

Start crop Point

End crop Point

0 -

10 -

0

500

1000

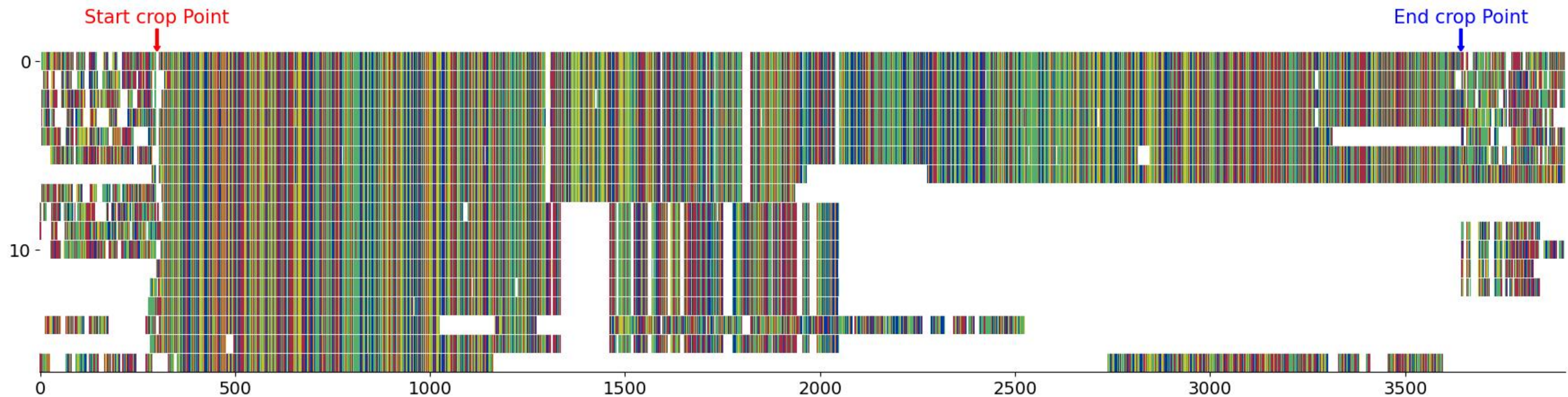
