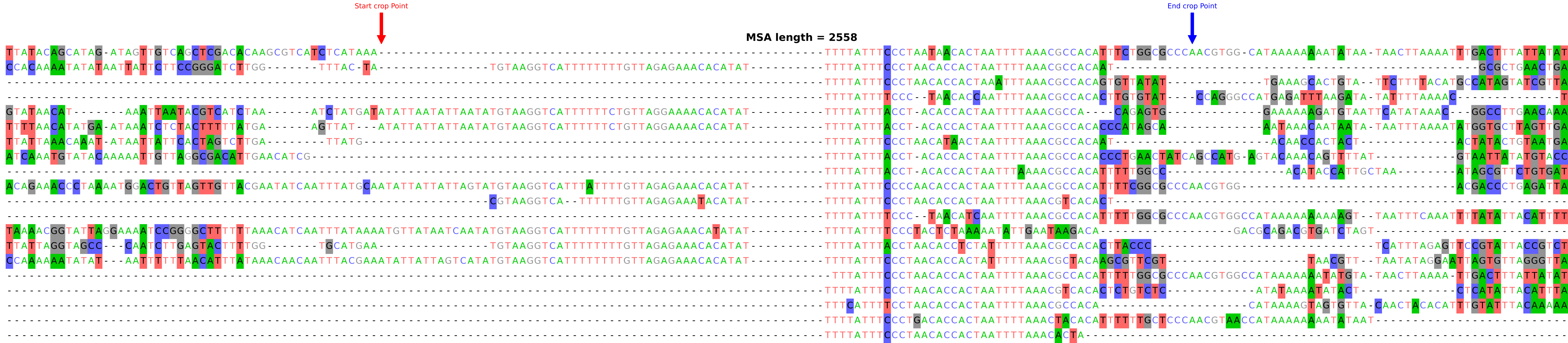
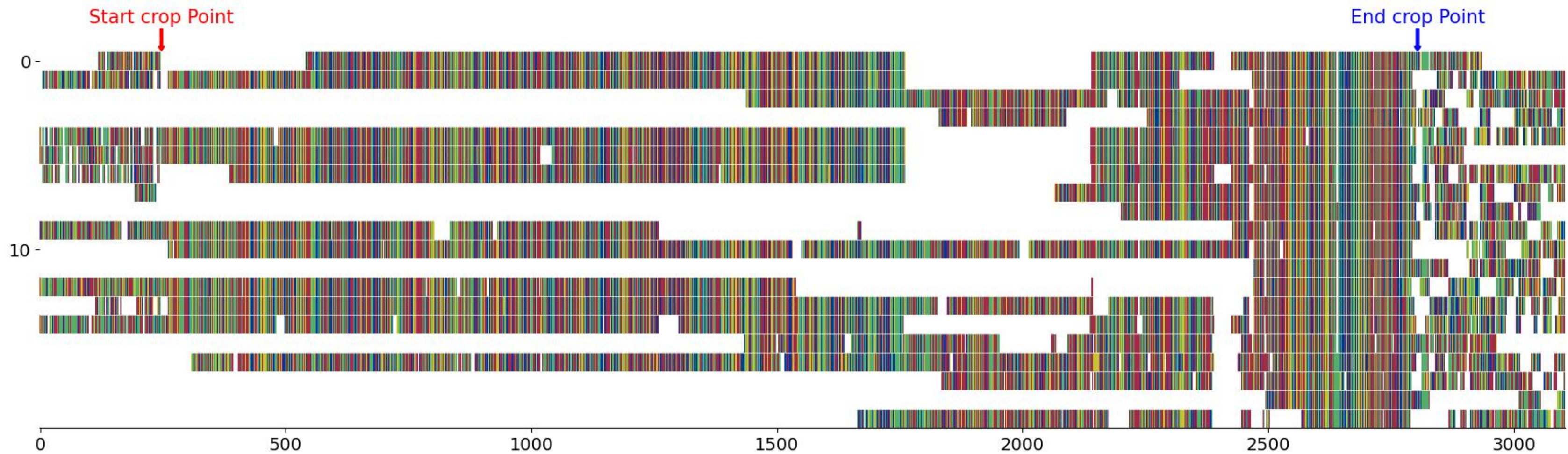




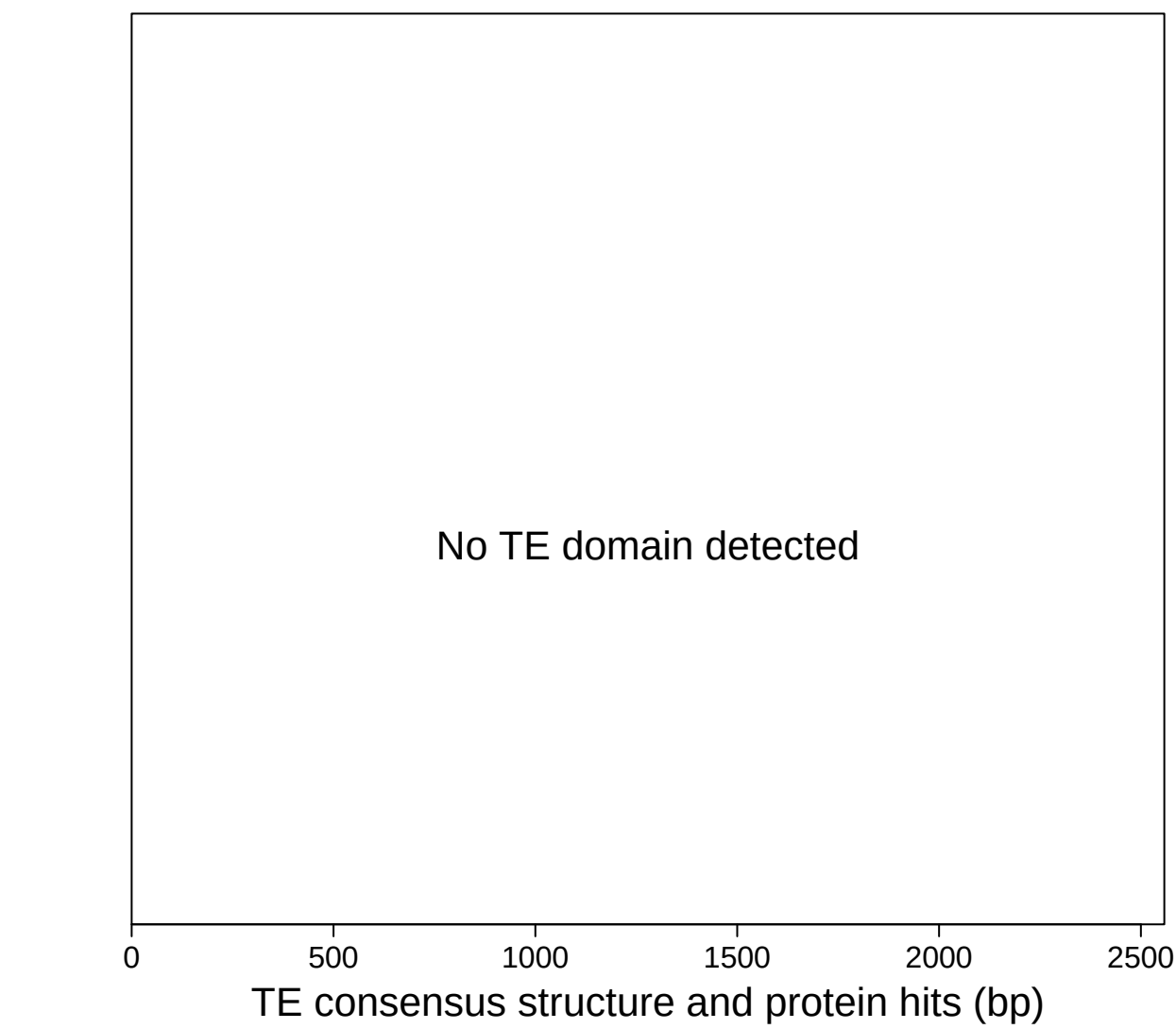
