

Start crop Point



End crop Point



MSA length = 2223



Start crop Point

End crop Point

0 -

10 -

0

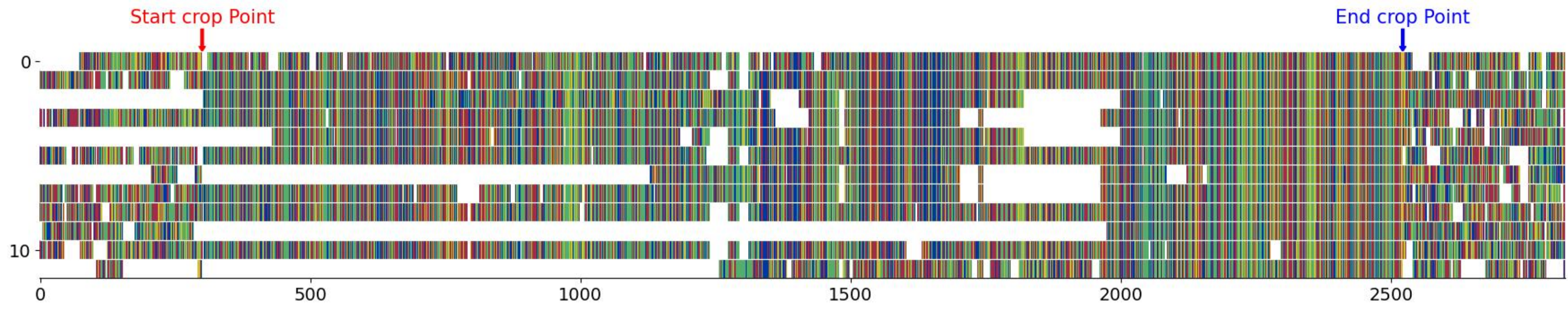
500

1000

1500

