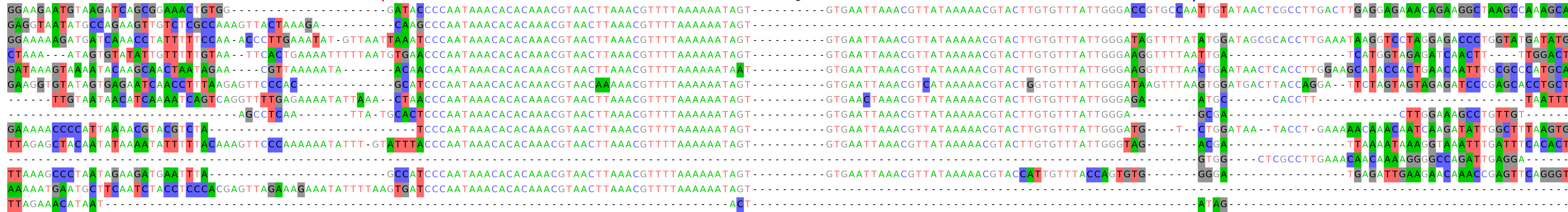


Start crop Point

End crop Point

MSA length = 2294



Start crop Point

End crop Point

0 -

10 -

0

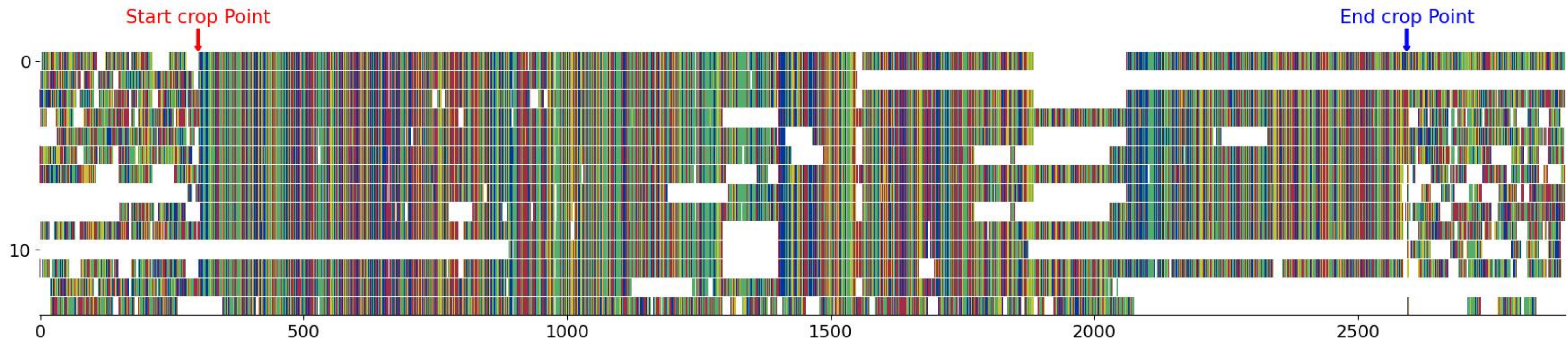