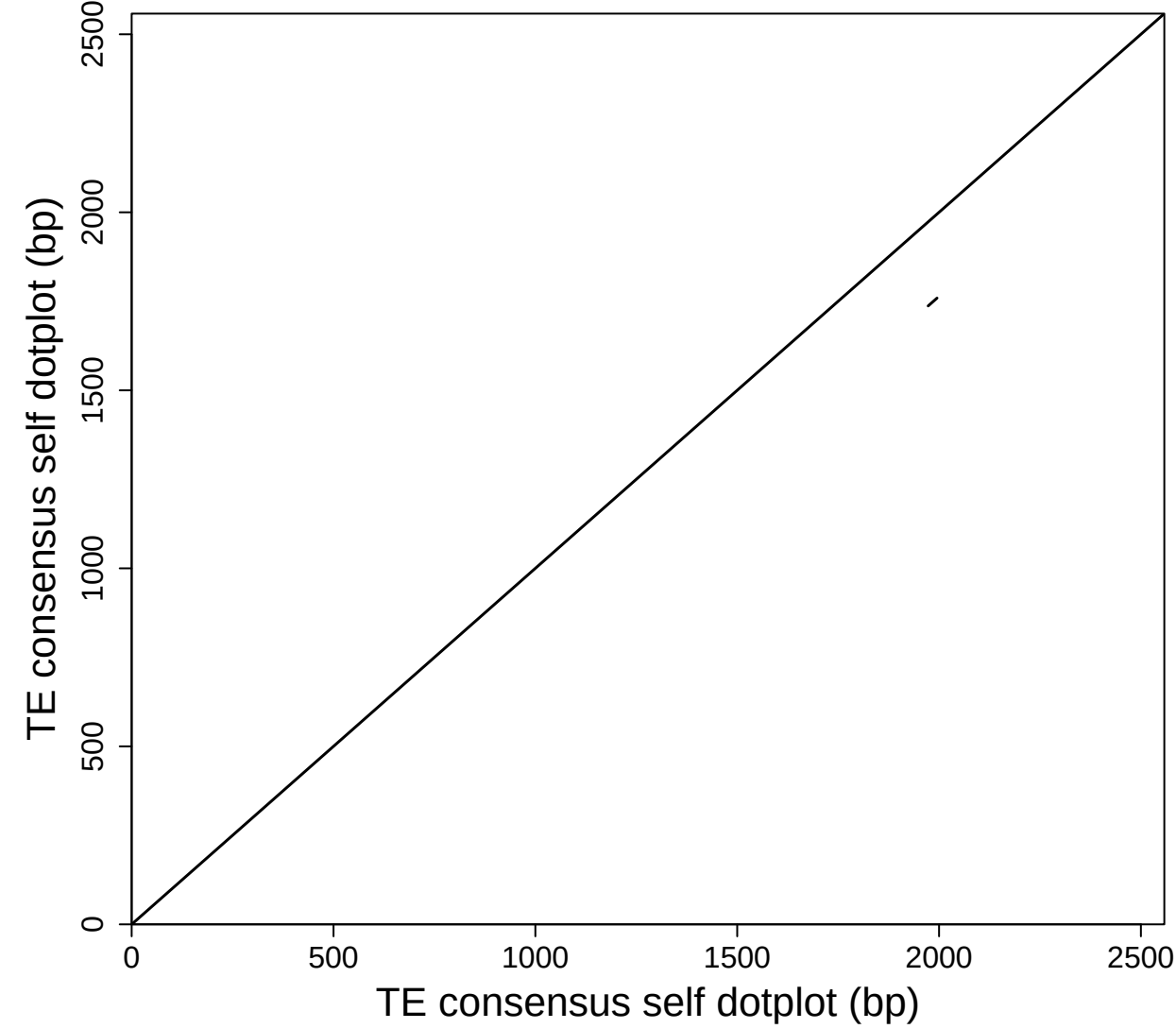
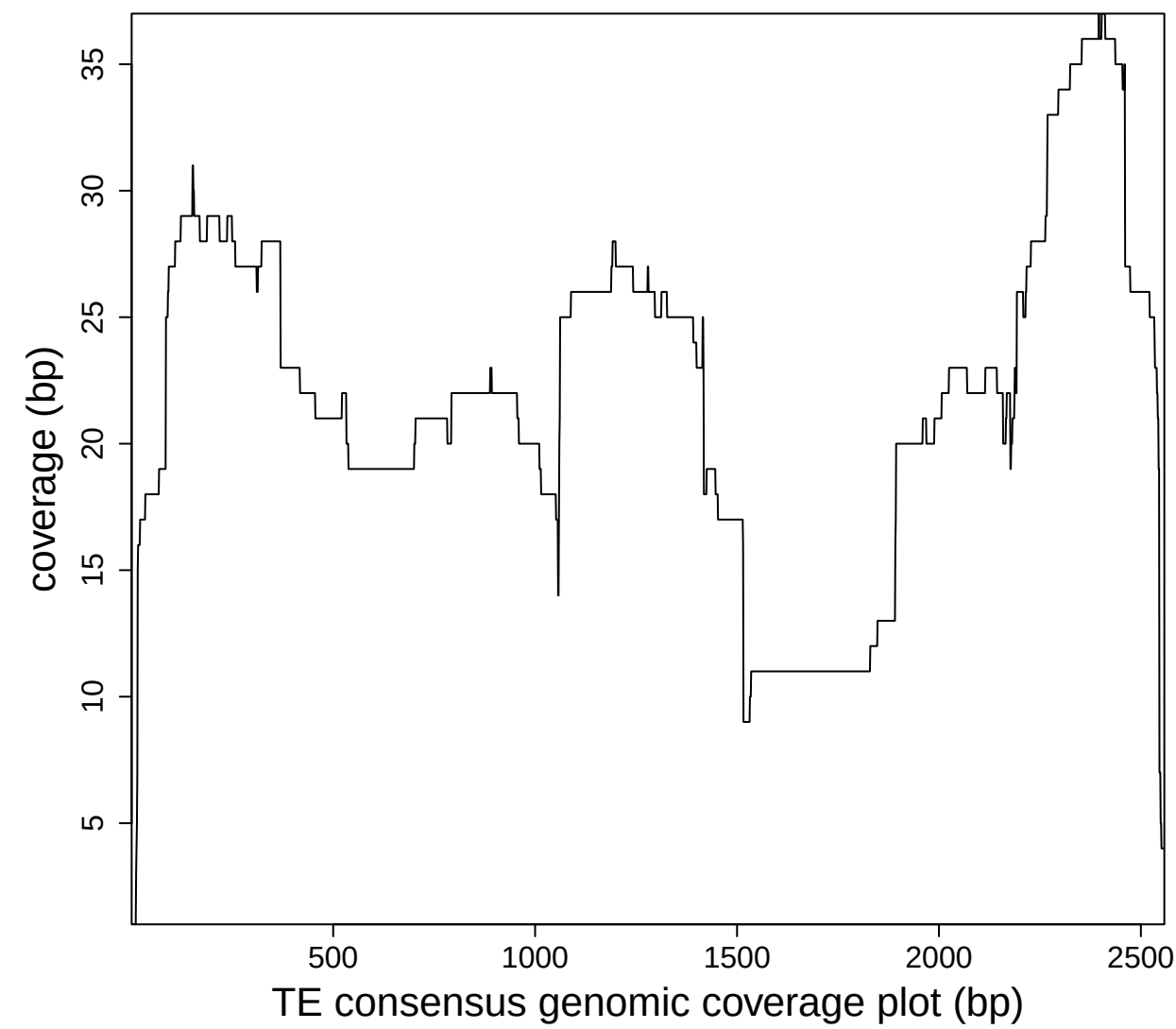
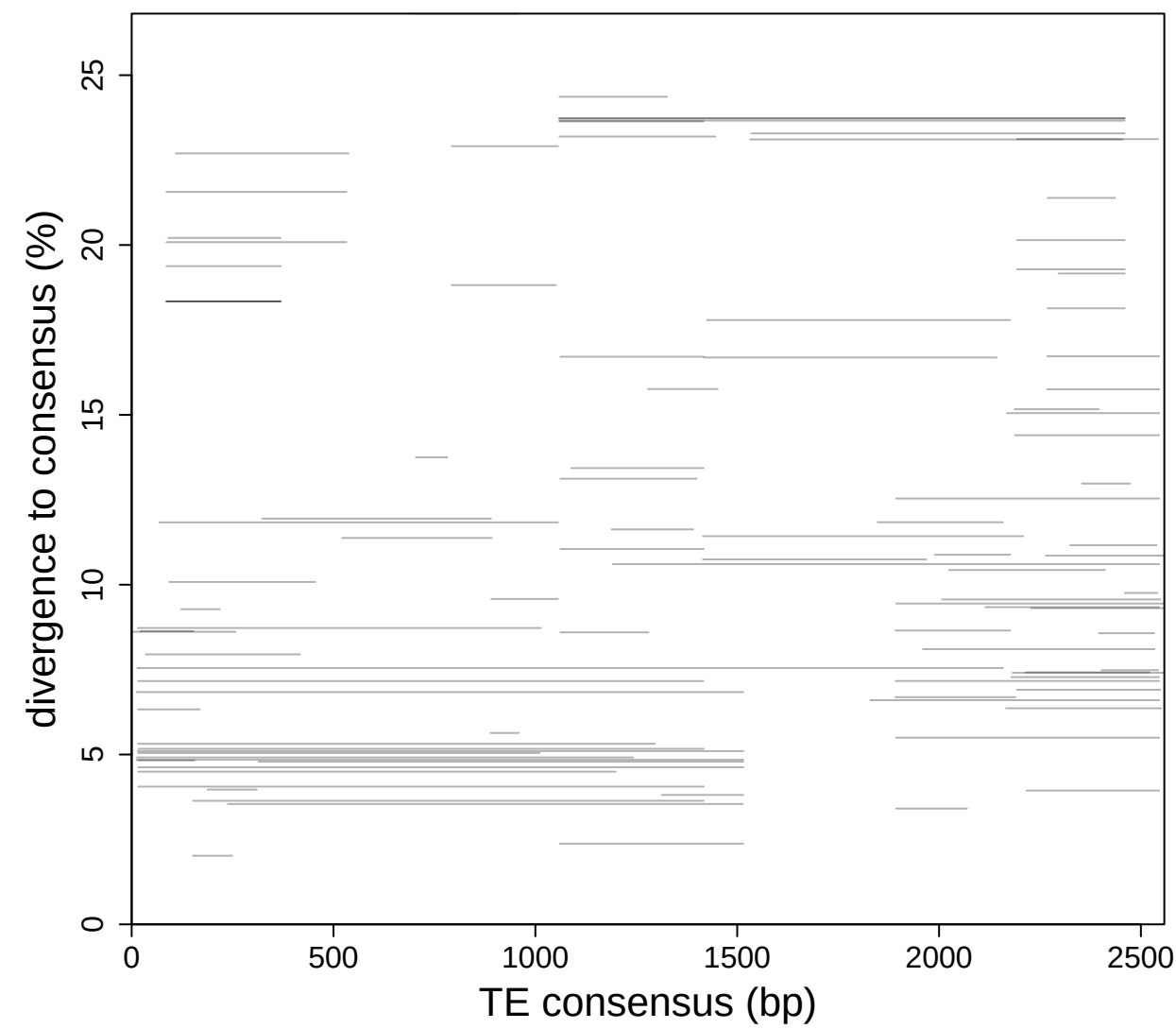
Start crop Point

End crop Point



