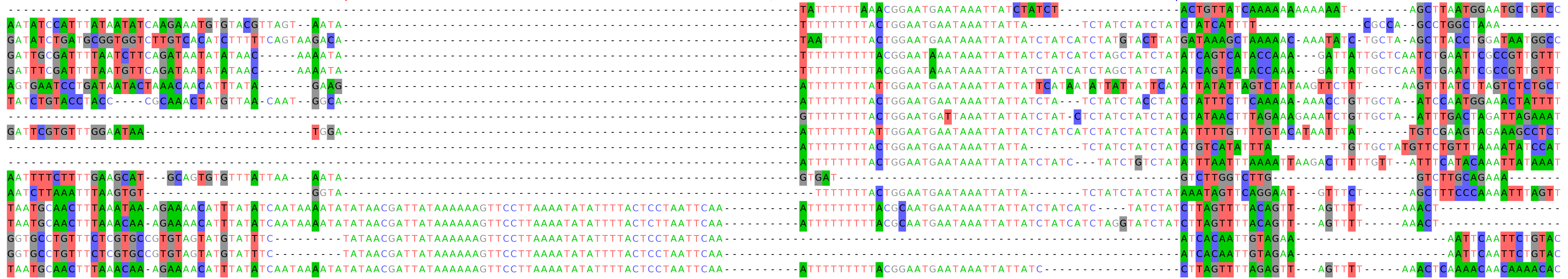
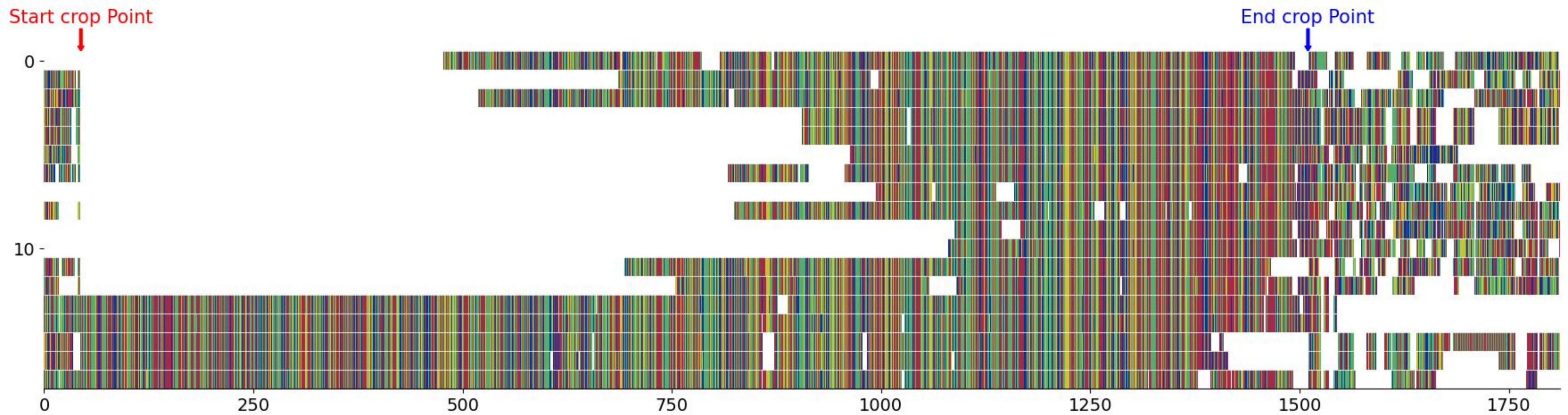


End crop Point

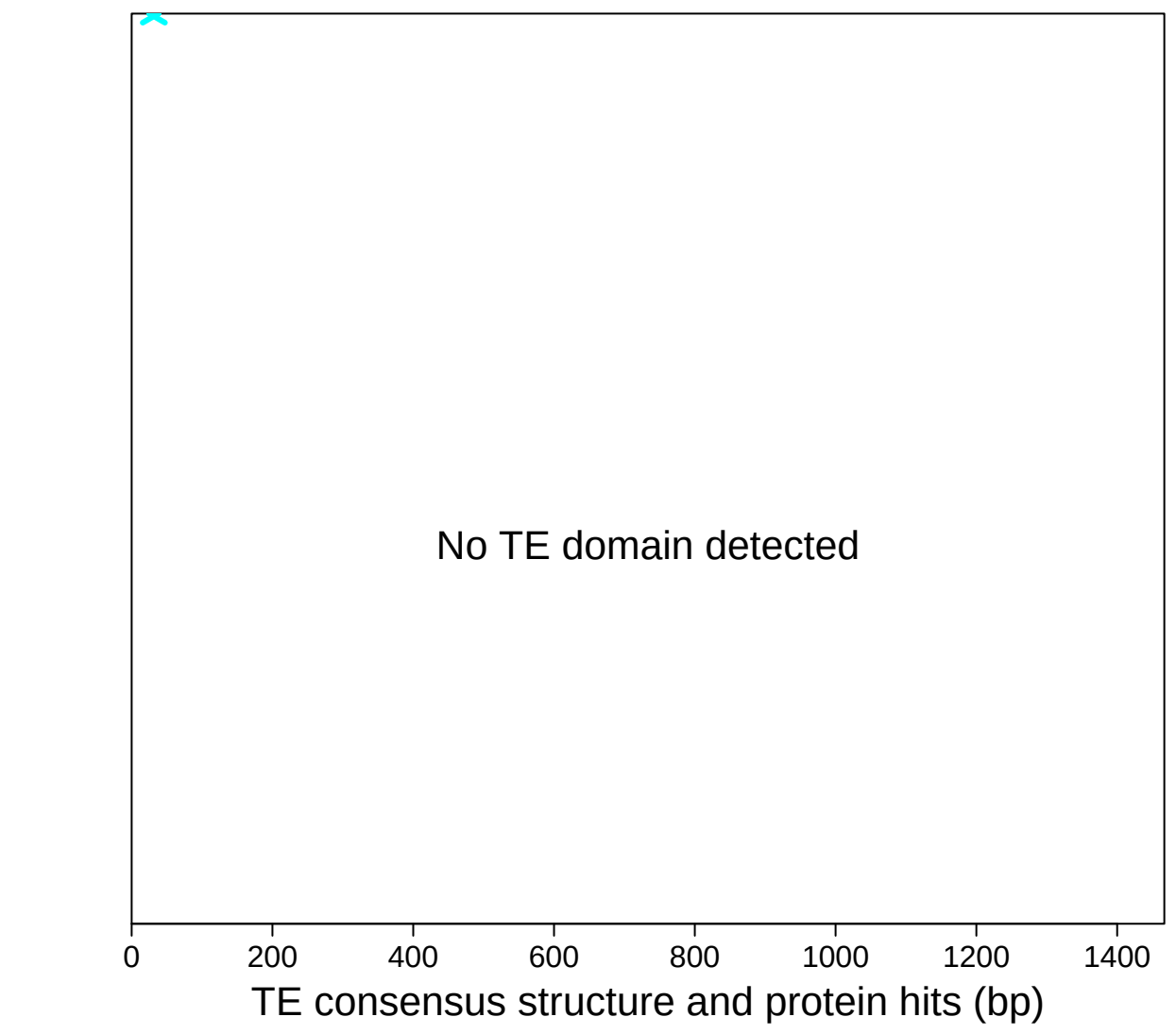
