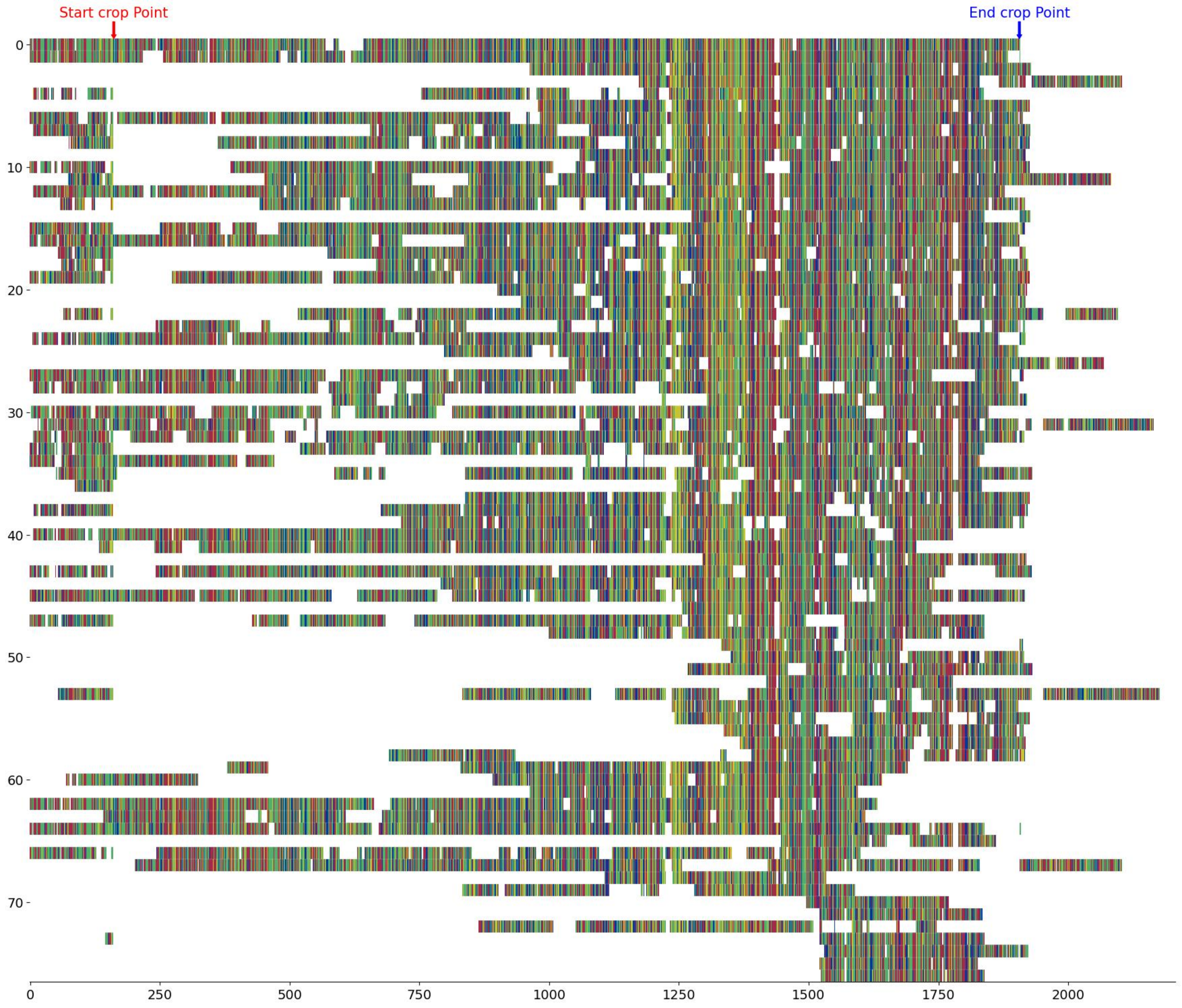


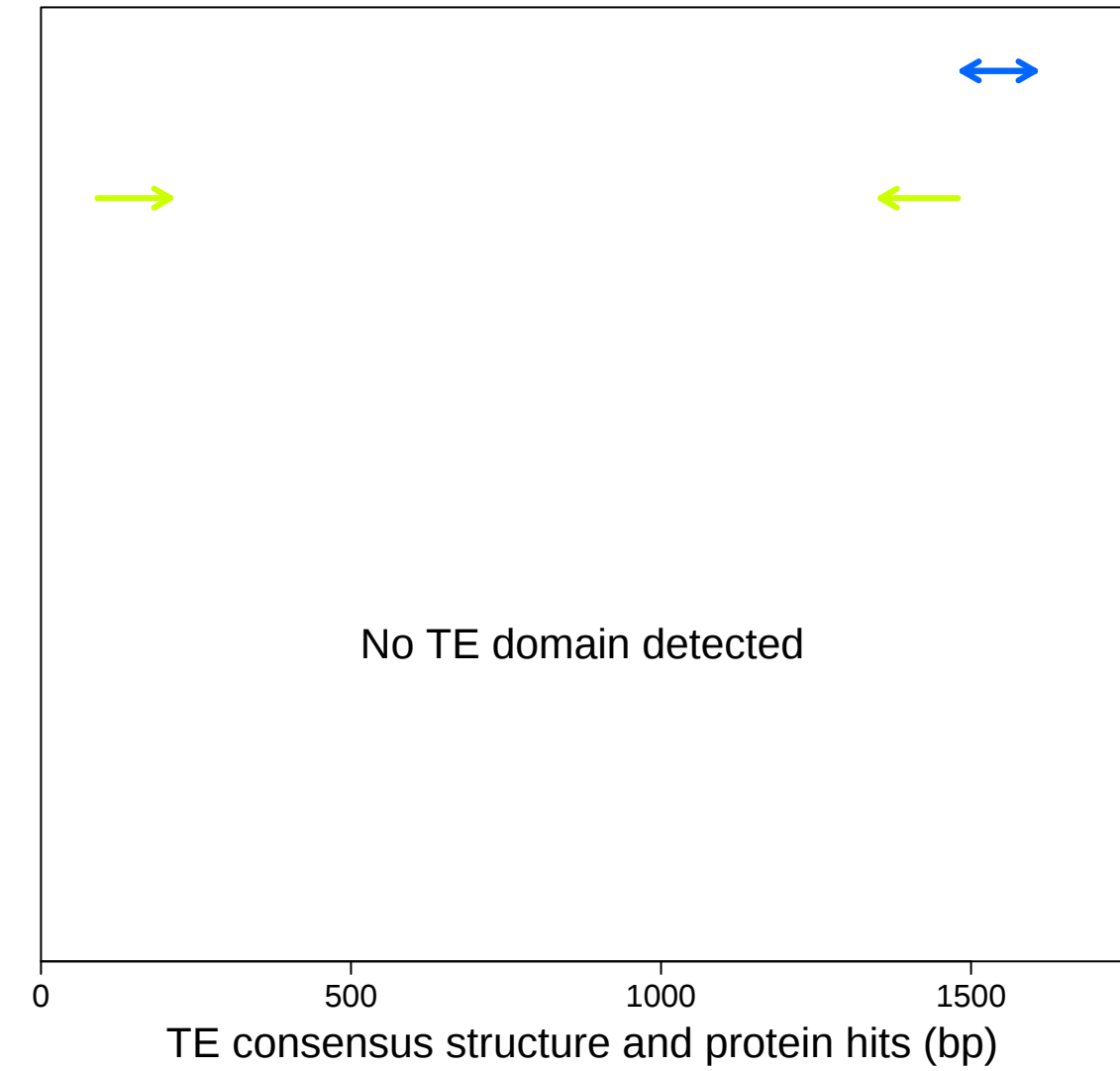
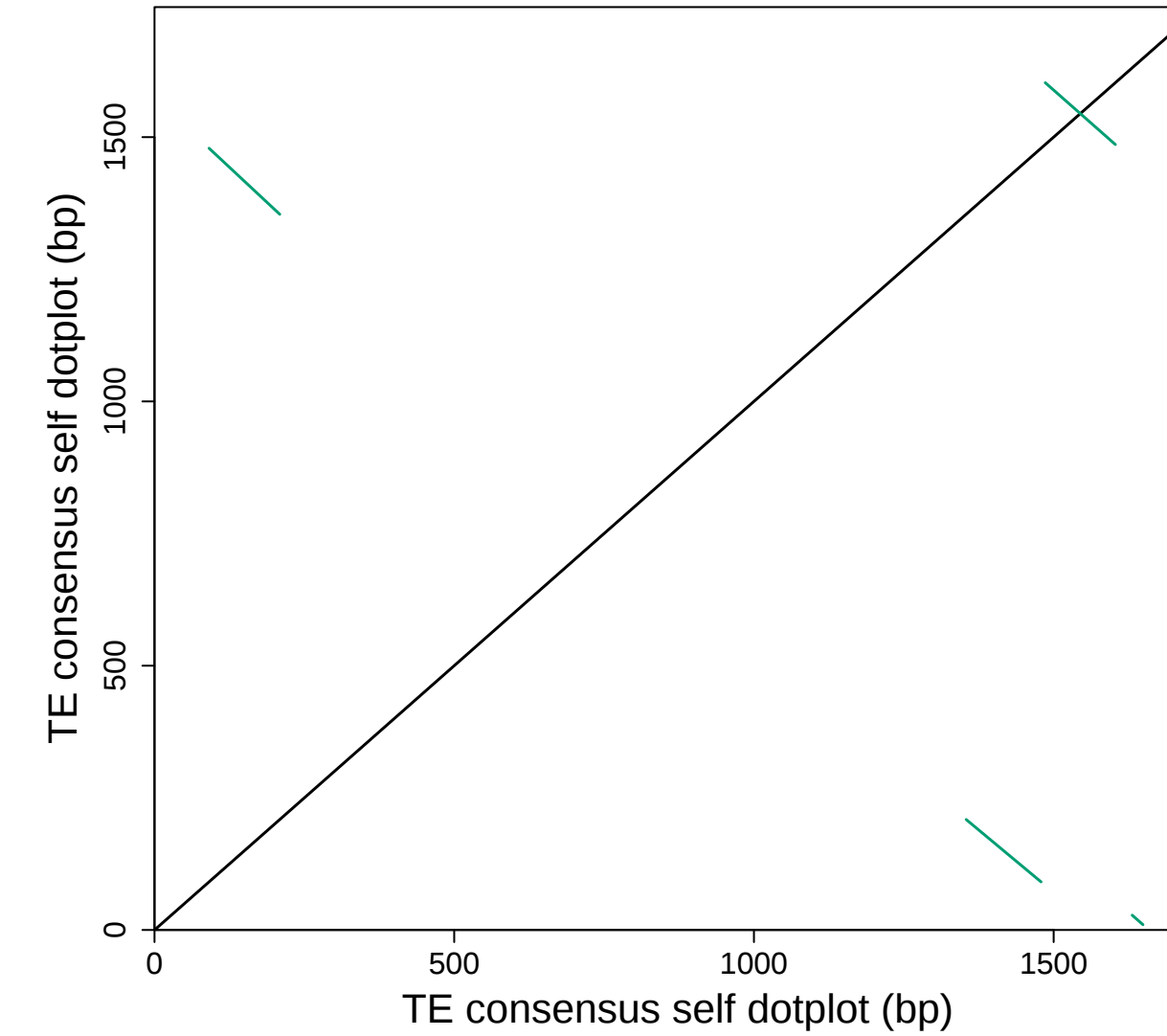
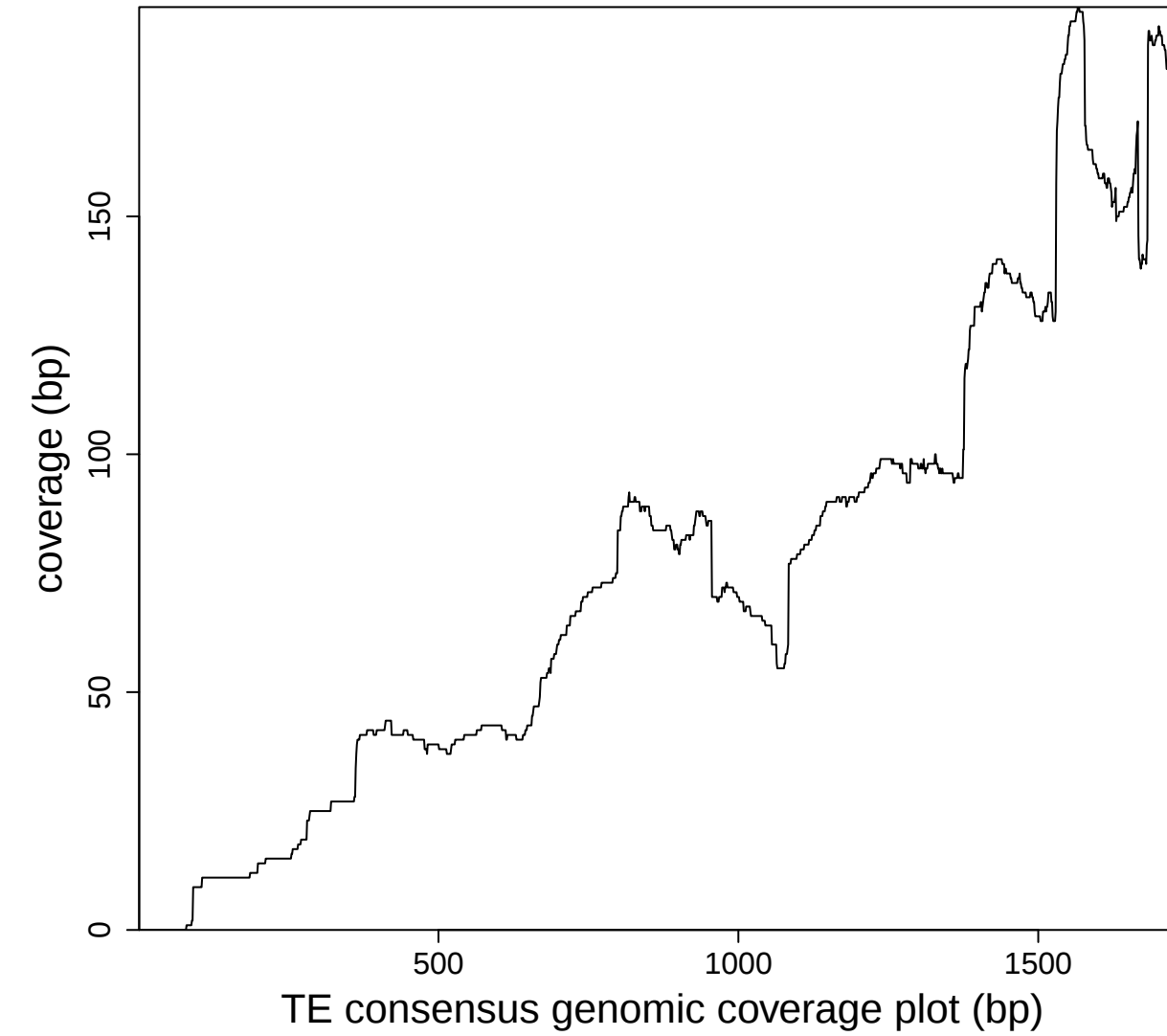