500

1000

1500

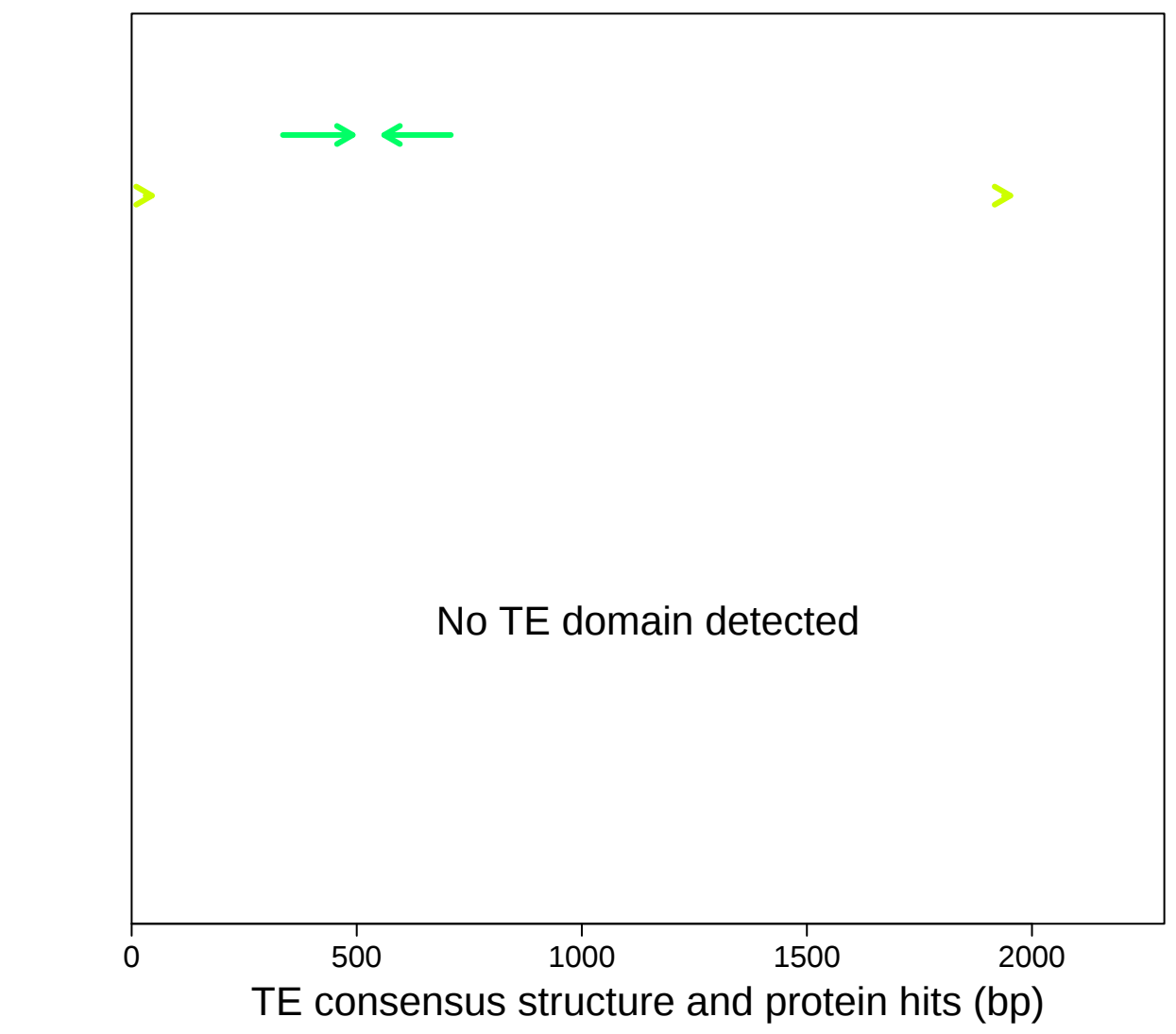
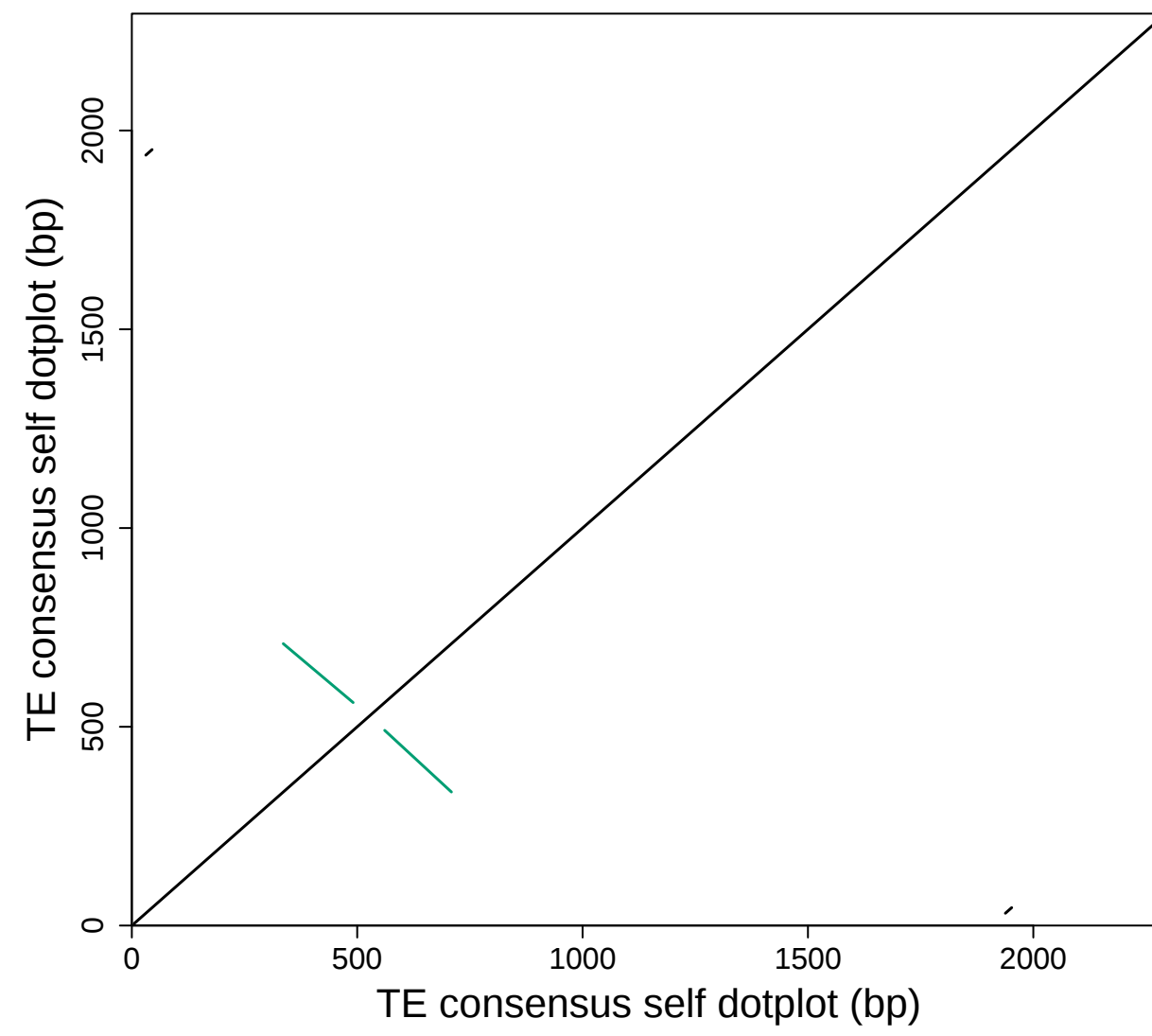
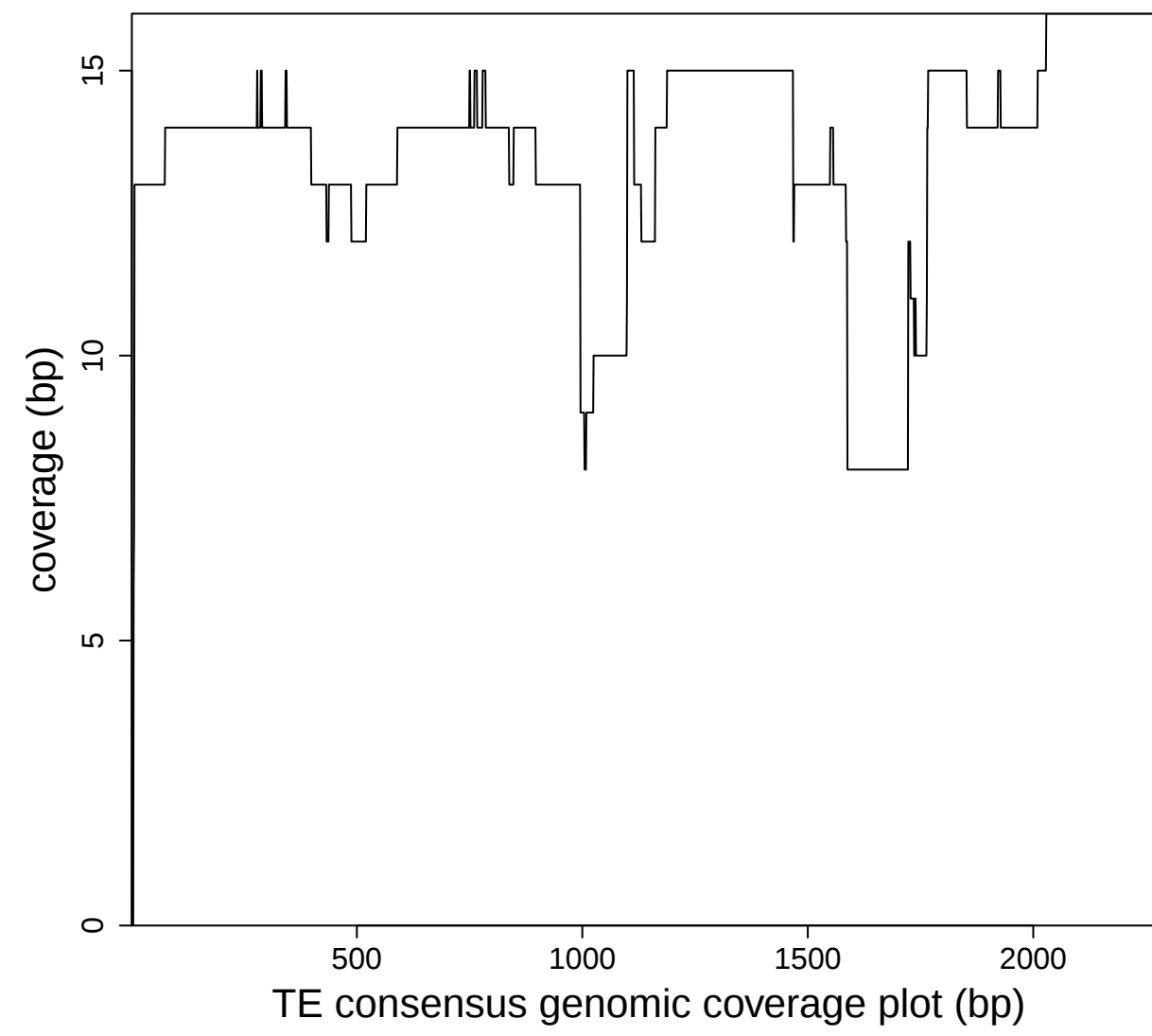
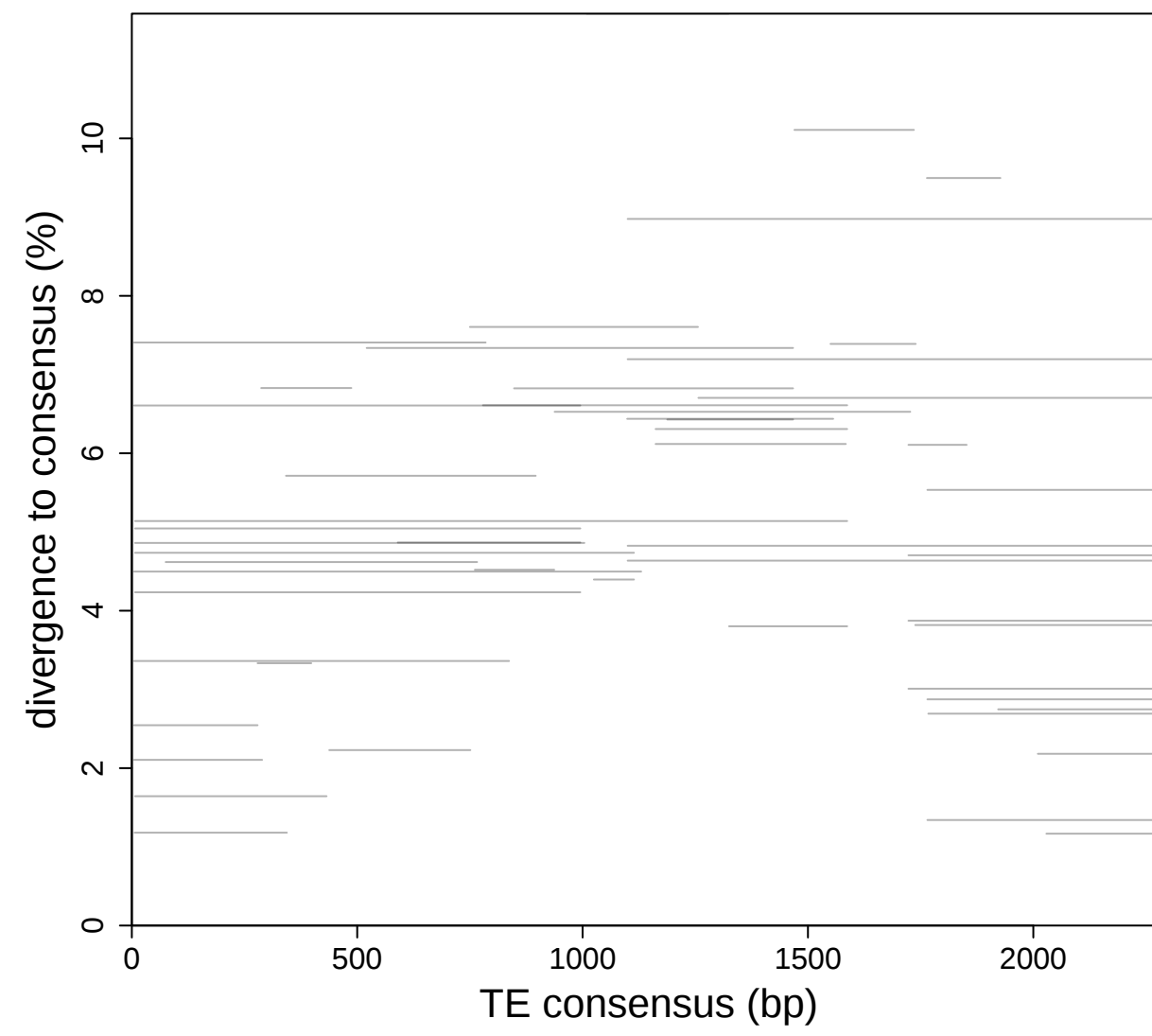
2000