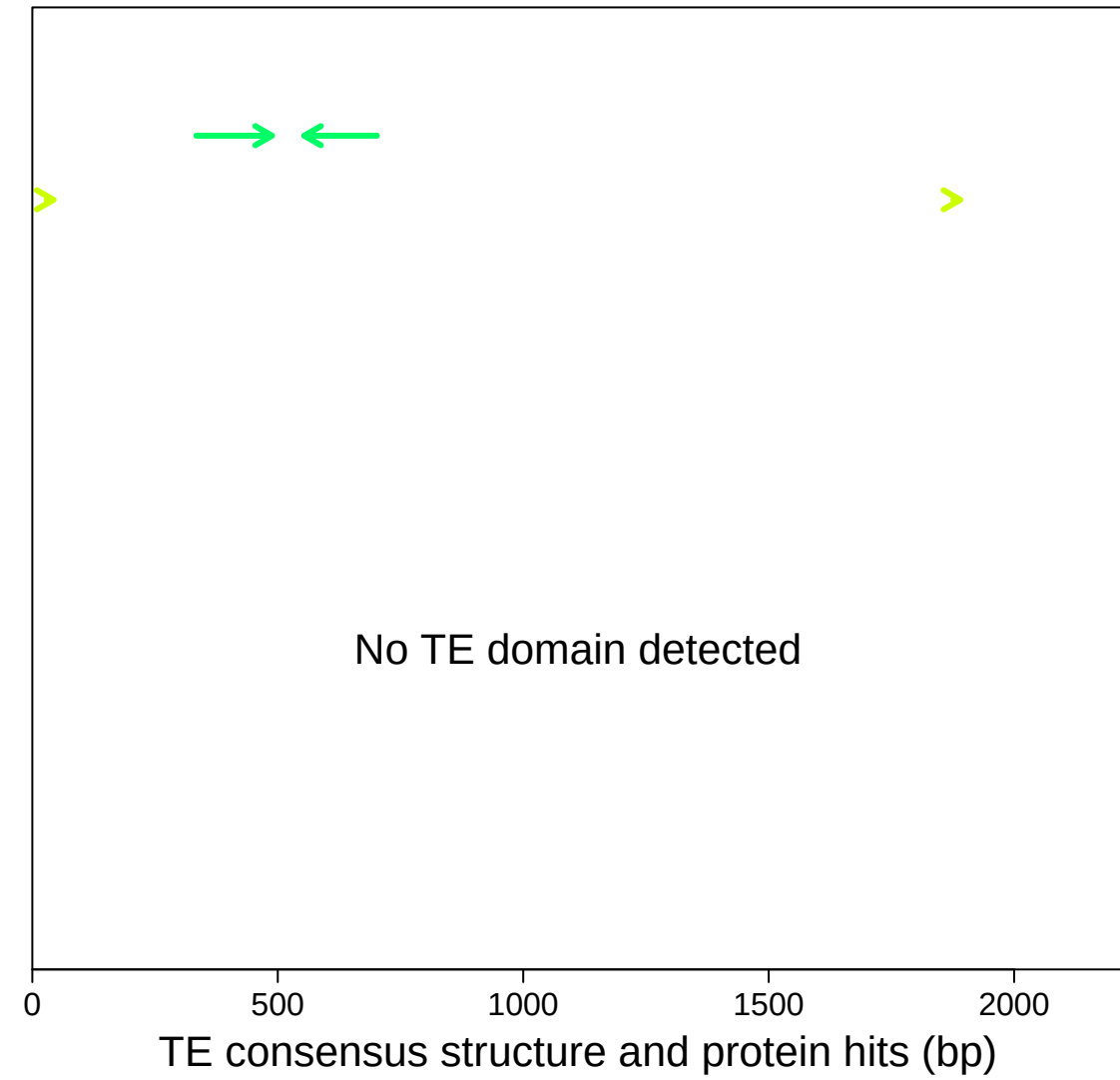
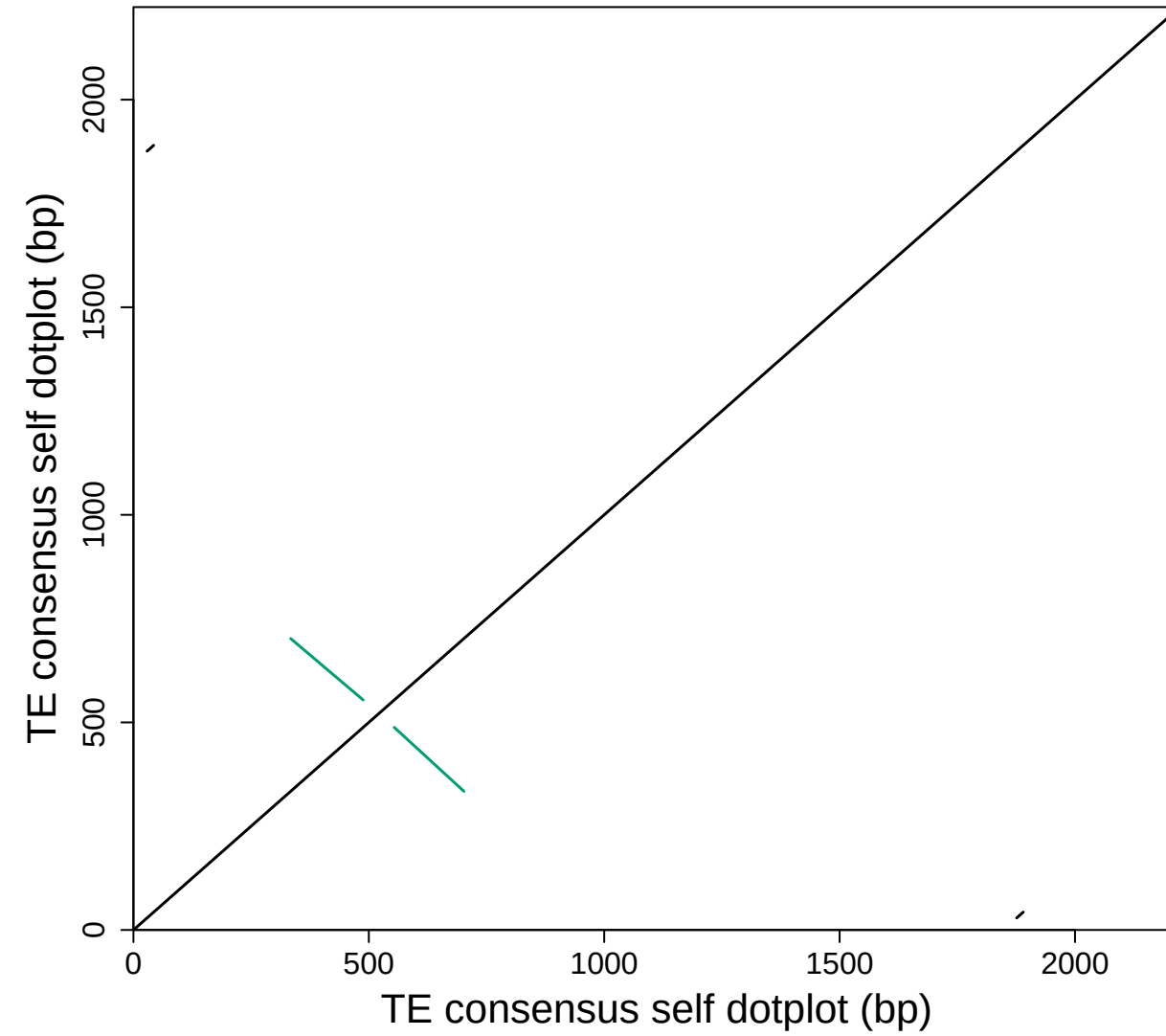
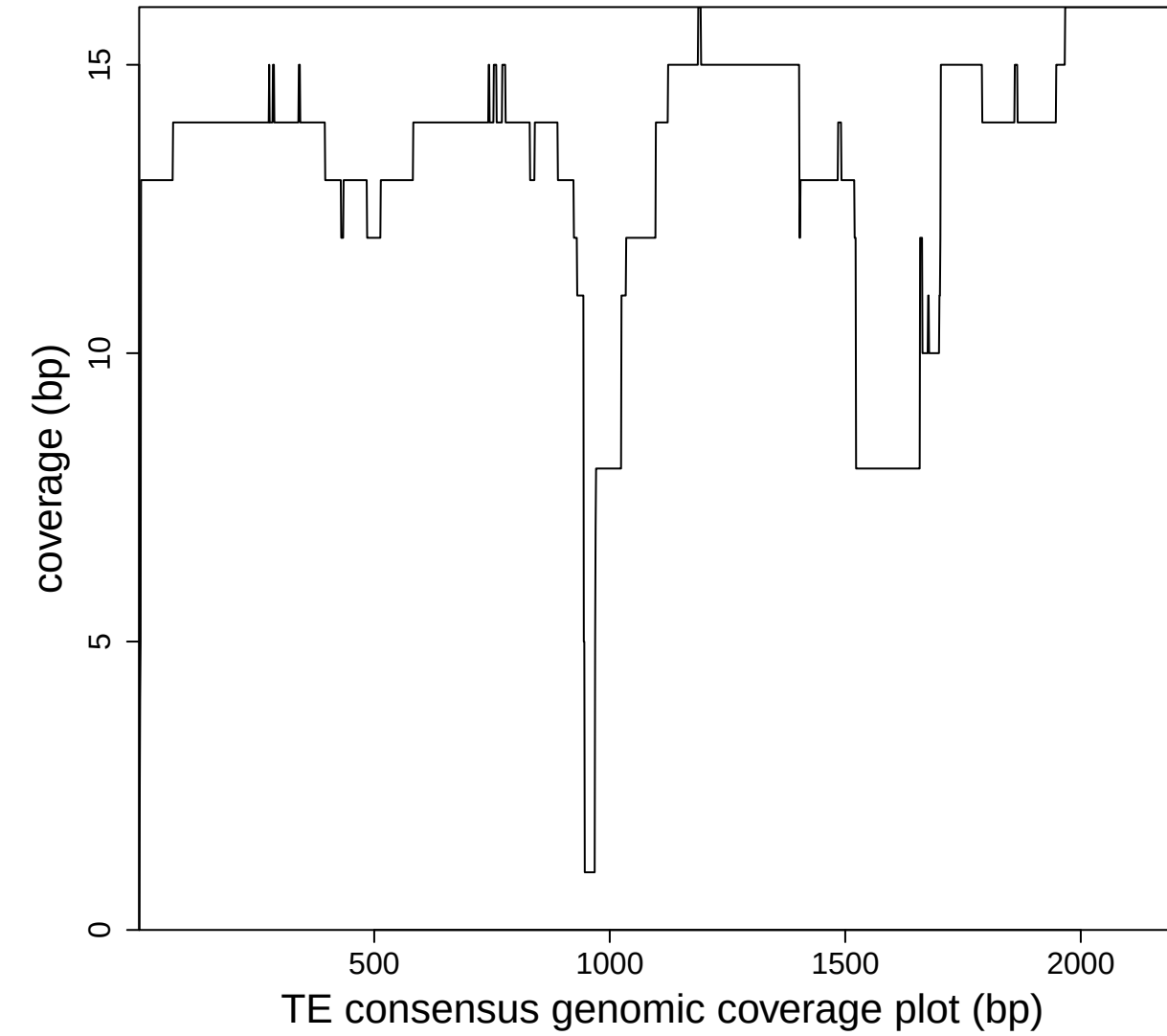
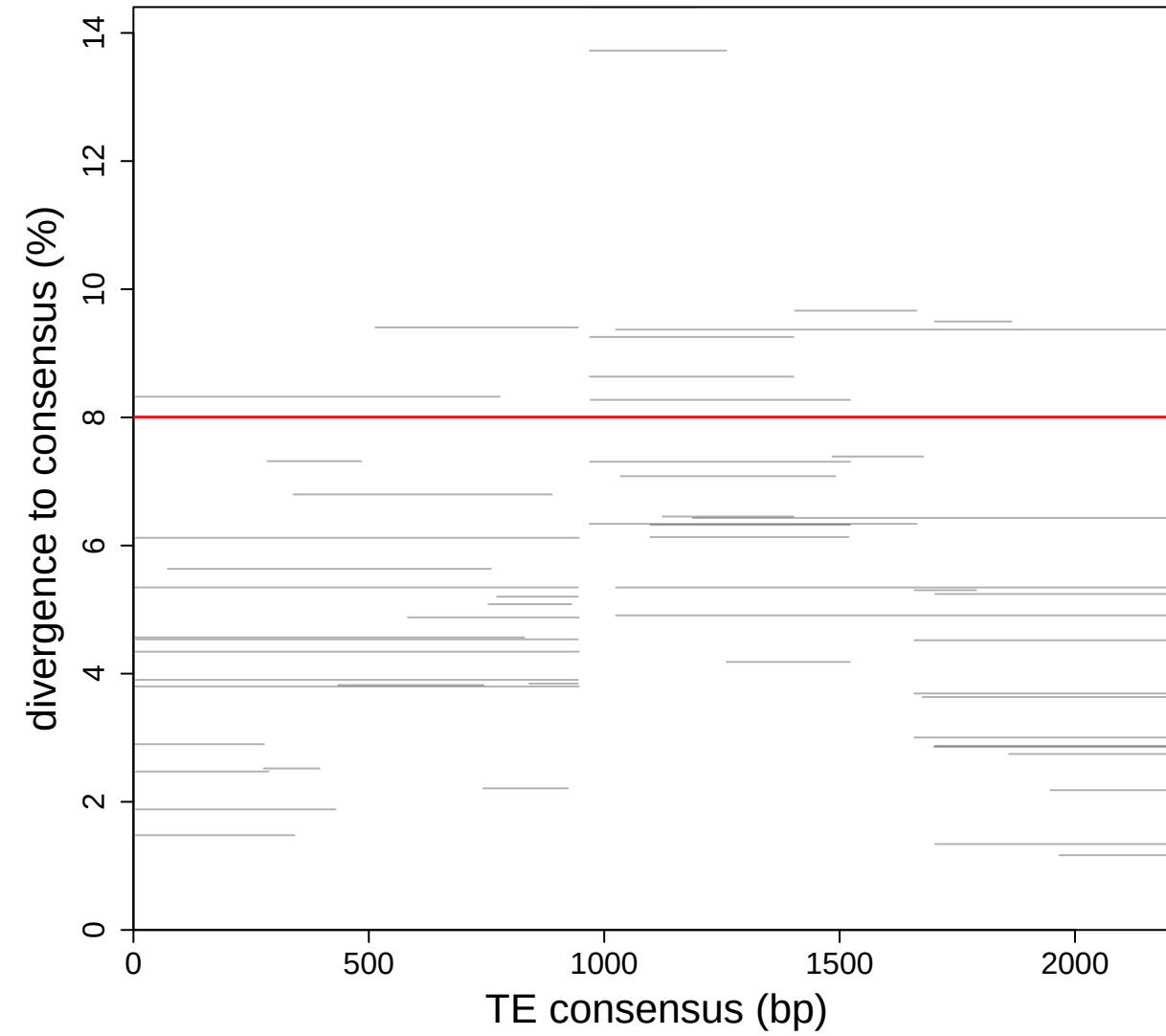
2000

