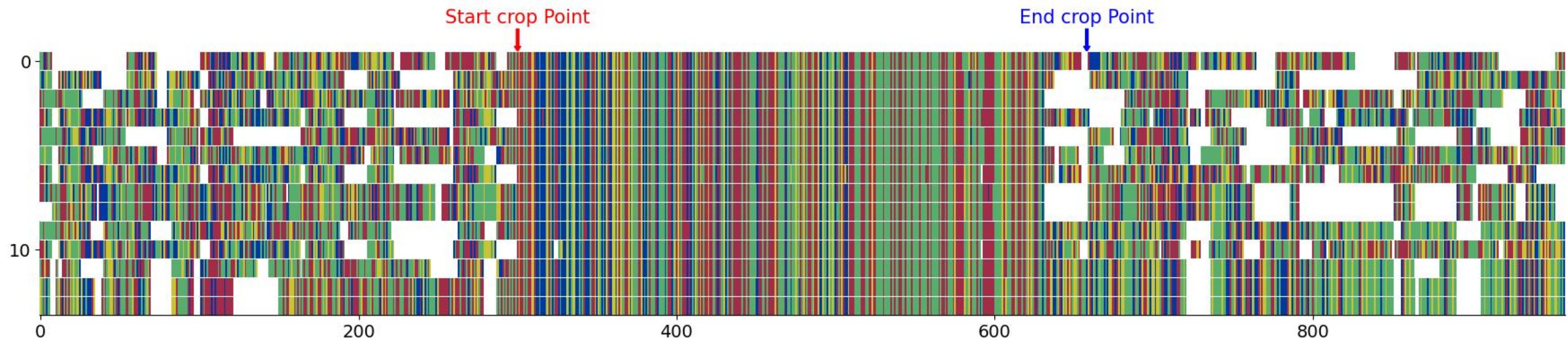
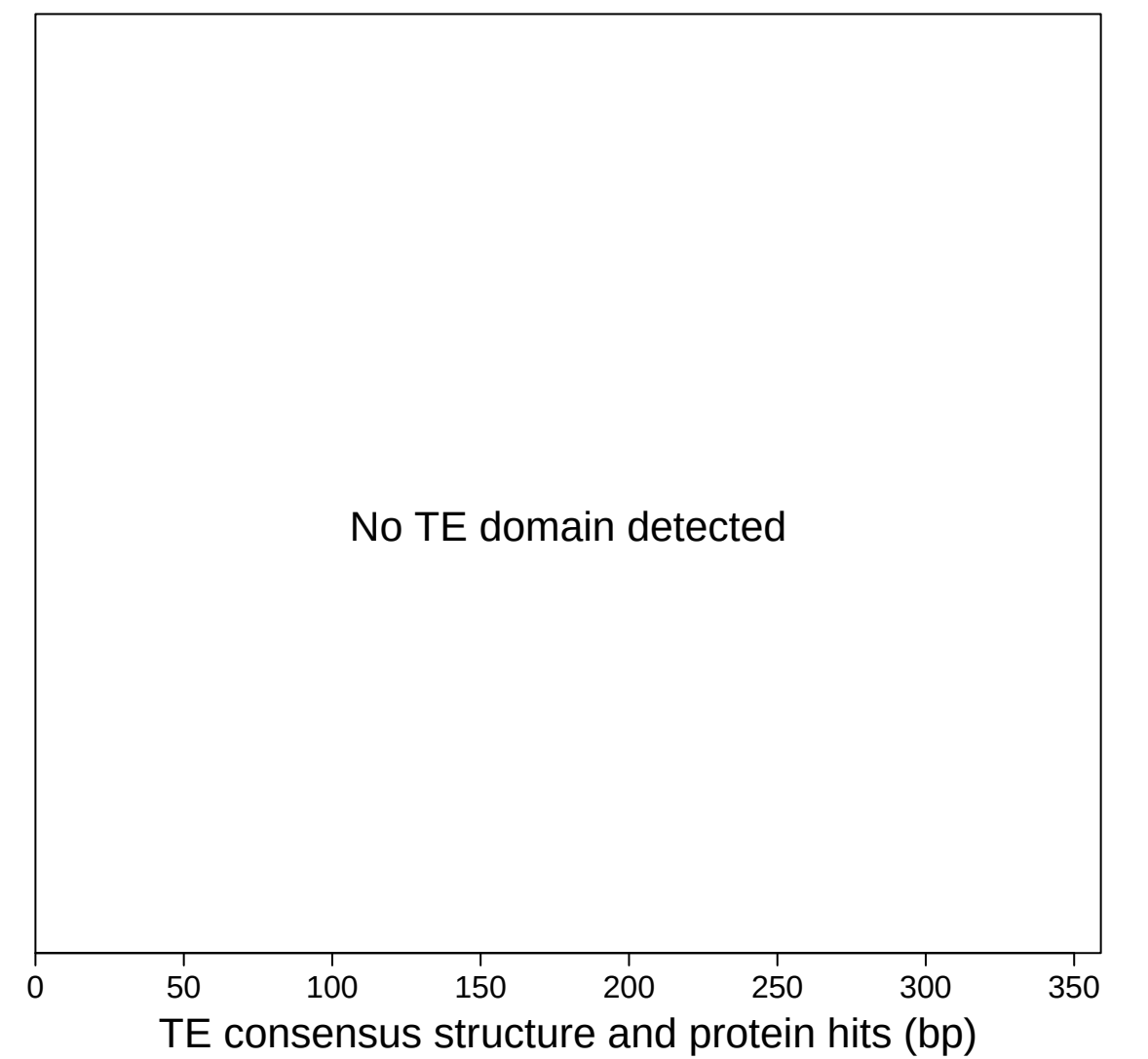
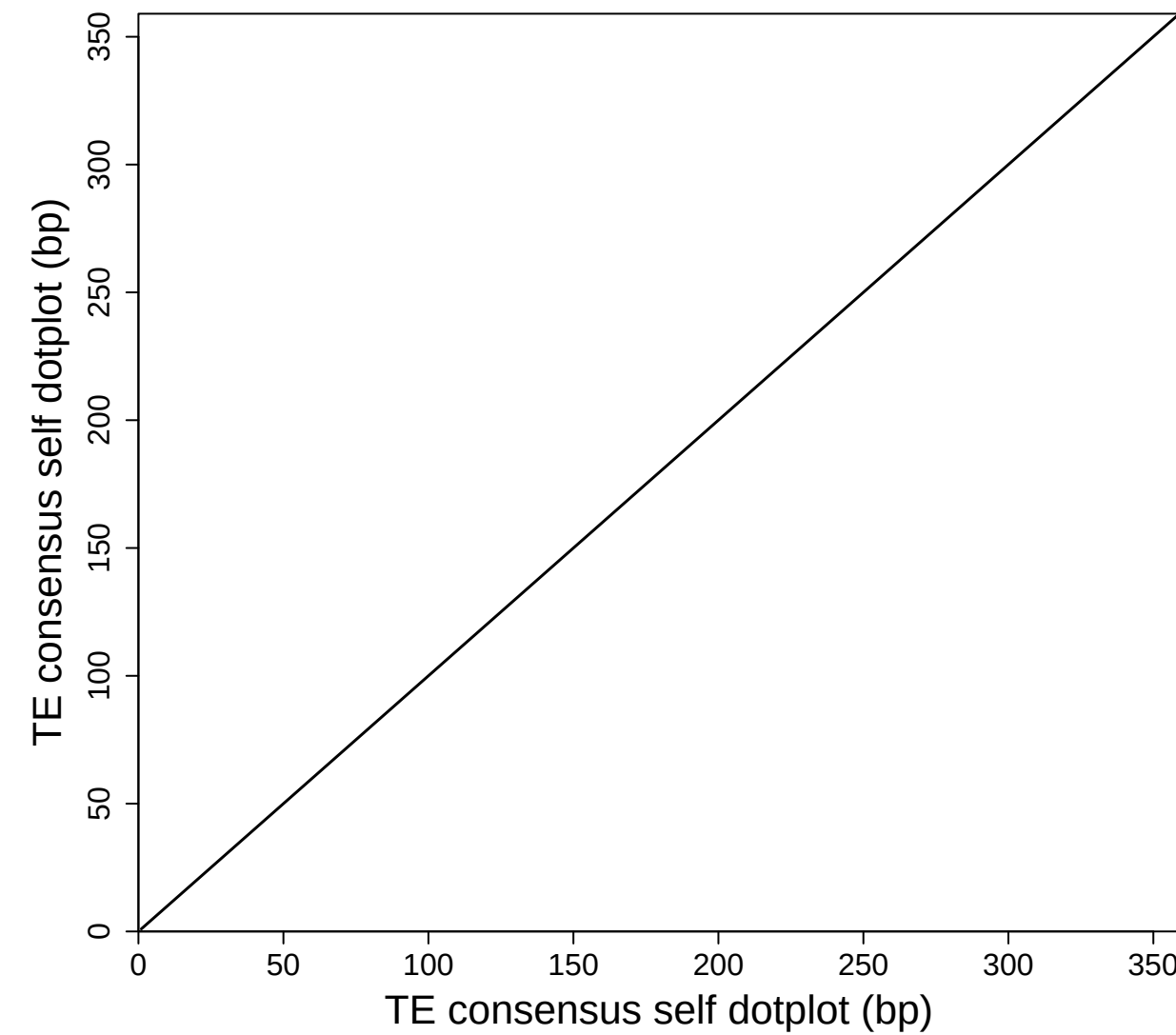