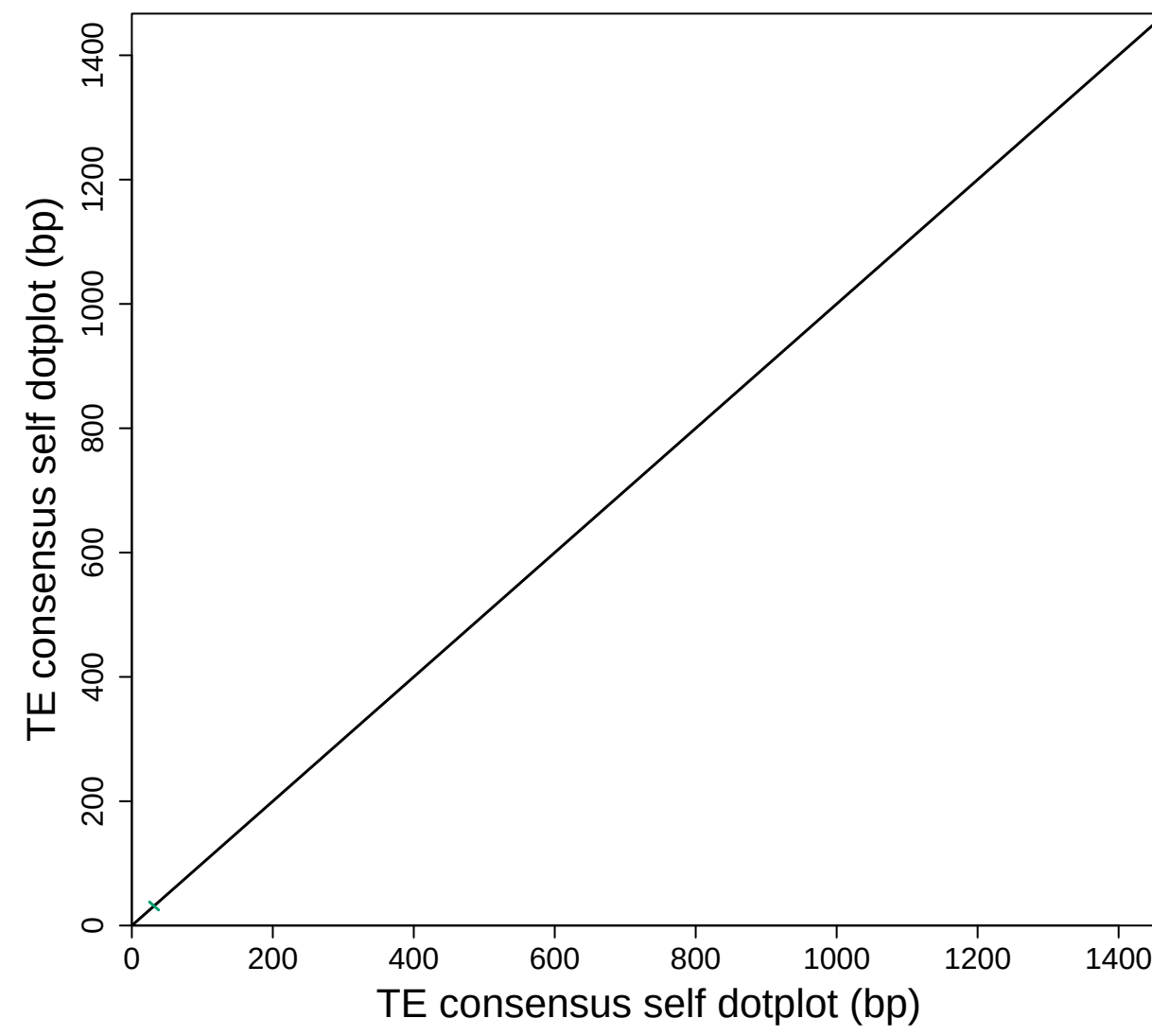
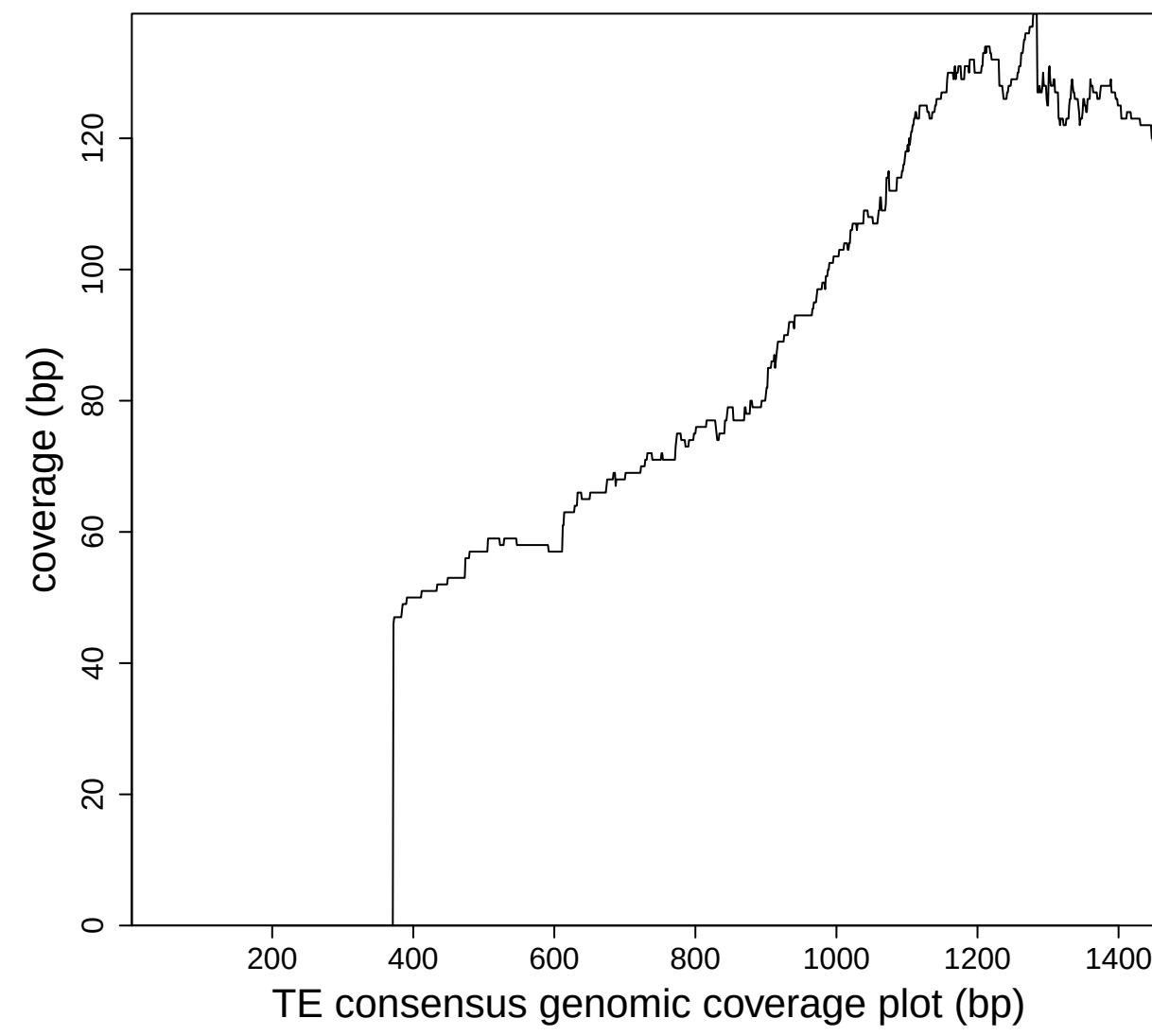
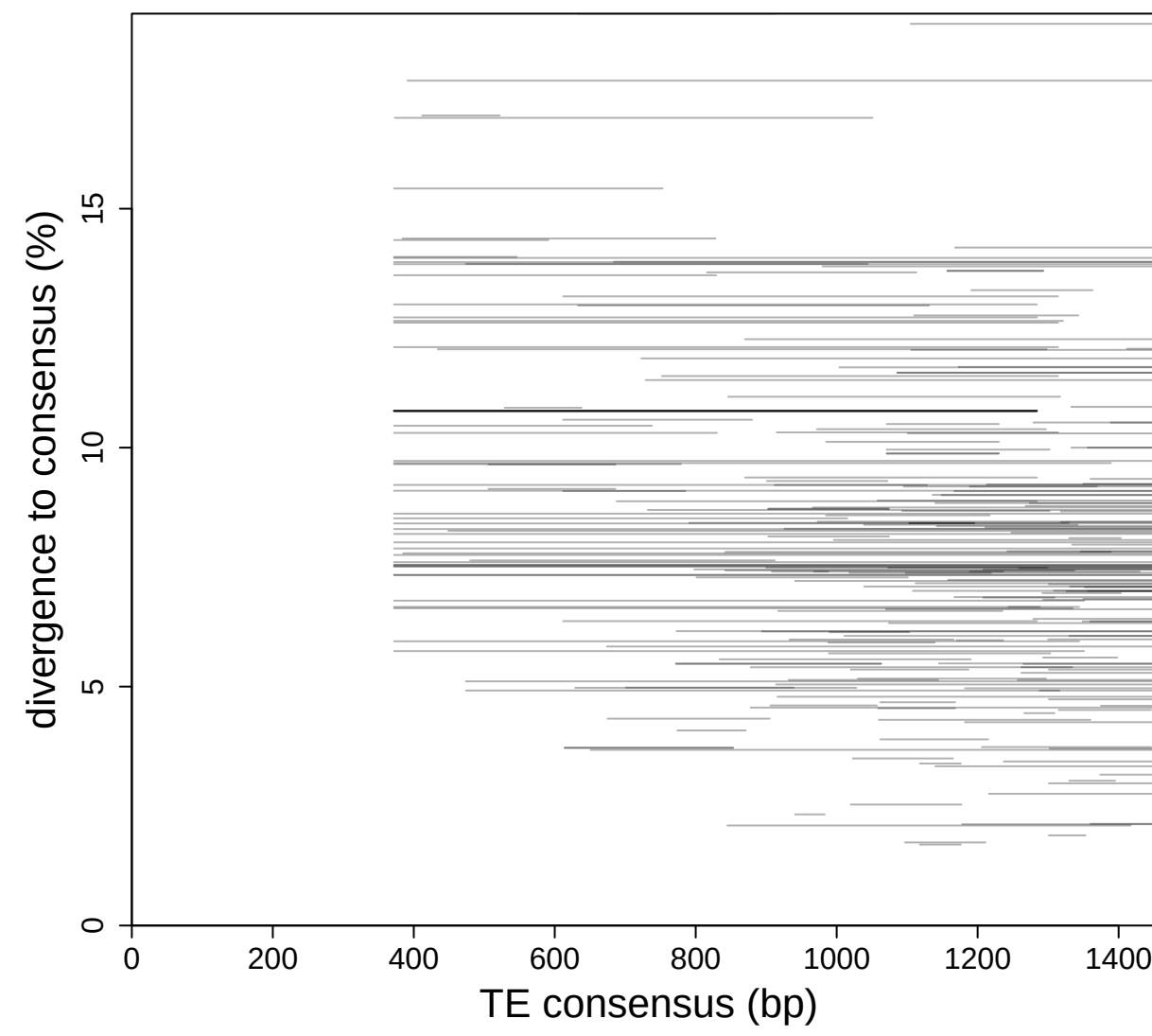
End crop Point





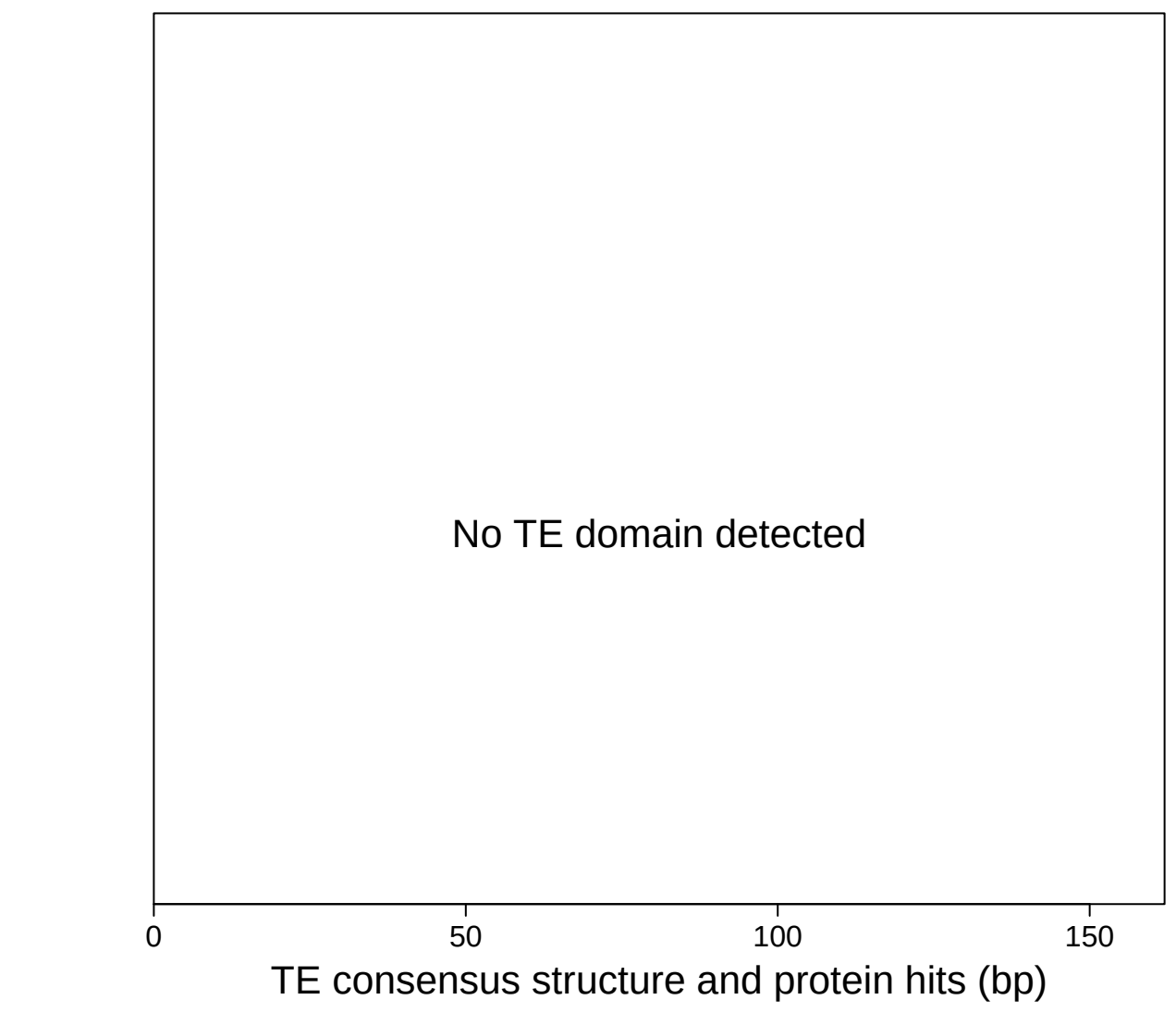