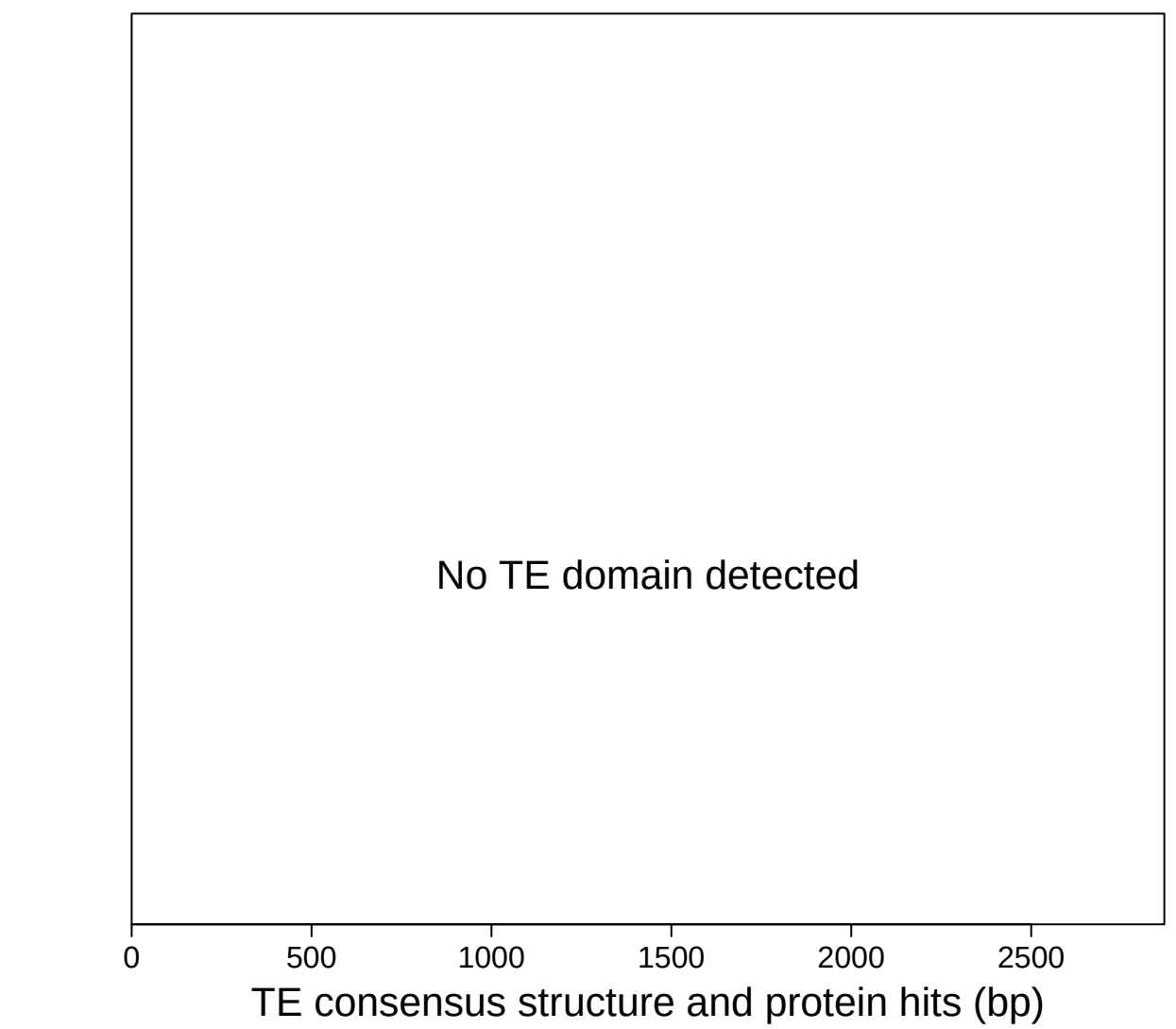
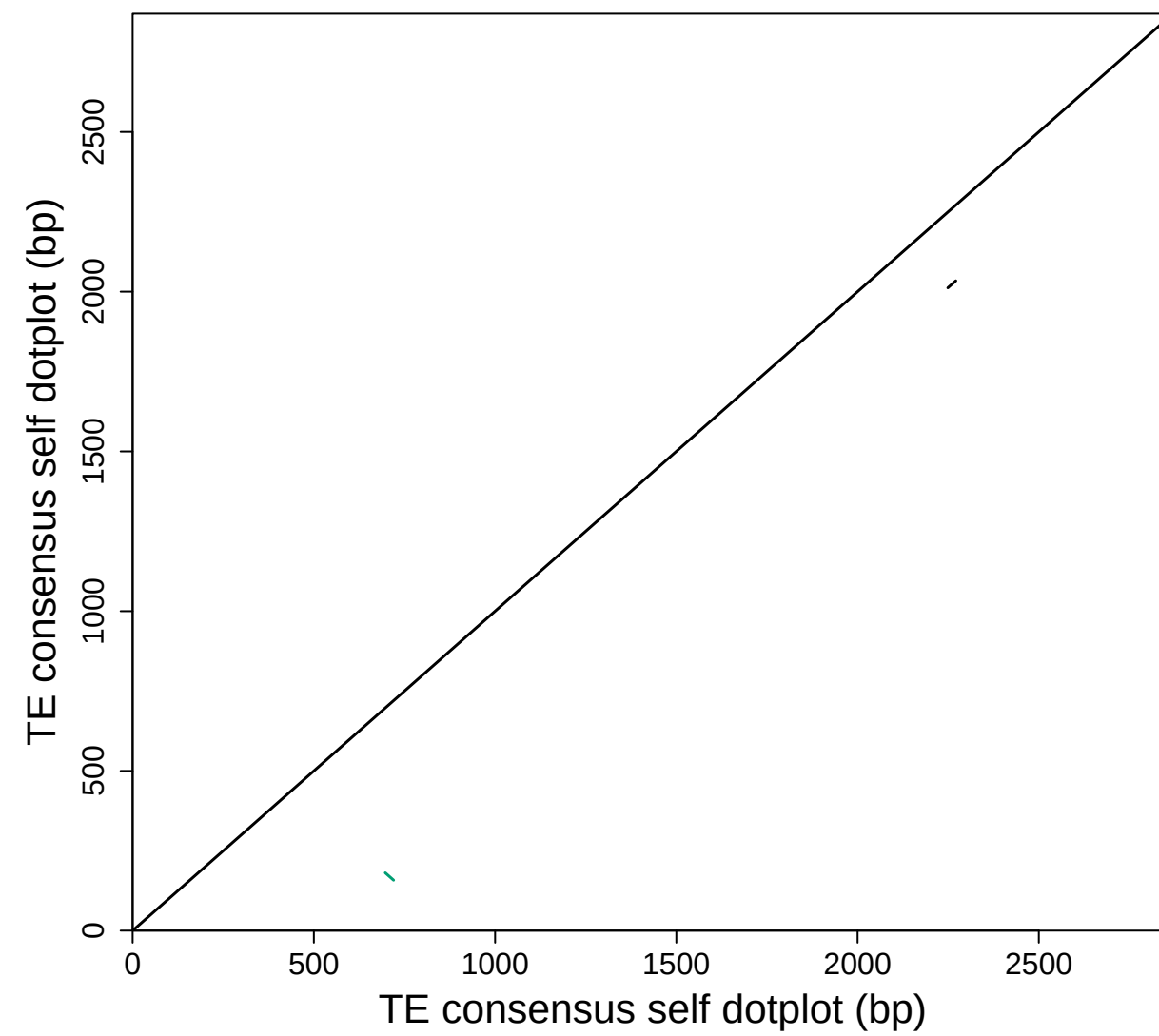
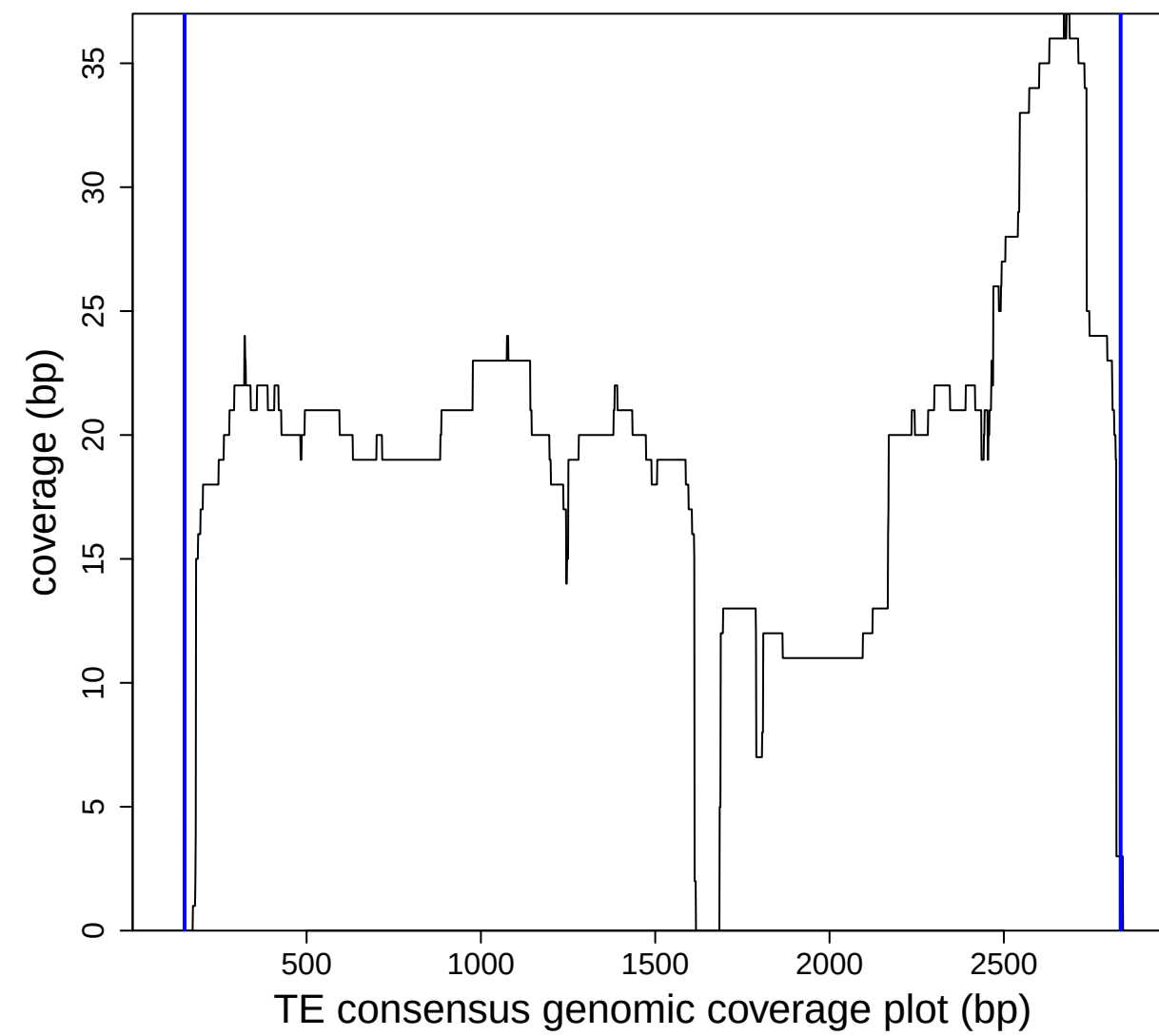