2500



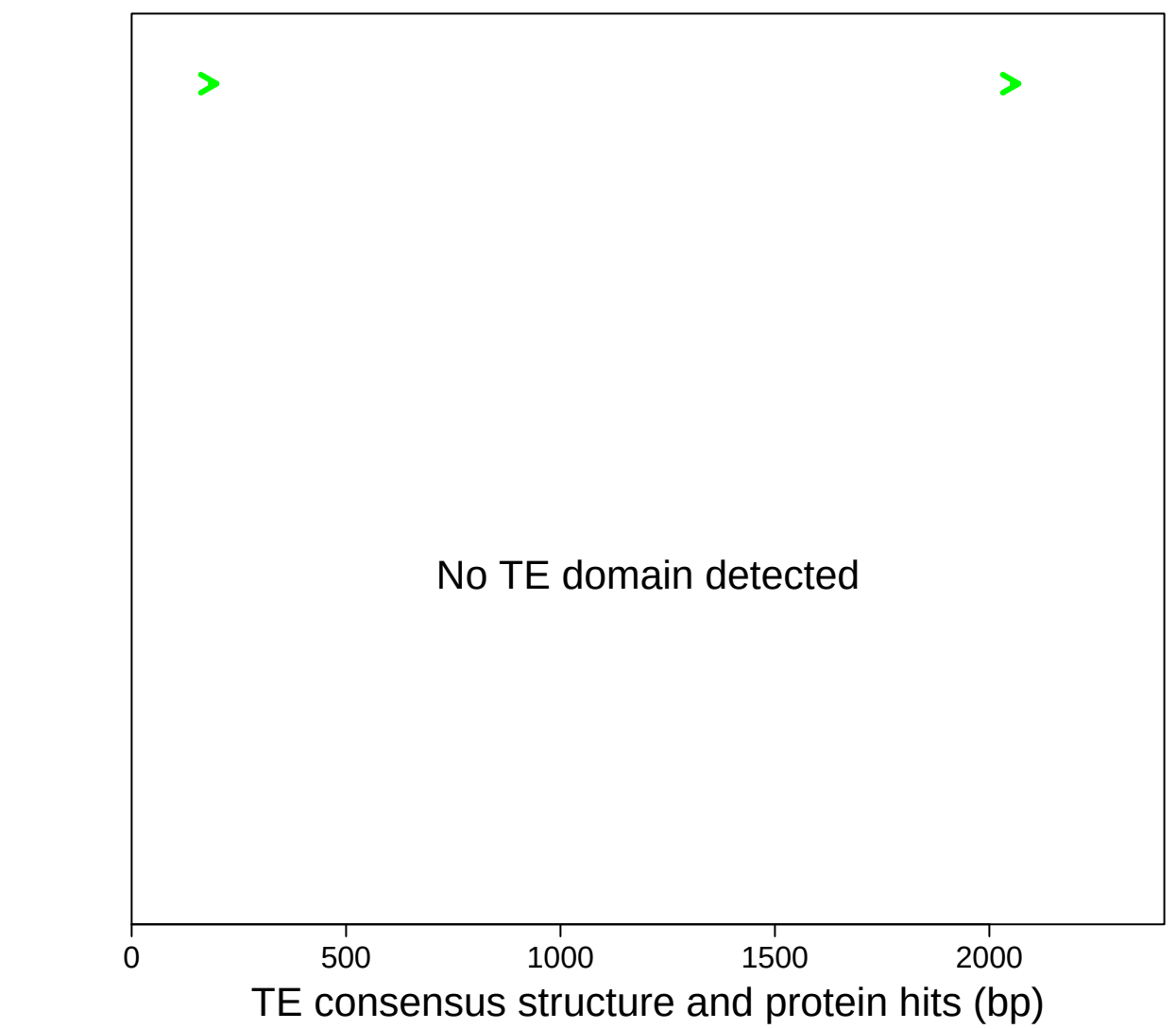
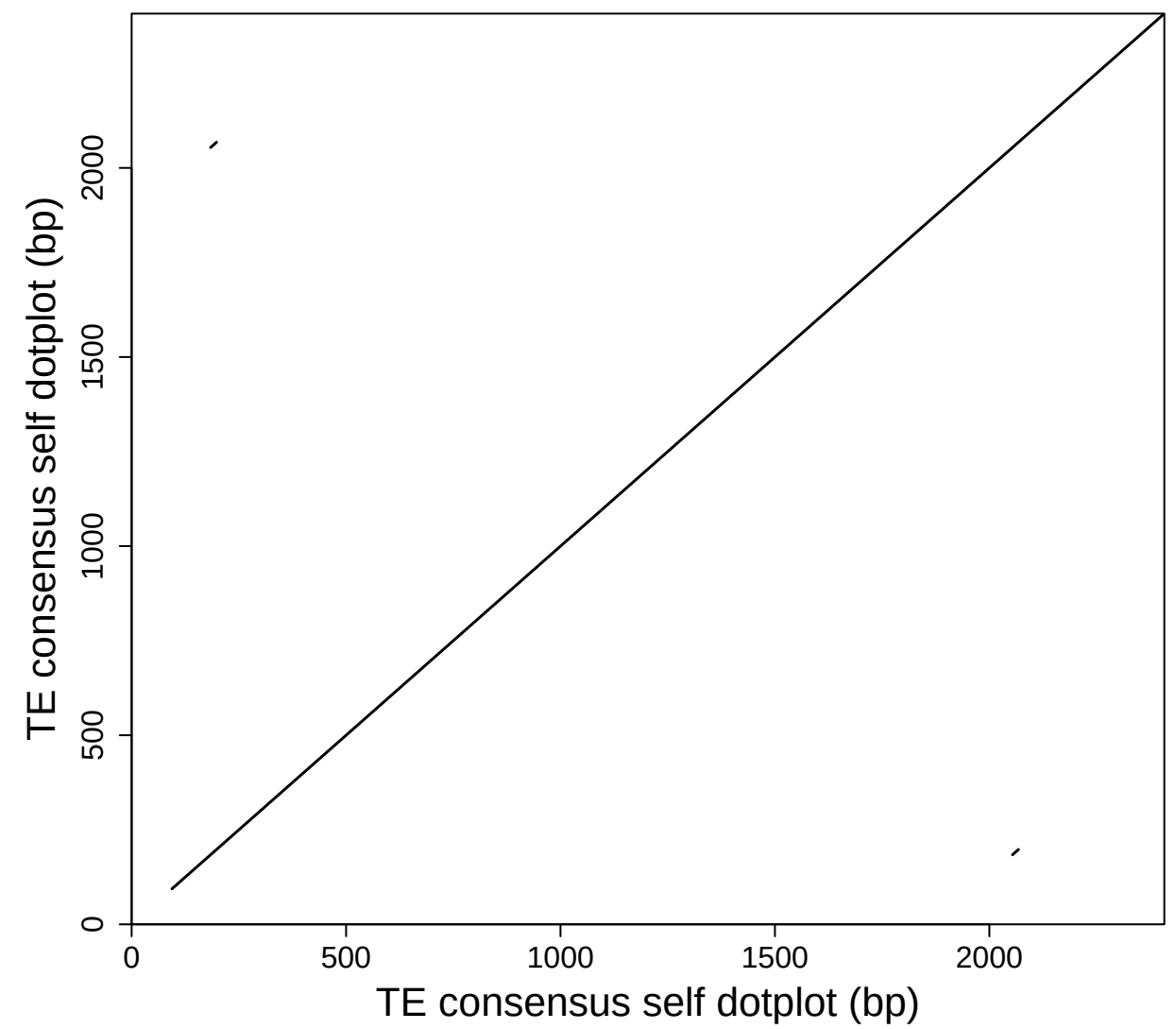
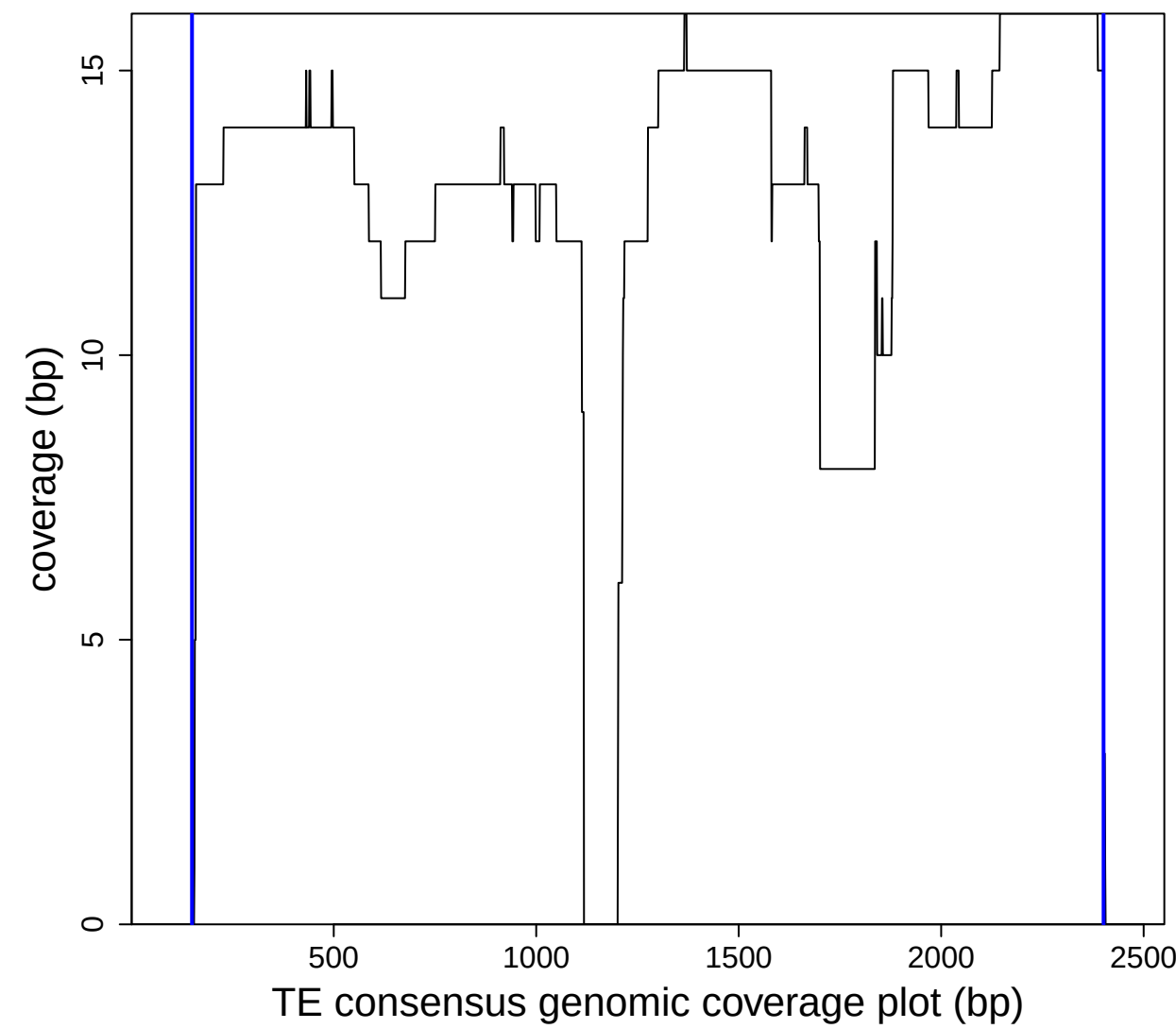