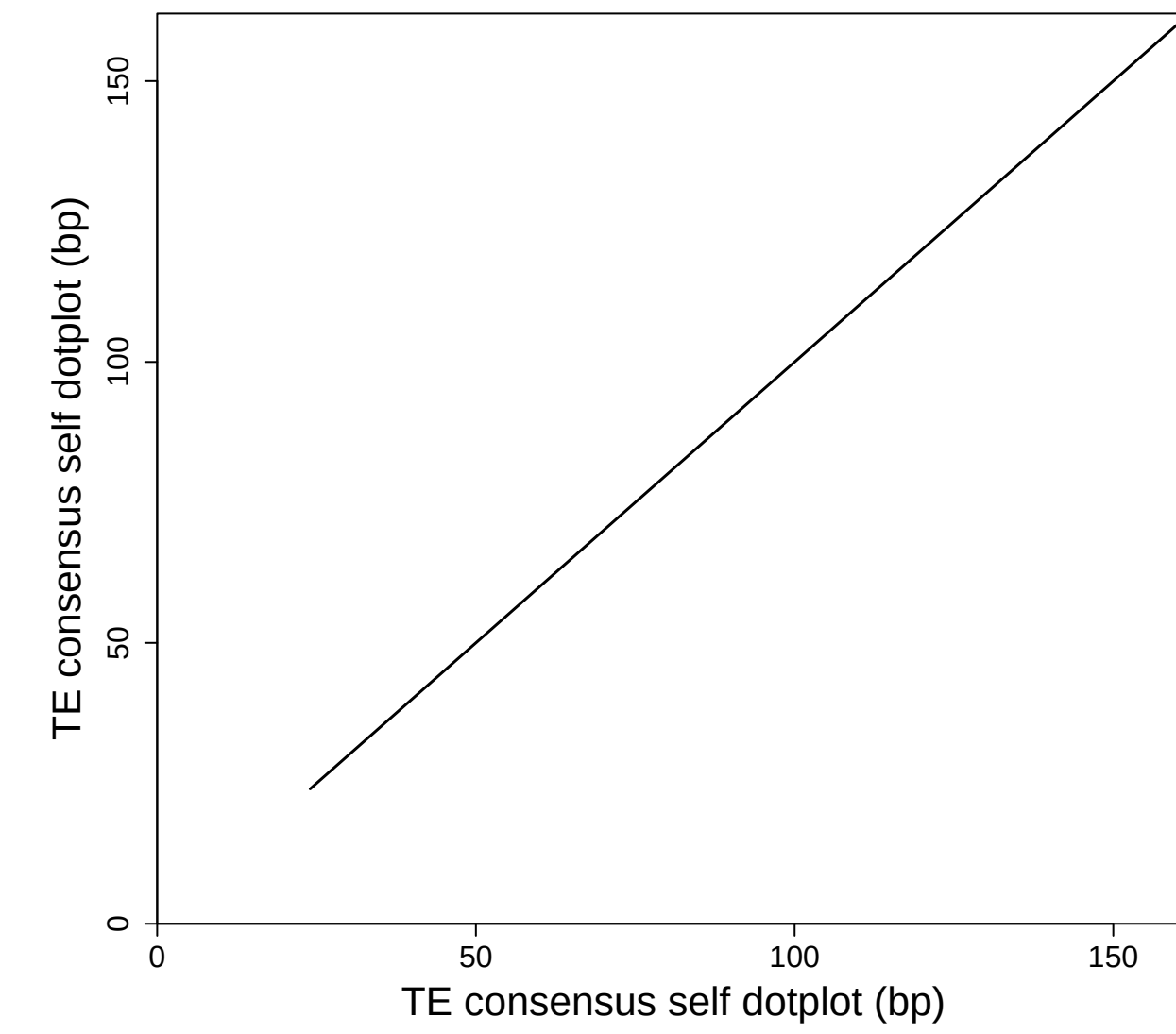
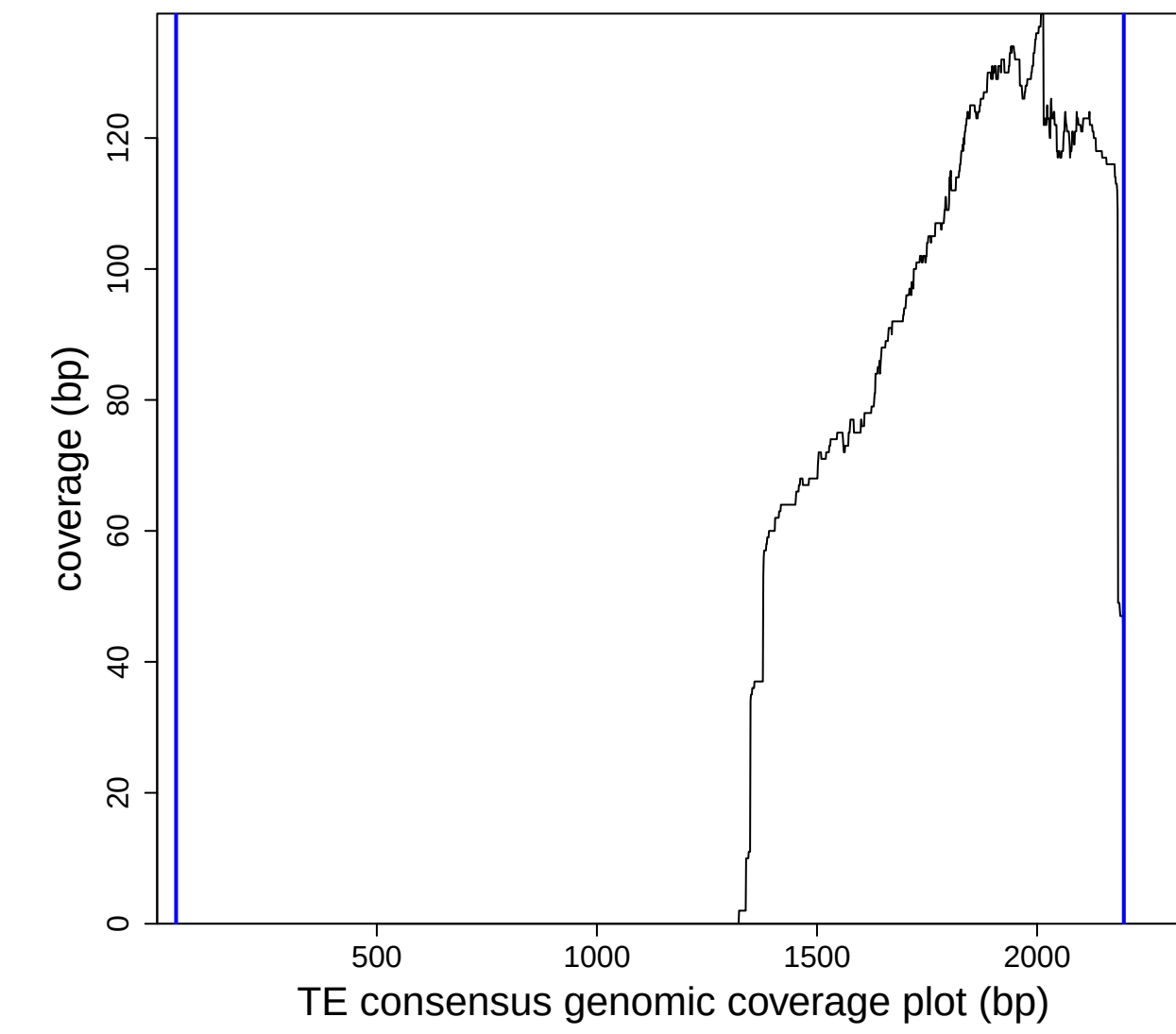
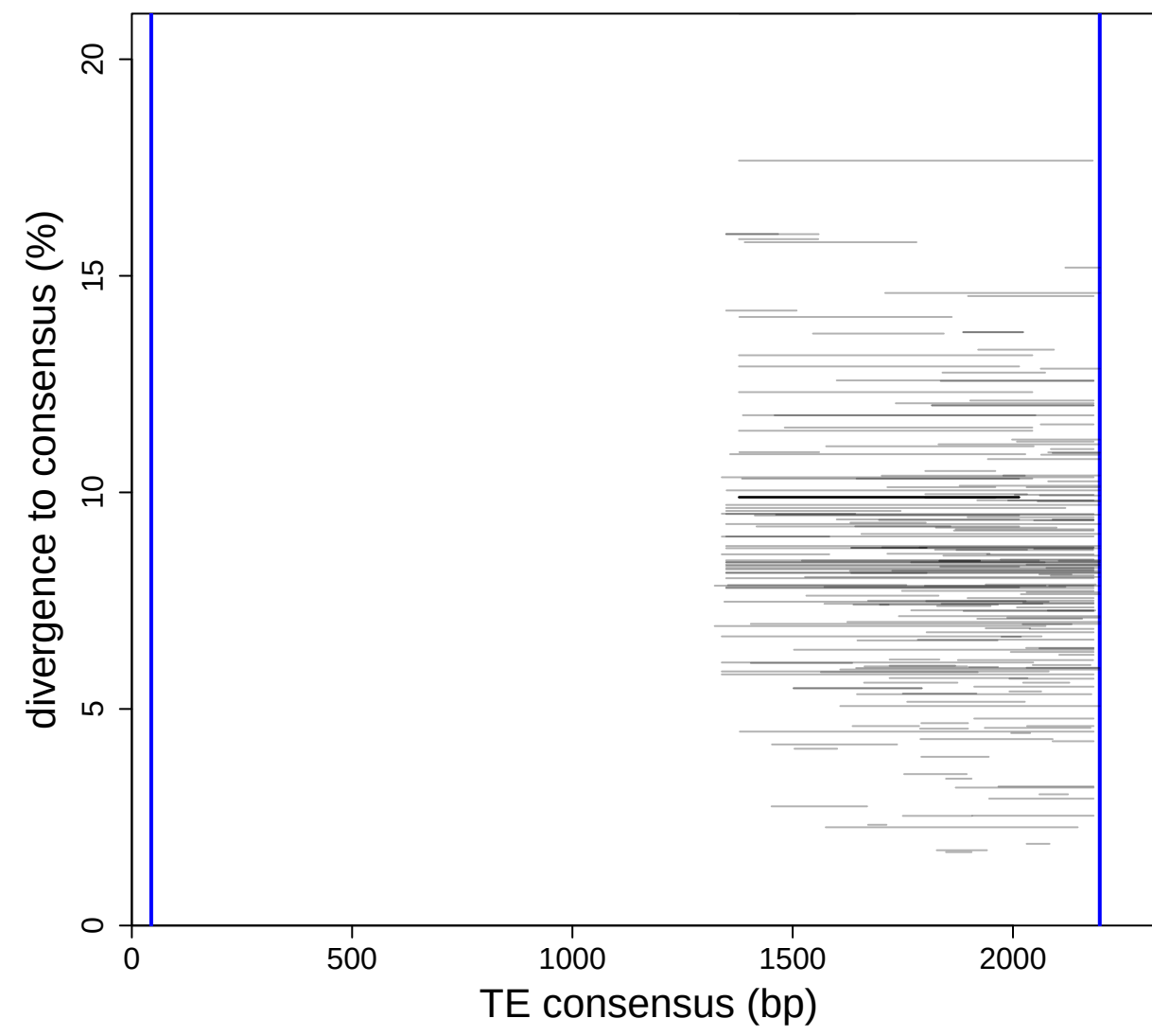
bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 1467bp; fragments: 239; full length: 0 (>=1320.3bp)

After TETrimmer 1467 bp

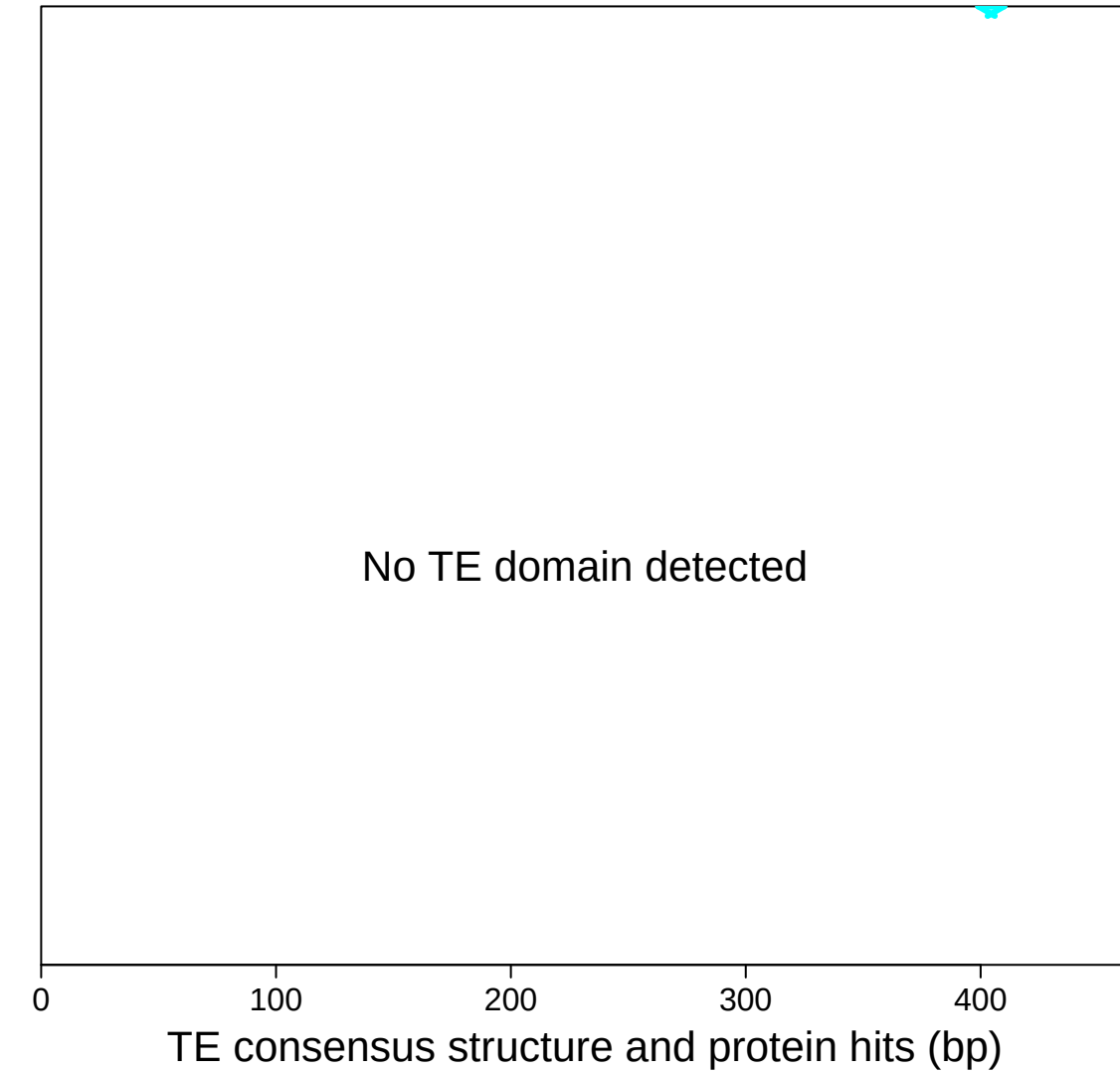