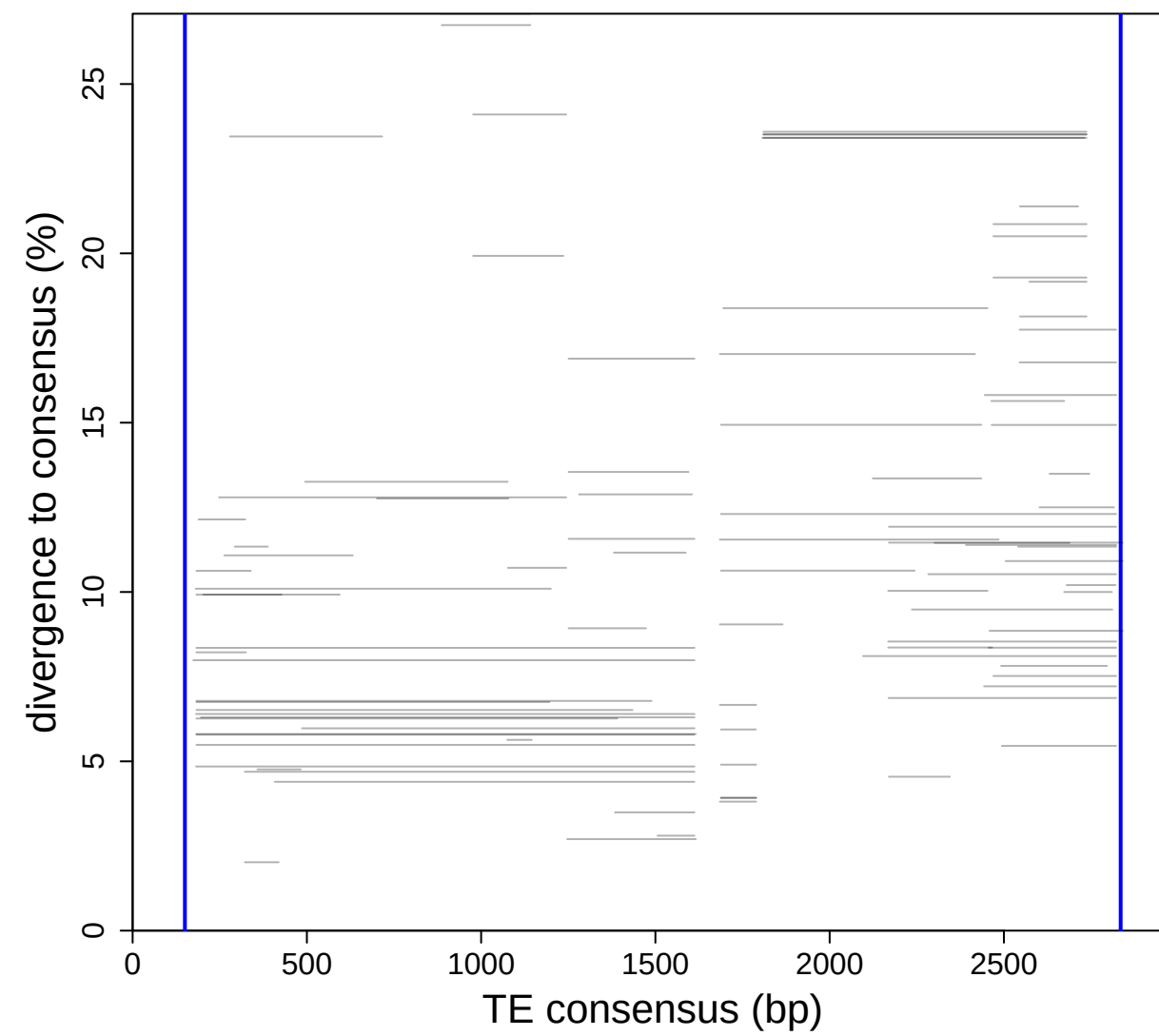
b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 2558bp; fragments: 102; full length: 0 (>=2302.2bp)

