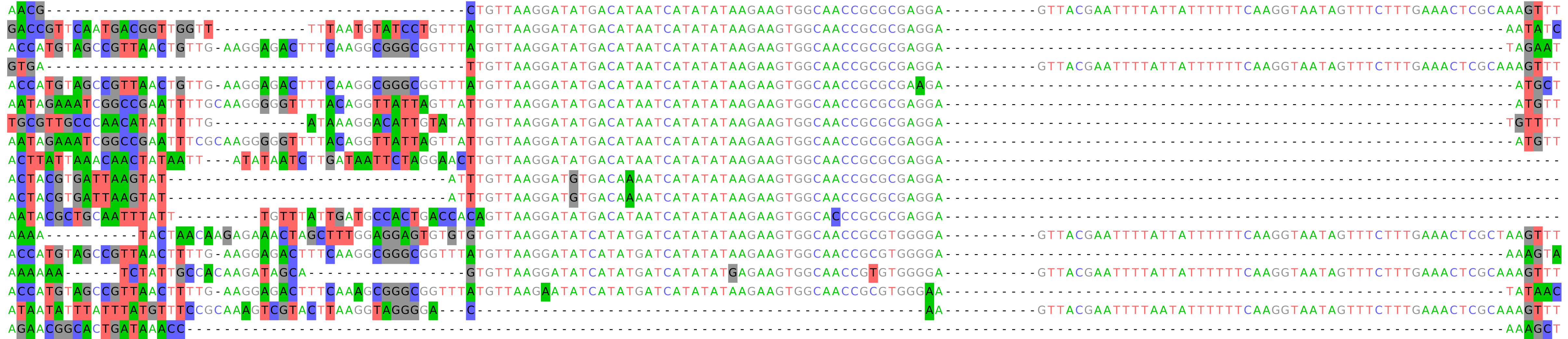
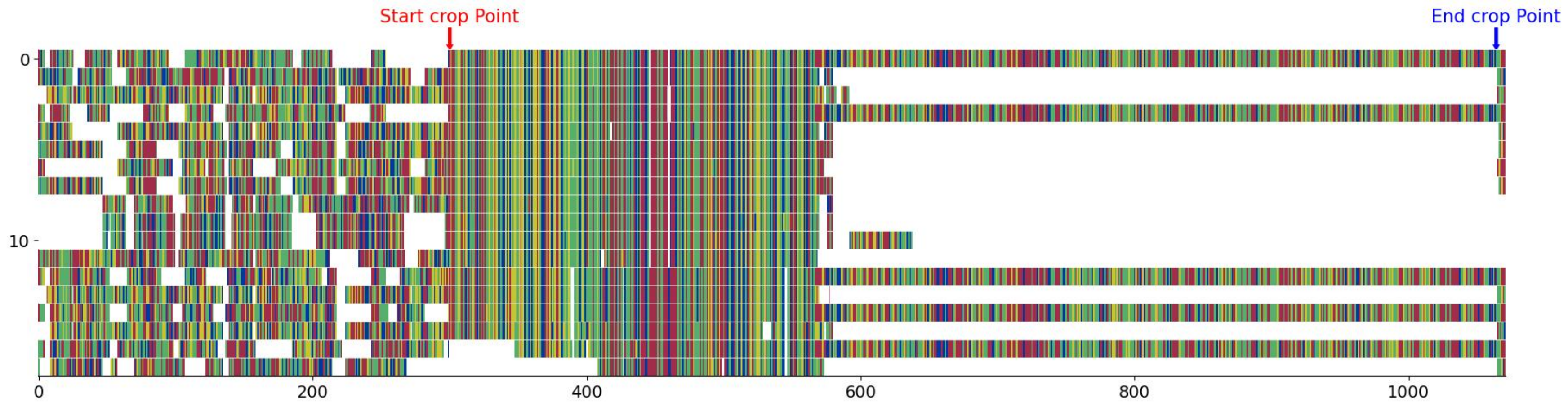


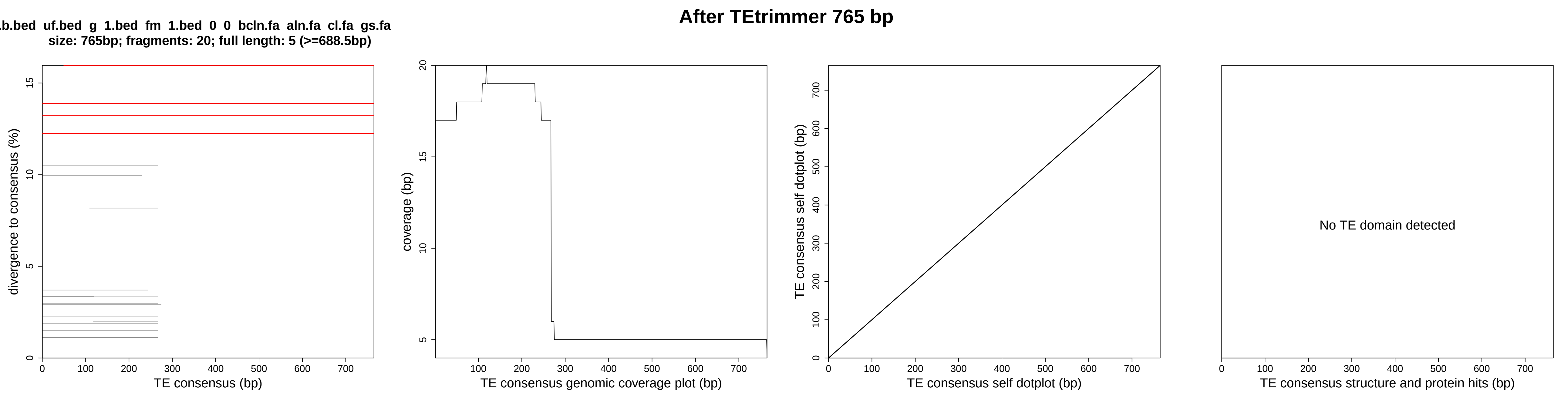