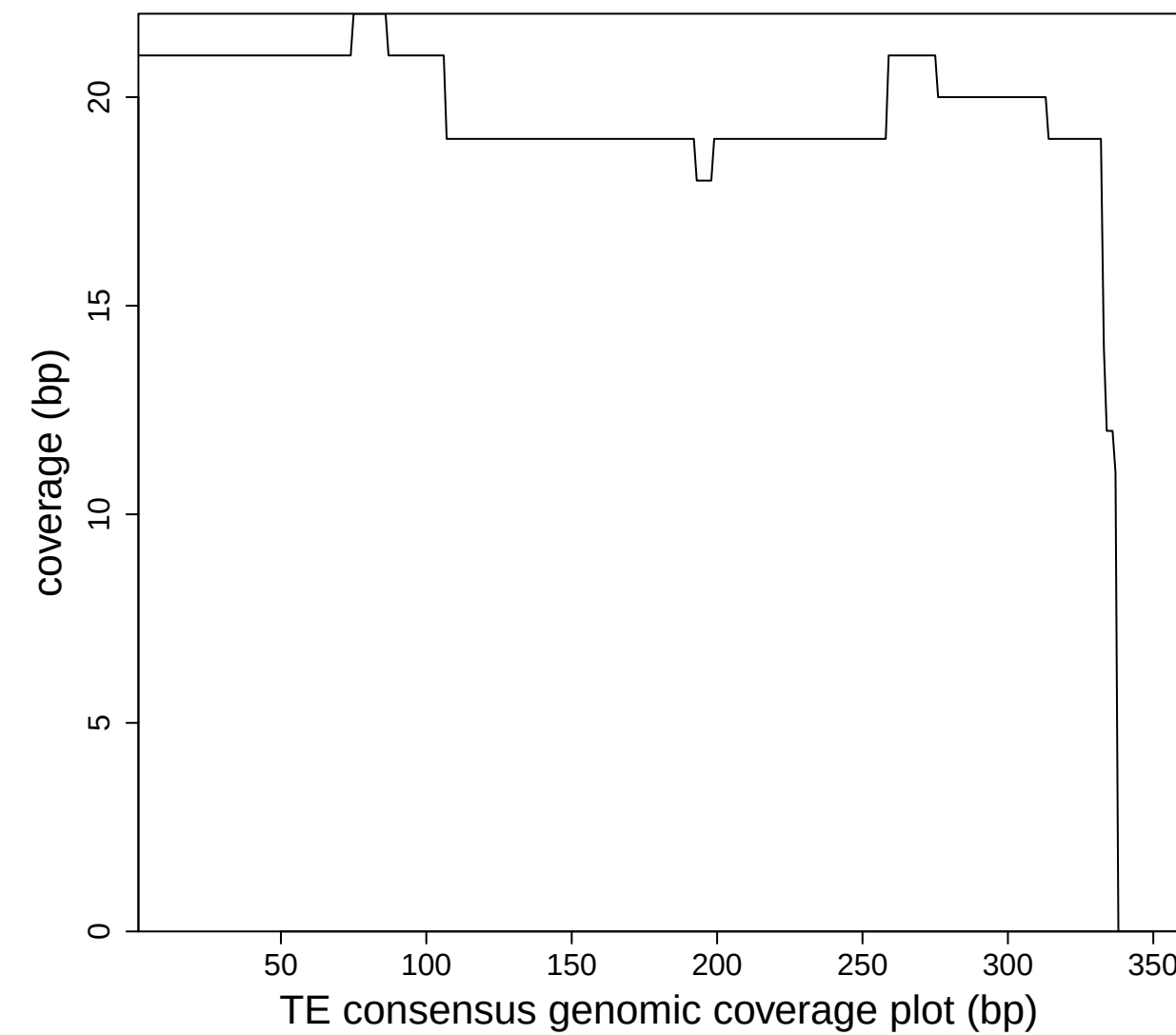
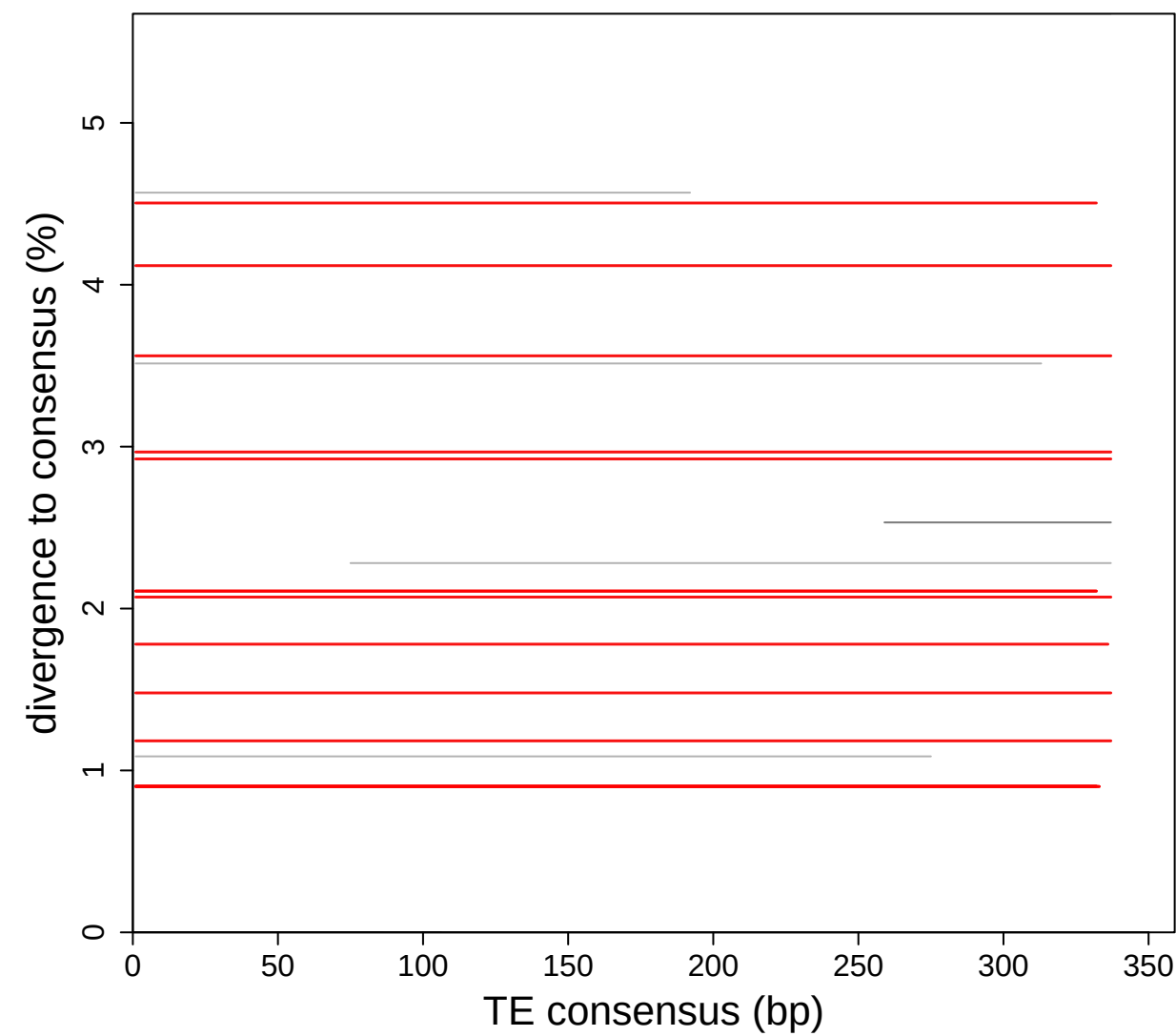




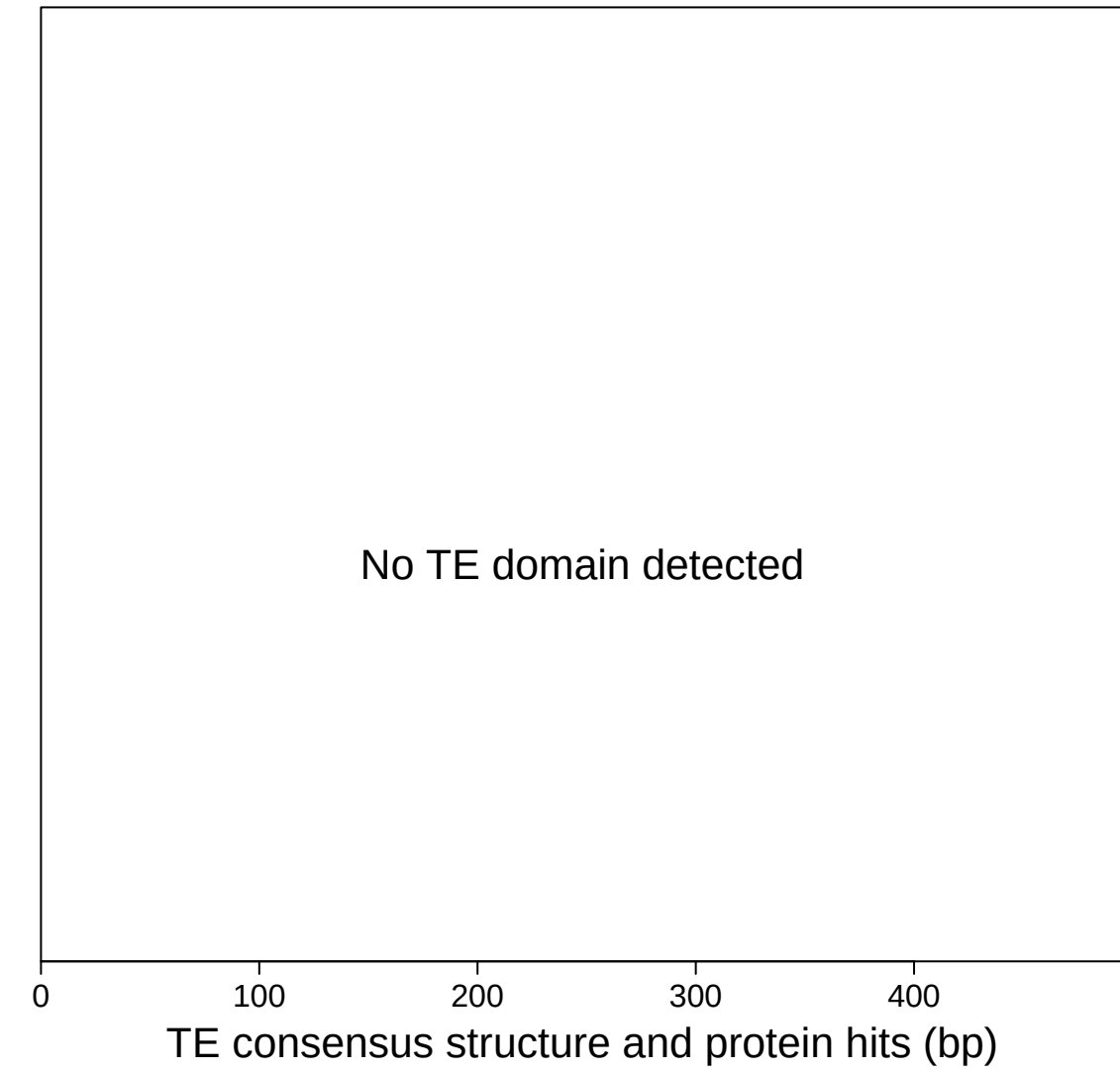
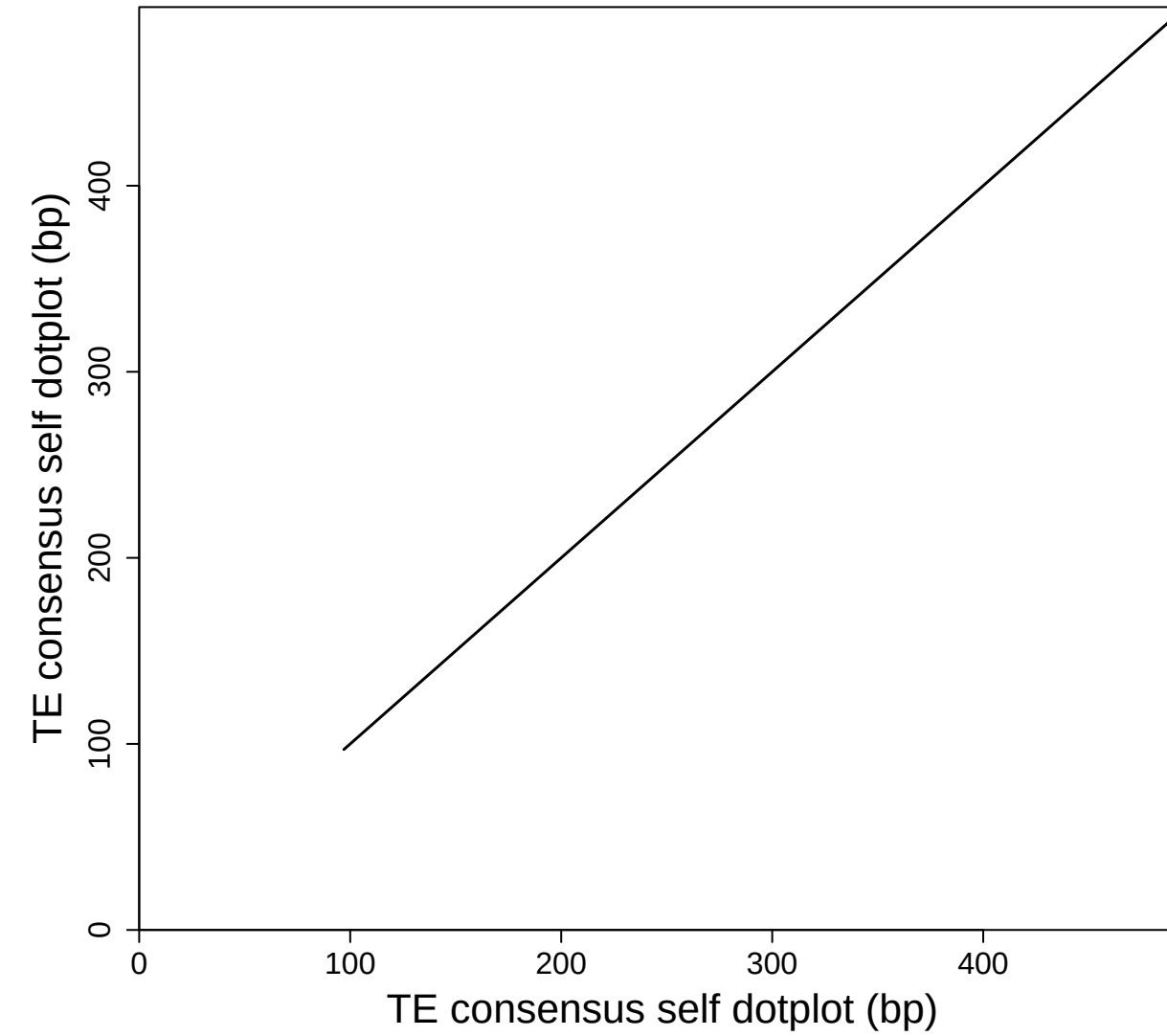
