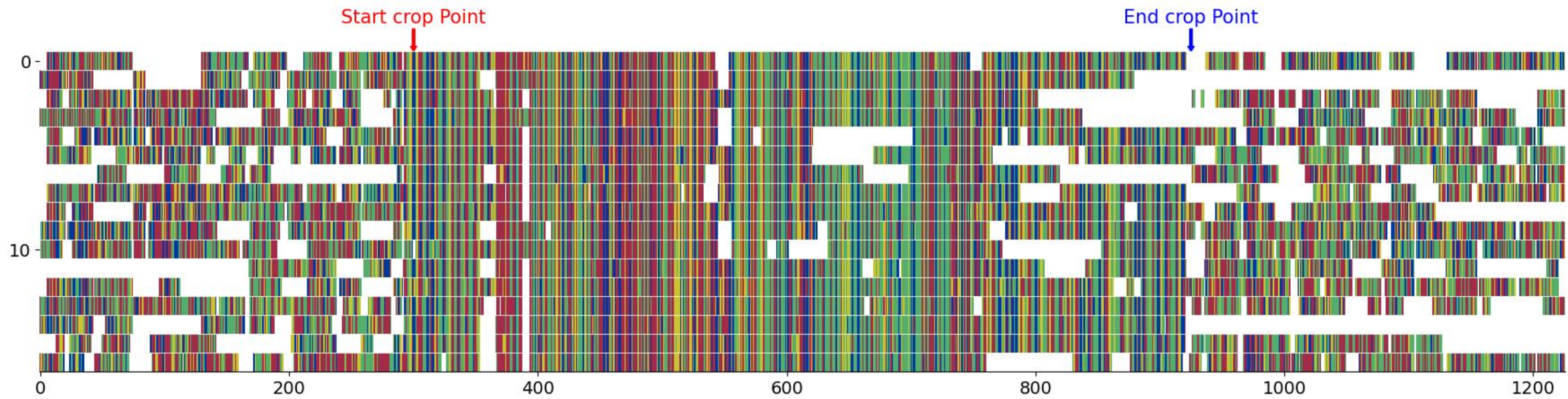
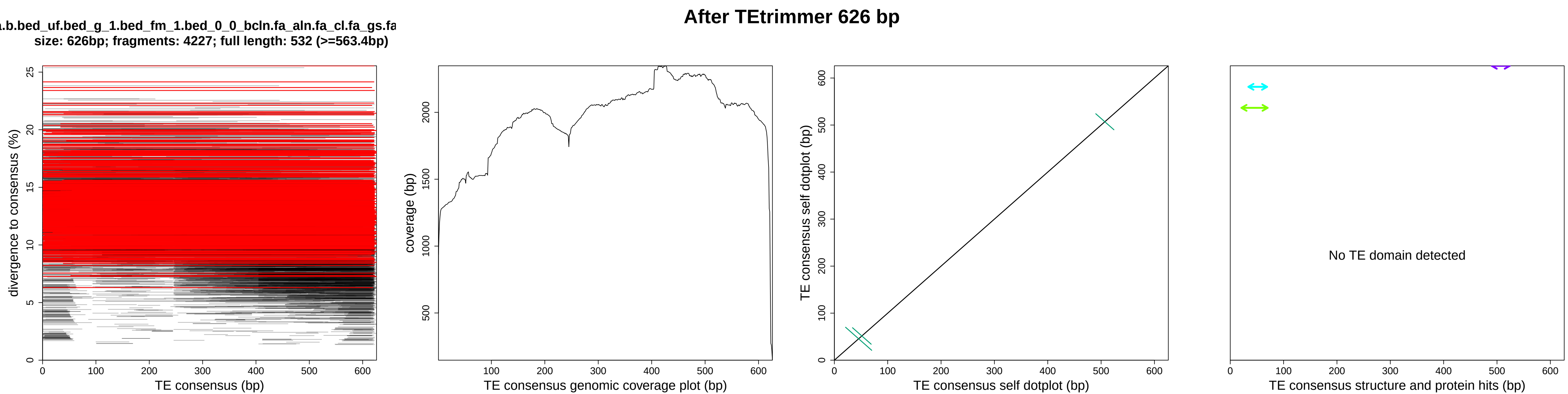
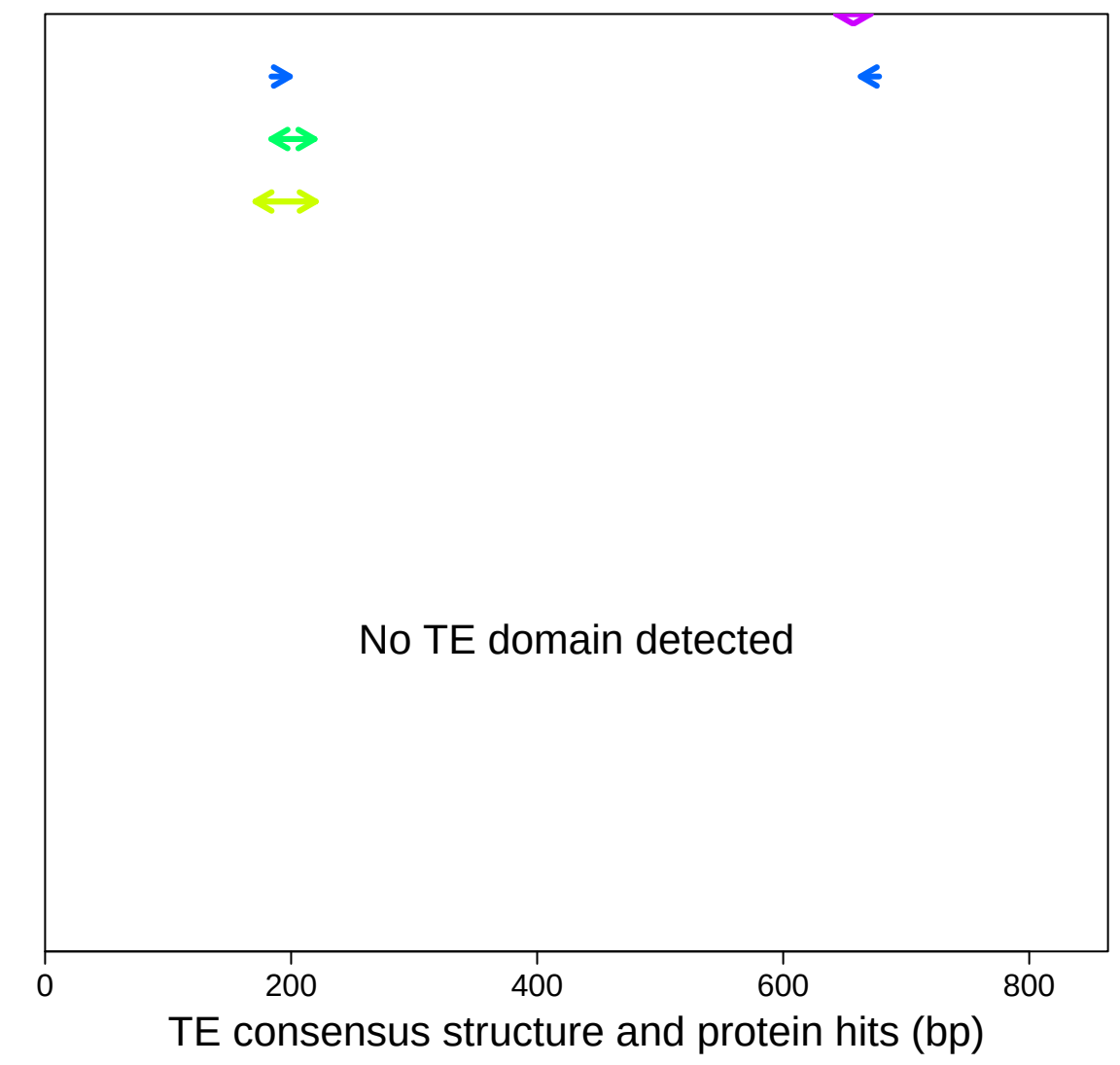