Start crop Point

End crop Point

MSA length = 765

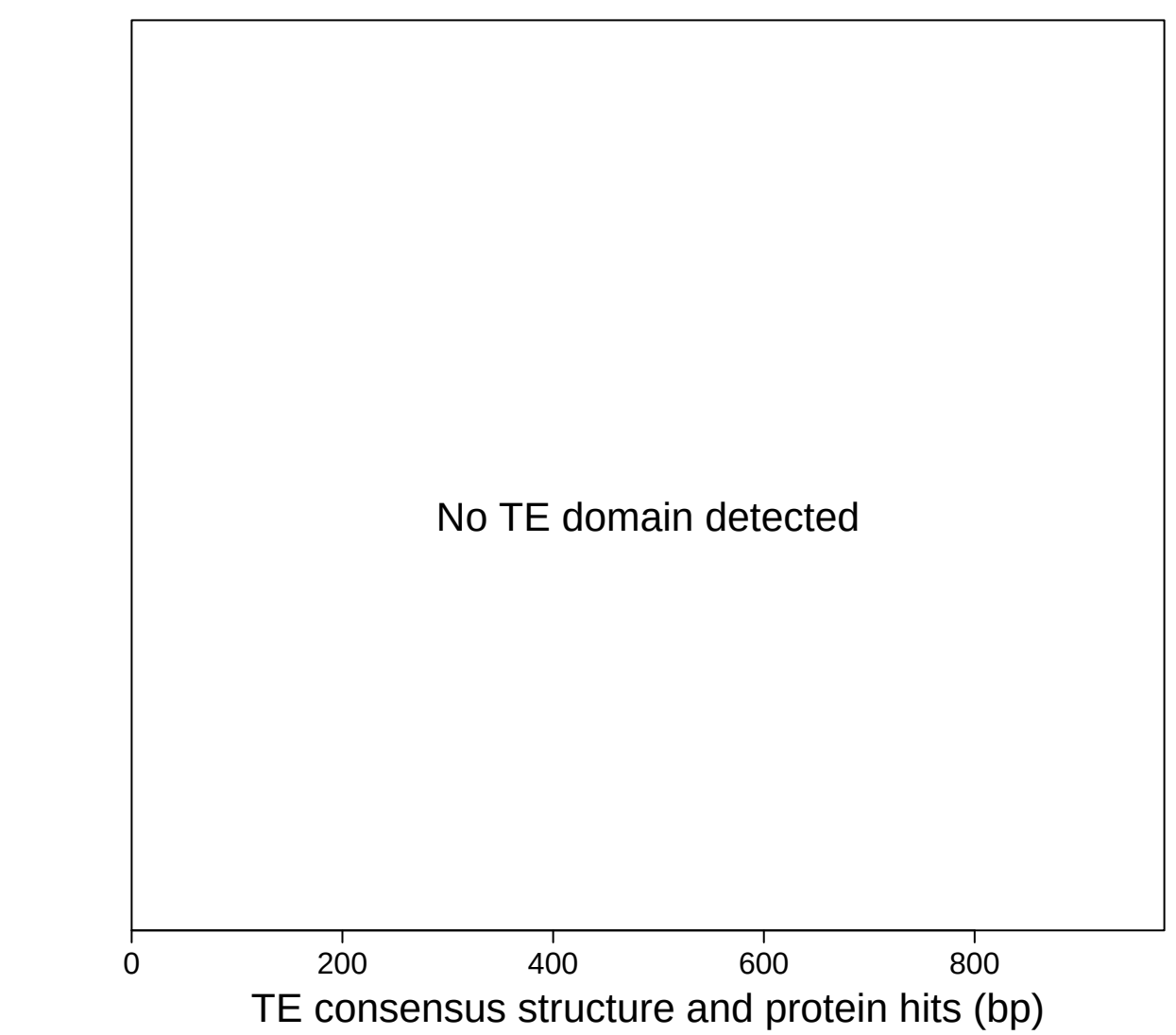
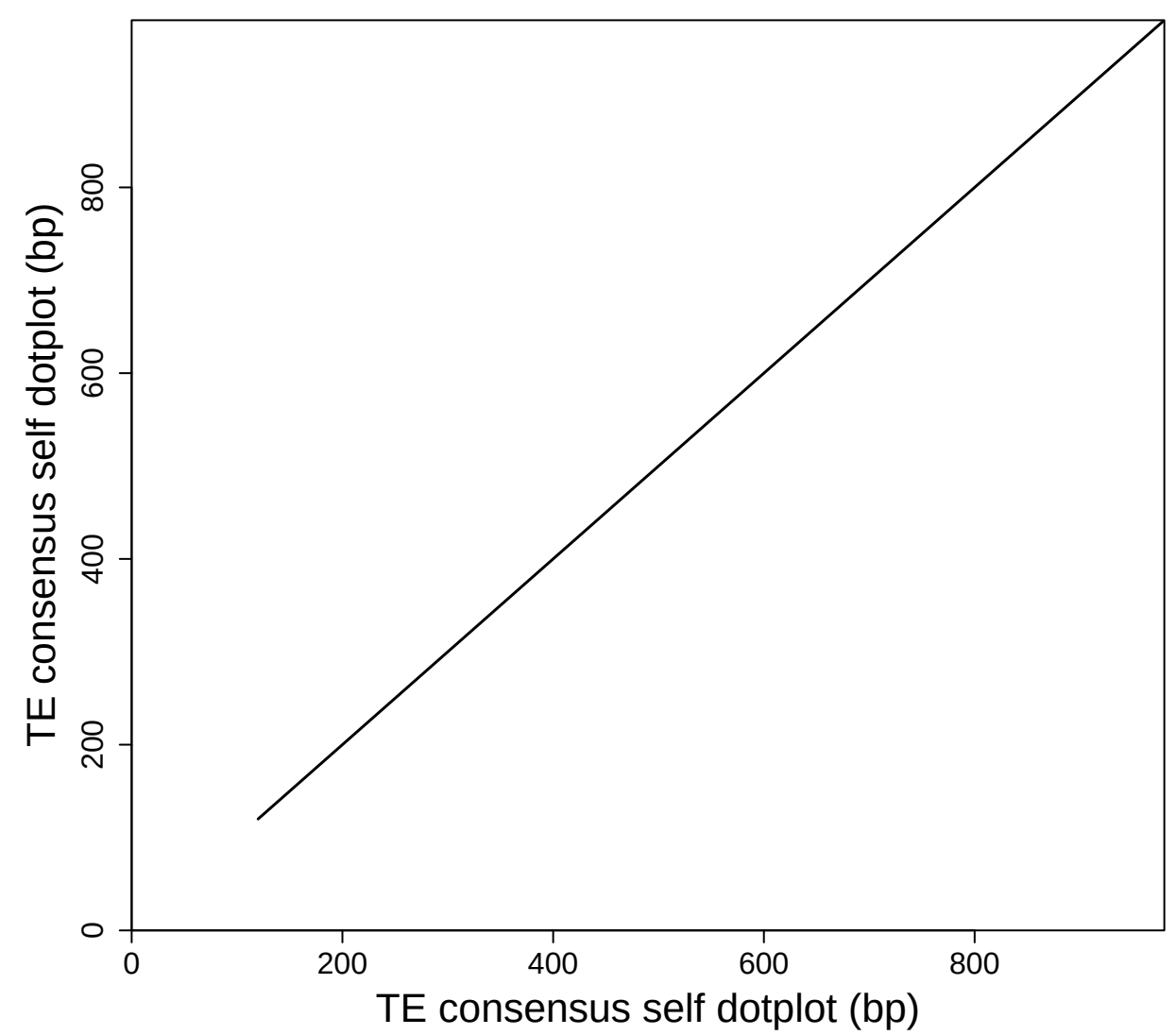