2500



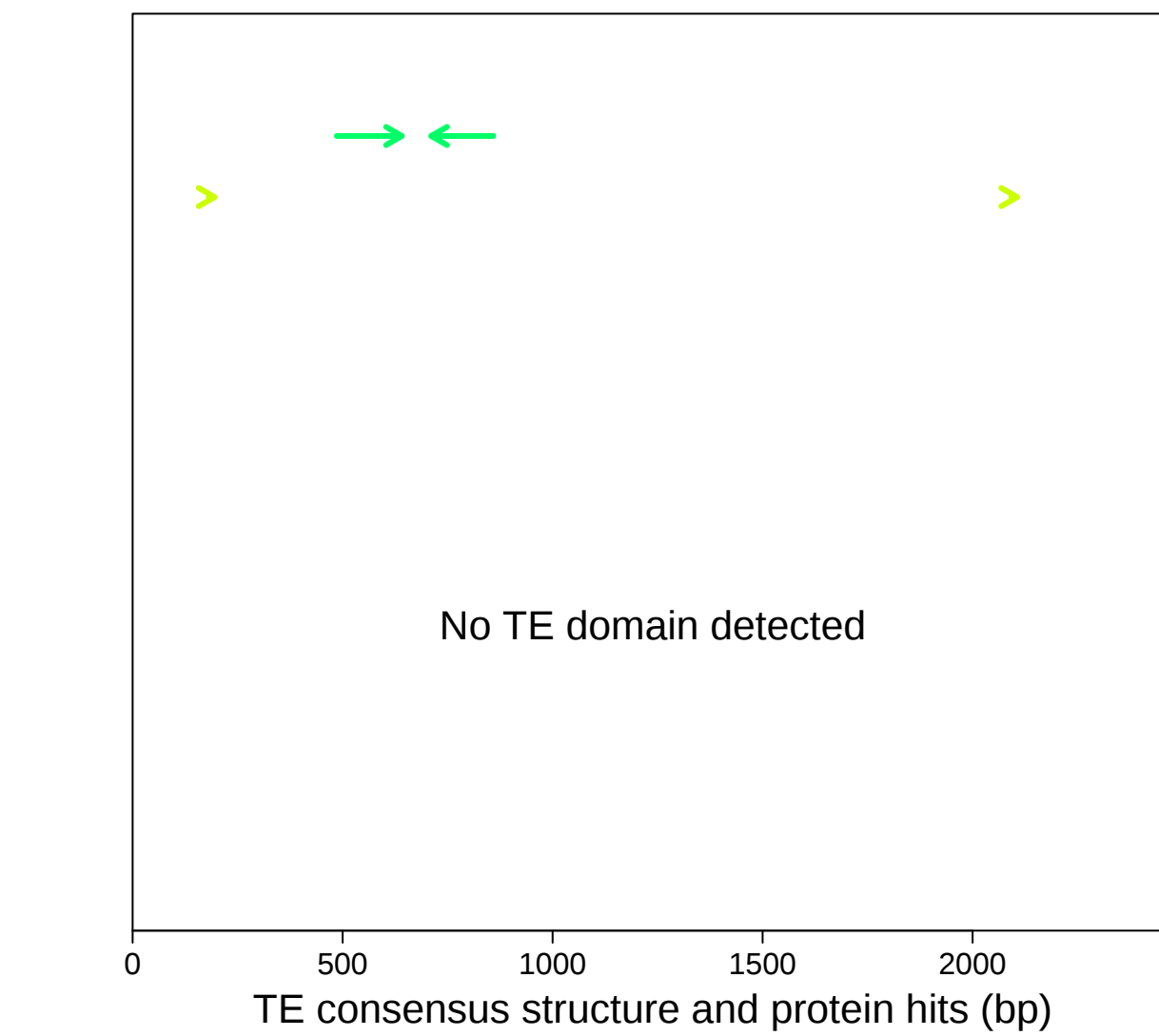
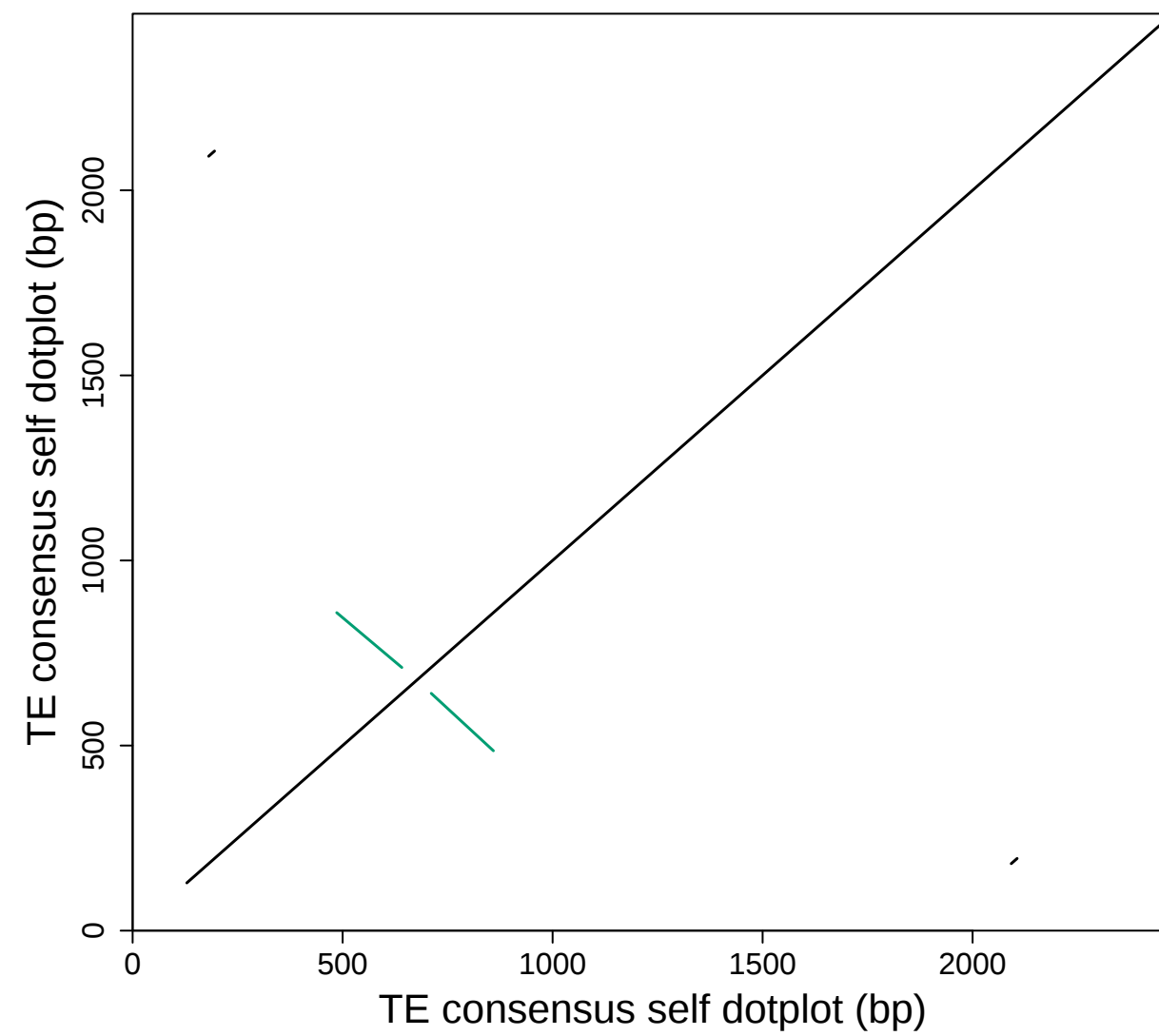
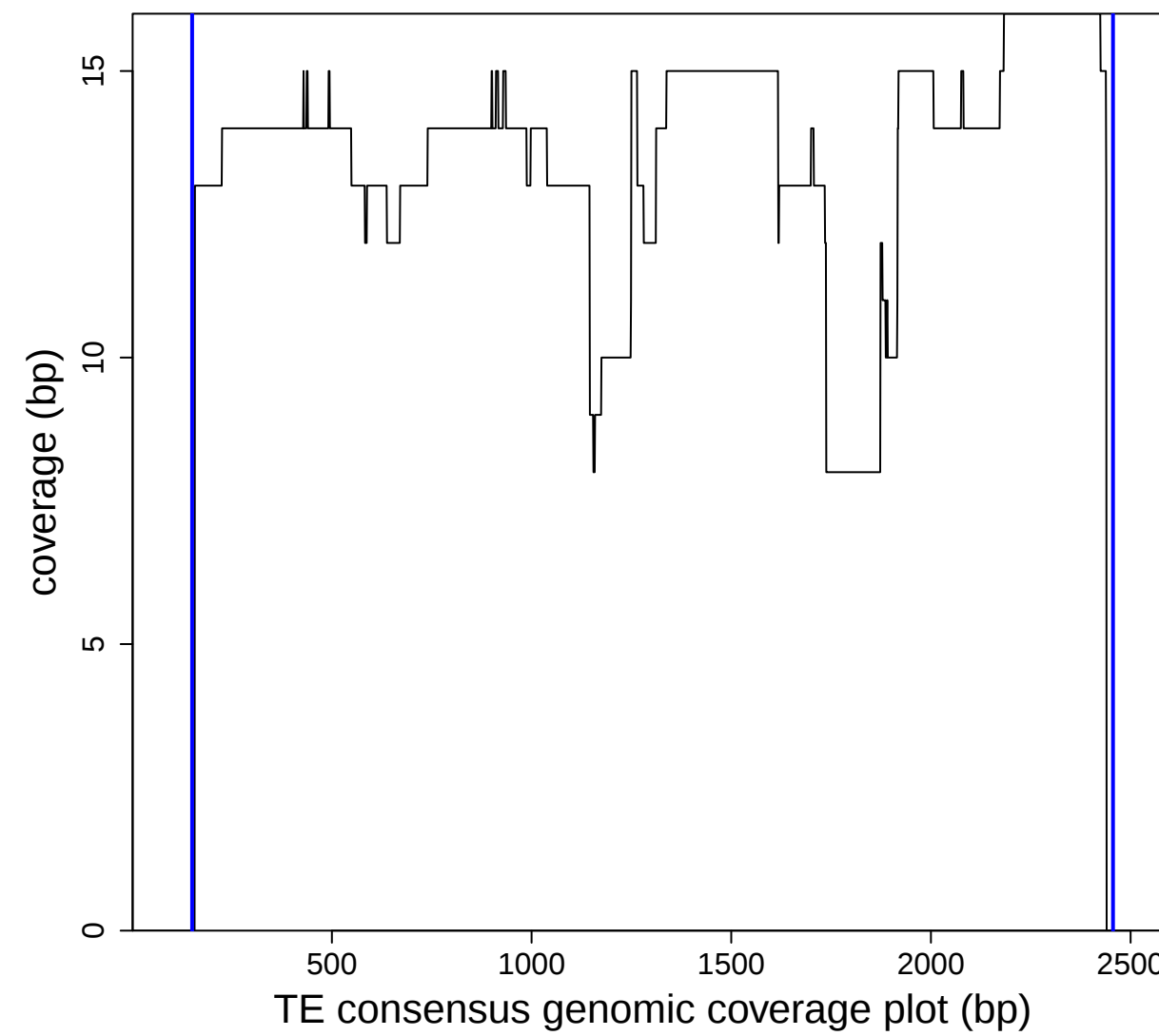