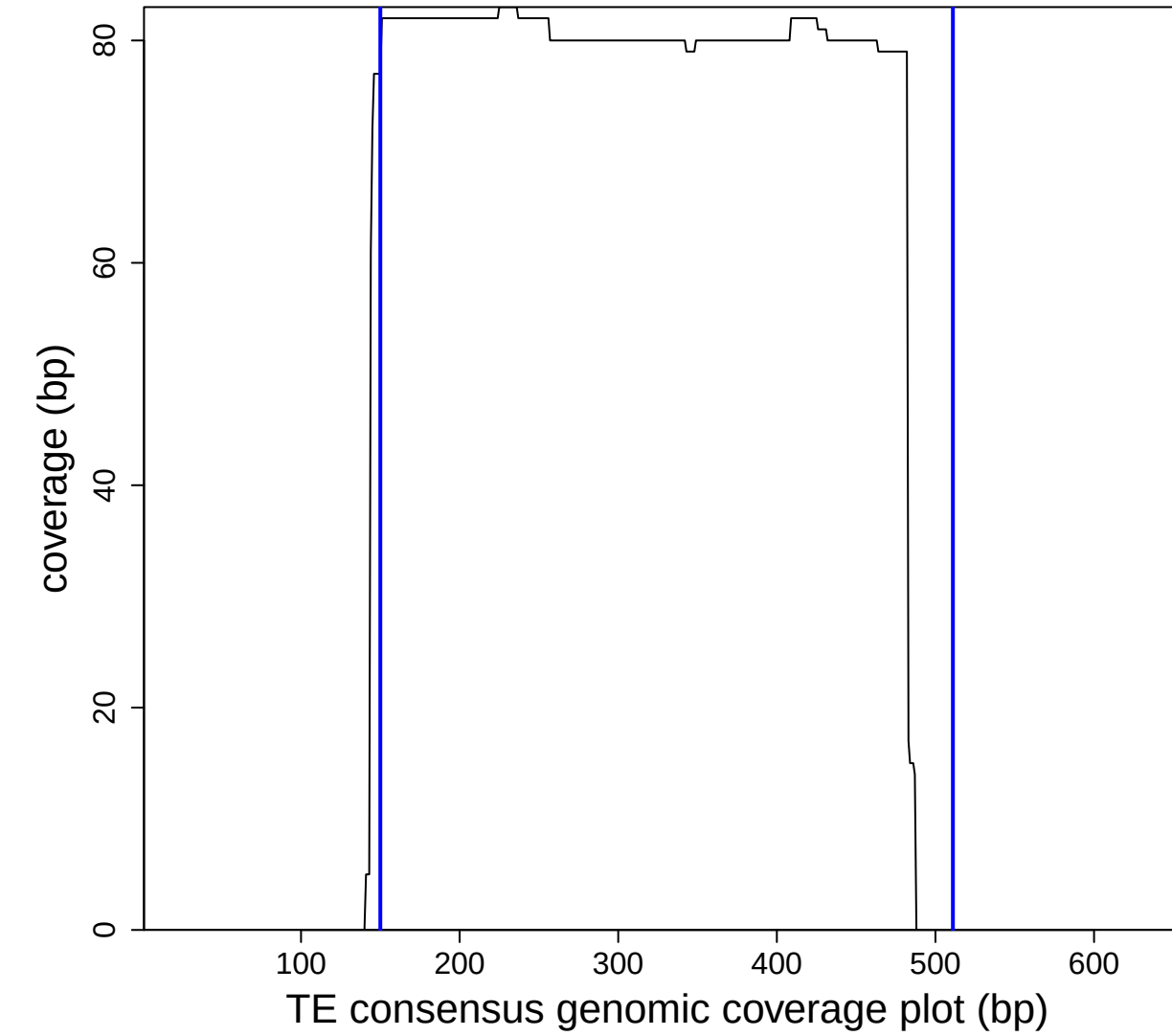
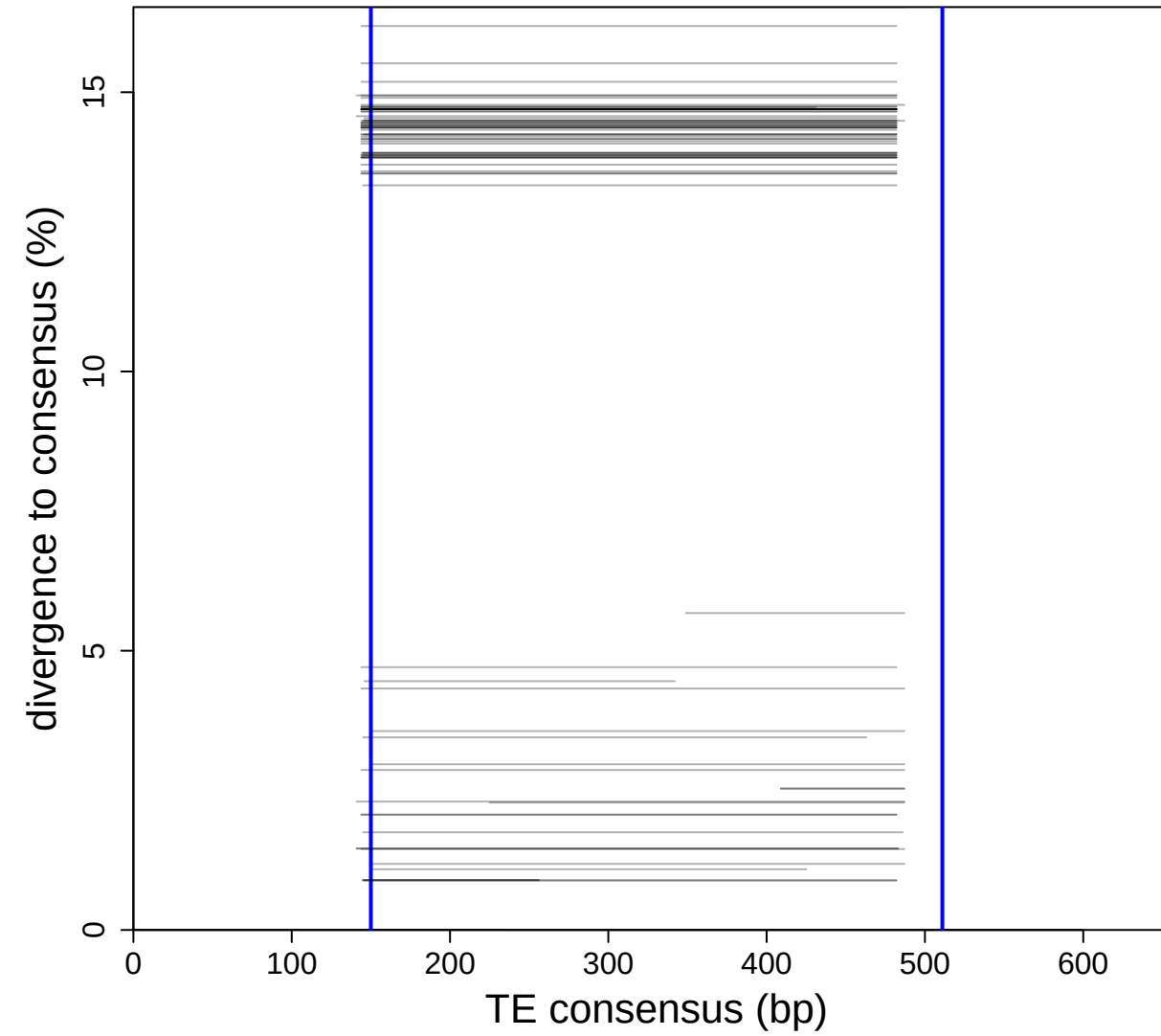
After TEtrimmer 359 bp