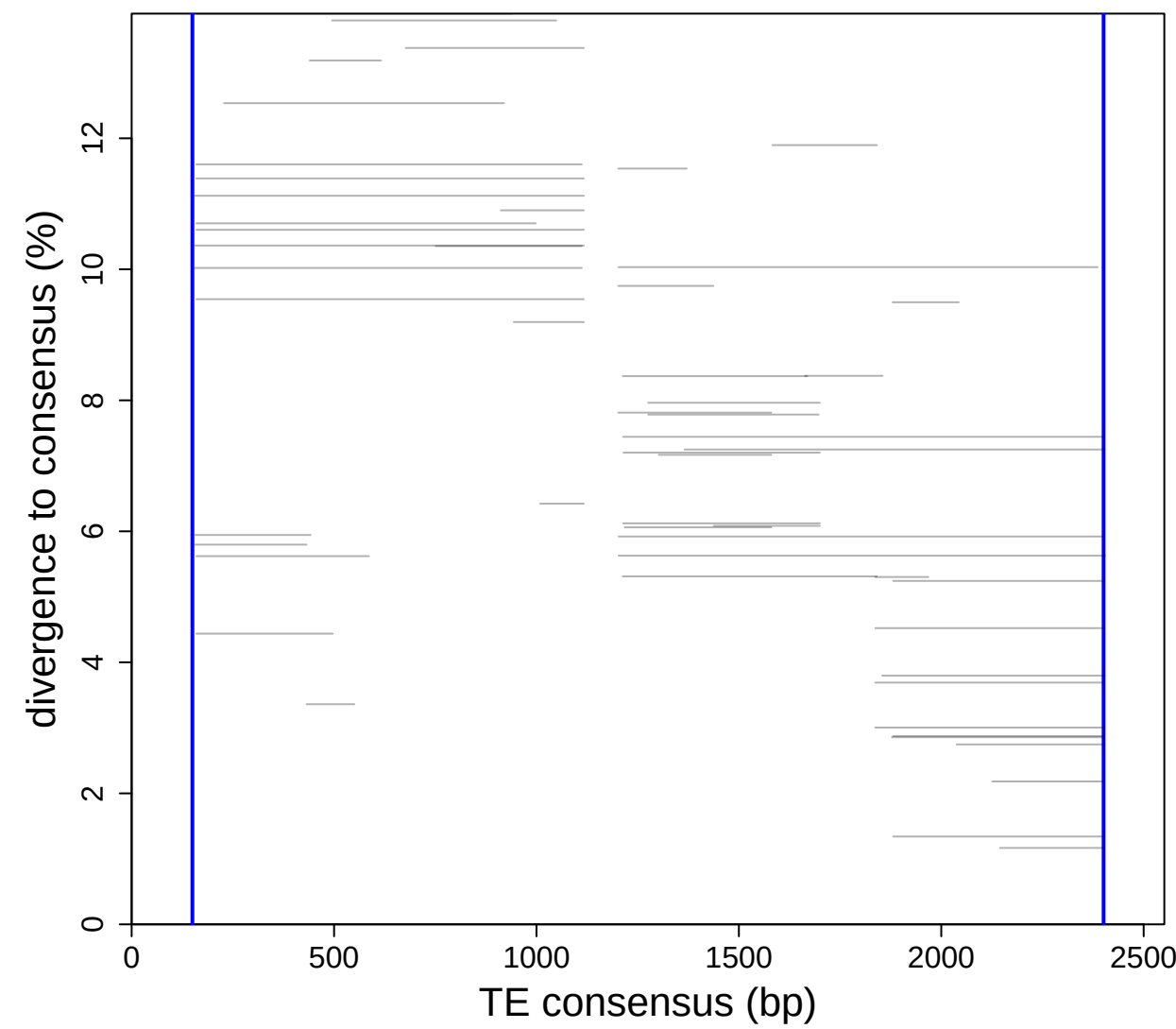
b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 2223bp; fragments: 55; full length: 1 (>=2000.7bp)