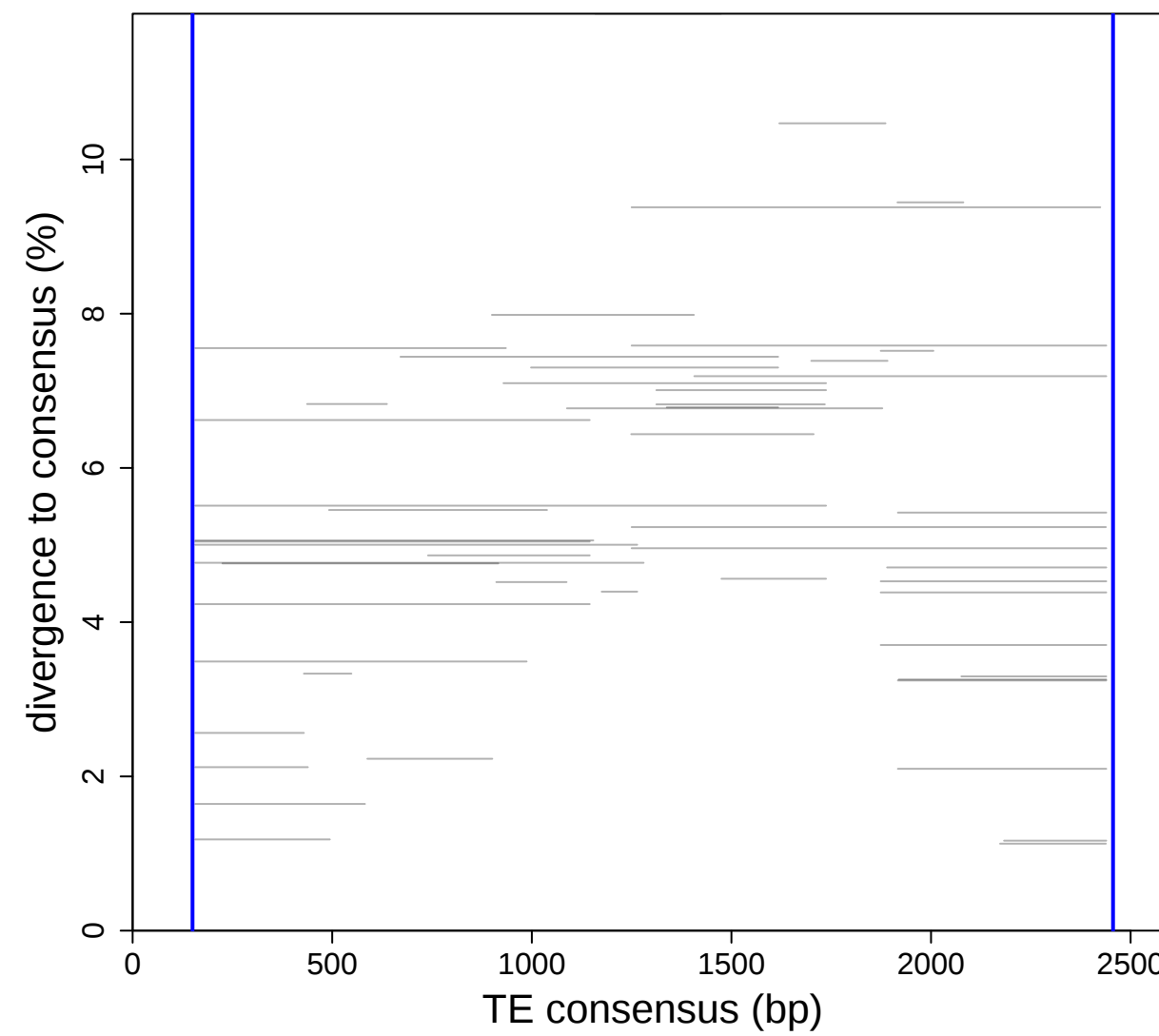
b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 2294bp; fragments: 52; full length: 0 (>=2064.6bp)