b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 359bp; fragments: 25; full length: 15 (≥ 323.1 bp)

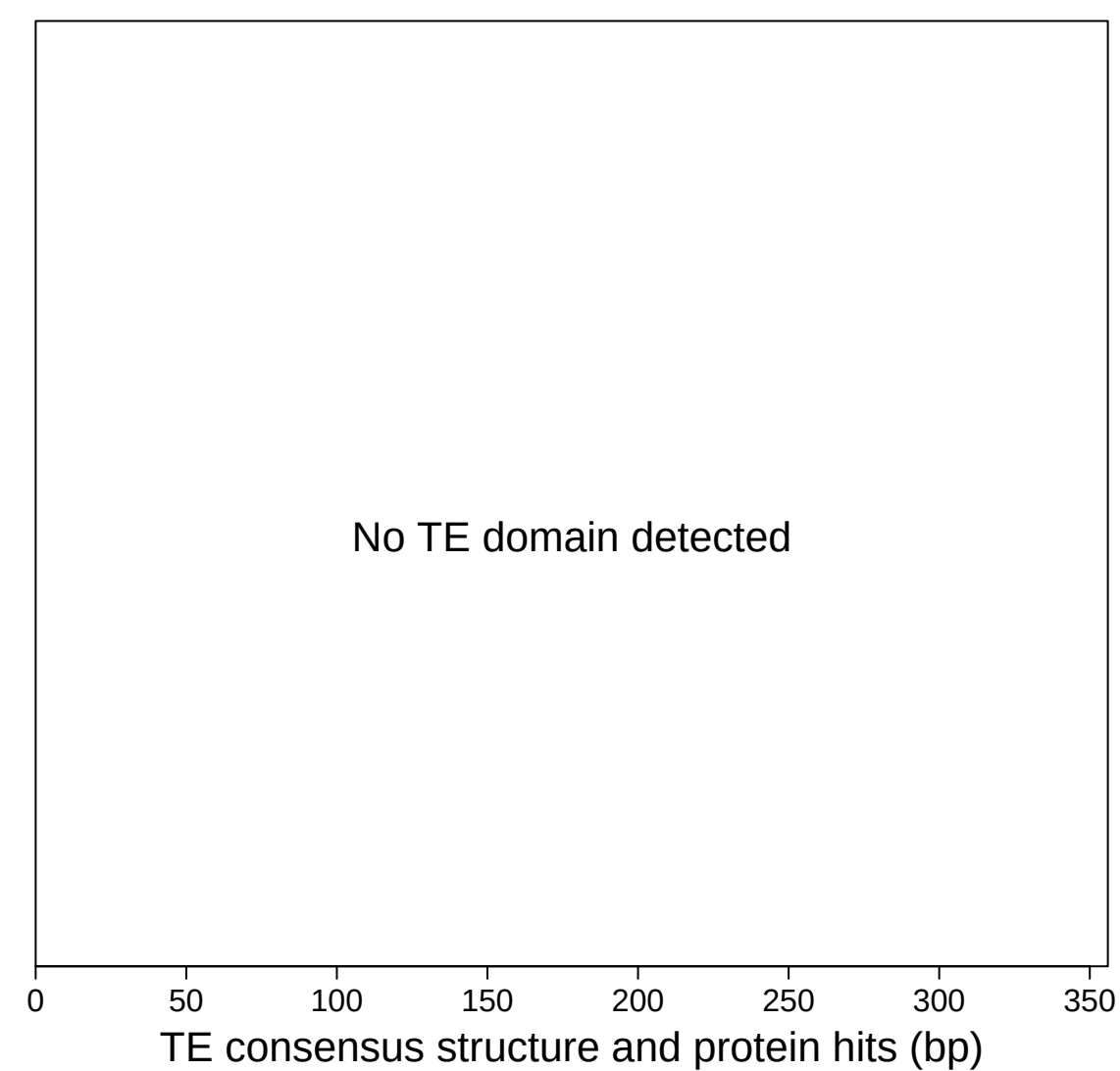