After TEtrimmer 2223 bp

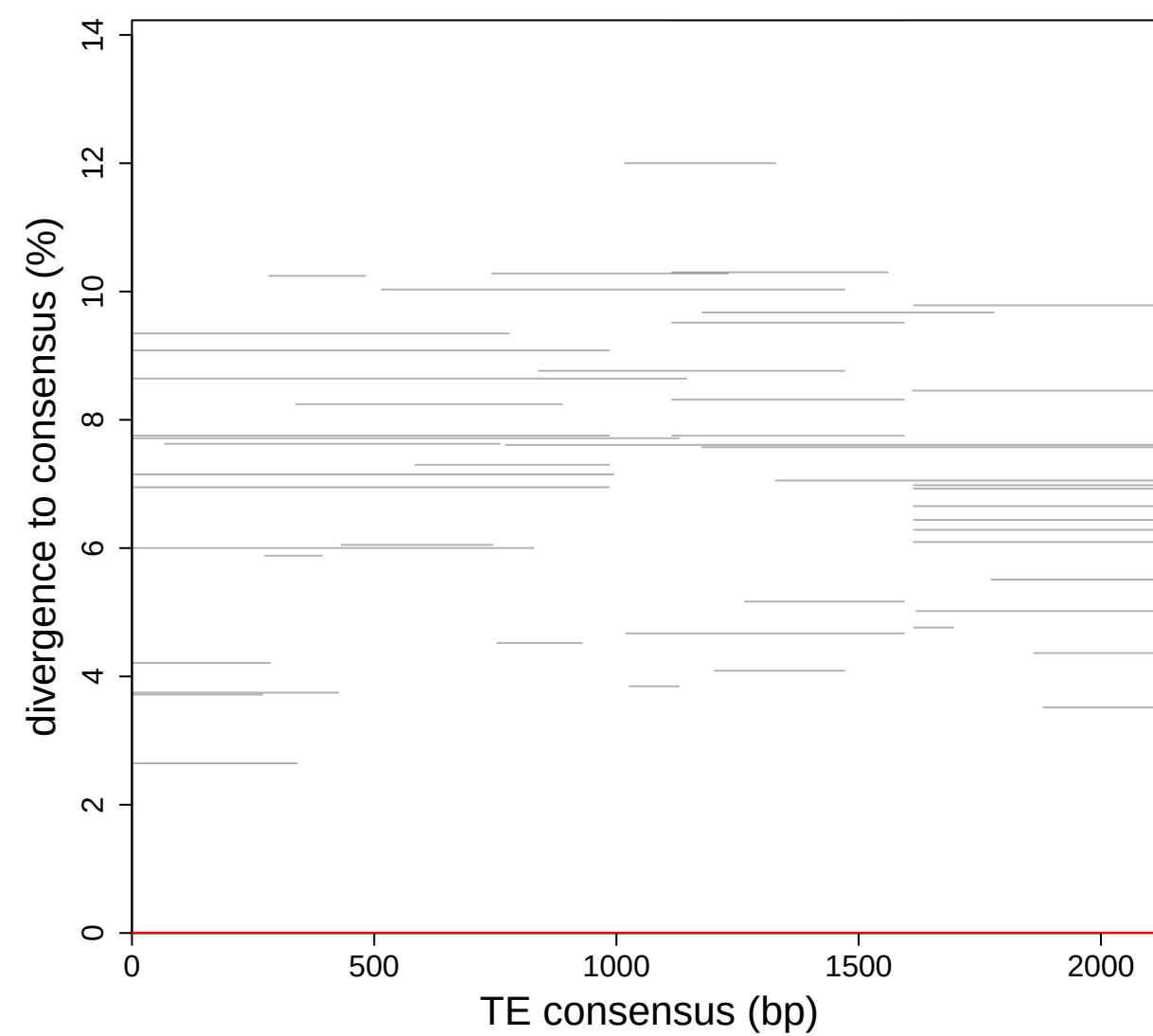


.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 2551bp; fragments: 54; full length: 0 (>=2295.9bp)