After TEtrimmer 2294 bp

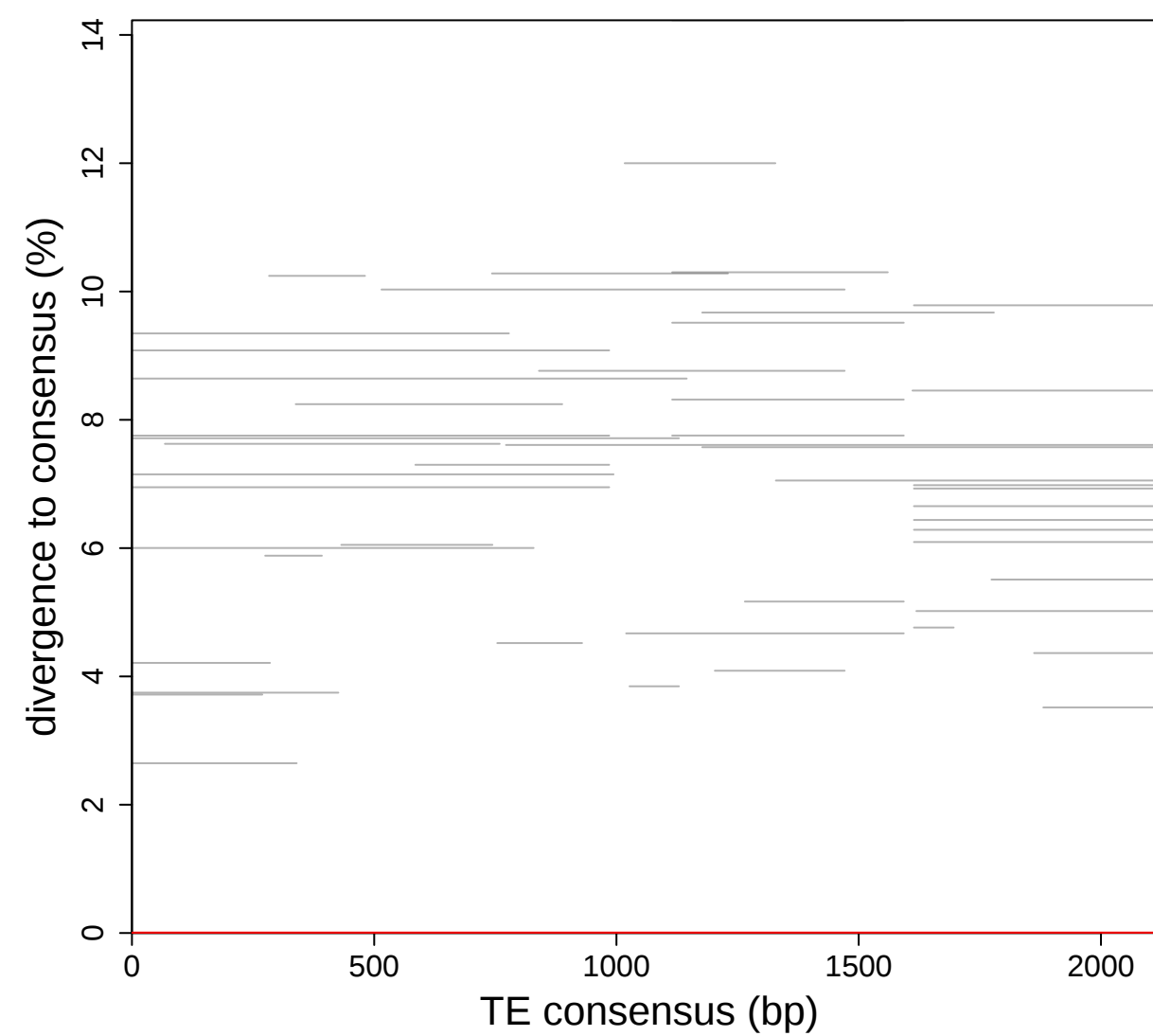


.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 2606bp; fragments: 52; full length: 0 (>=2345.4bp)

