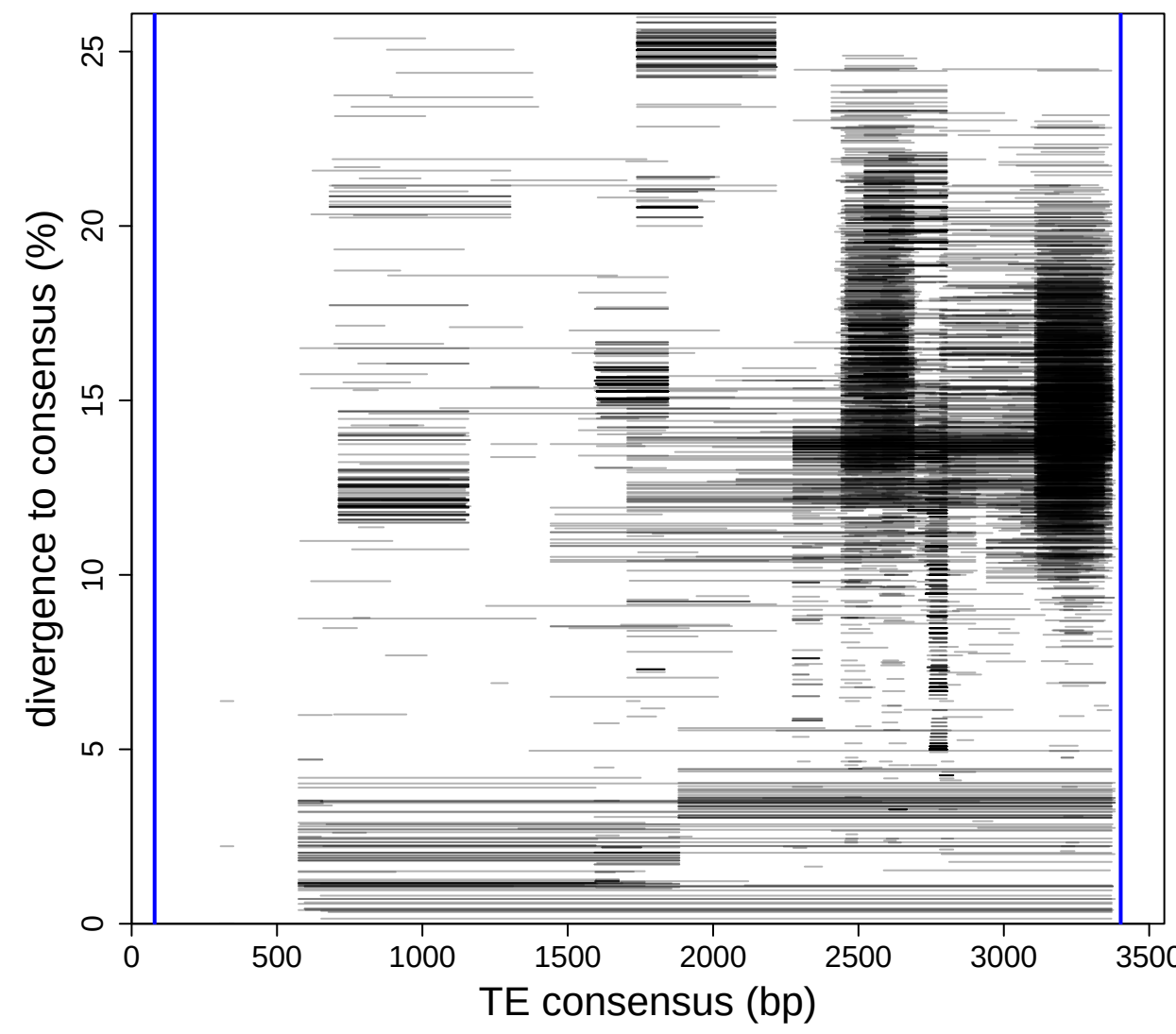
l.b.bed uf.bed g_3.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 3314bp; fragments: 