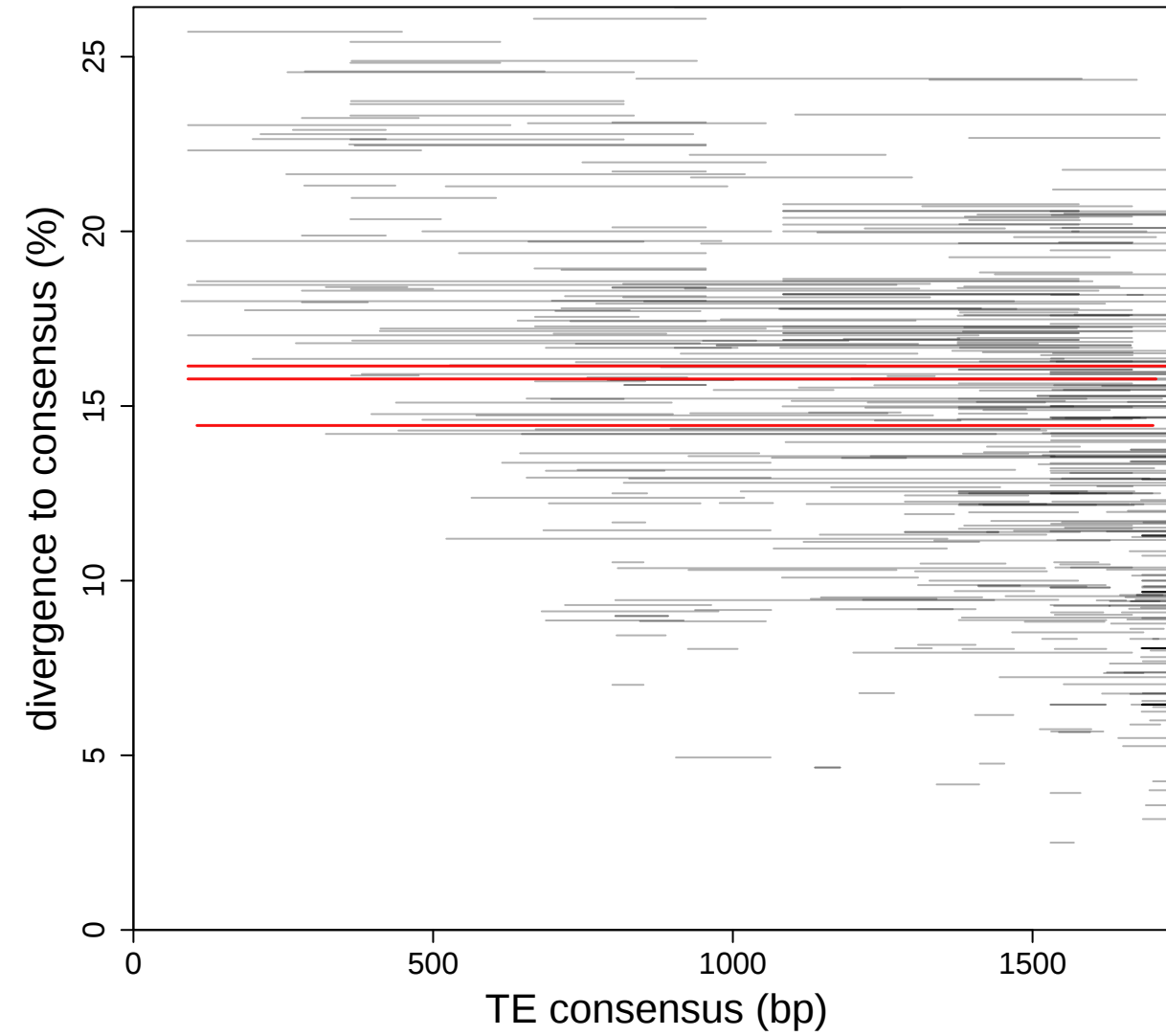
AAAAA
TCT
TTG
CAA
TTG
TTC
GAA

-ATTGTTCCAAATGTTTTGACAGCGA-CTGGGCAGGCACAGCTAGC--GTACCTATGTAAAAAGAA-