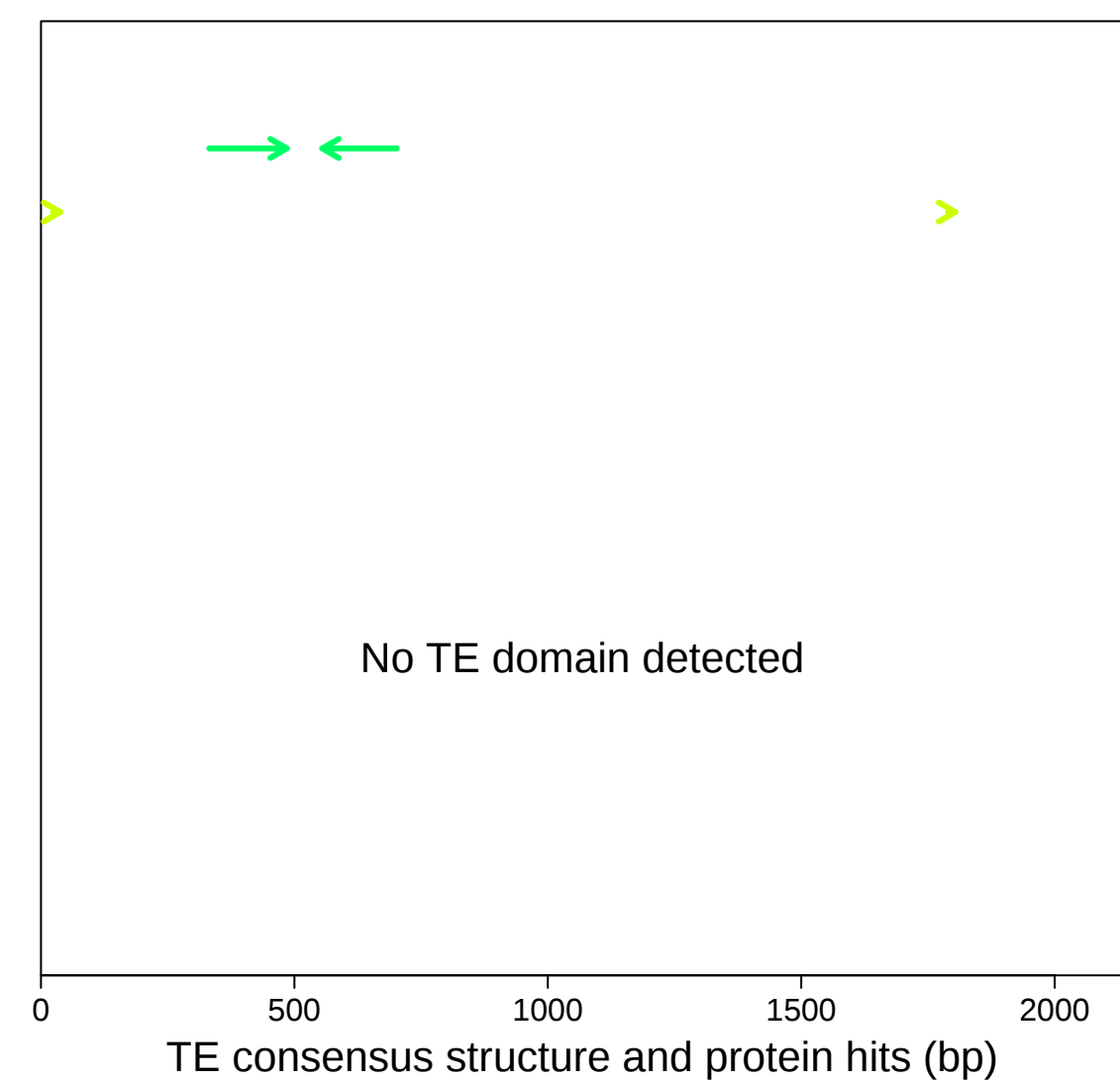
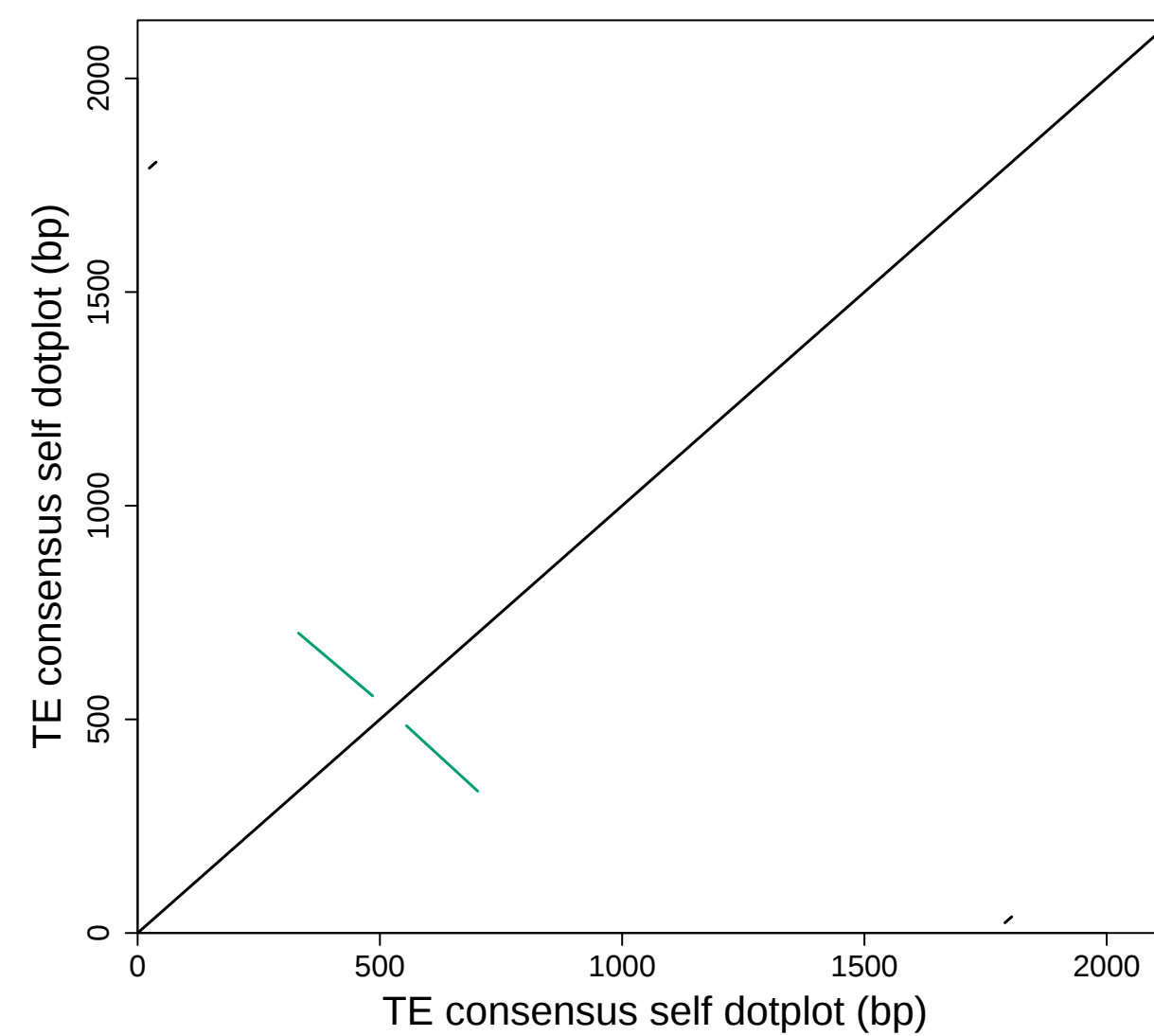