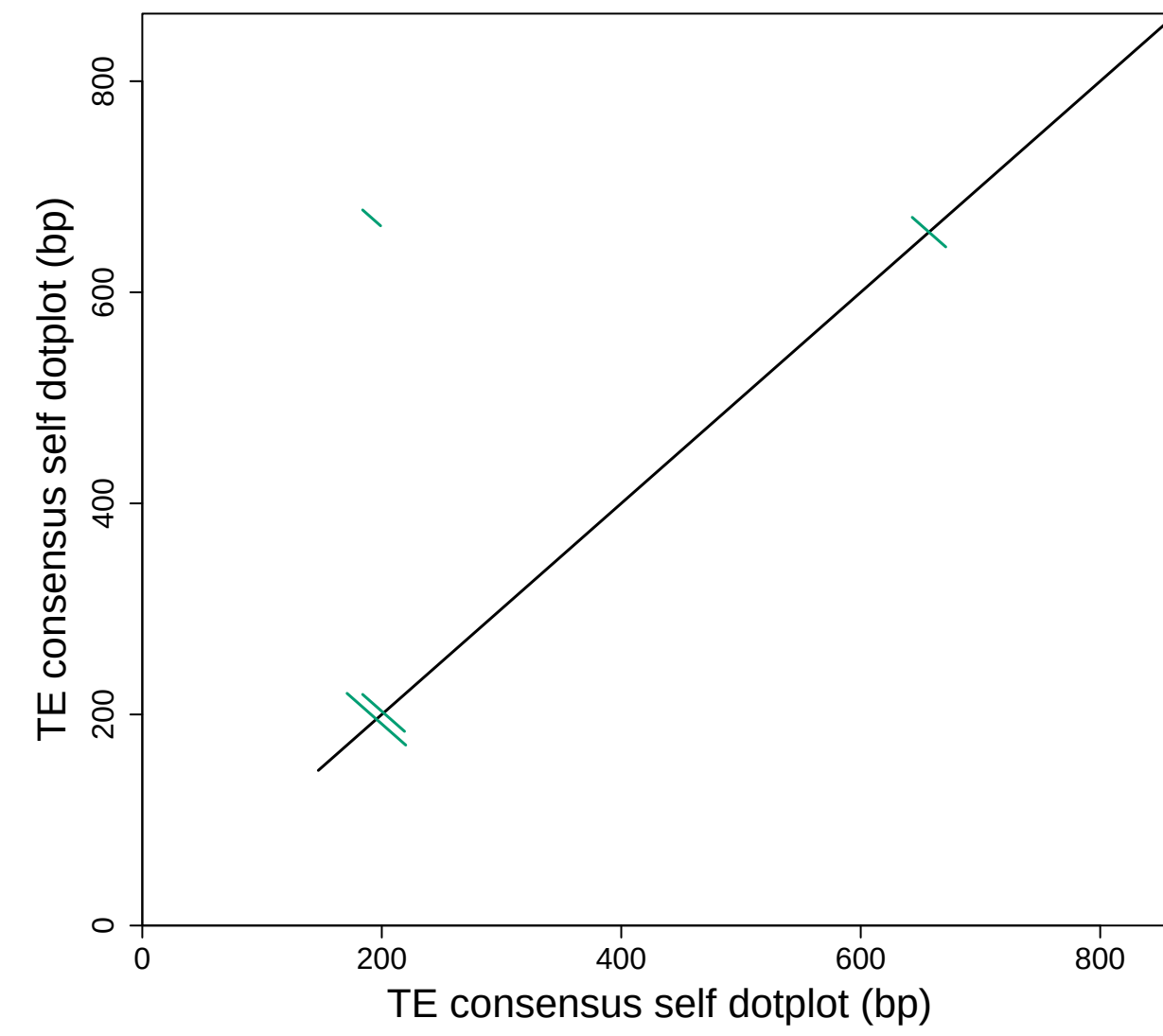
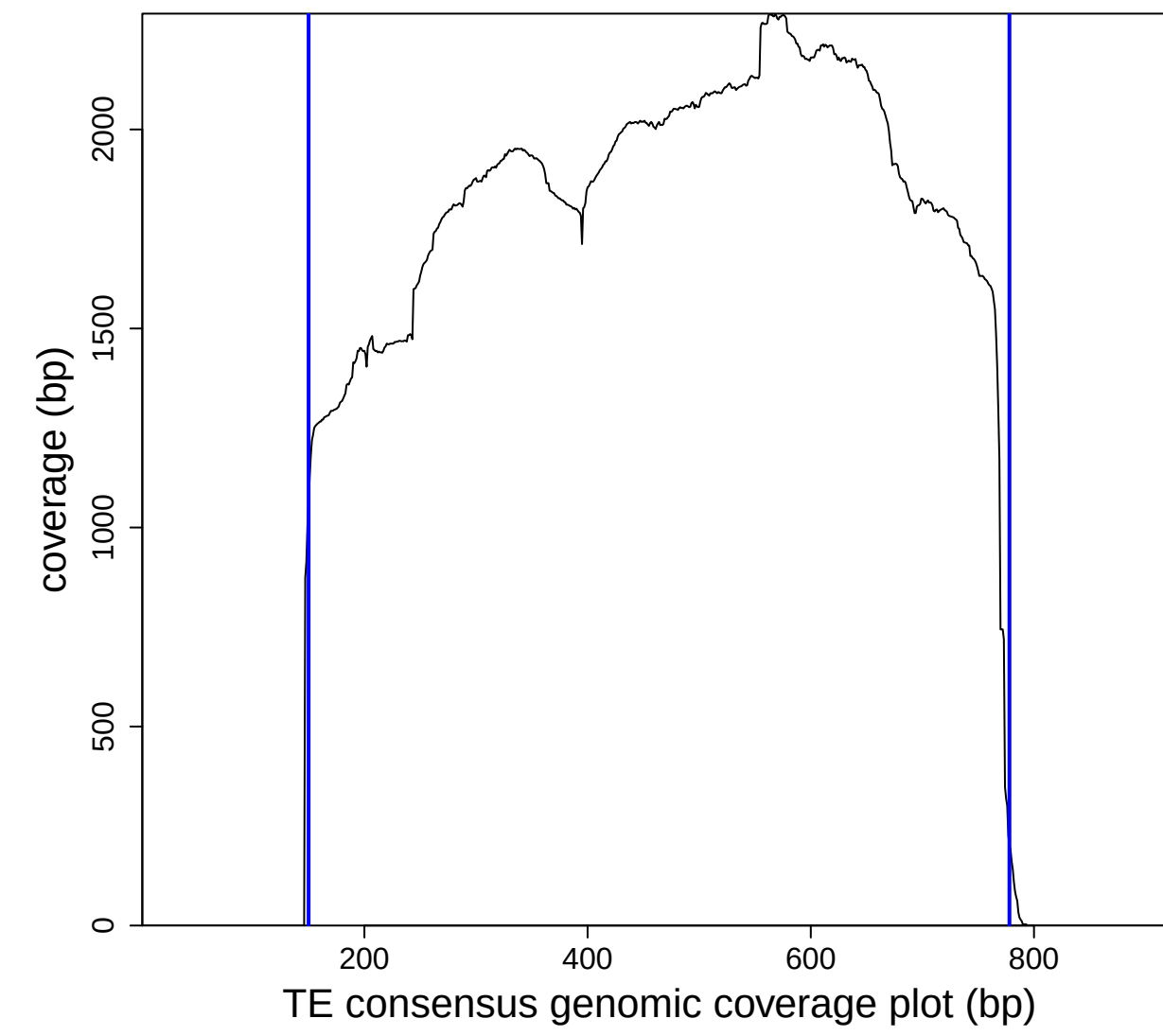
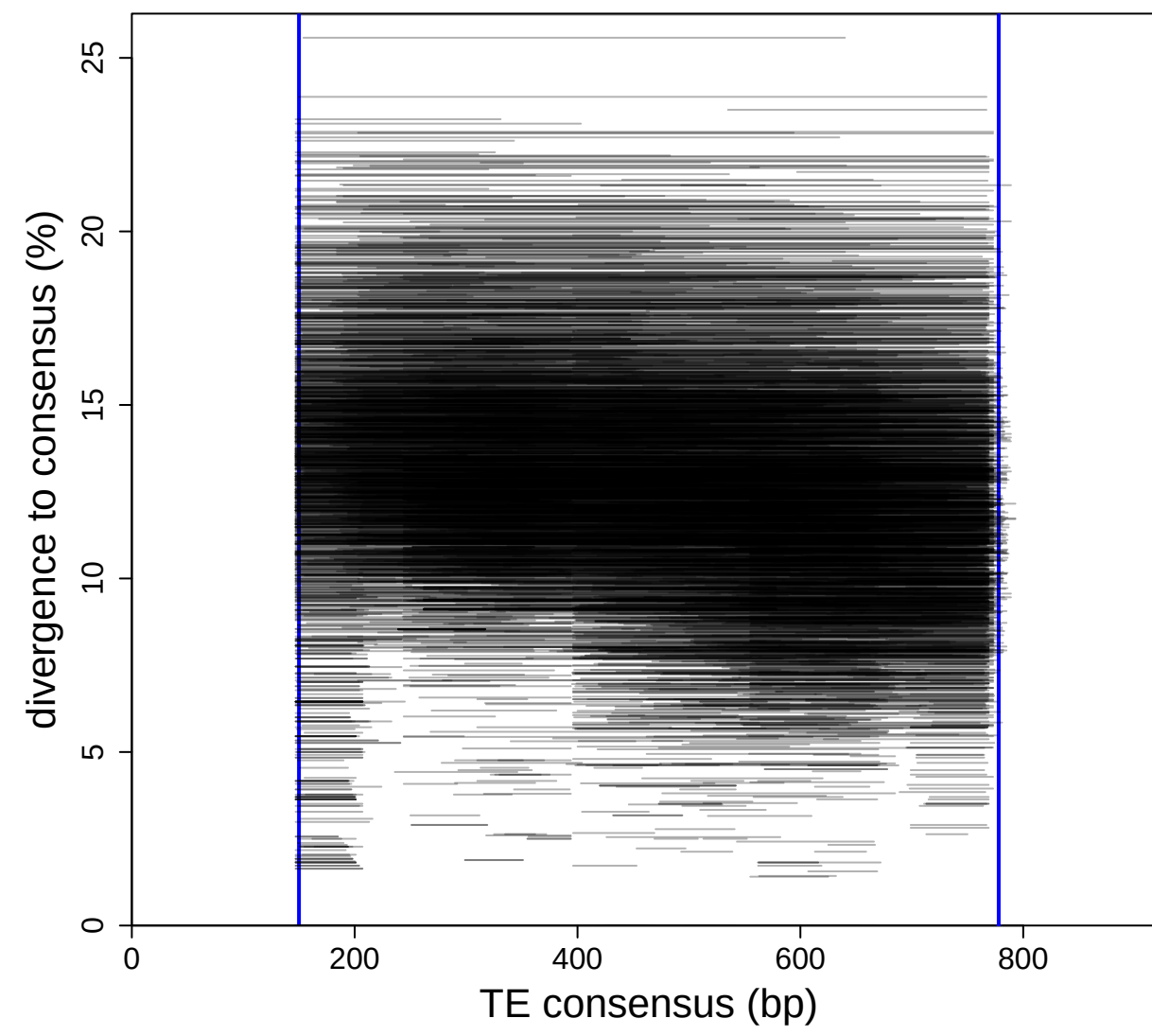