5959; full length: 7 (≥ 2982.6 bp)

After TEtrimmer 3314 bp



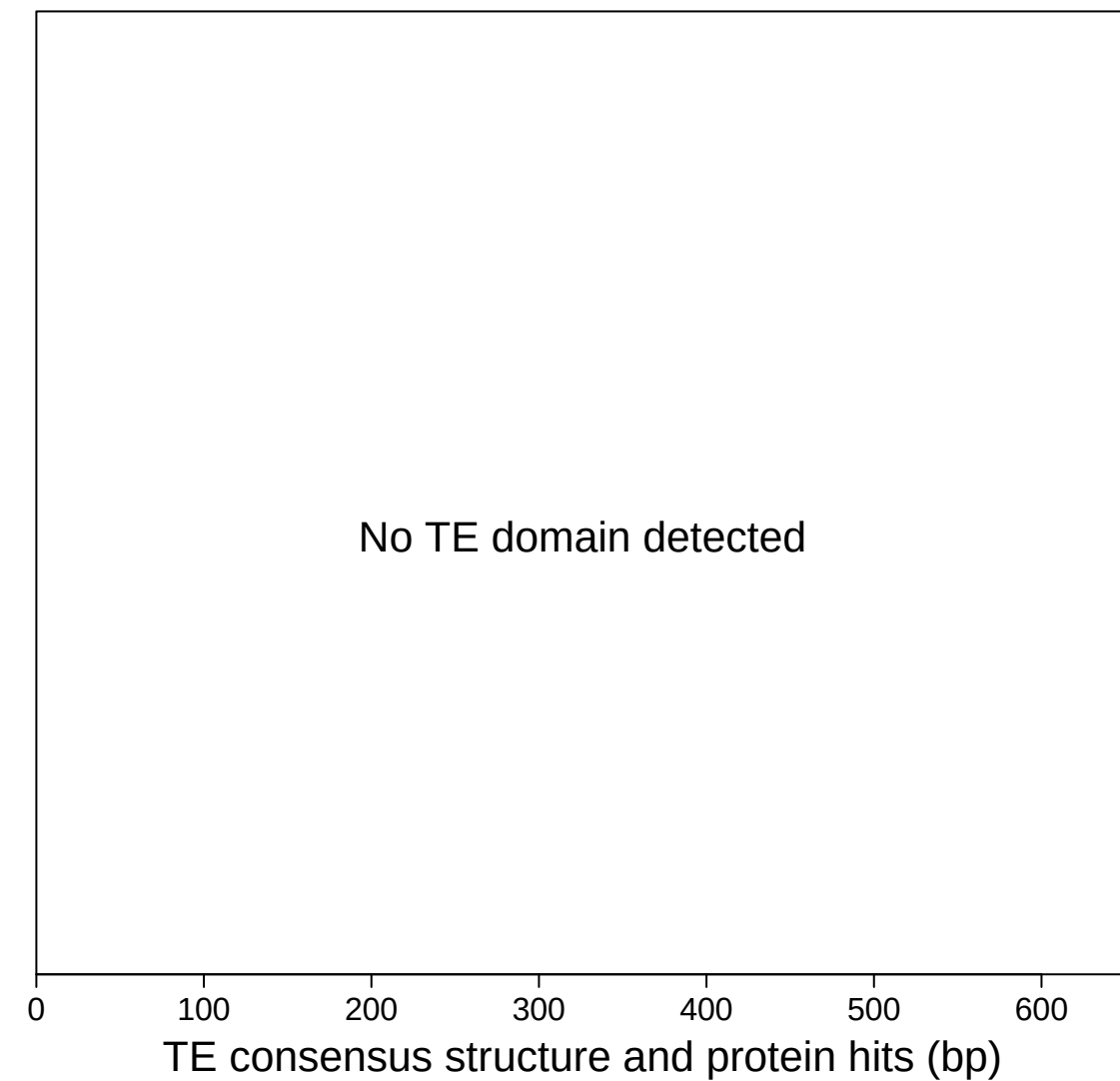