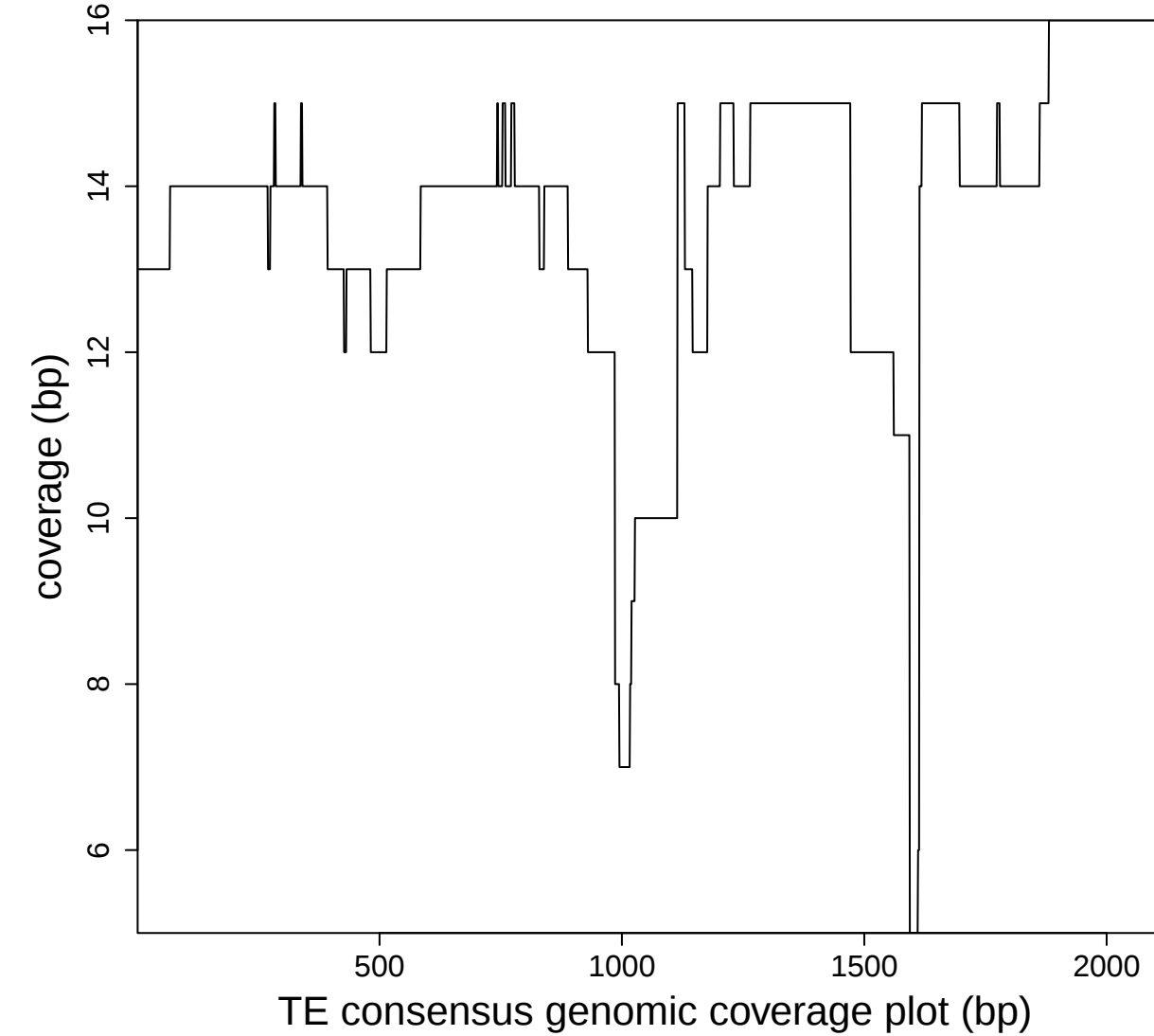
After TEtrimmer Extended plot Blue lines are boundaries