After TEtrimmer 2558 bp

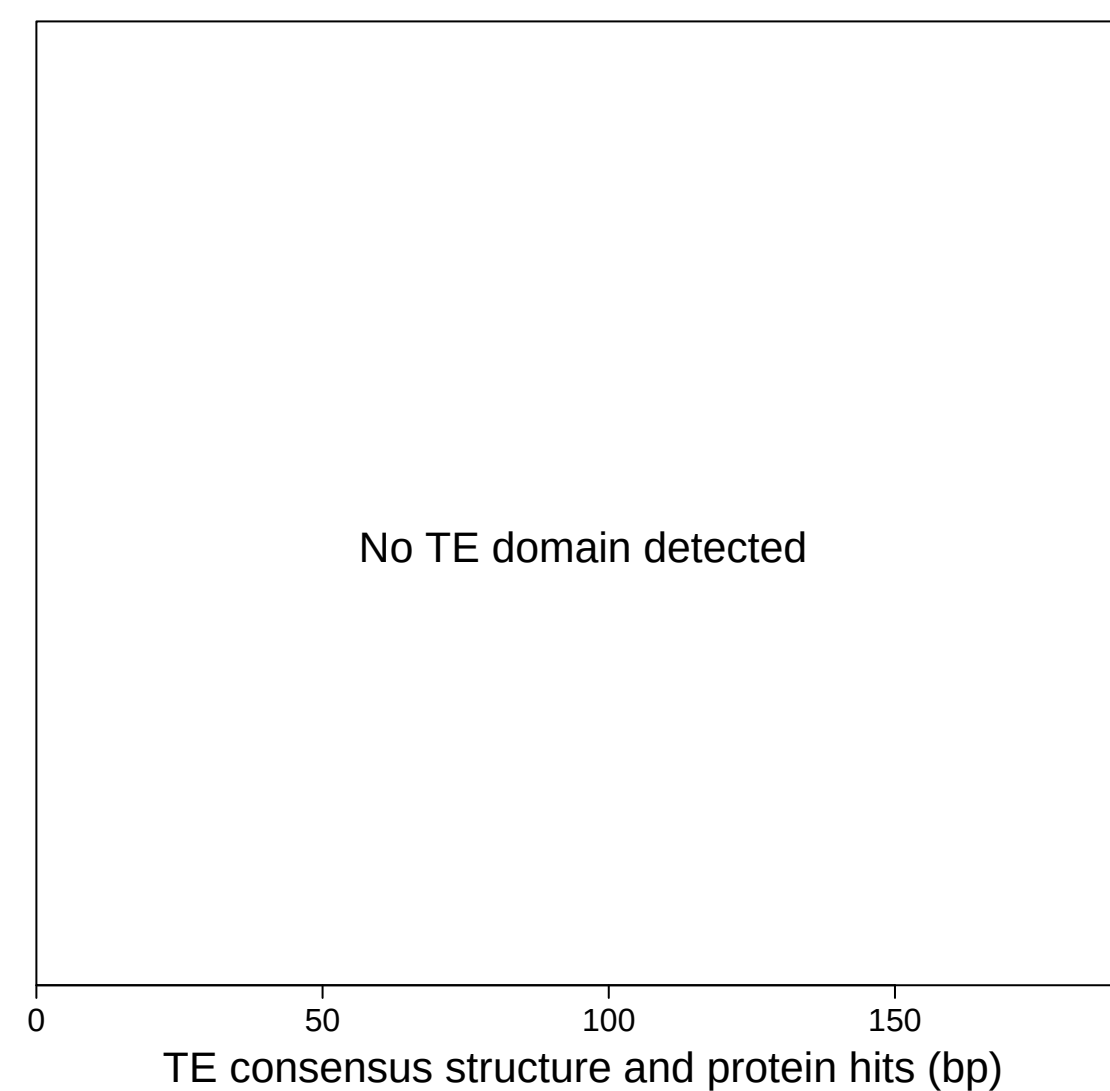
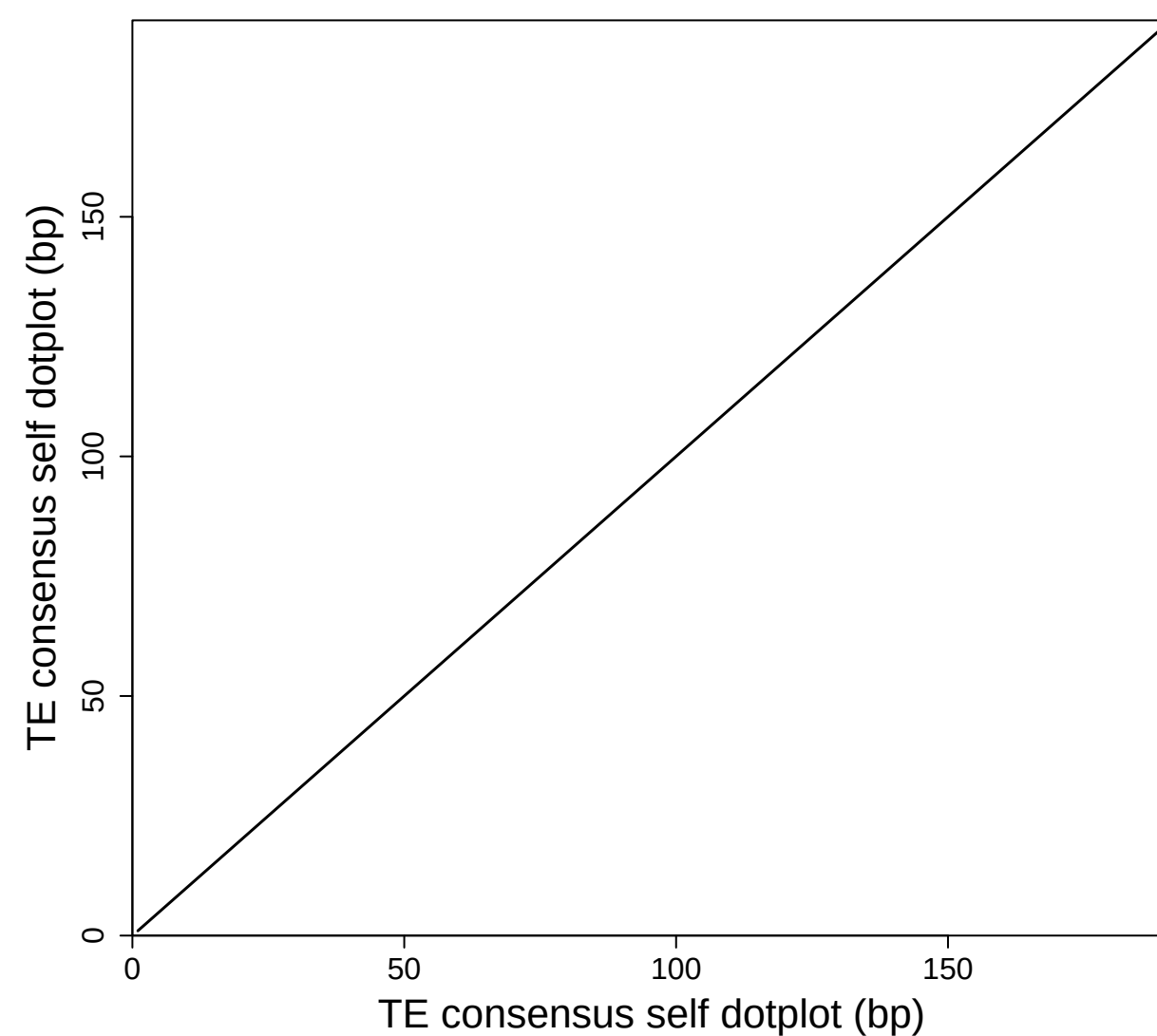
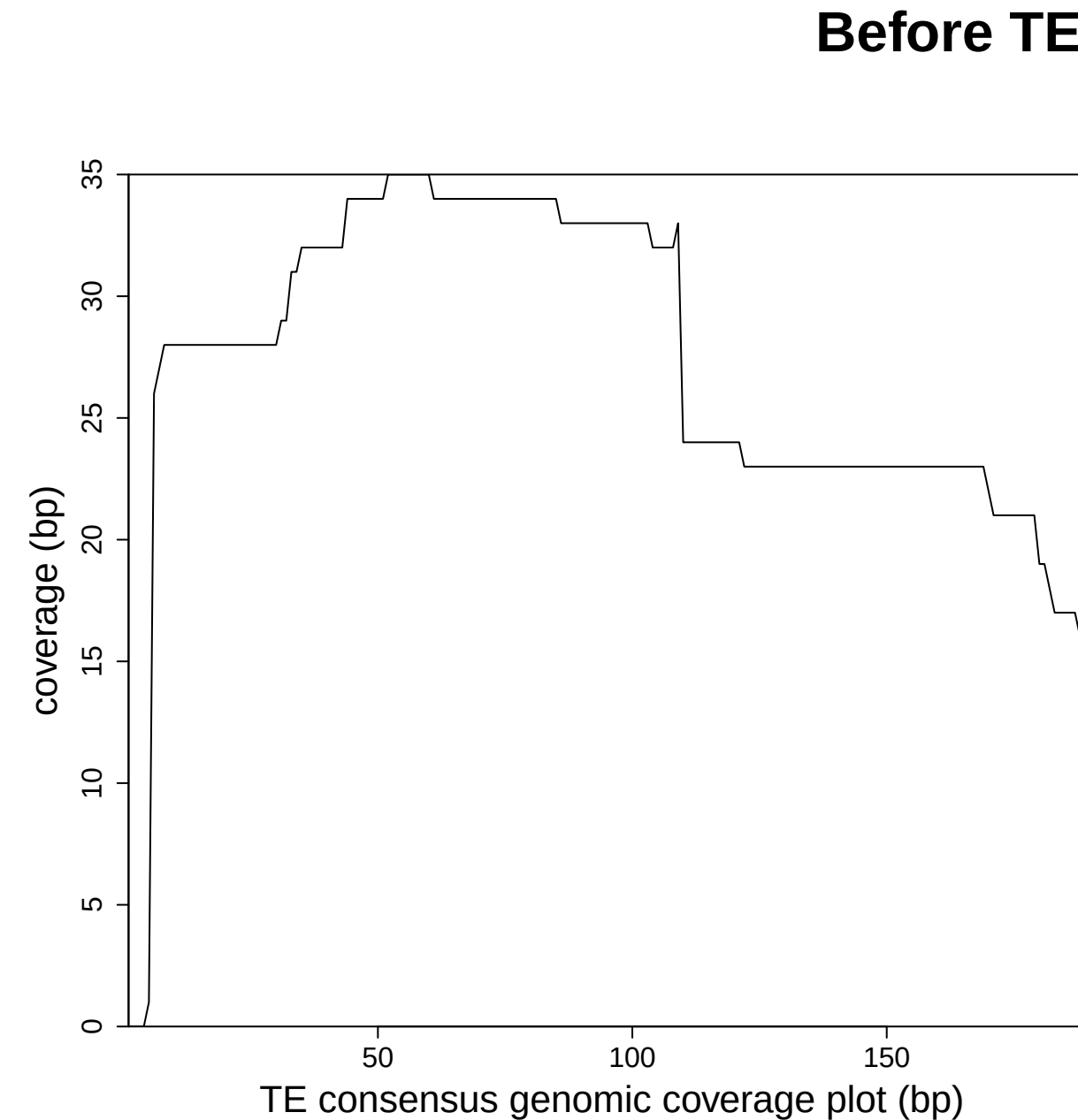