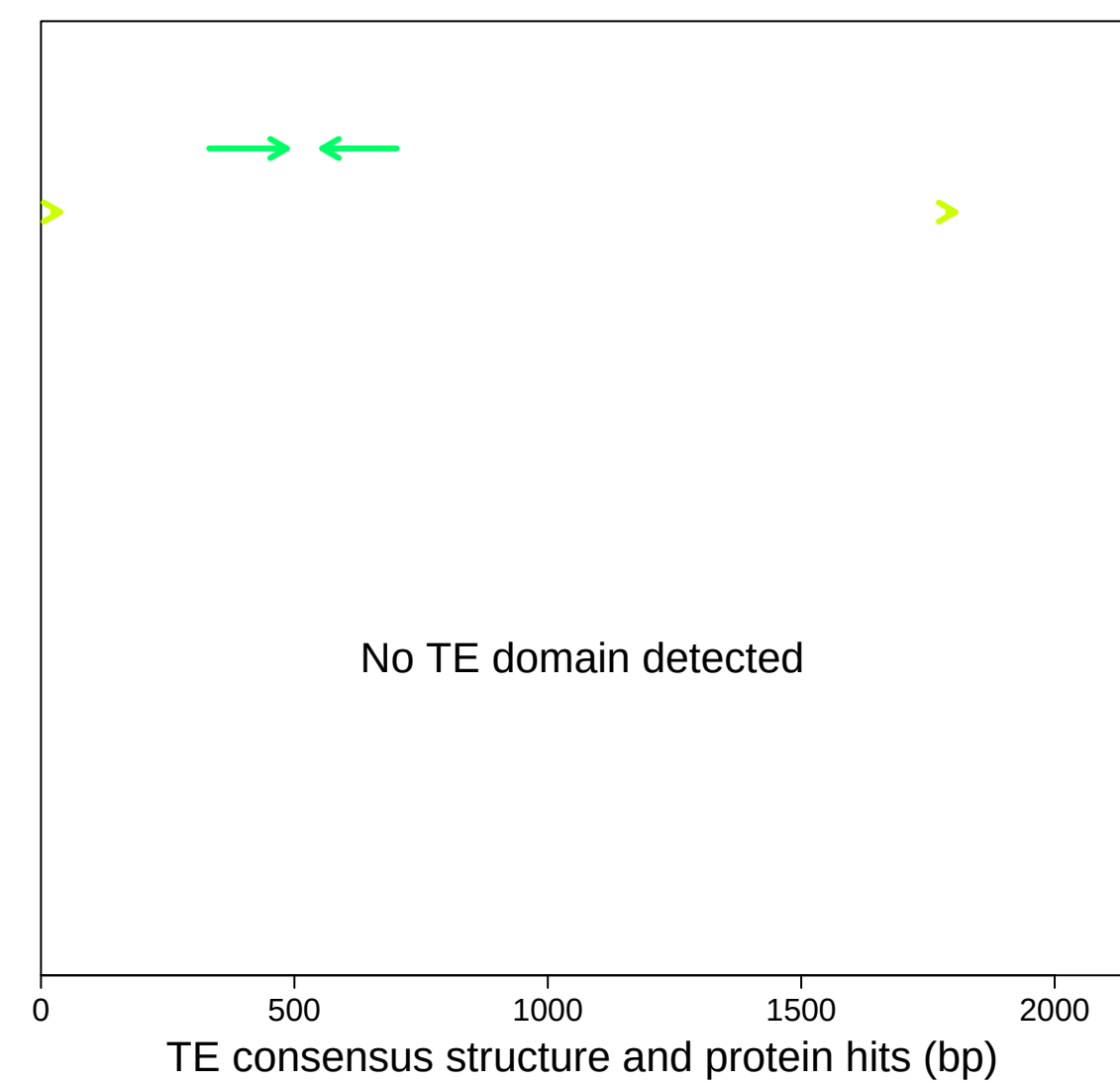
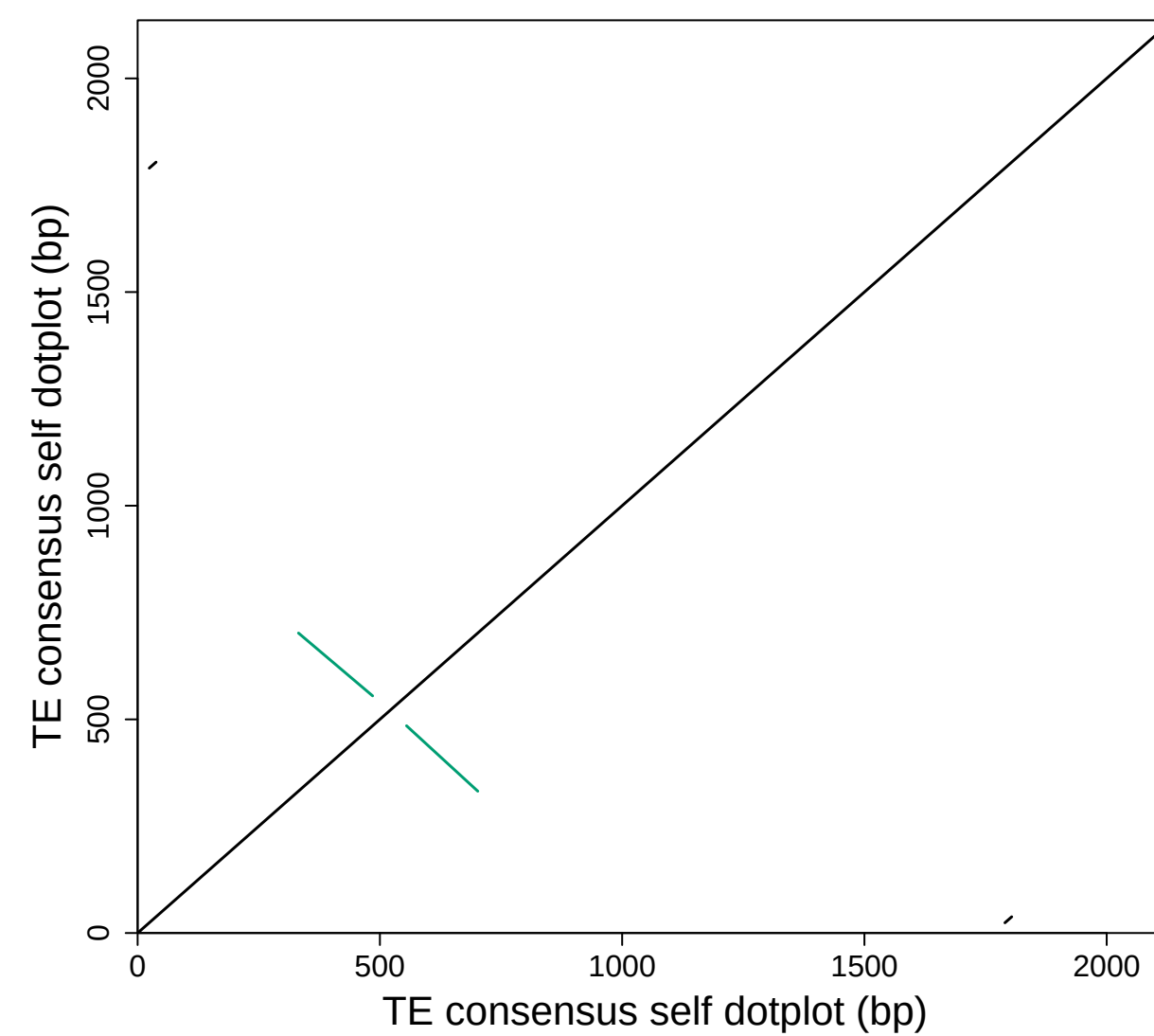