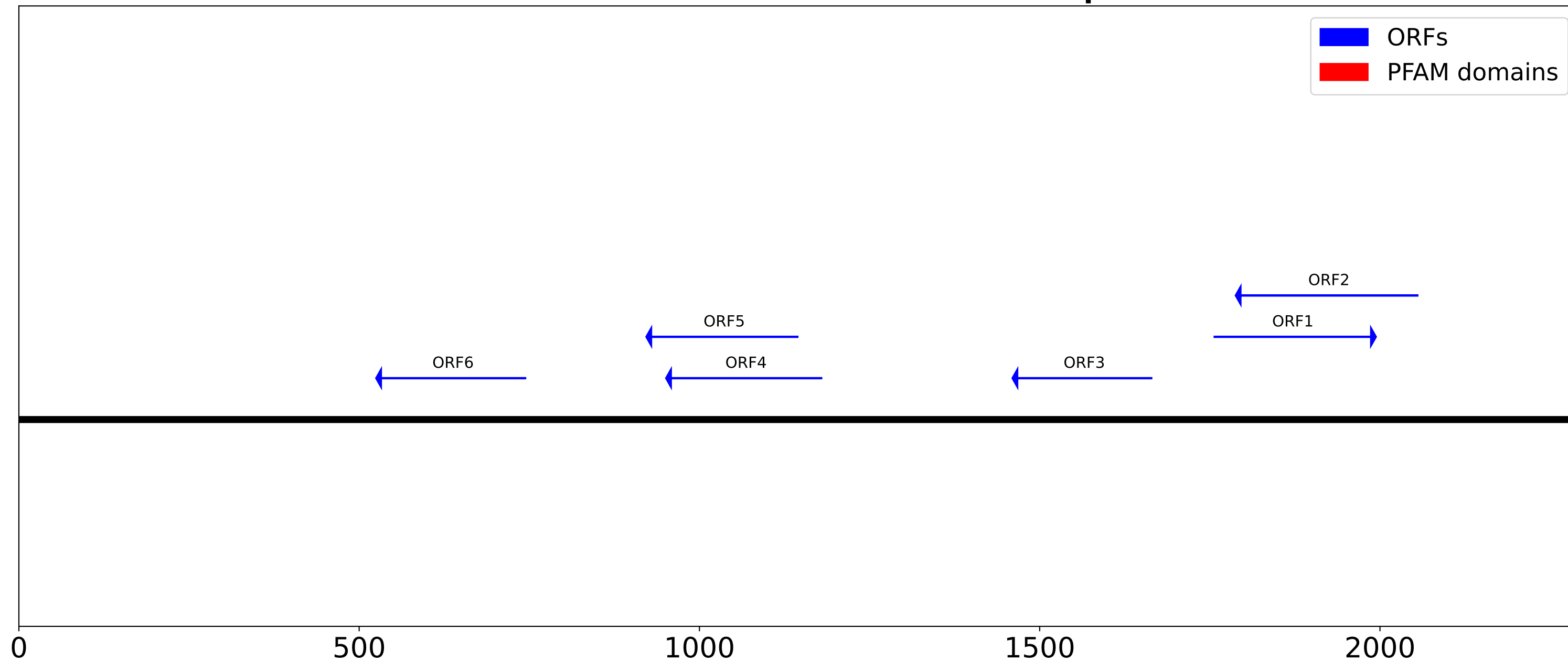
TE: ltr_1_family_571
size: 2136bp; fragments: 50; full length: 1 (>=1922.4bp)



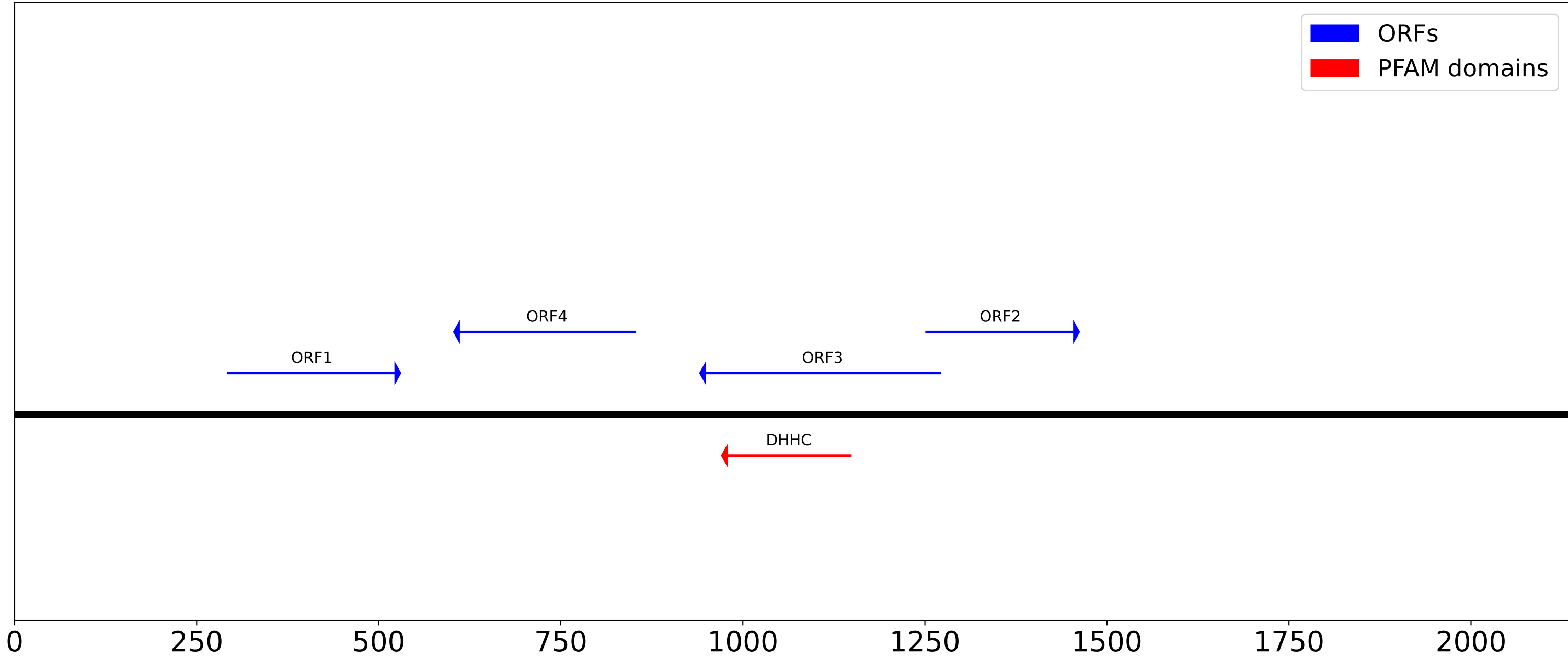
Before TEtrimmer 2136 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