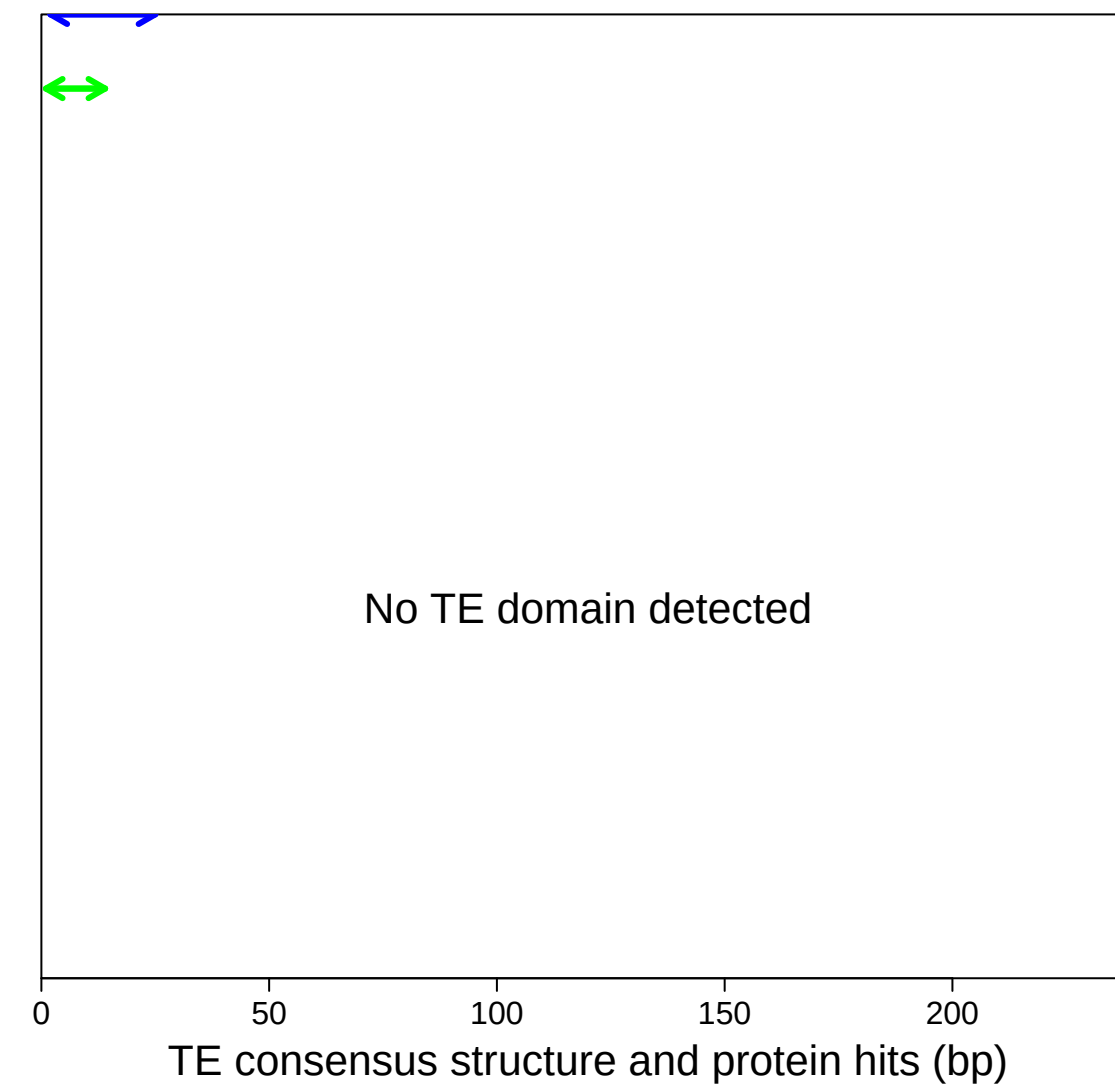
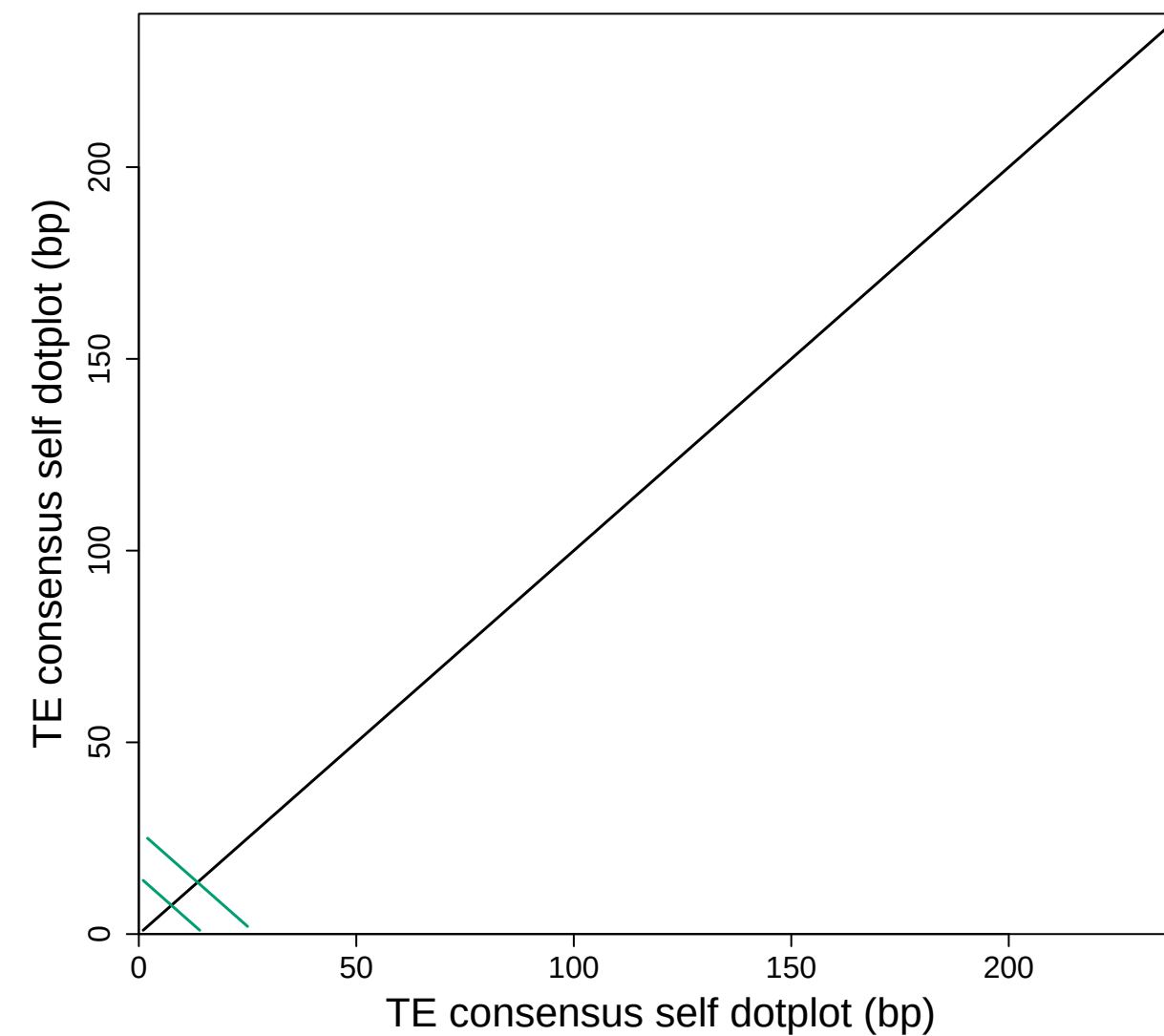