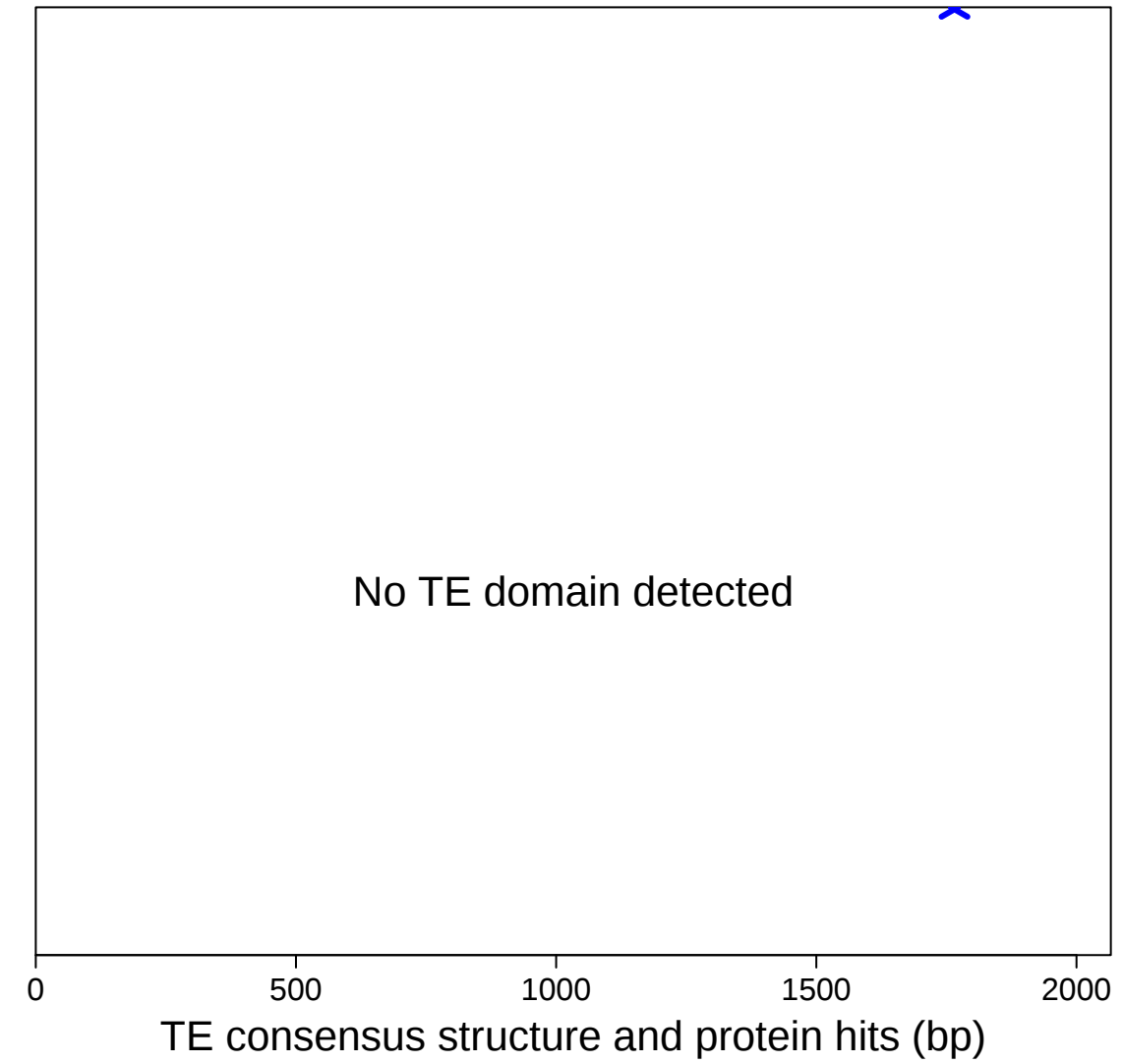
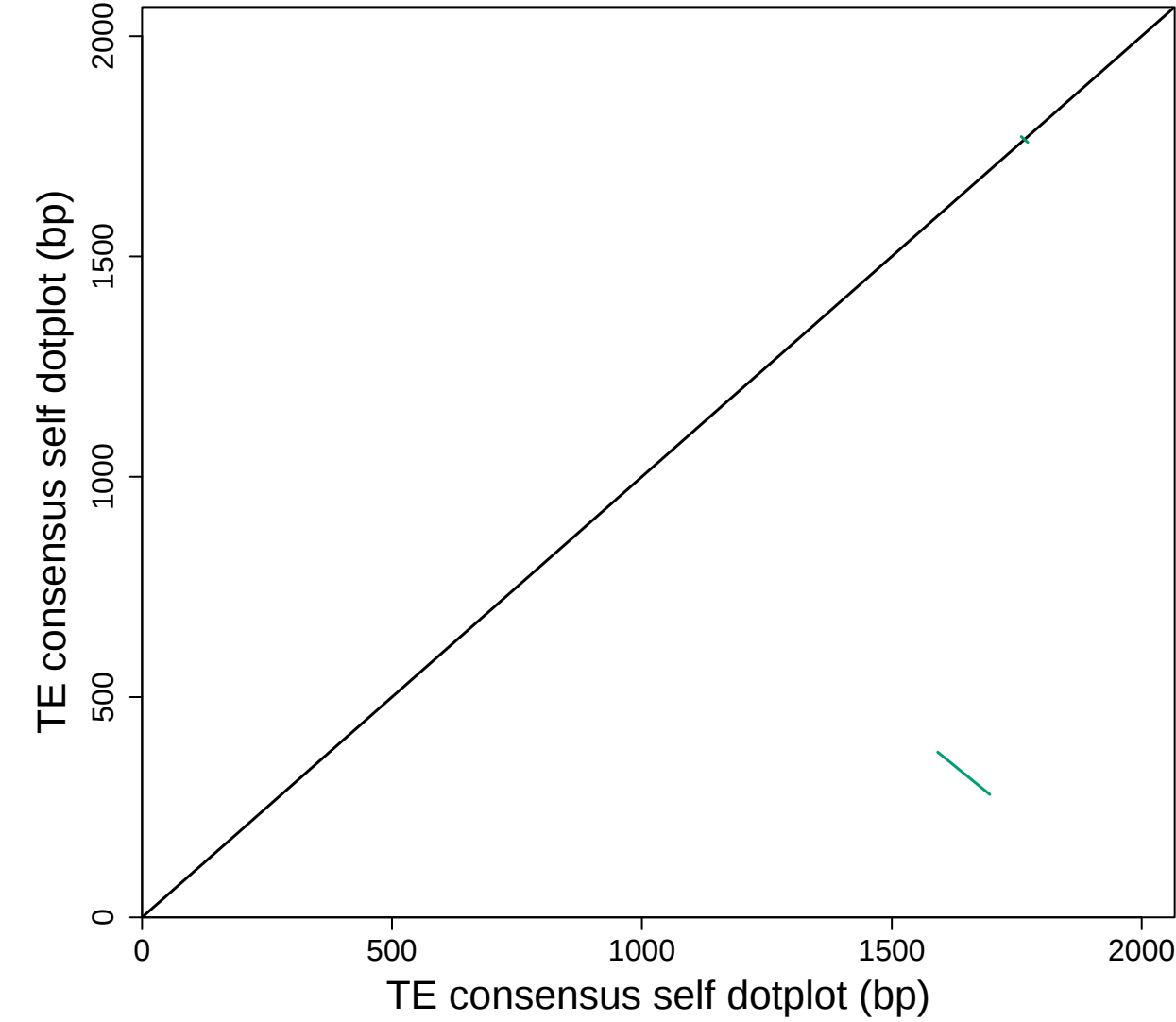
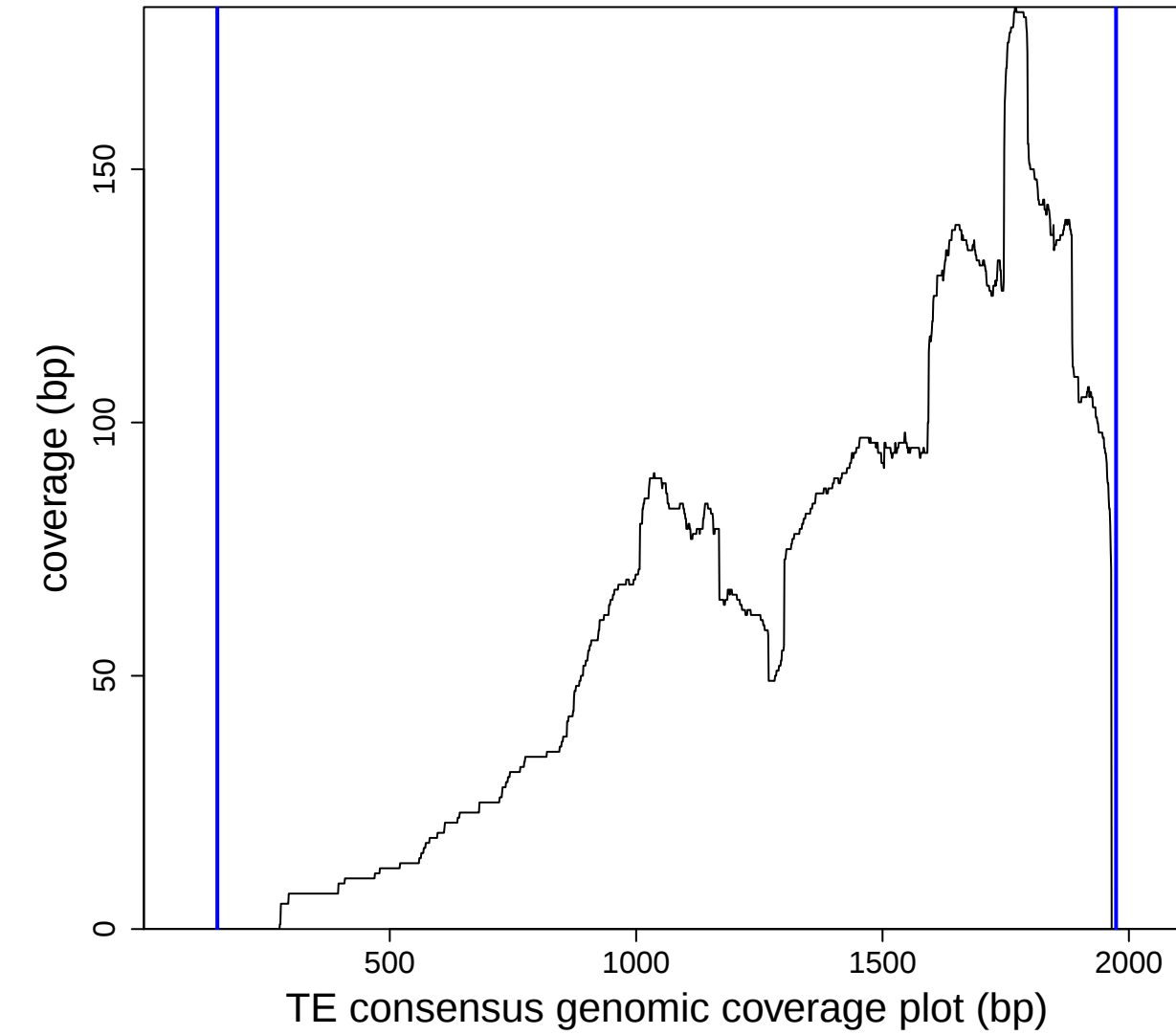
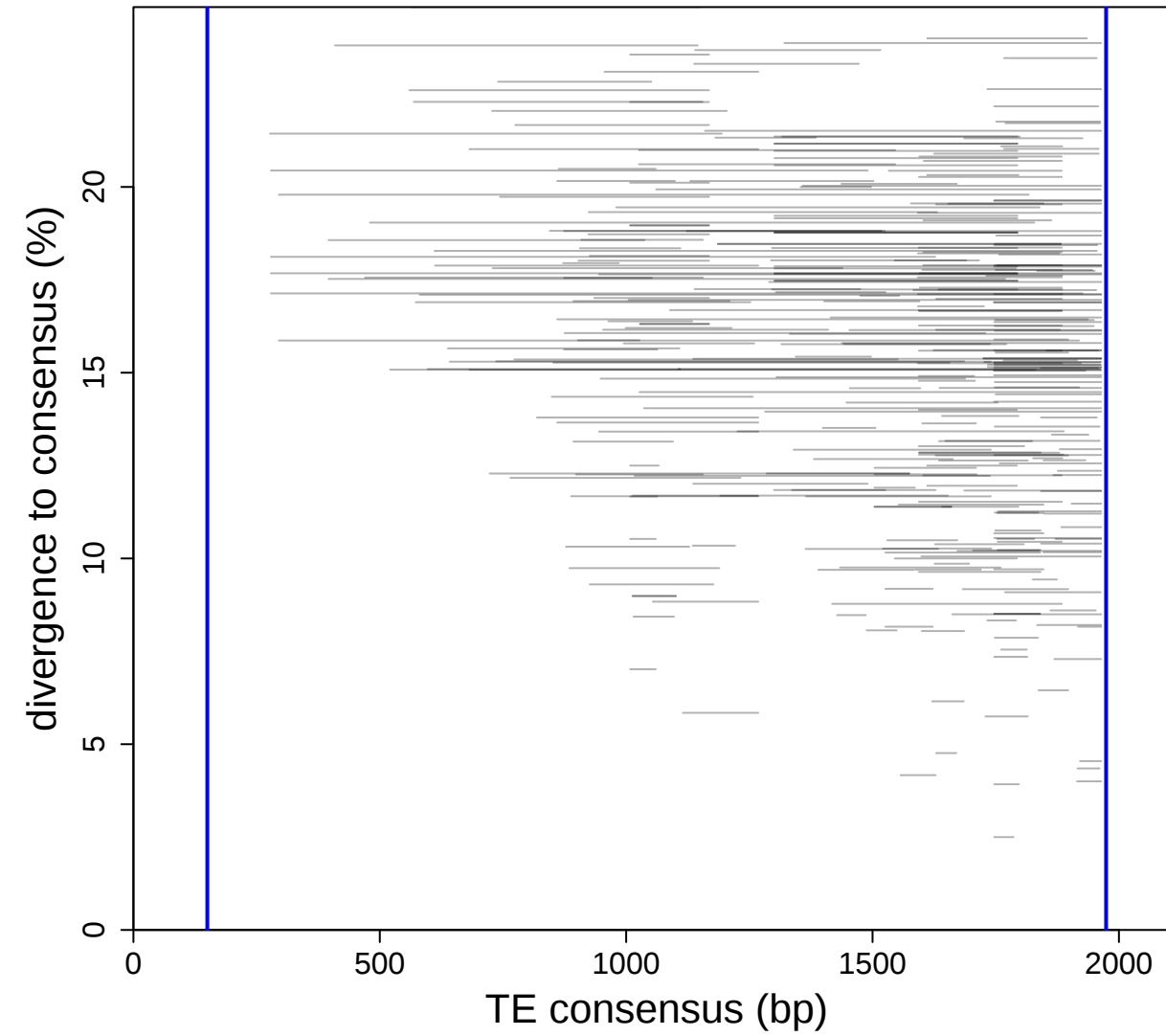