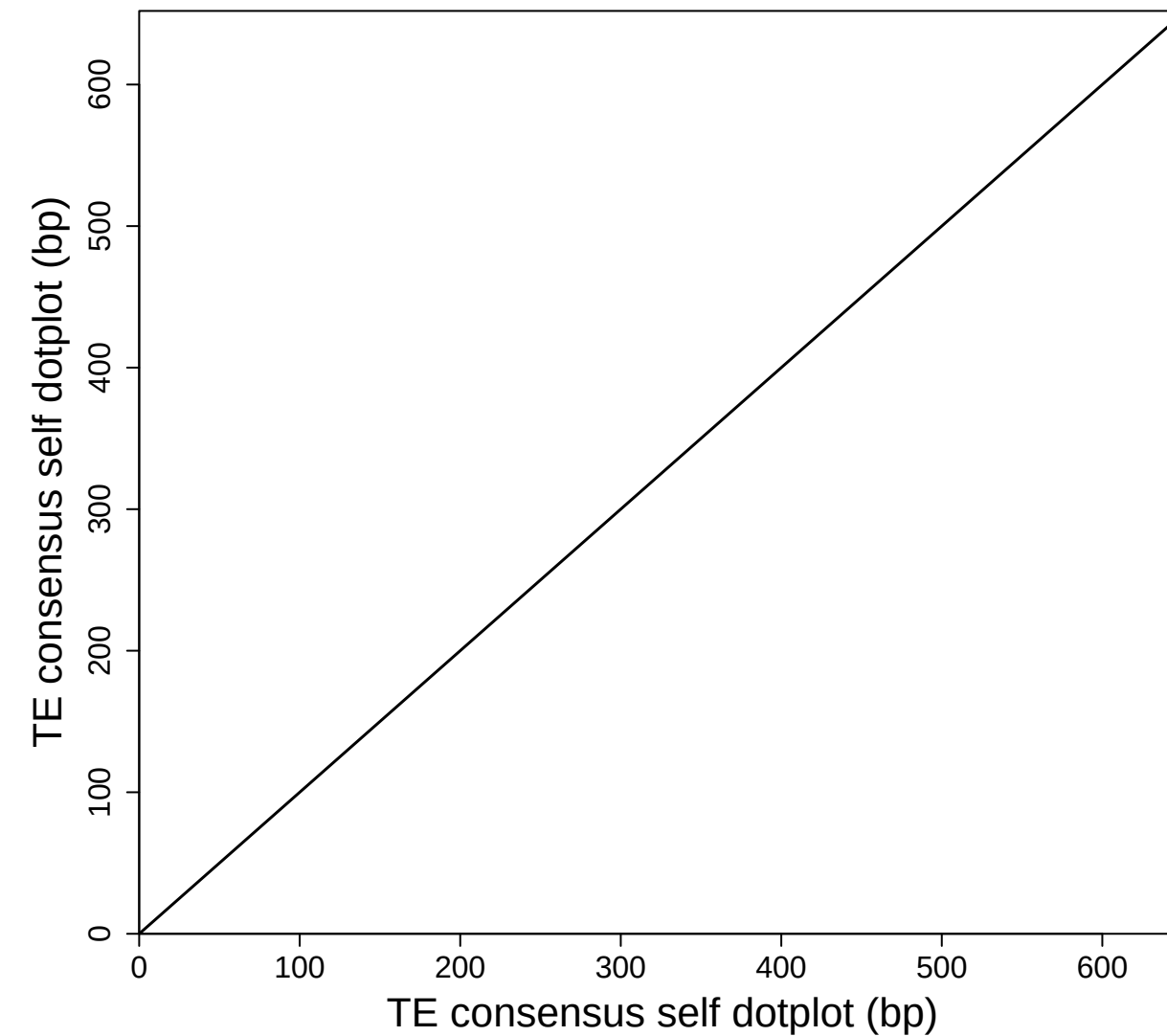
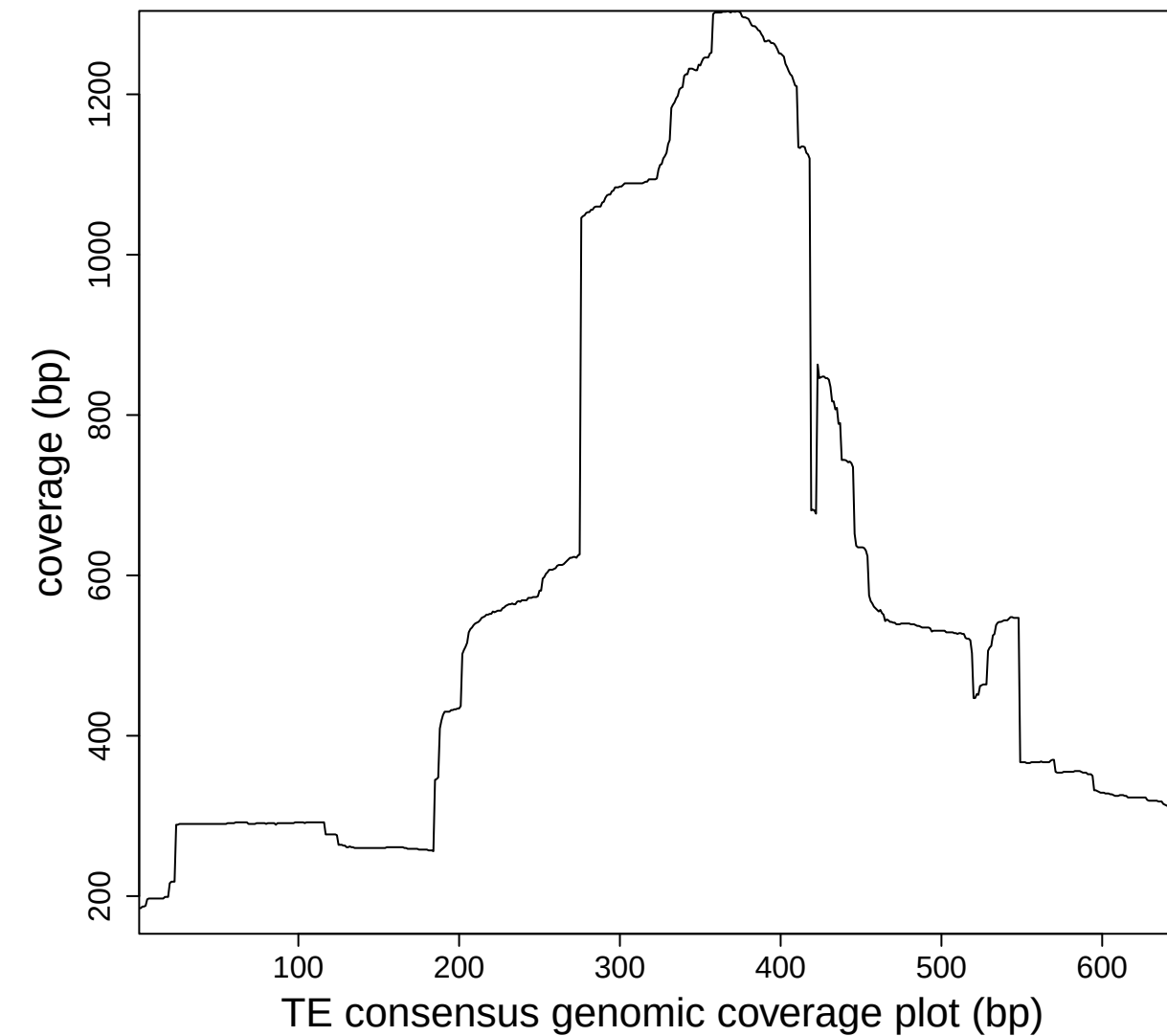
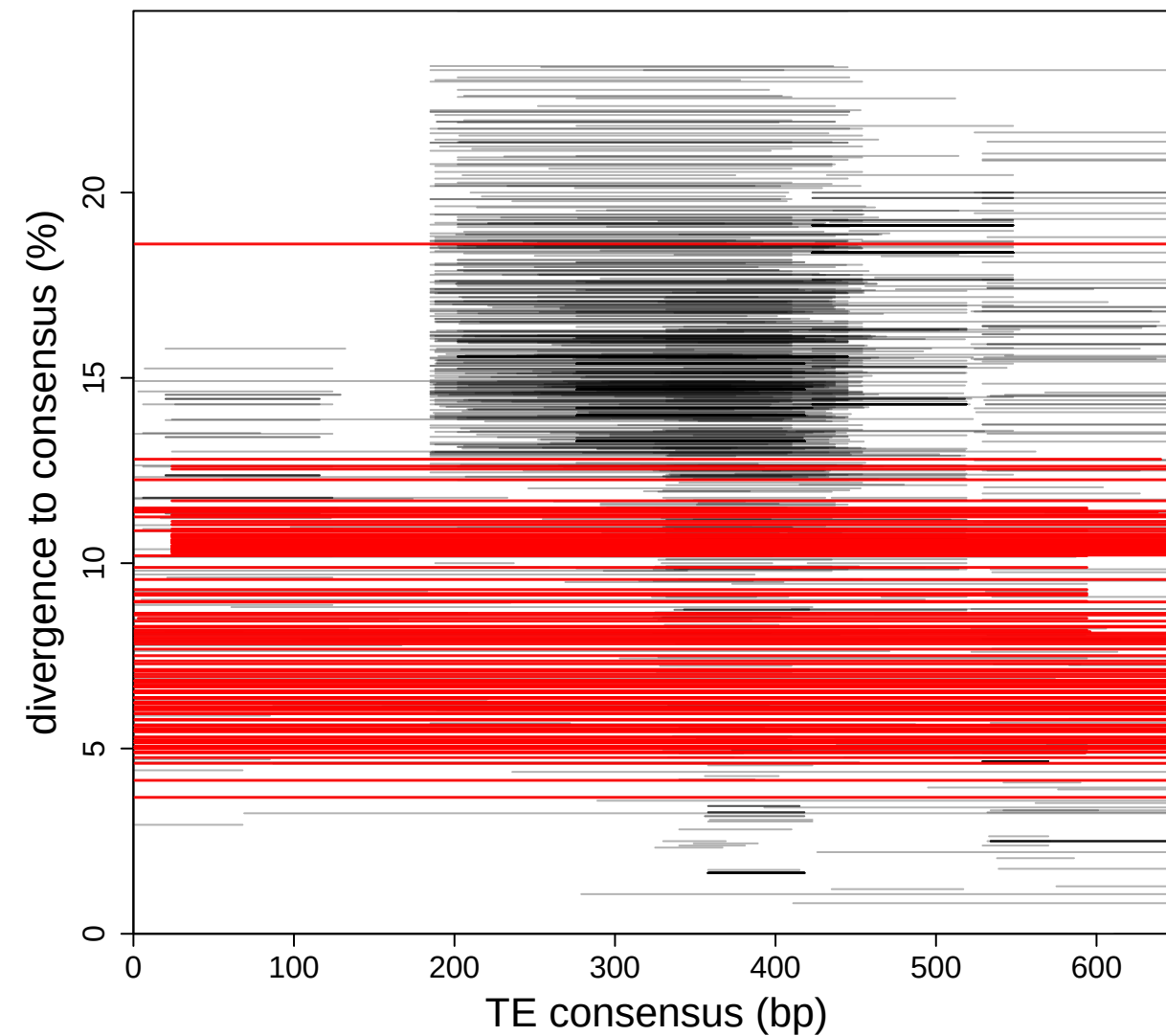