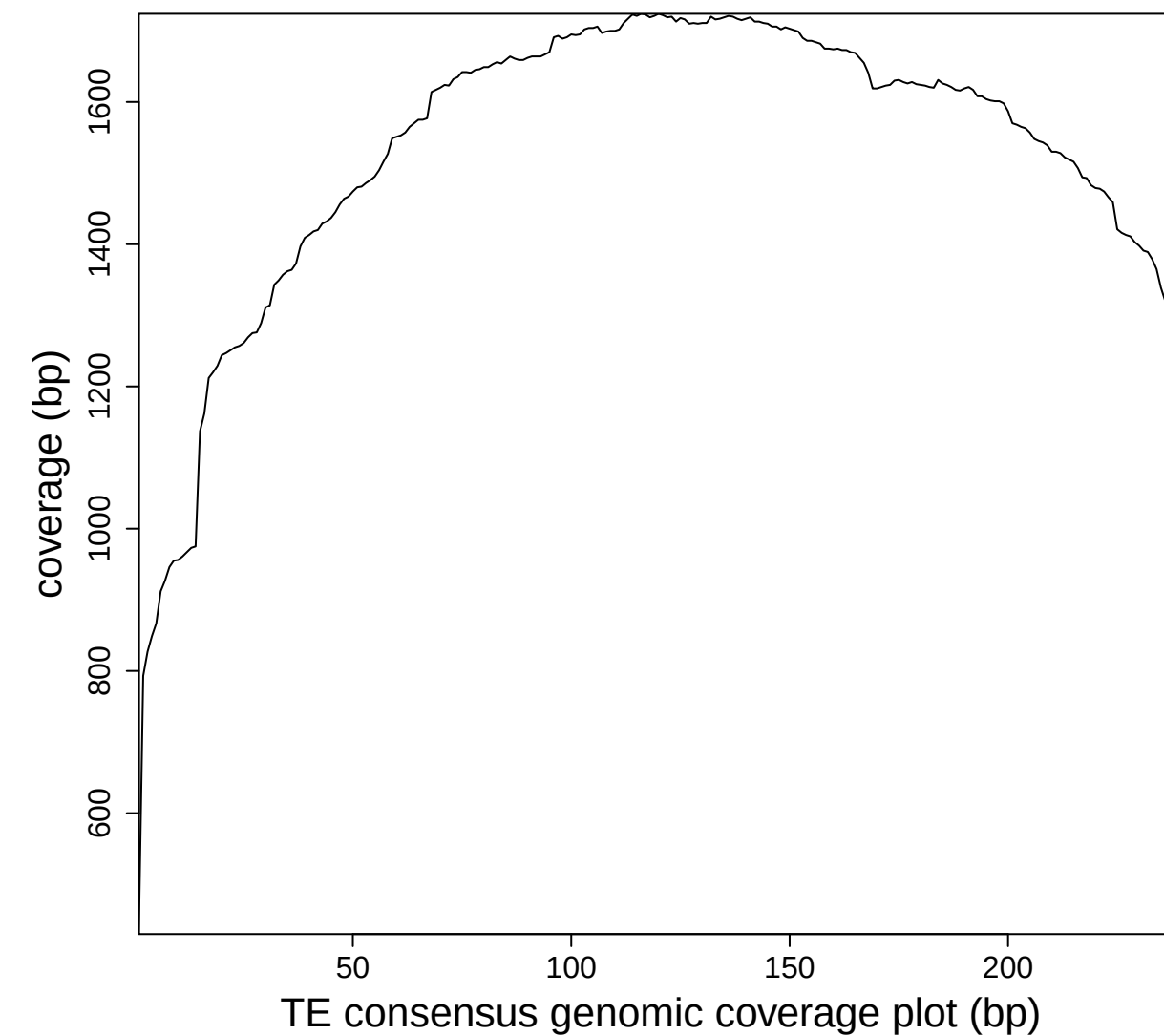
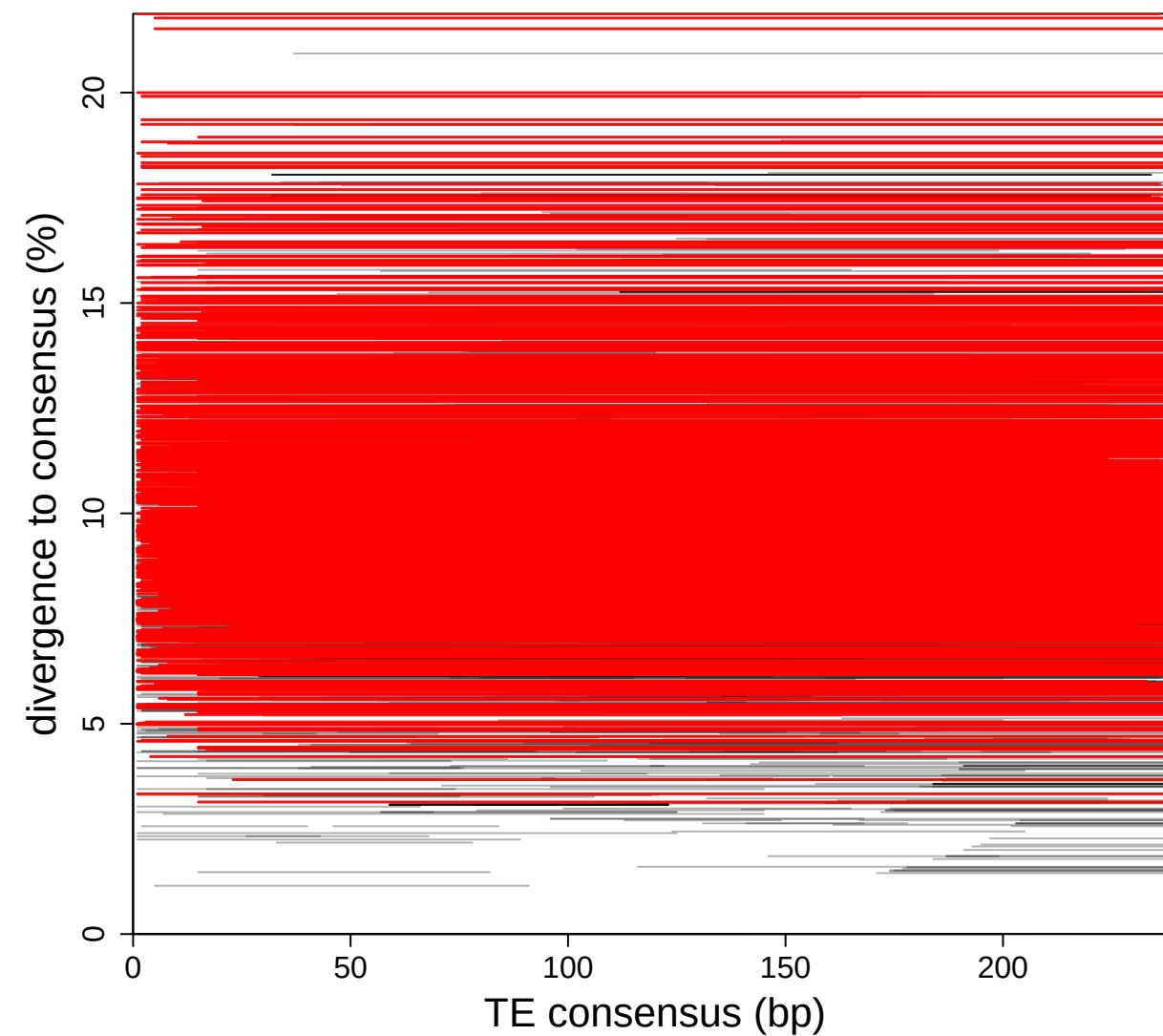


8.fasta.b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