ta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 1746bp; fragments: 450; full length: 3 (>=1571.4bp)

After TEtrimmer 1746 bp

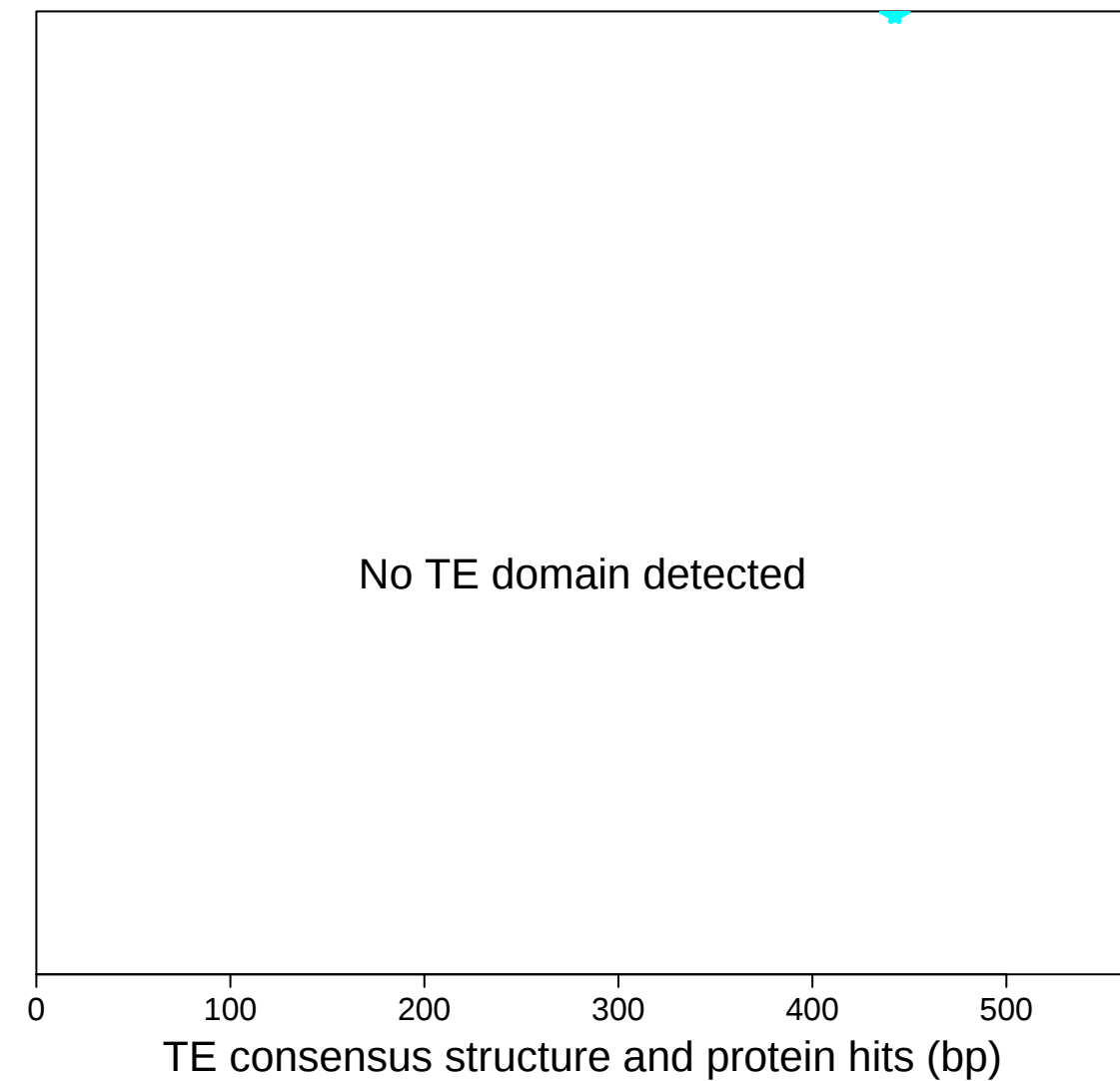