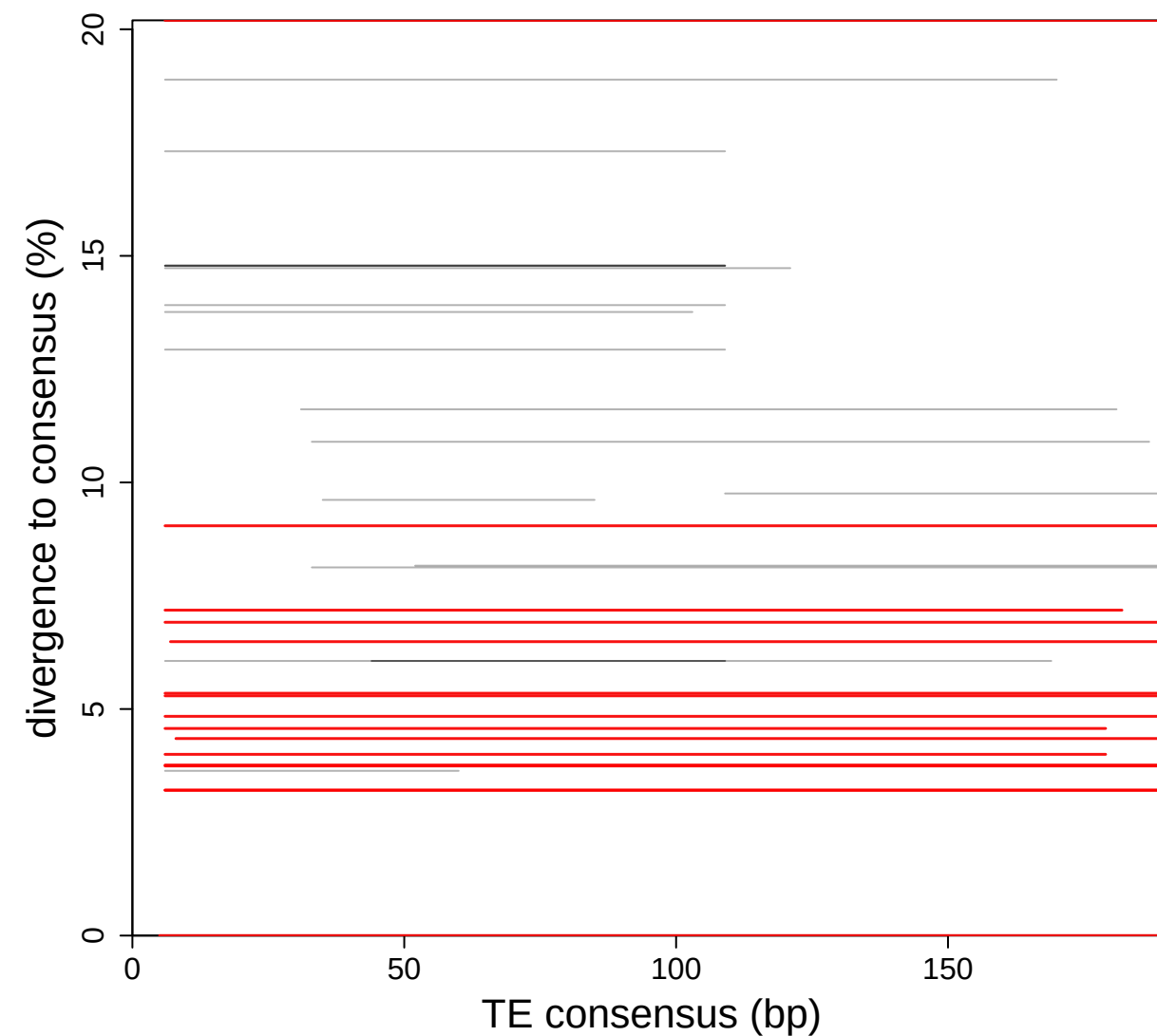


.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 2985bp; fragments: 98; full length: 0 (>=2686.5bp)