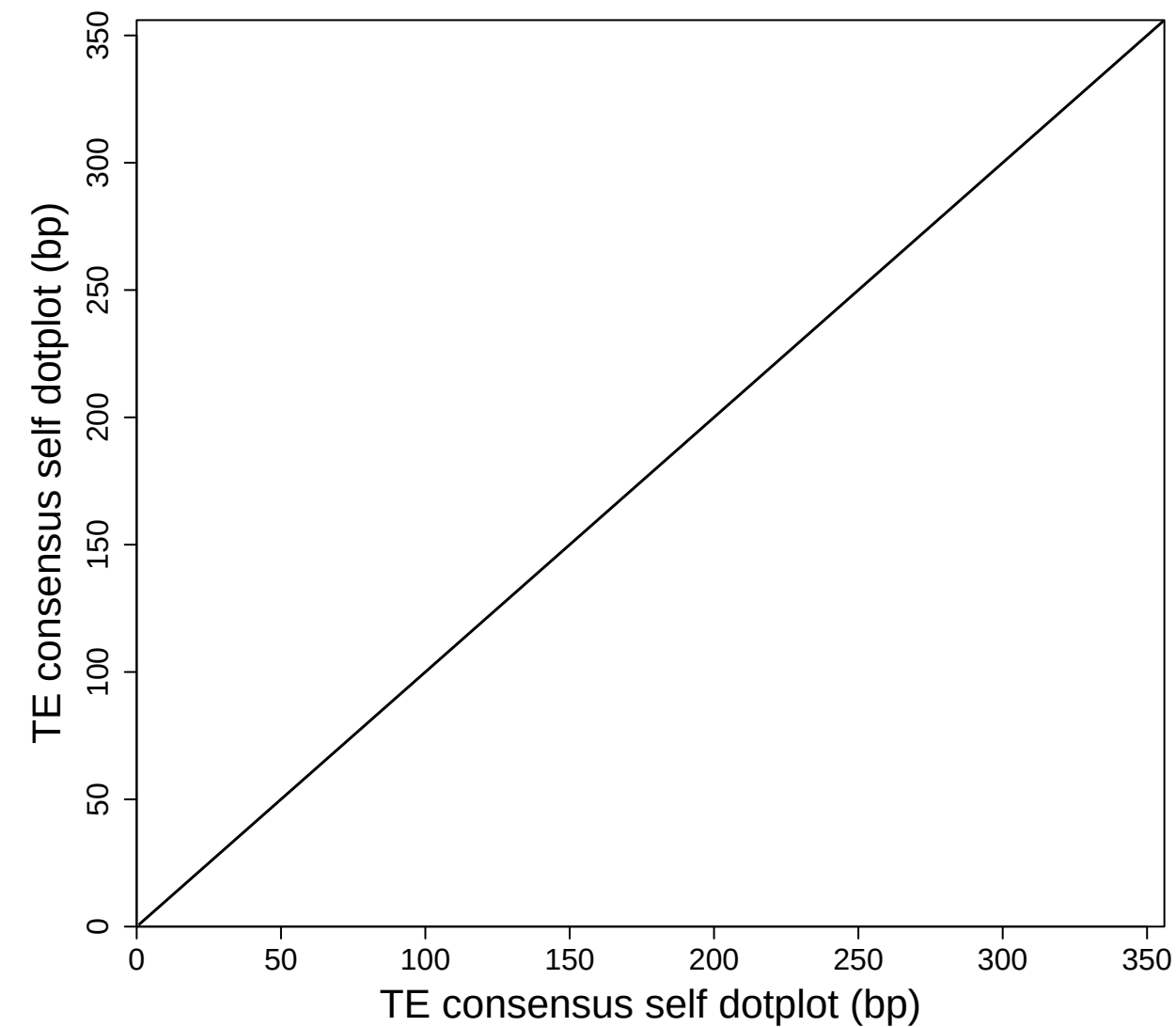
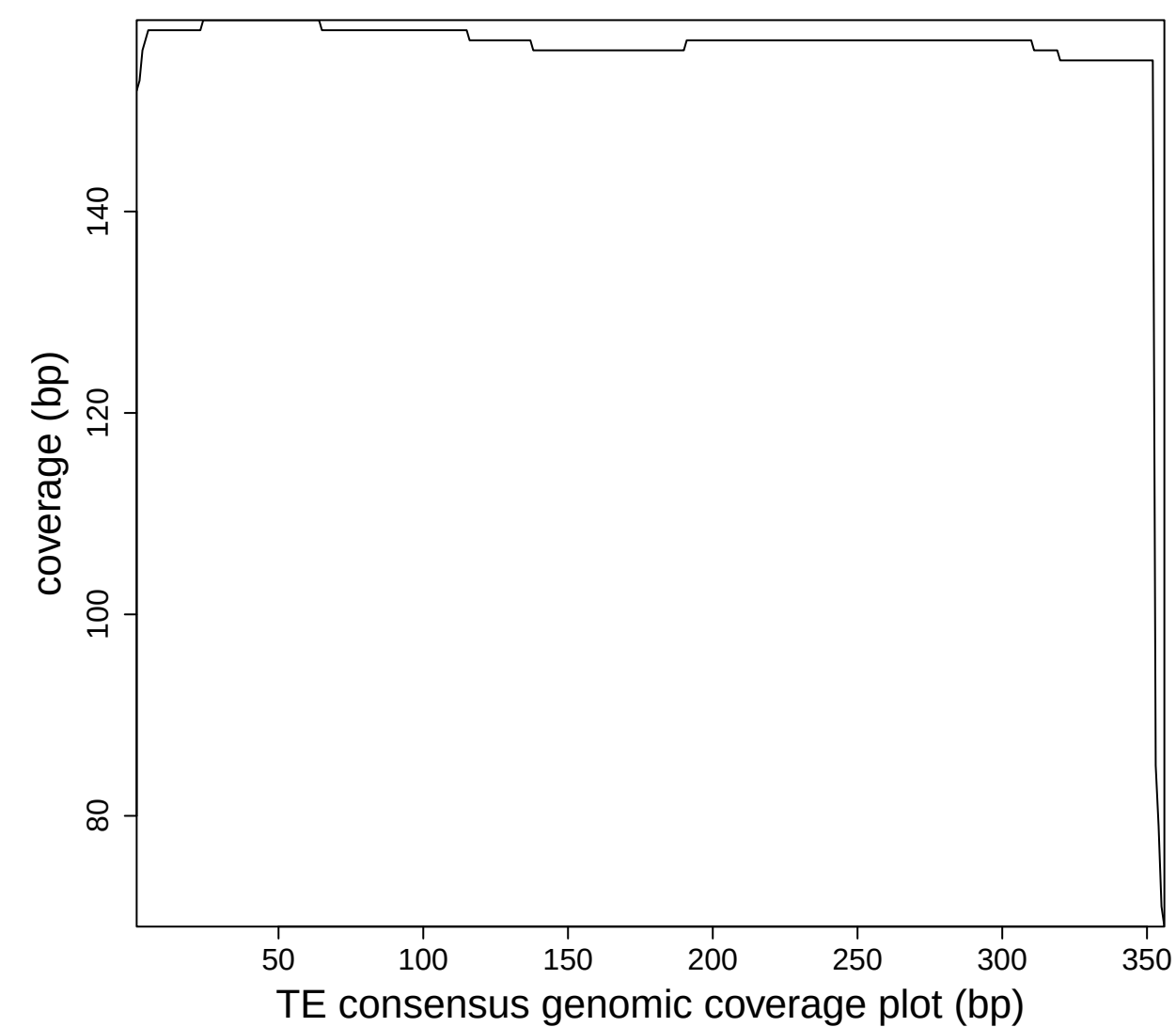
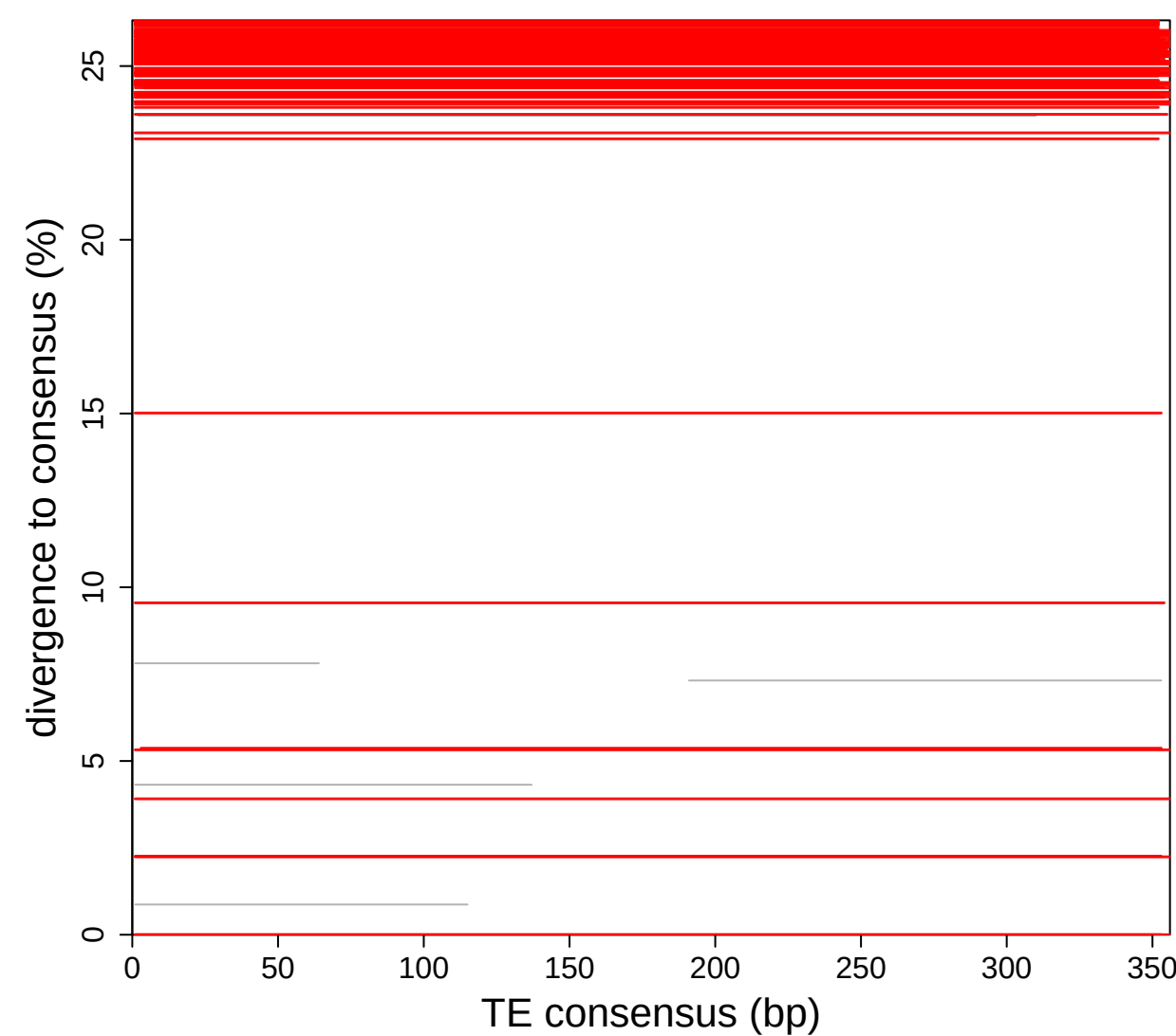