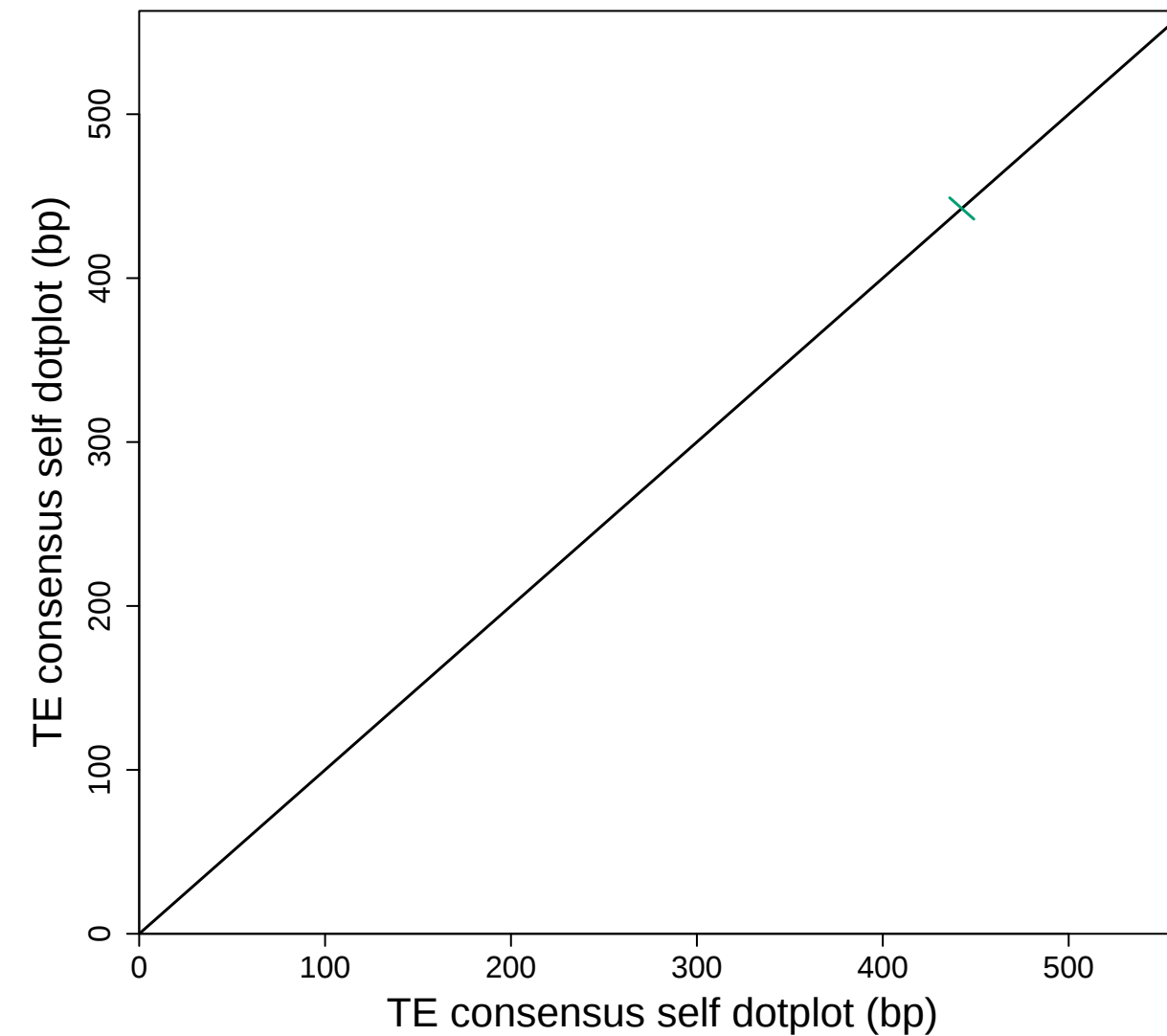
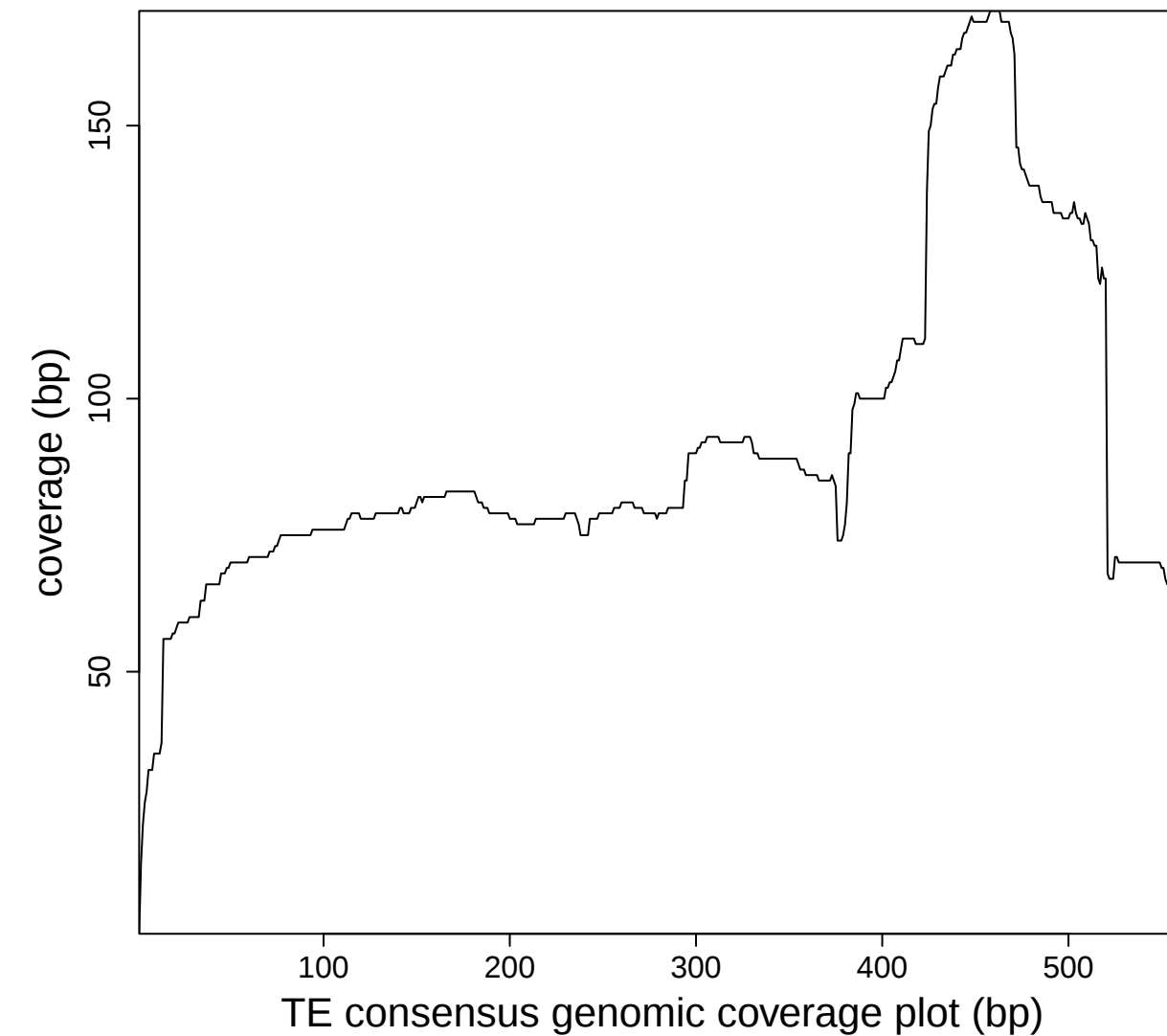
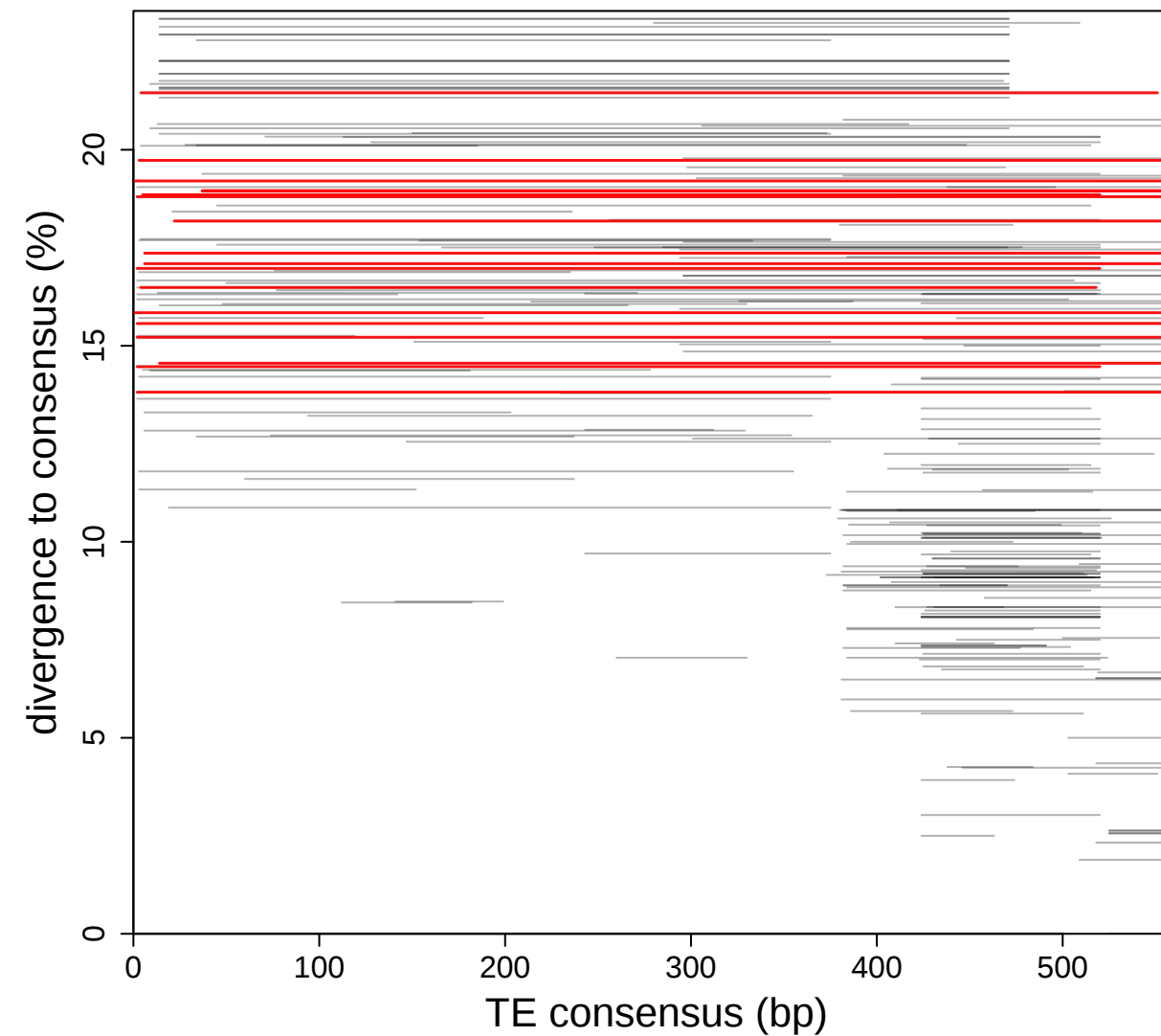