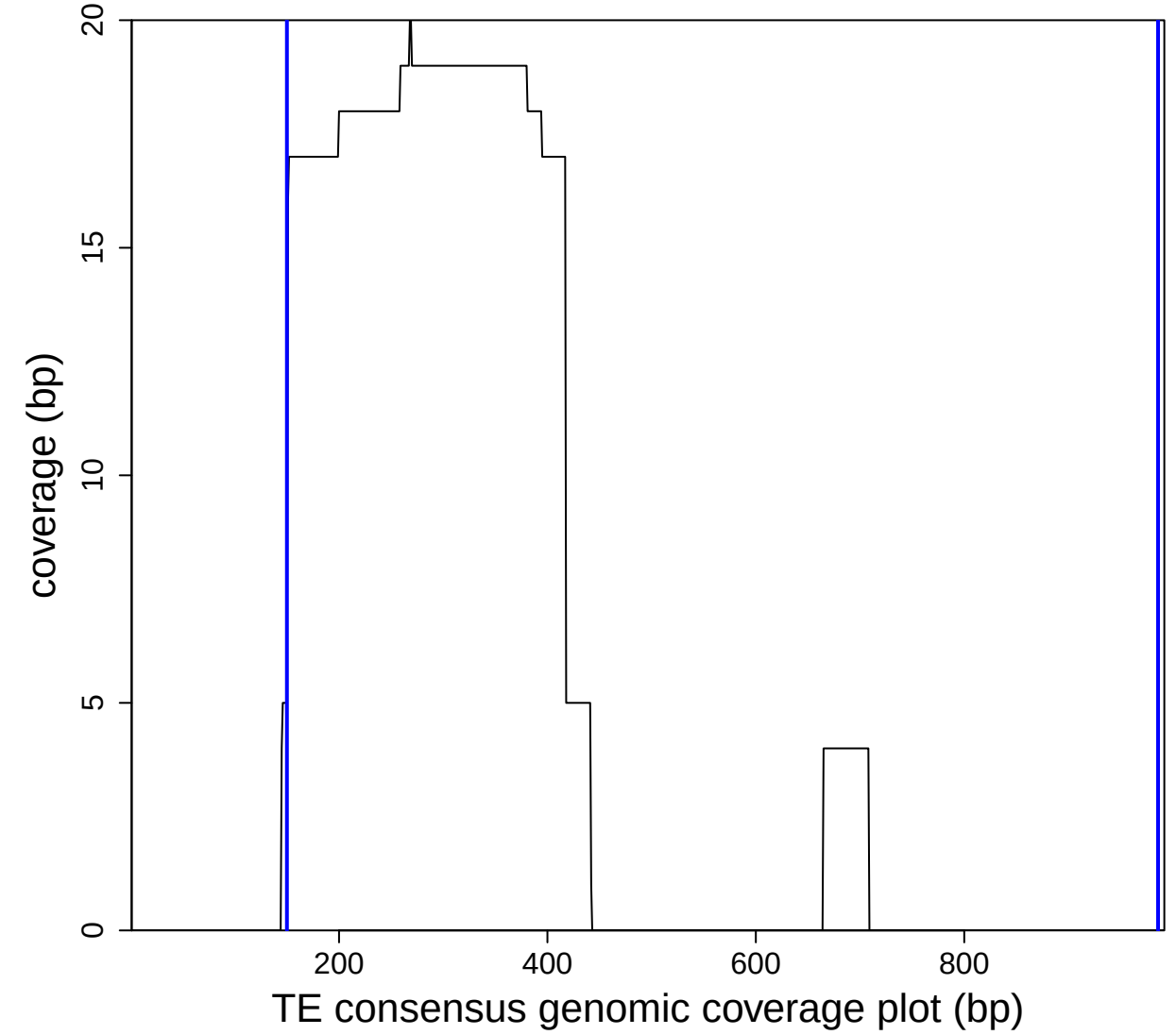
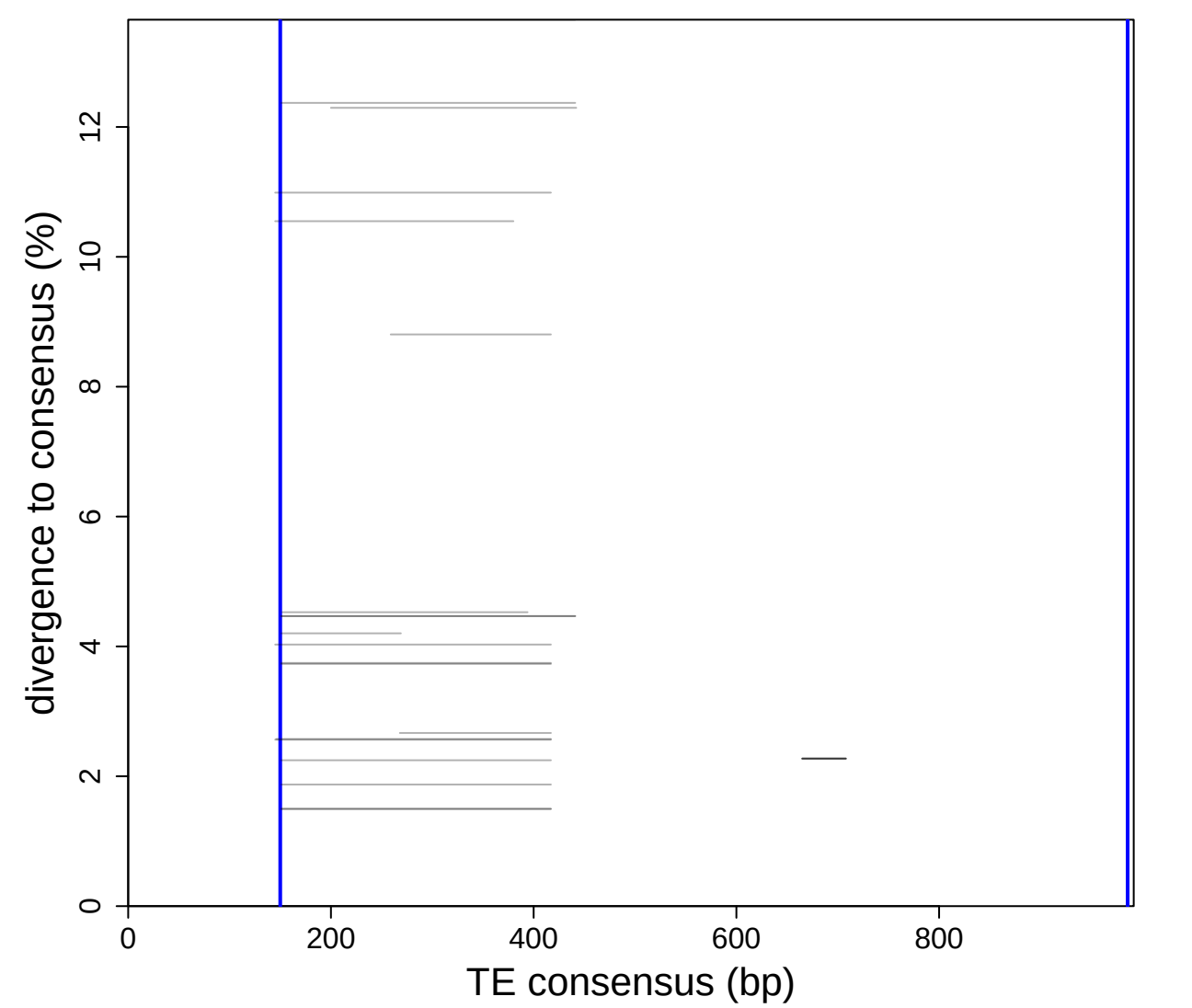