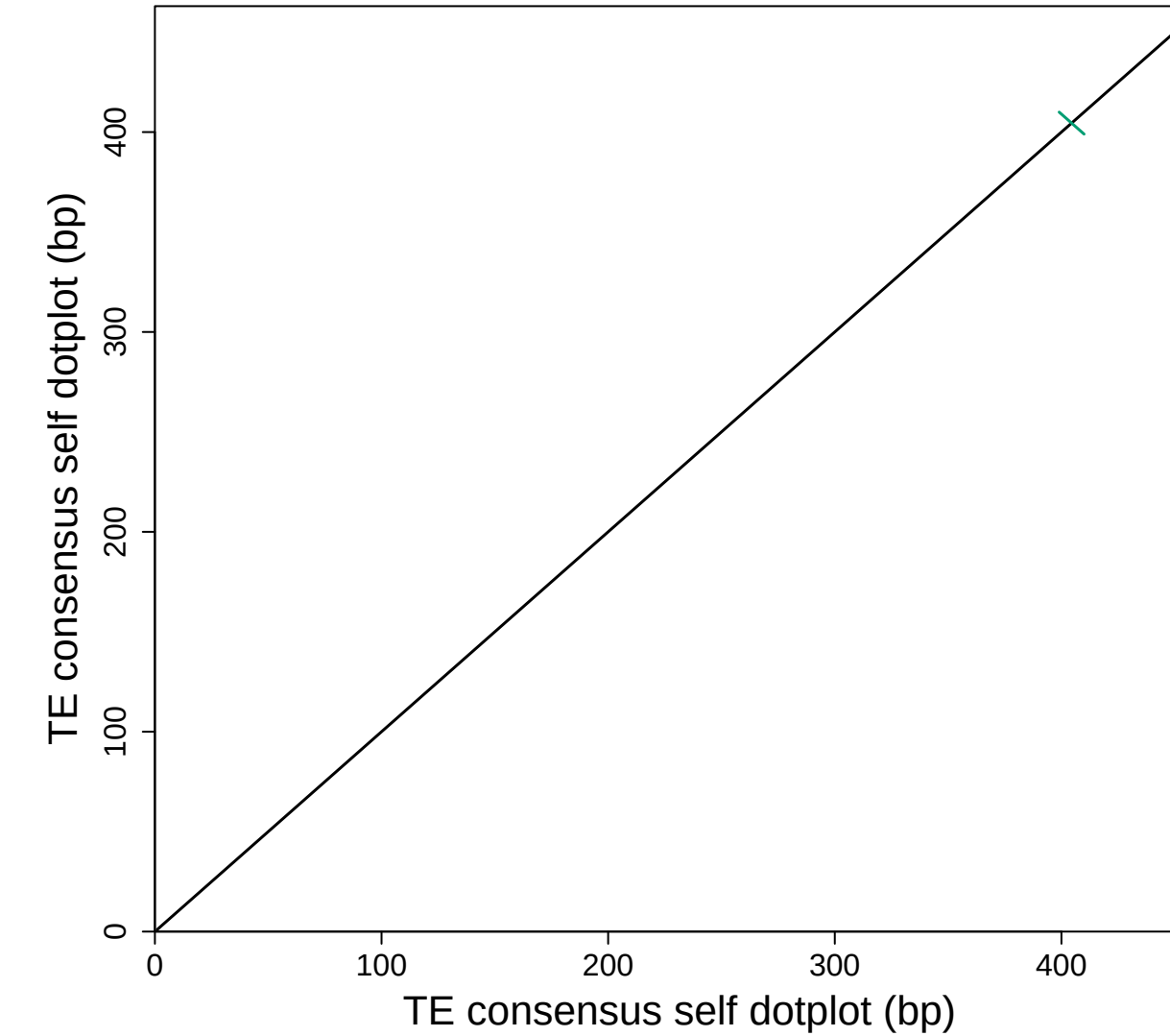
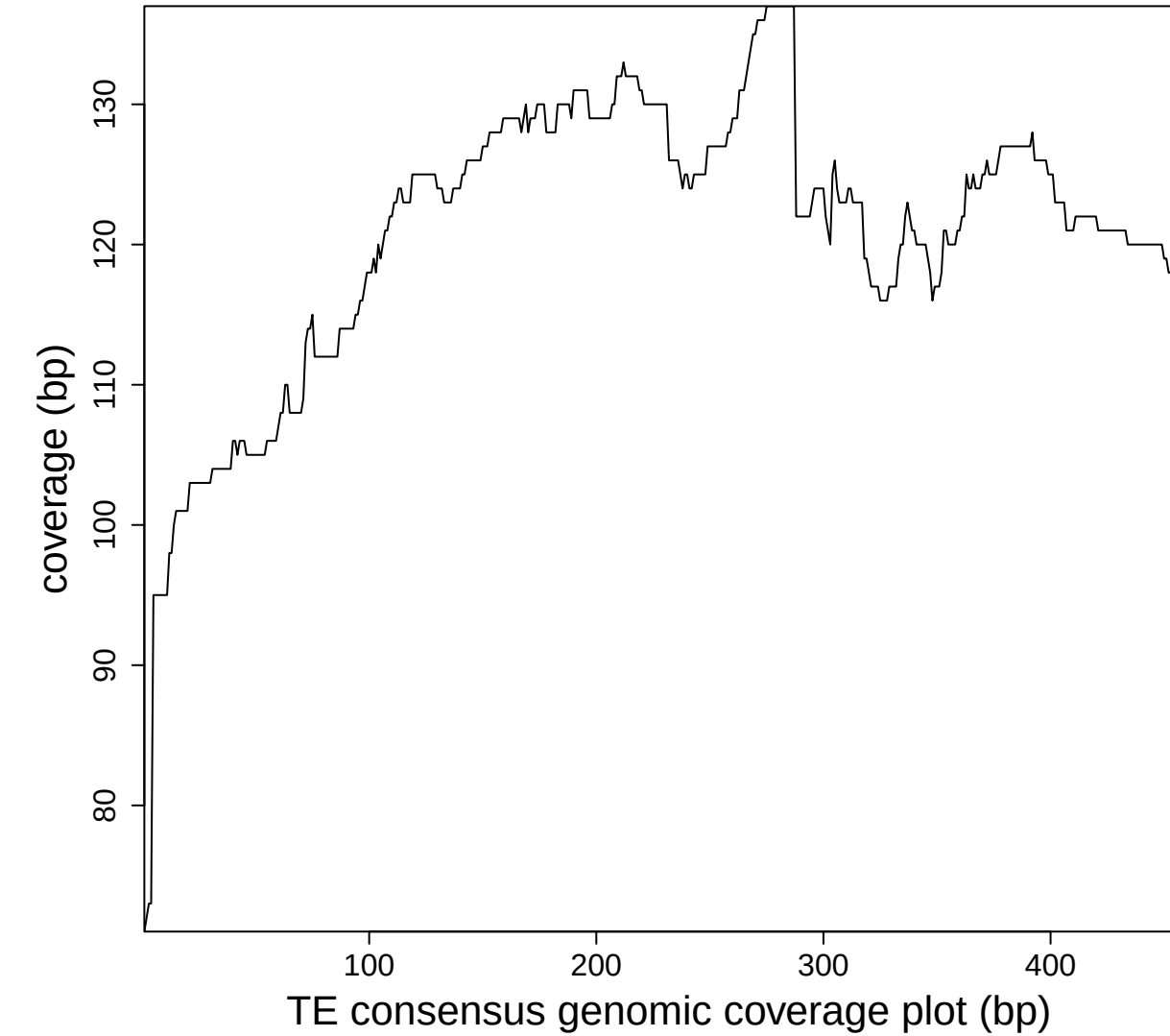
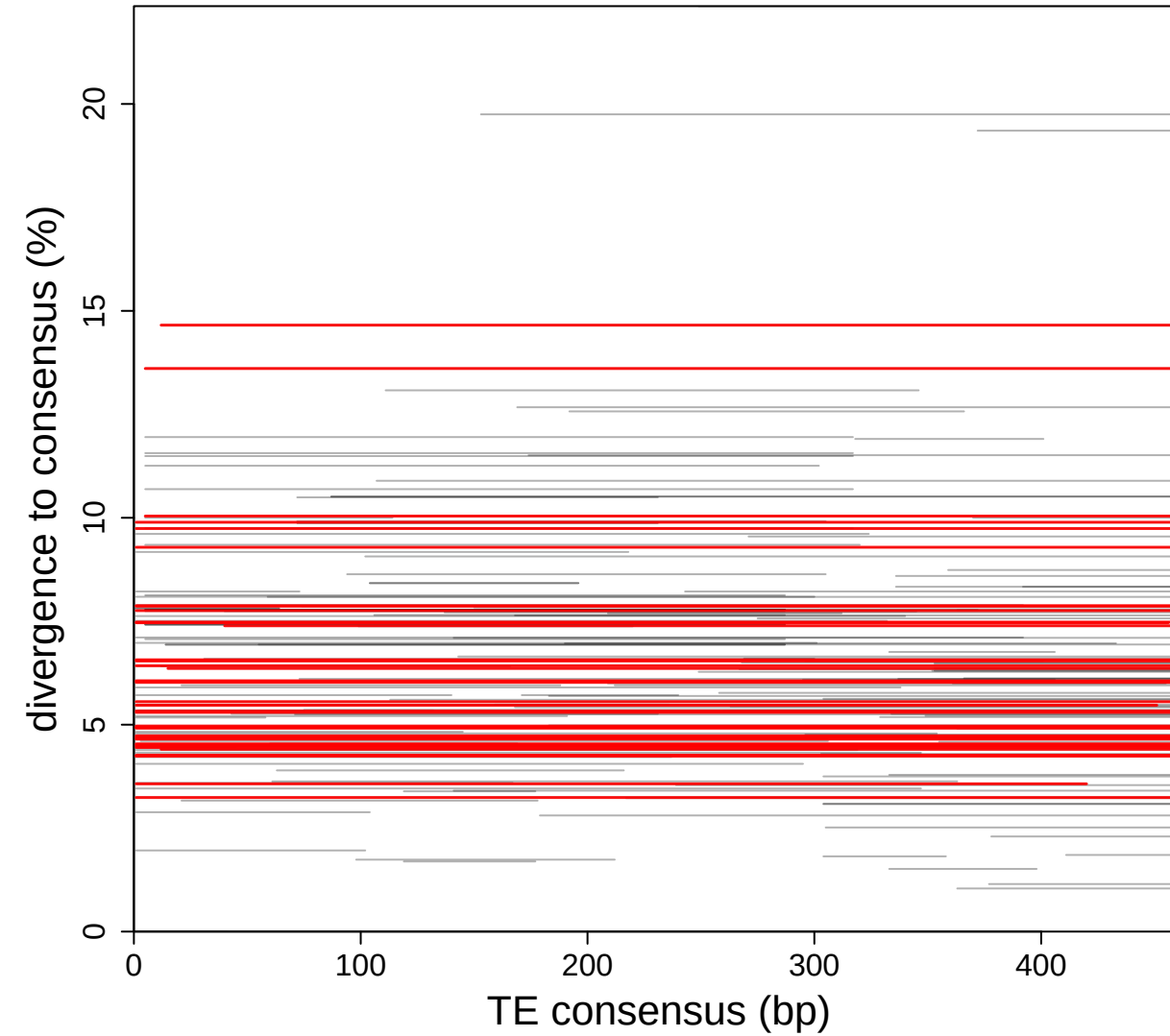


26.