0.fasta.b.bed uf.bed g 3.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 3552bp; fragments: 5695; full length: 0 (>=3196.8bp)

After TEtrimmer Extended plot Blue lines are boundaries

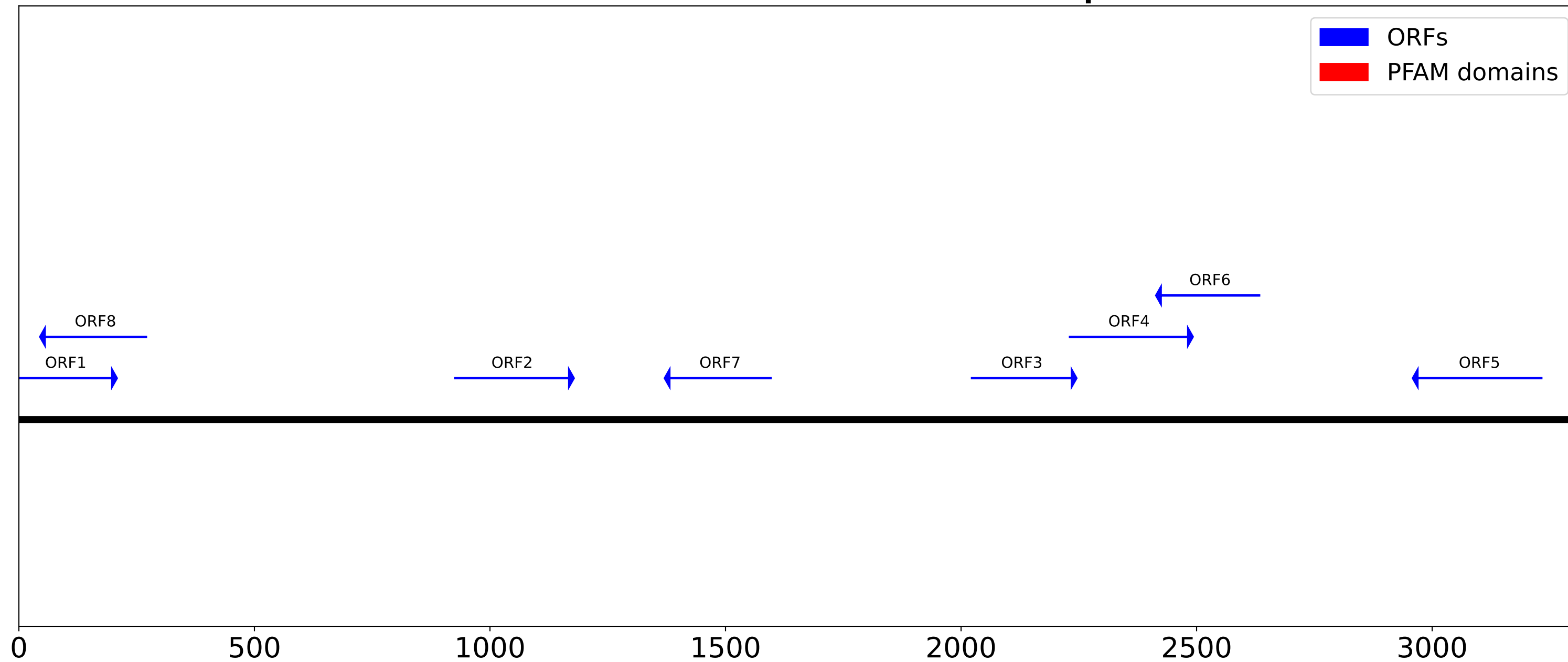


TE: rnd_1_family_920