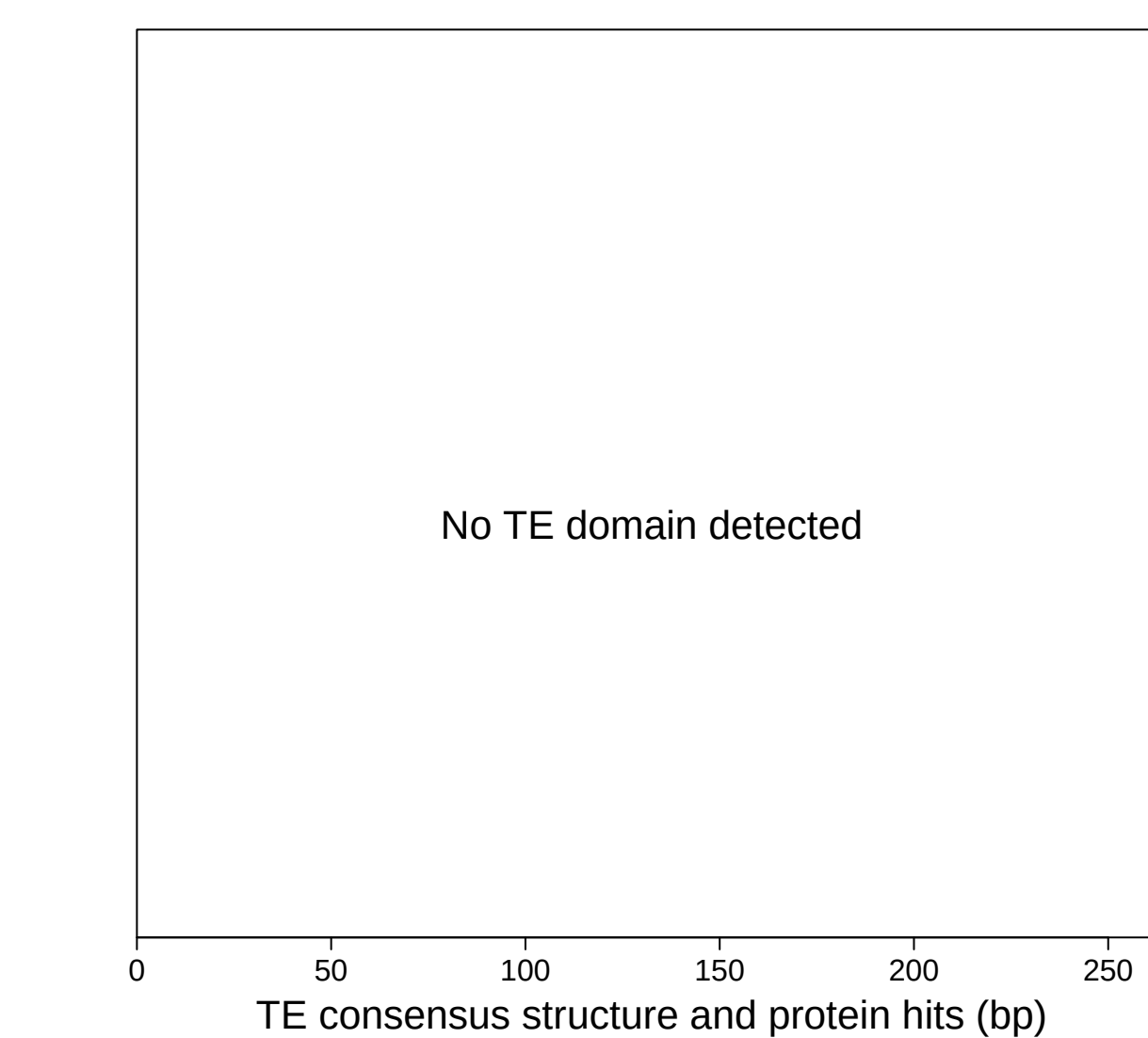
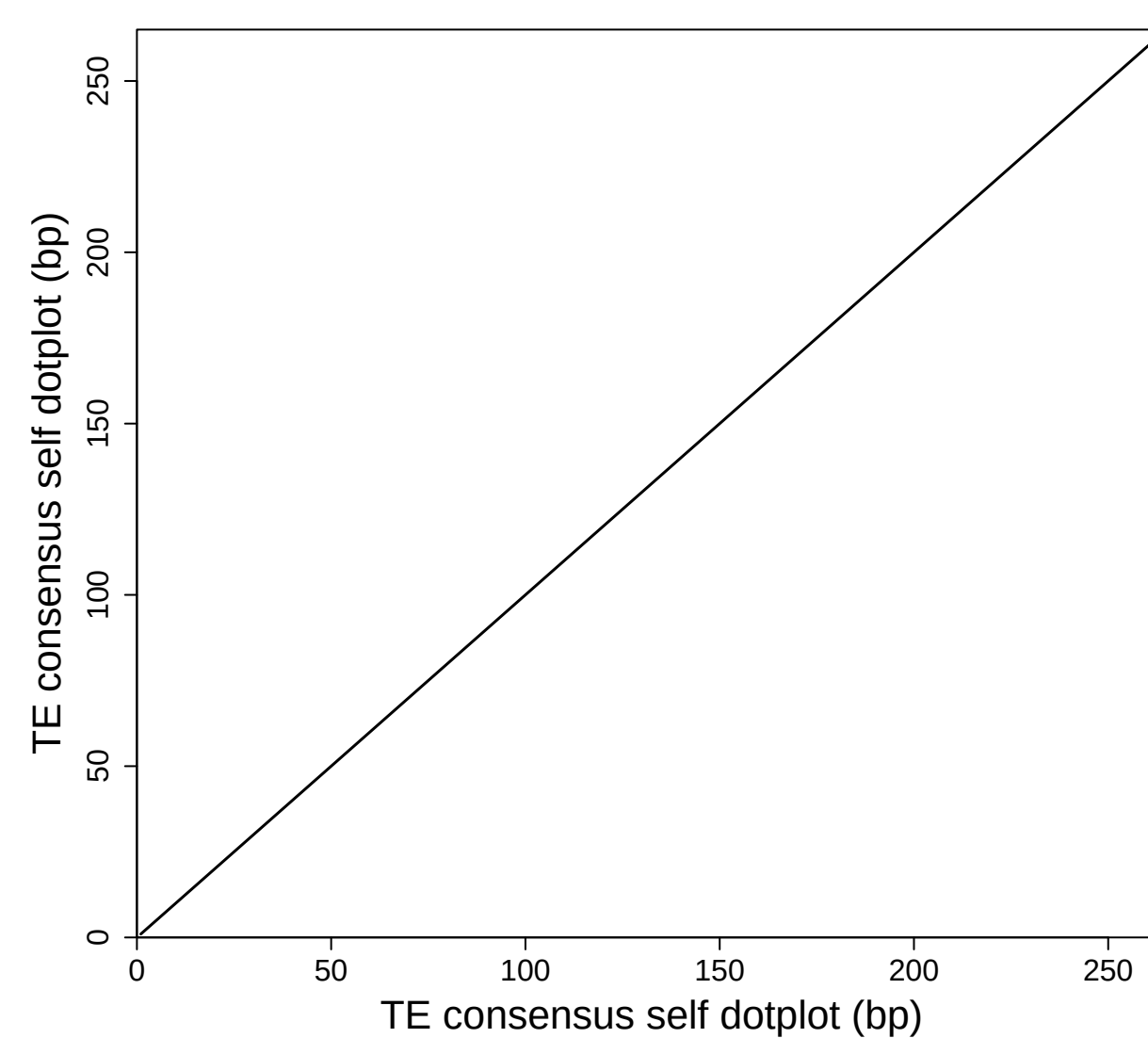
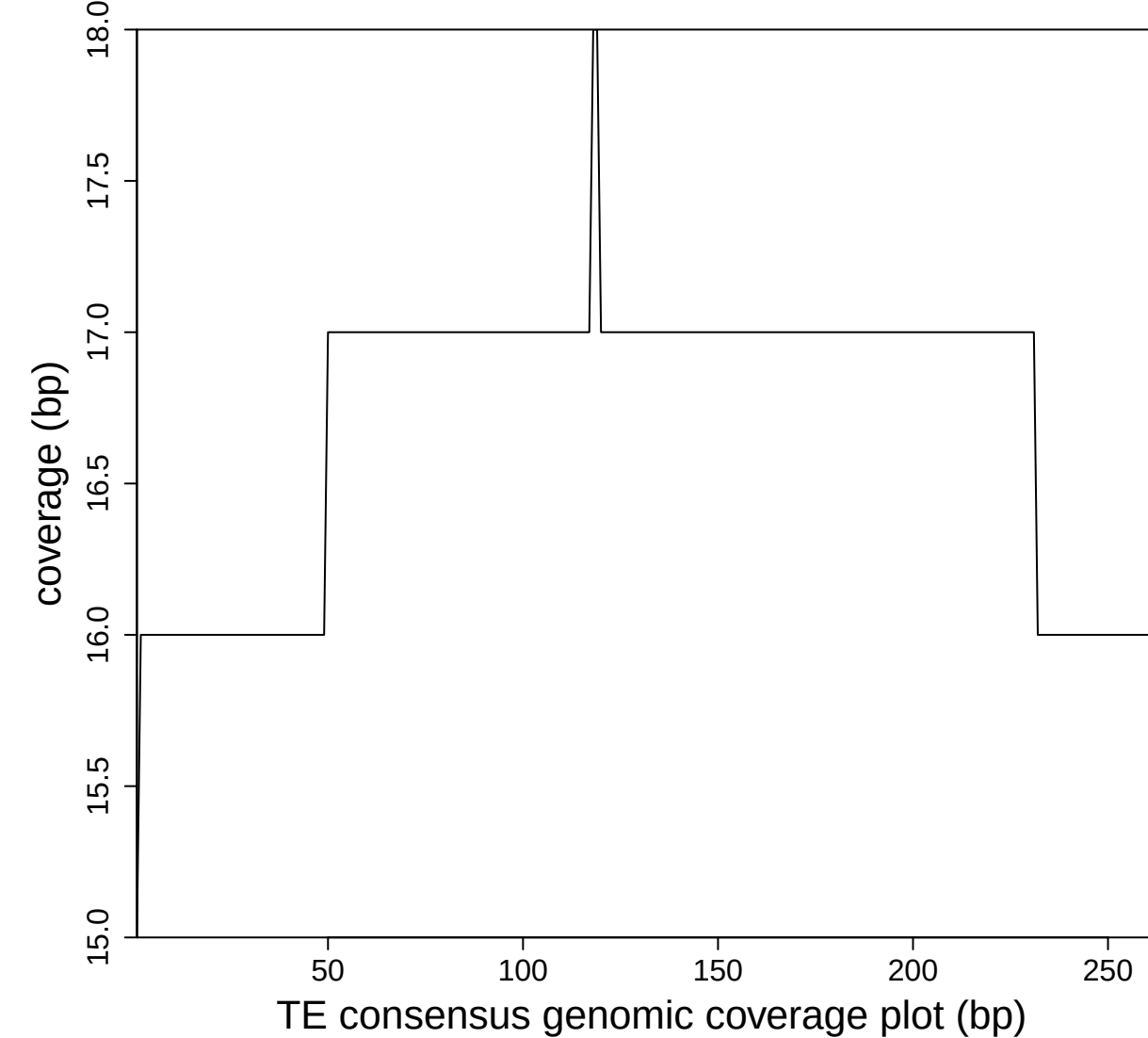