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 2347bp; fragments: 229; full length: 0 (>=2112.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

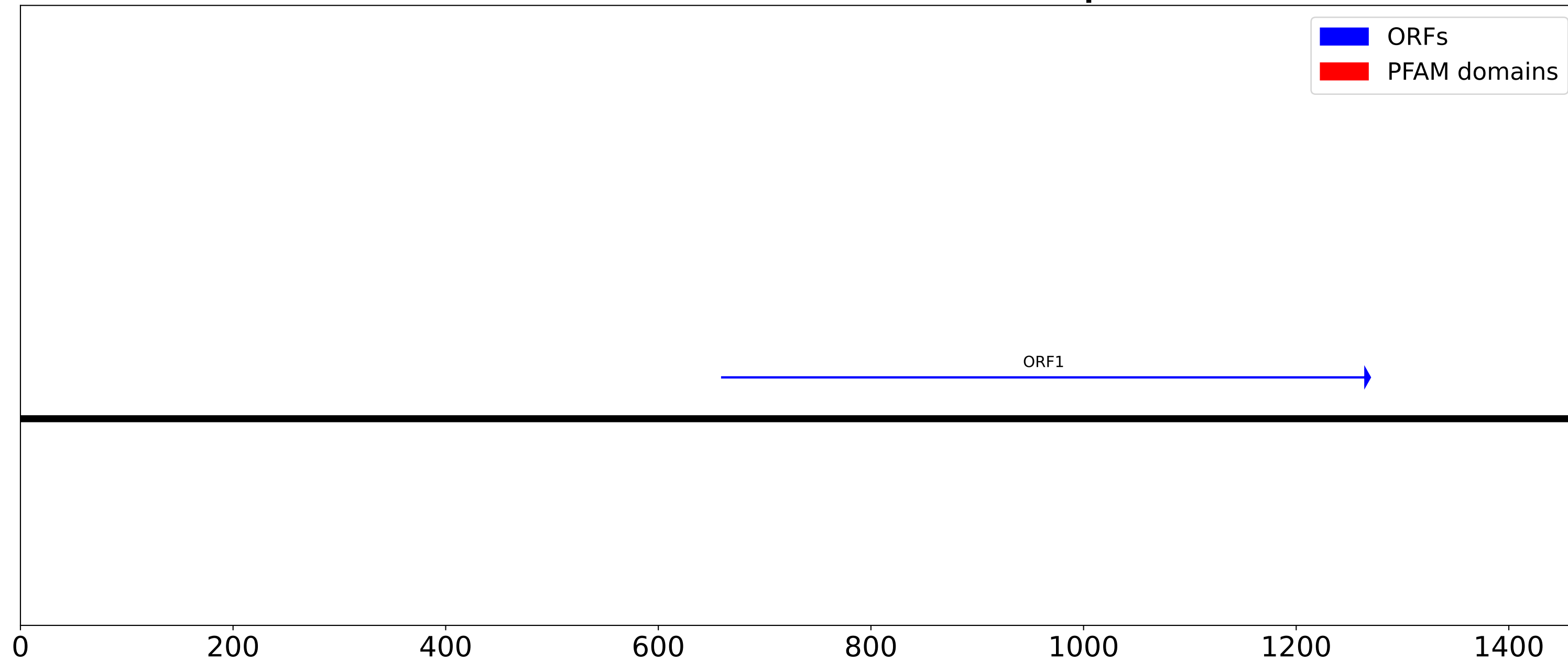