 928bp; fragments: 3842; full length: 0 (>=835.2bp)

After TEtrimmer Extended plot Blue lines are boundaries

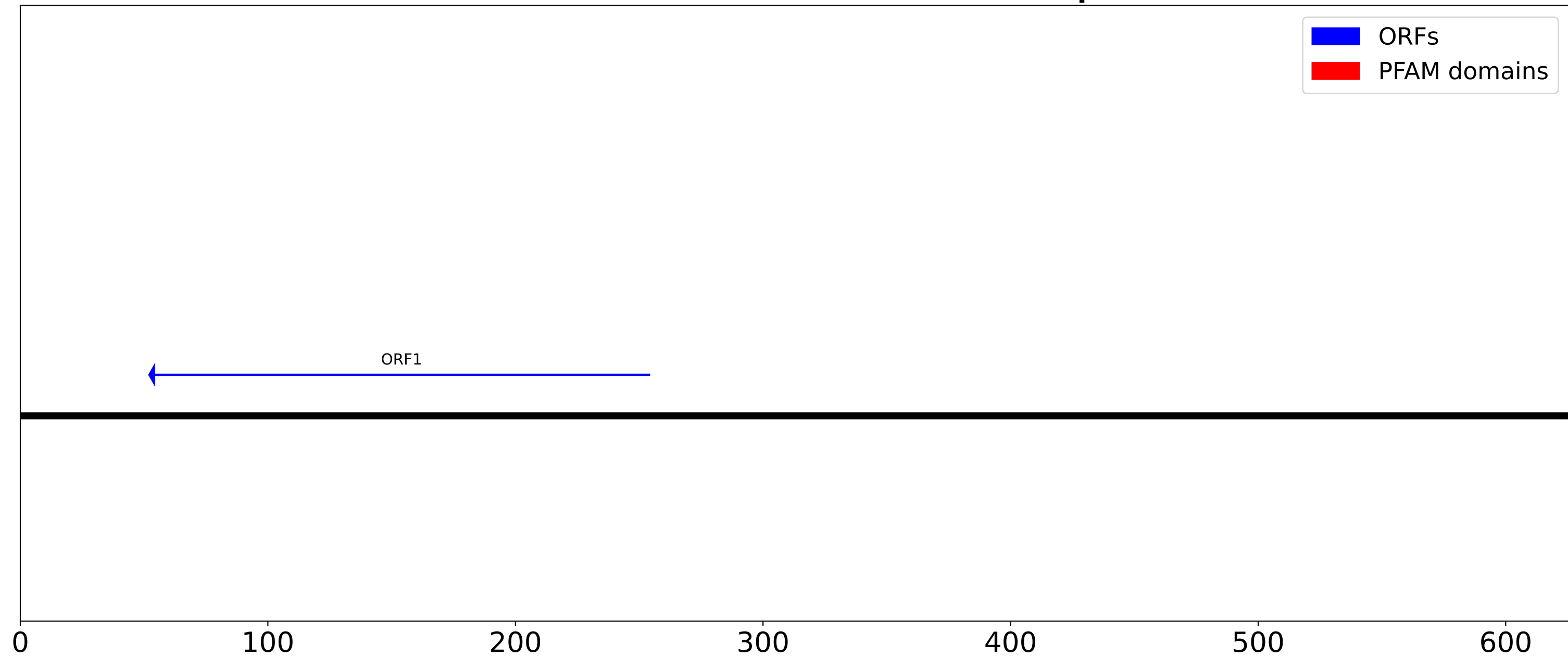


TE: rnd_1_family_248
size: 240bp; fragments: 2036; full