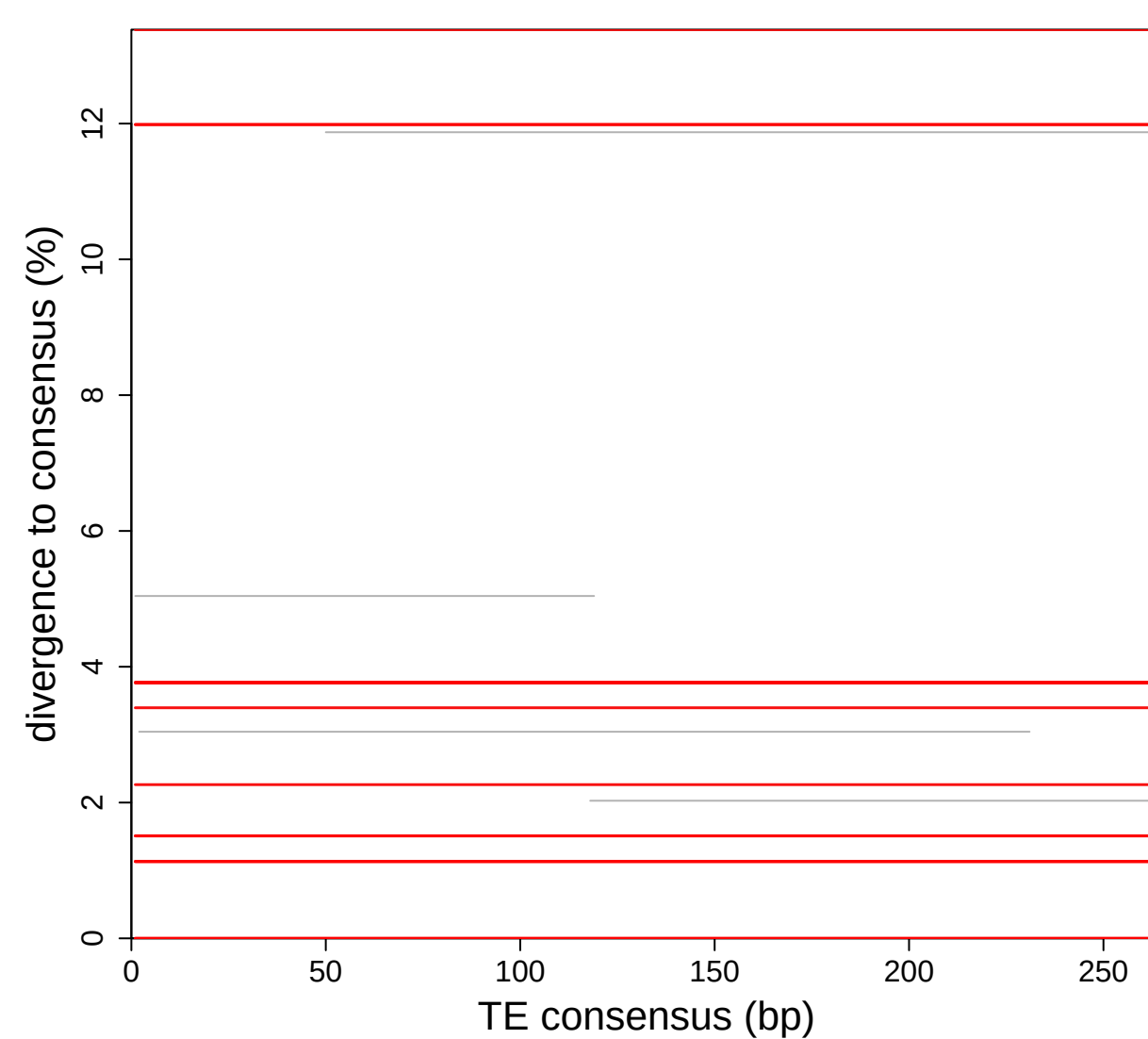


.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 992bp; fragments: 24; full length: 0 (>=892.8bp)