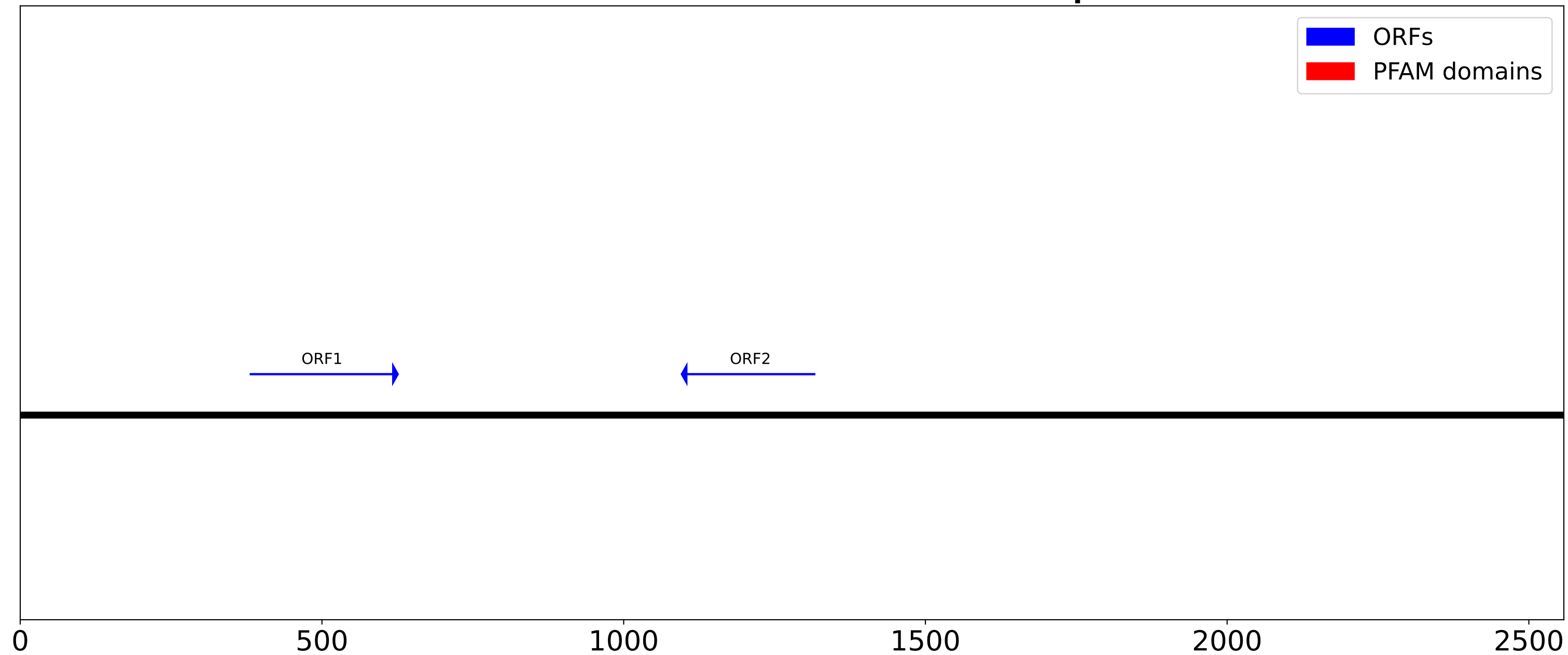
After TEtrimmer Extended plot Blue lines are boundaries



TE: ltr_1_family_392
size: 191bp; fragments: 36; full length: 16 (≥ 171.9 bp)



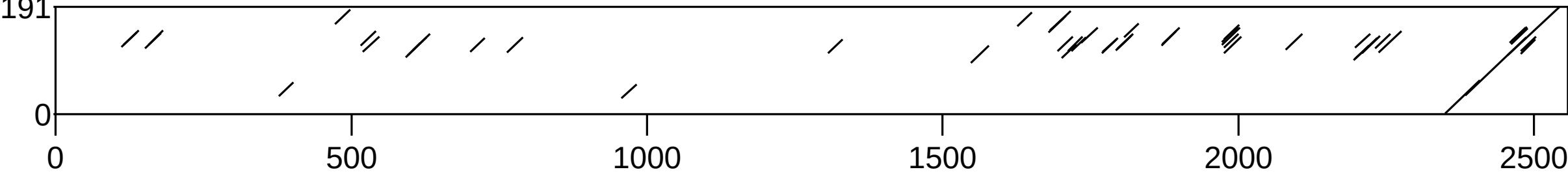
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)