21.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 2124bp; fragments: 337; full length: 0 (>=1911.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

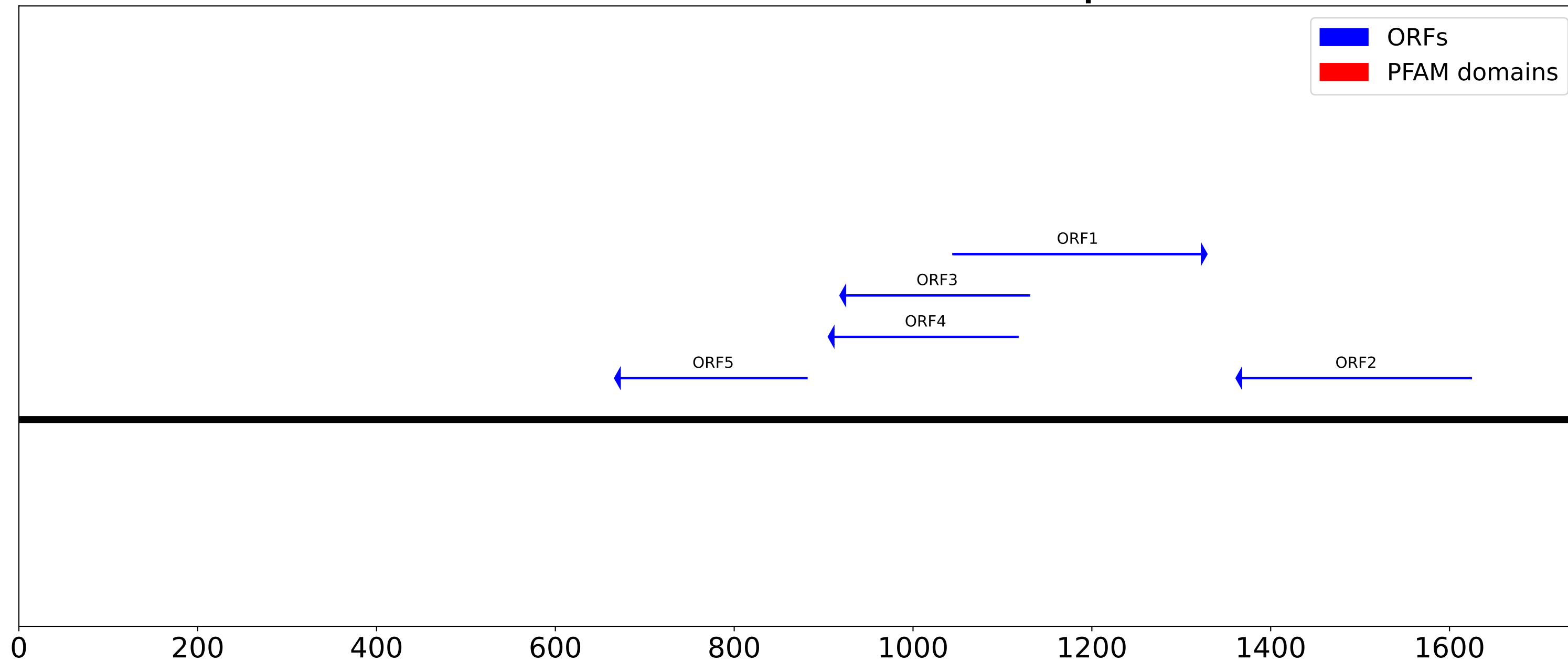


TE: rnd_5_family_16921
size: 563bp; fragments: 228; full length: 18 (≥ 506.7 bp)

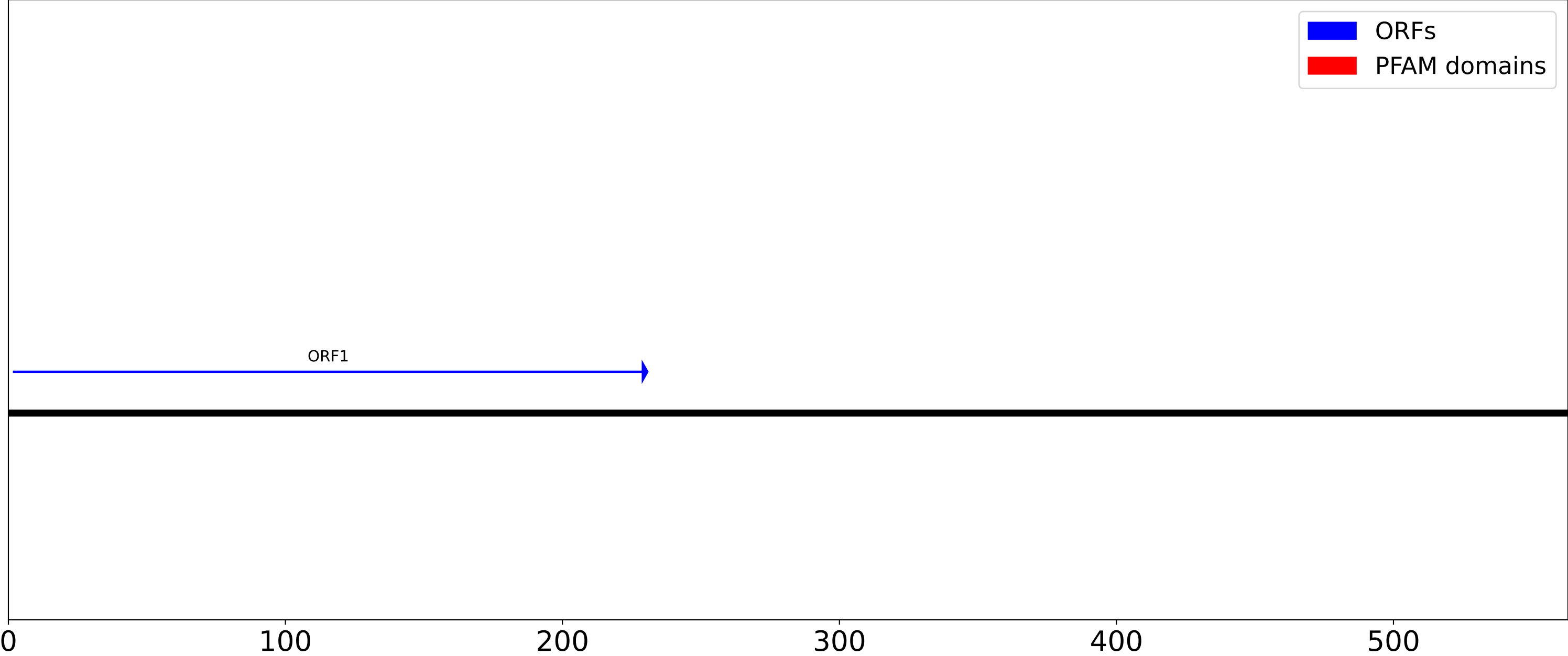