length: 850 (≥ 216 bp)

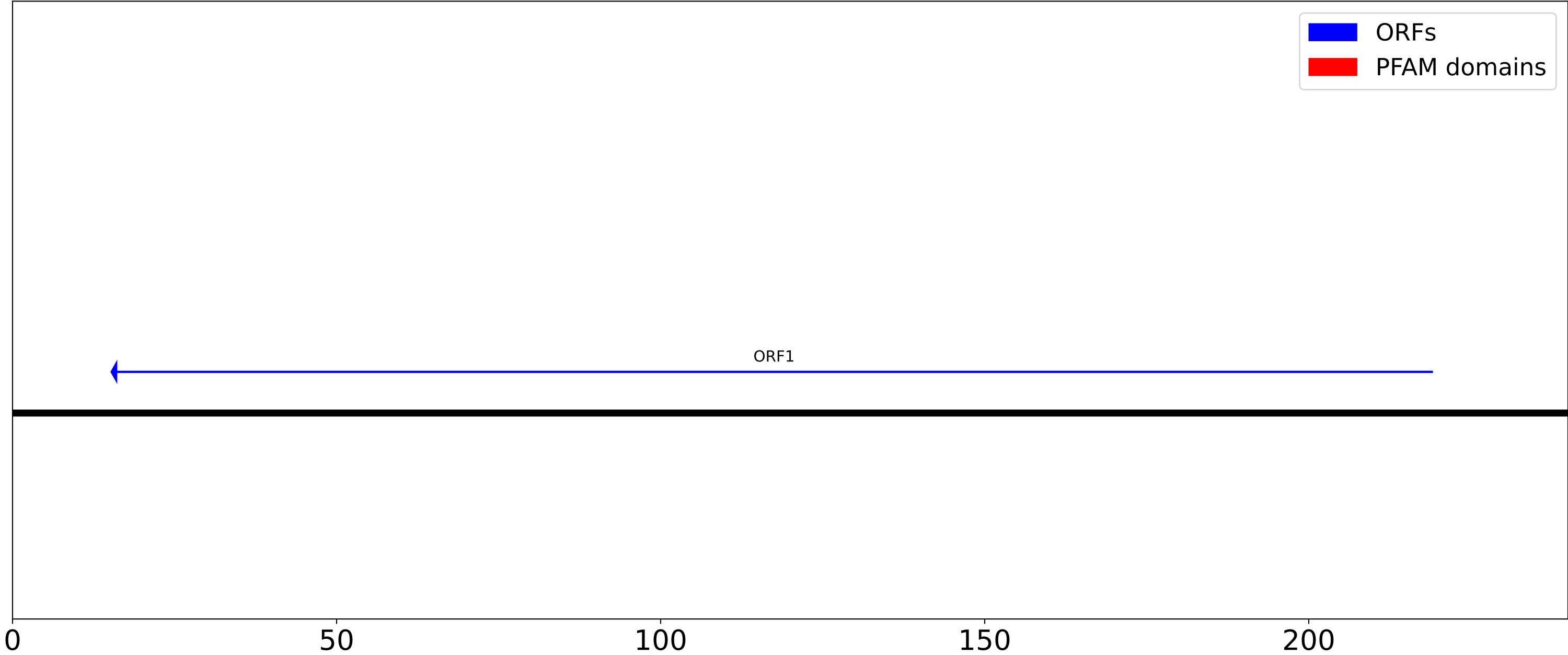


Before TEtrimmer 240 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

(window size = 25, threshold = 50.00 09/10/25)

TE consensus before TEtrimmer (bp)