size: 652bp; fragments: 1716; full length: 222 (≥ 586.8 bp)

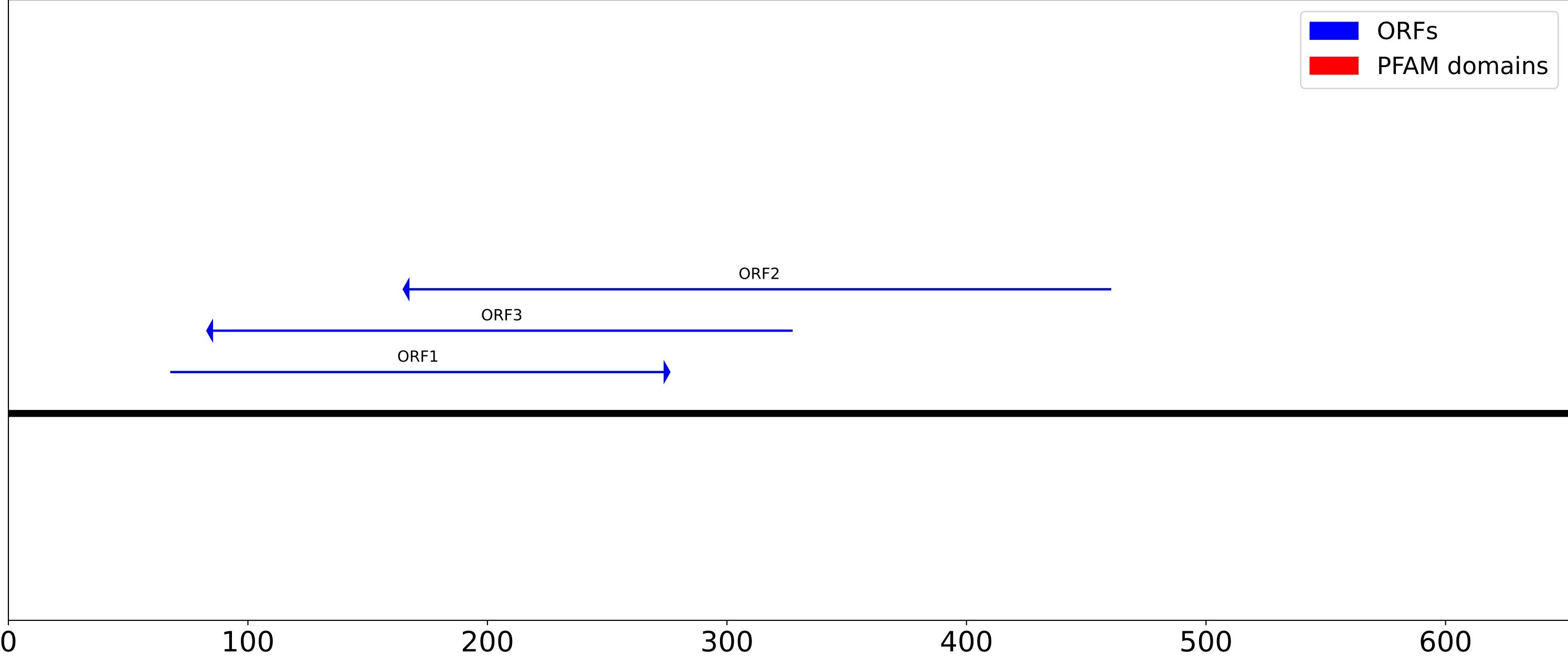
Before TEtrimmer 652 bp