TE: rnd_5_family_10926
size: 463bp; fragments: 214; full length: 44 (≥ 416.7 bp)

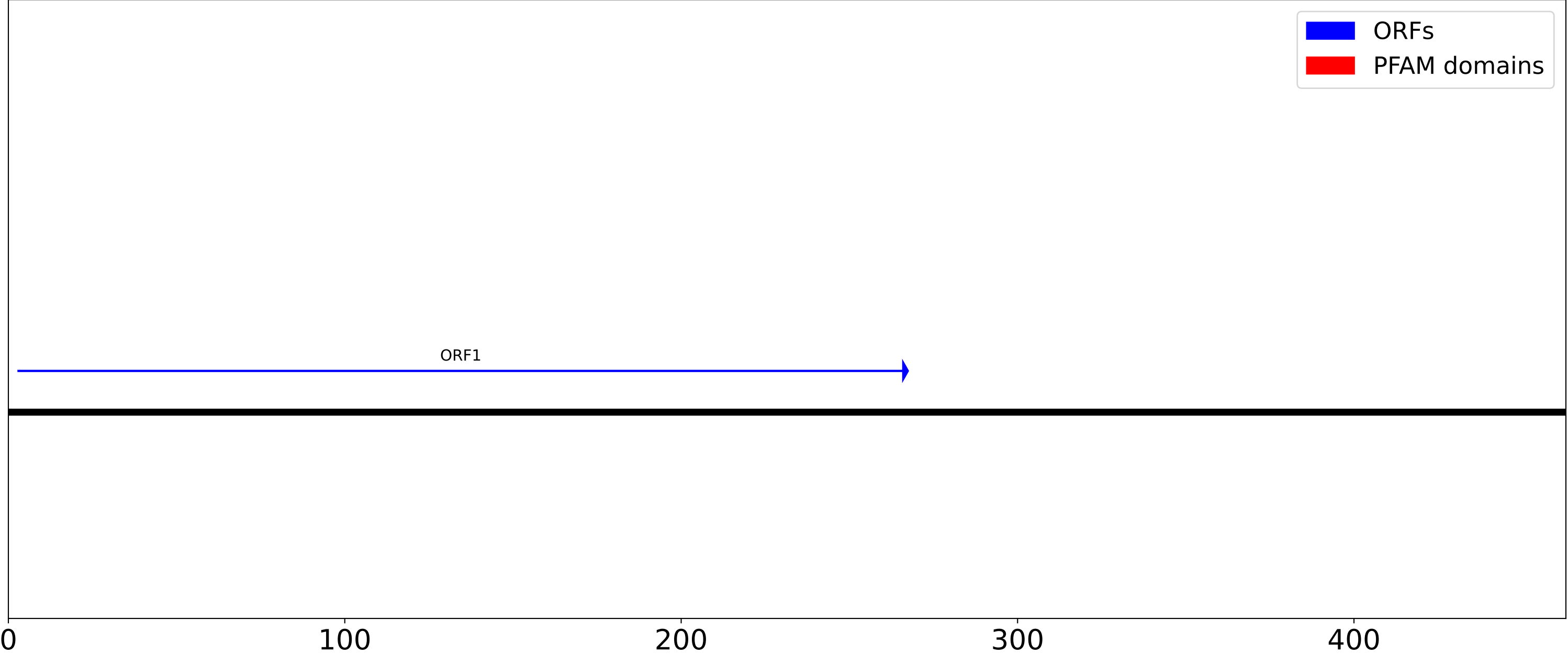