.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 661bp; fragments: 86; full length: 0 (>=594.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

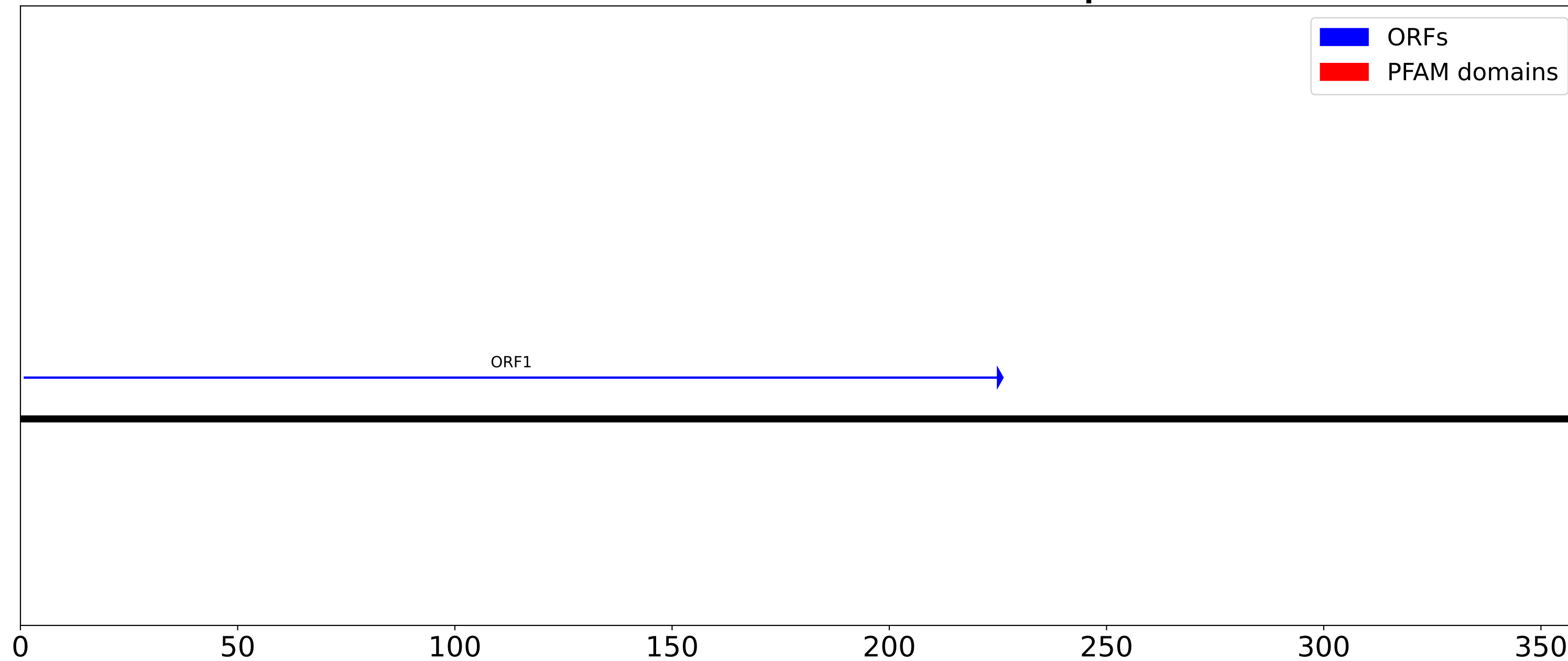


TE: ltr_1_family_269
size: 356bp; fragments: 160; full length: 154 (≥ 320.4 bp)



Before TEtrimmer 356 bp

After TEtrimmer ORF and PFAM domain plot



Dotplot

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