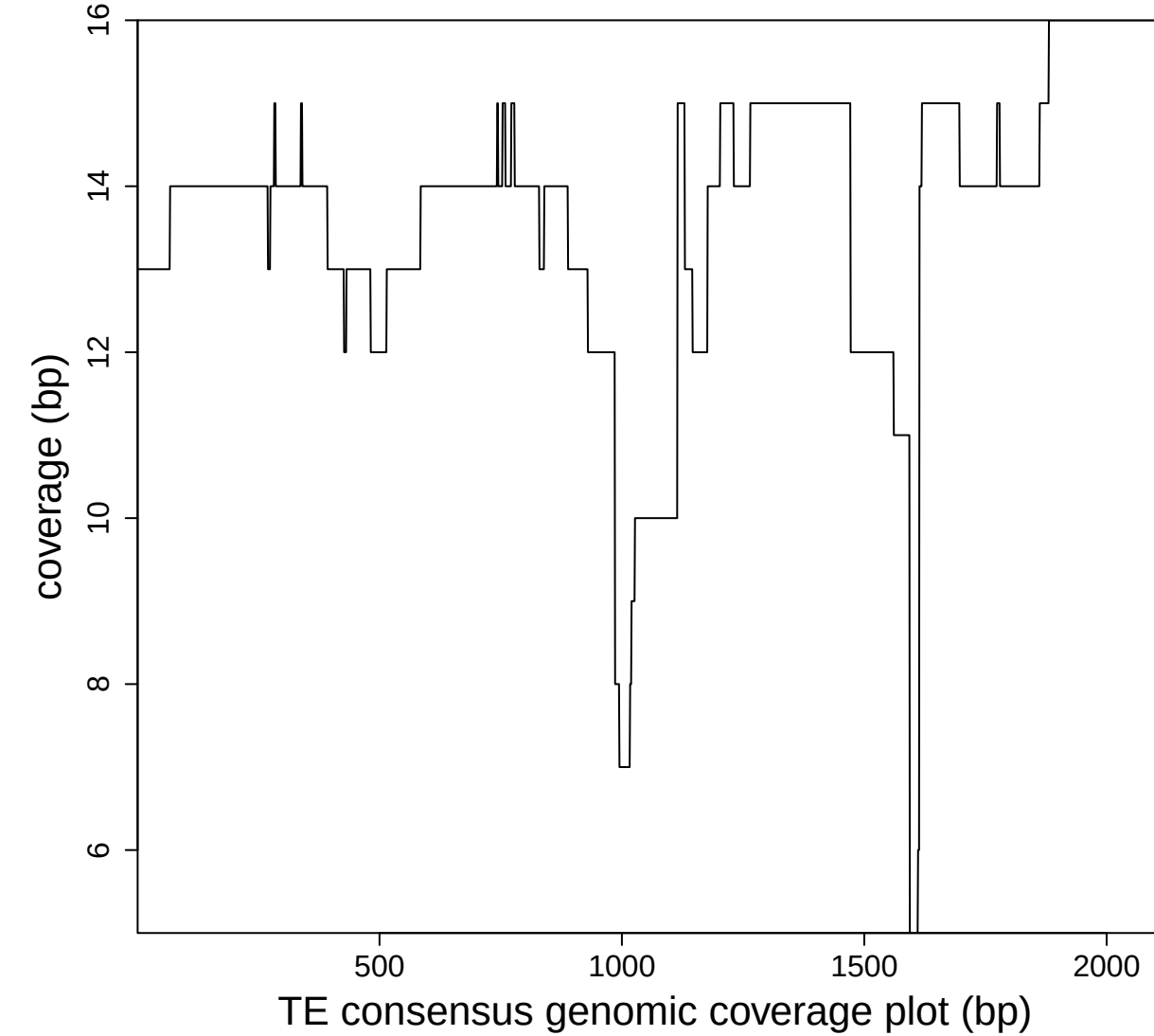
After TEtrimmer Extended plot Blue lines are boundaries