Before TEtrimmer 463 bp

After TEtrimmer ORF and PFAM domain plot



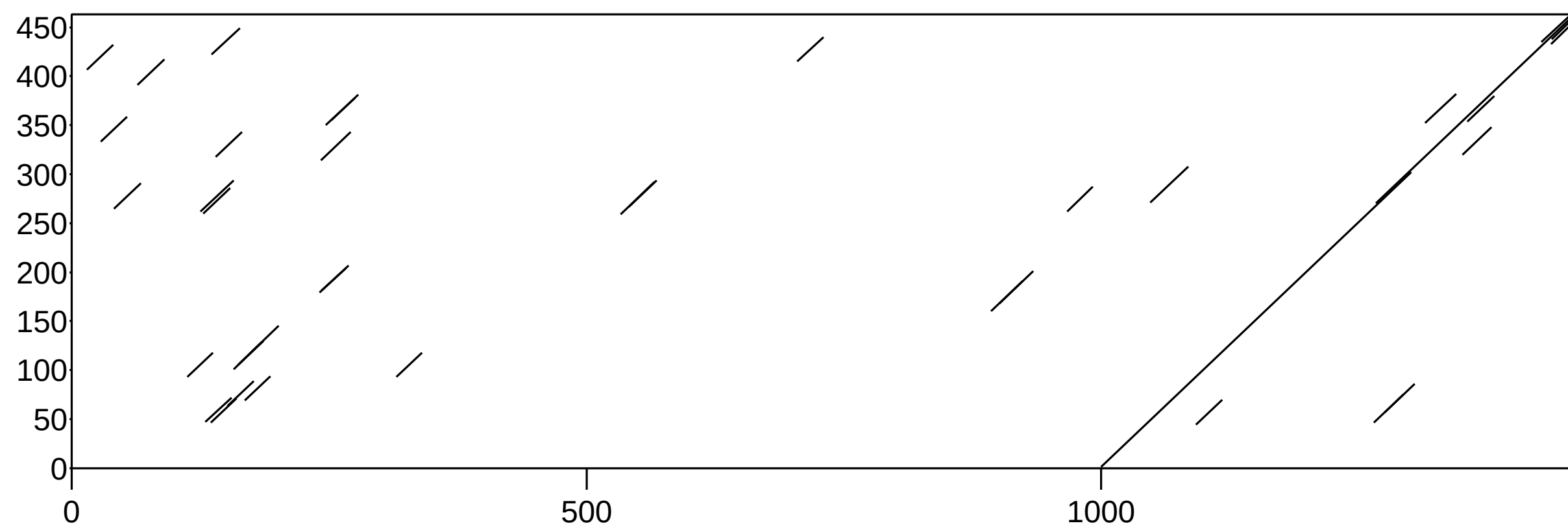
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)