Before TEtrimmer 563 bp

After TEtrimmer ORF and PFAM domain plot



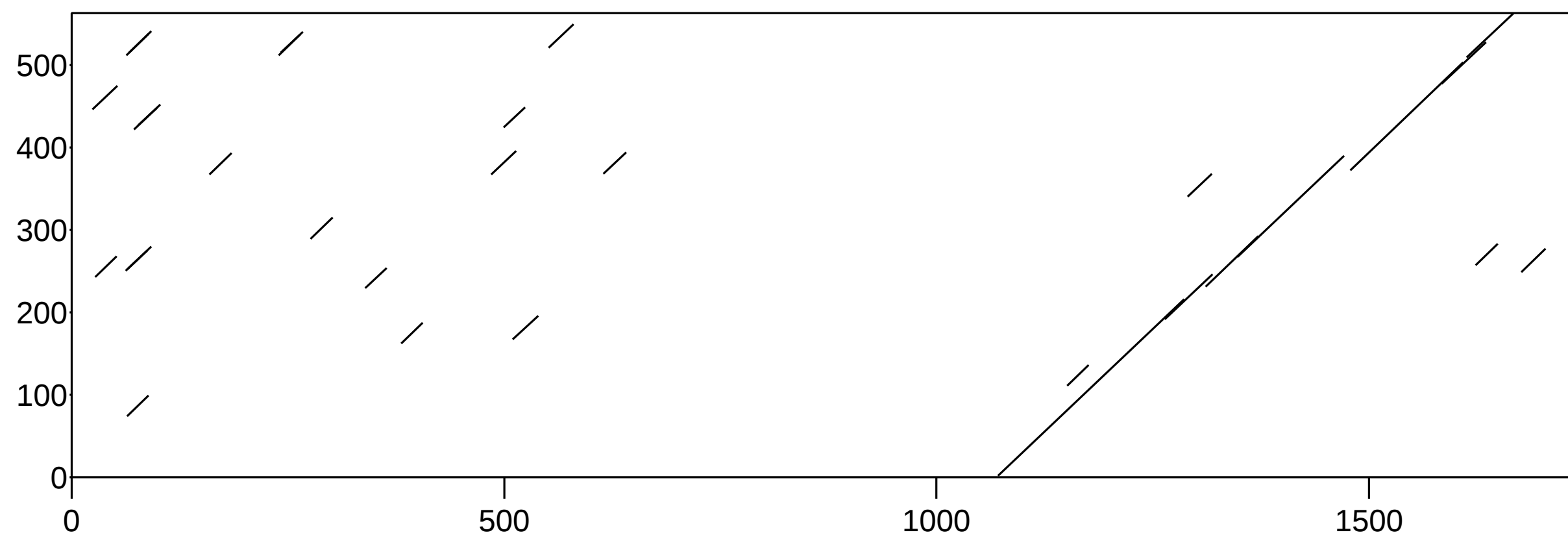
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)