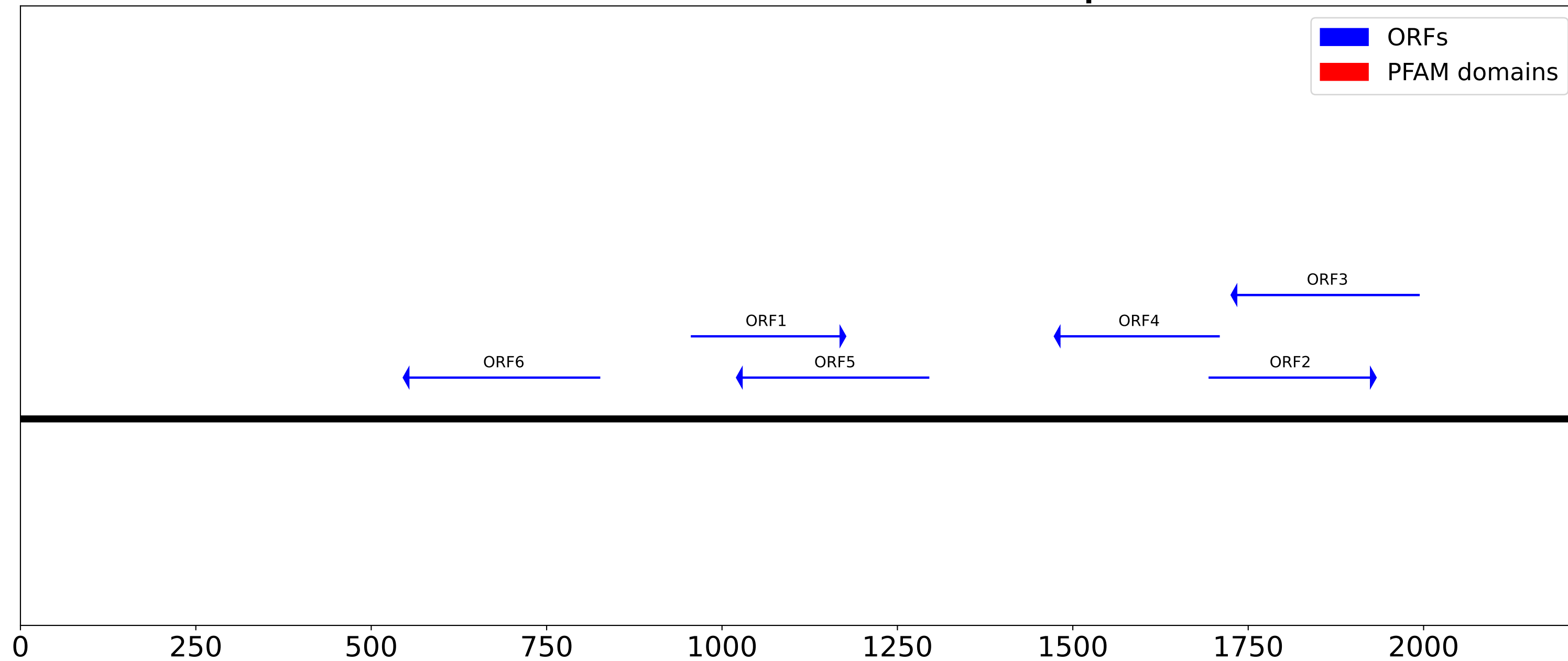
TE: ltr_1_family_571
size: 2136bp; fragments: 50; full length: 1 (>=1922.4bp)



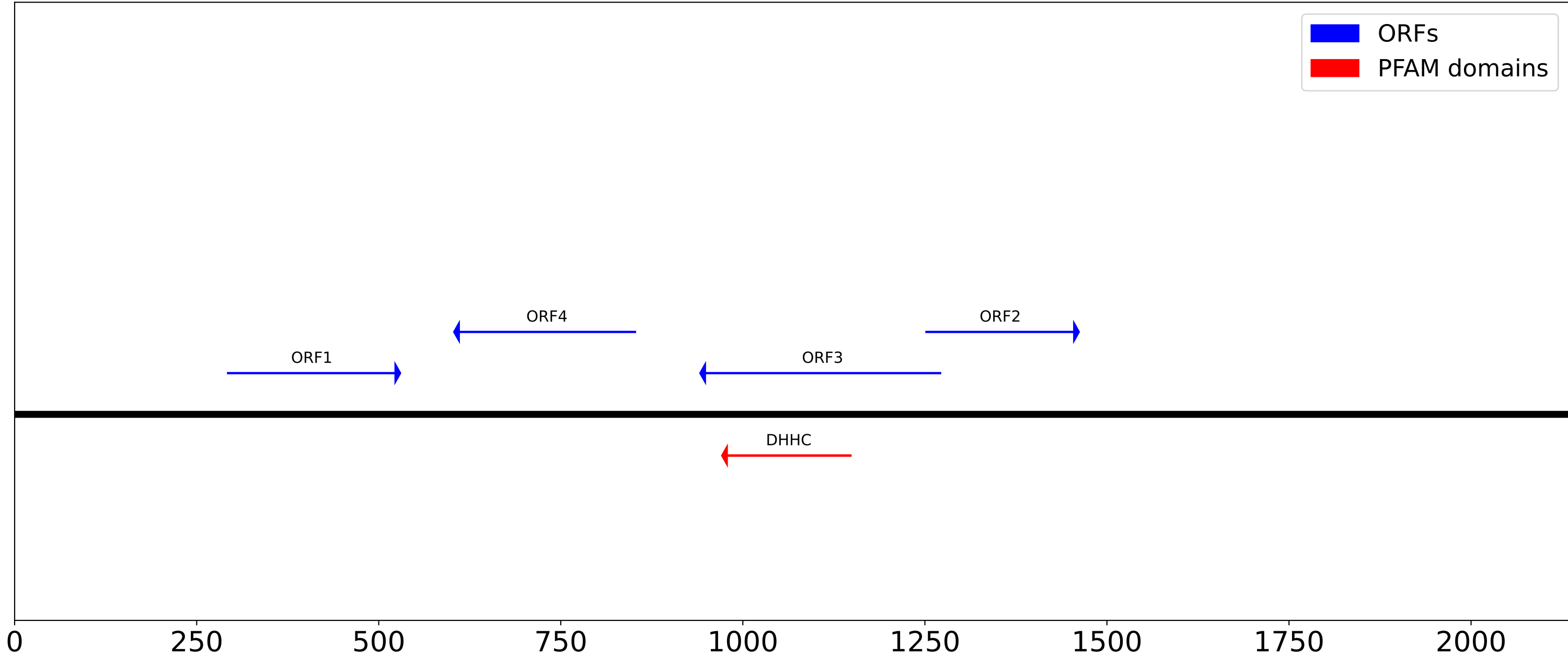
Before TEtrimmer 2136 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