After TEtrimmer Extended plot Blue lines are boundaries

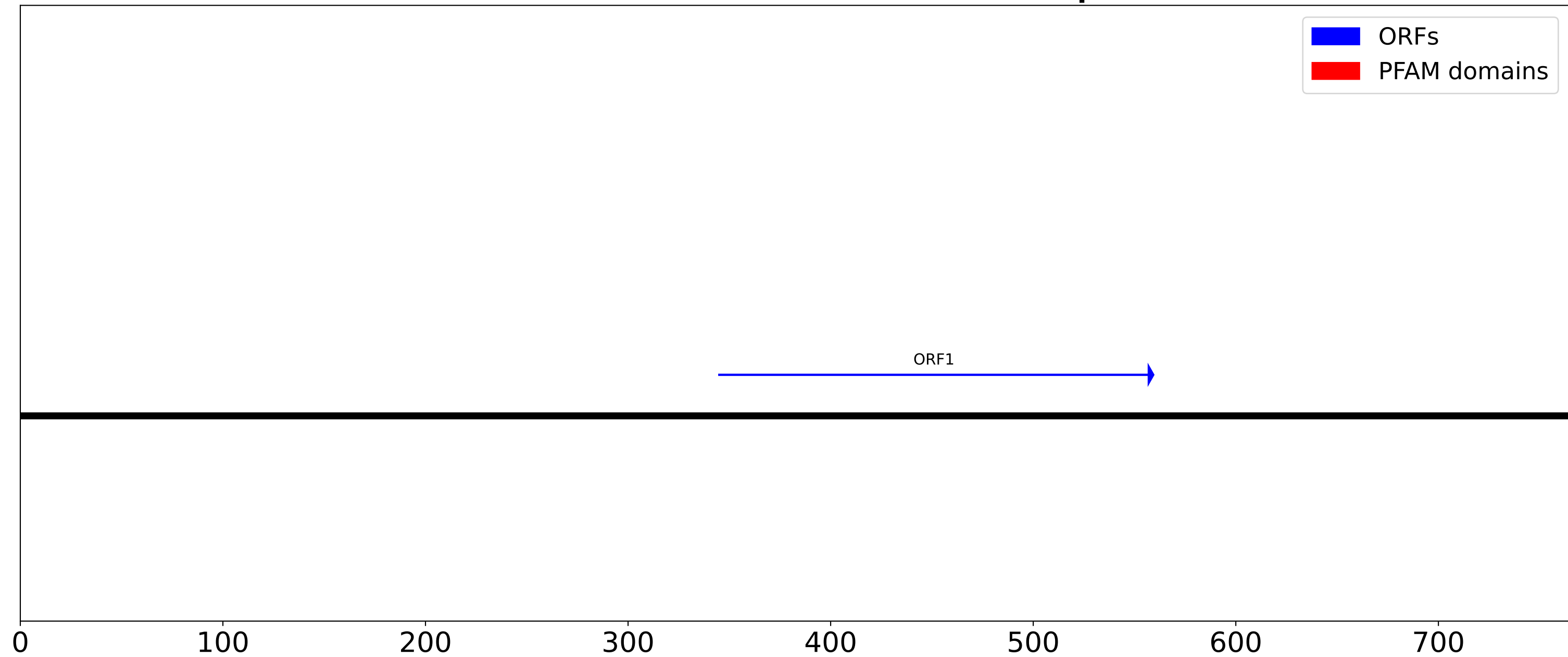


TE: ltr_1_family_395
size: 265bp; fragments: 18; full length: 14 (≥ 238.5 bp)



Before TEtrimmer 265 bp

After TEtrimmer ORF and PFAM domain plot



Dotplot

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

E consensus before TEtrimmer (bp)