After TEtrimmer ORF and PFAM domain plot



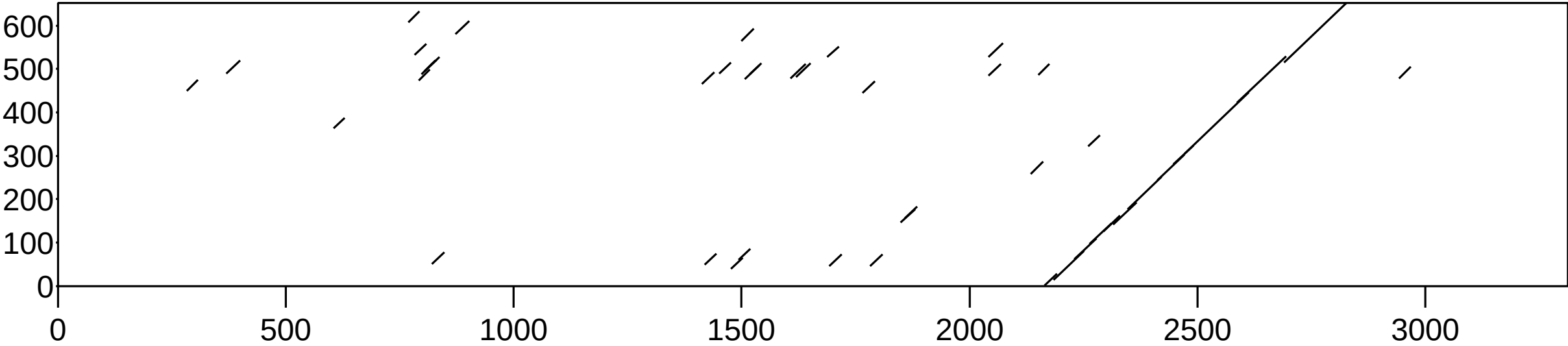
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)