1500

2000

2500

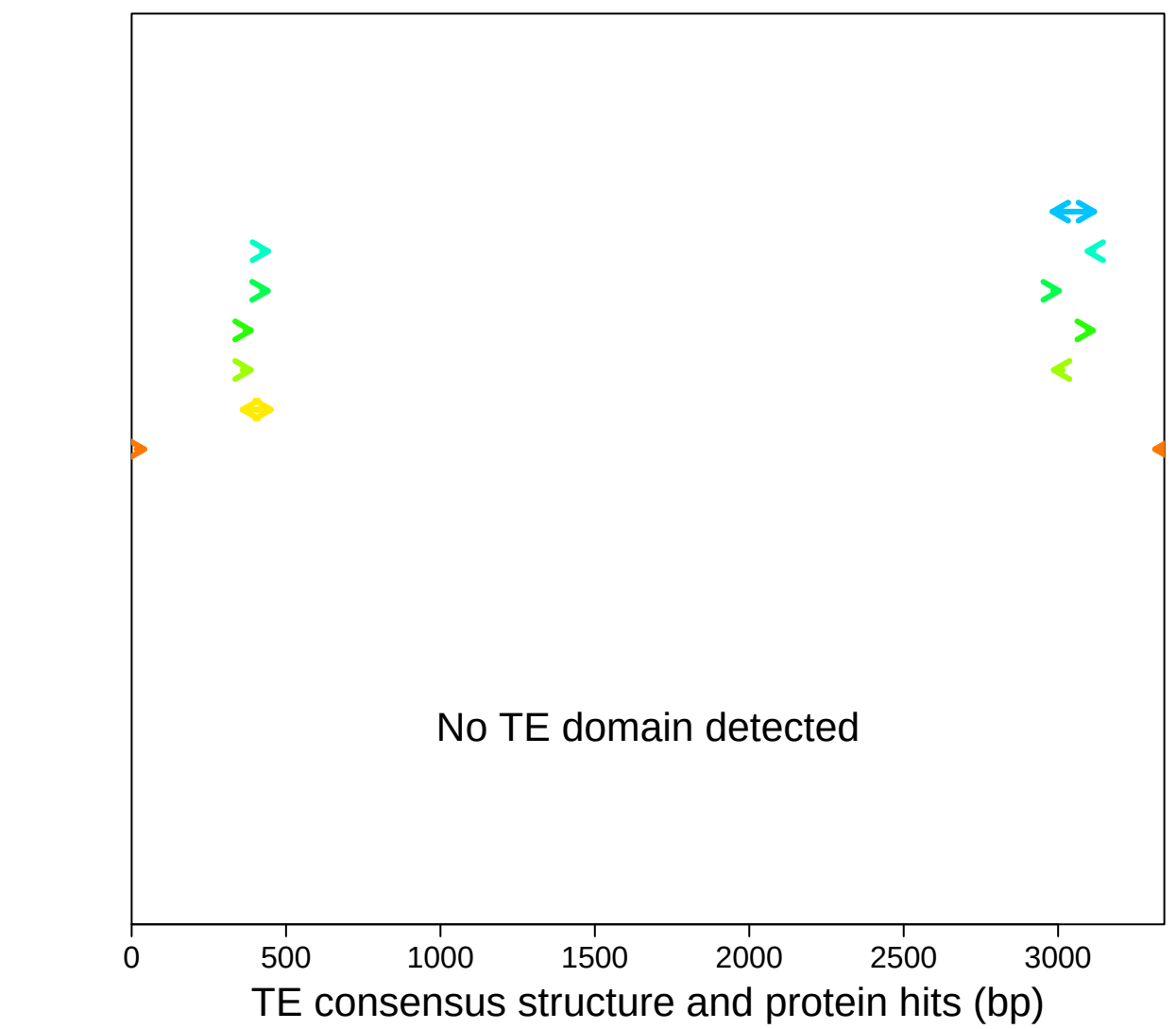
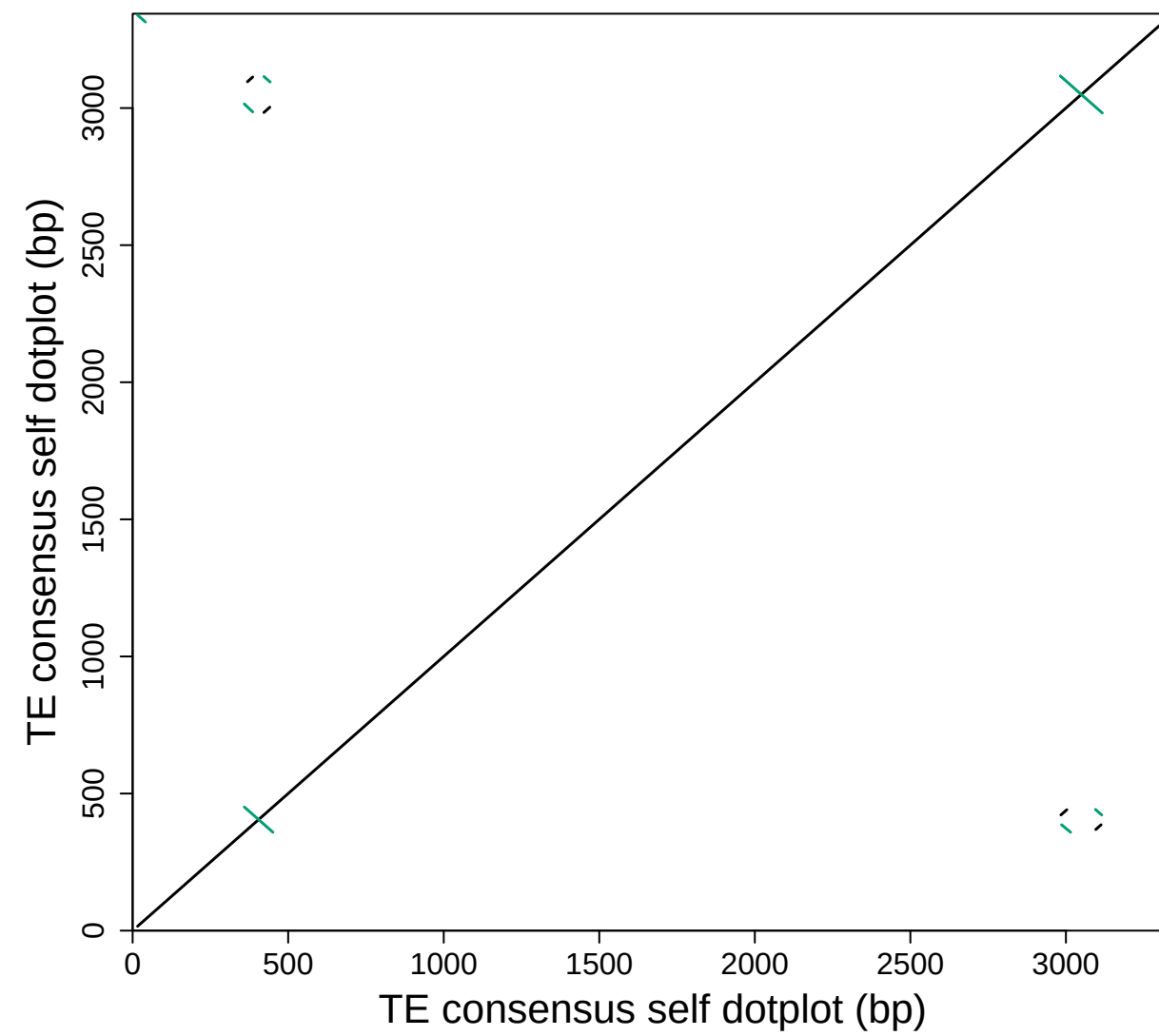
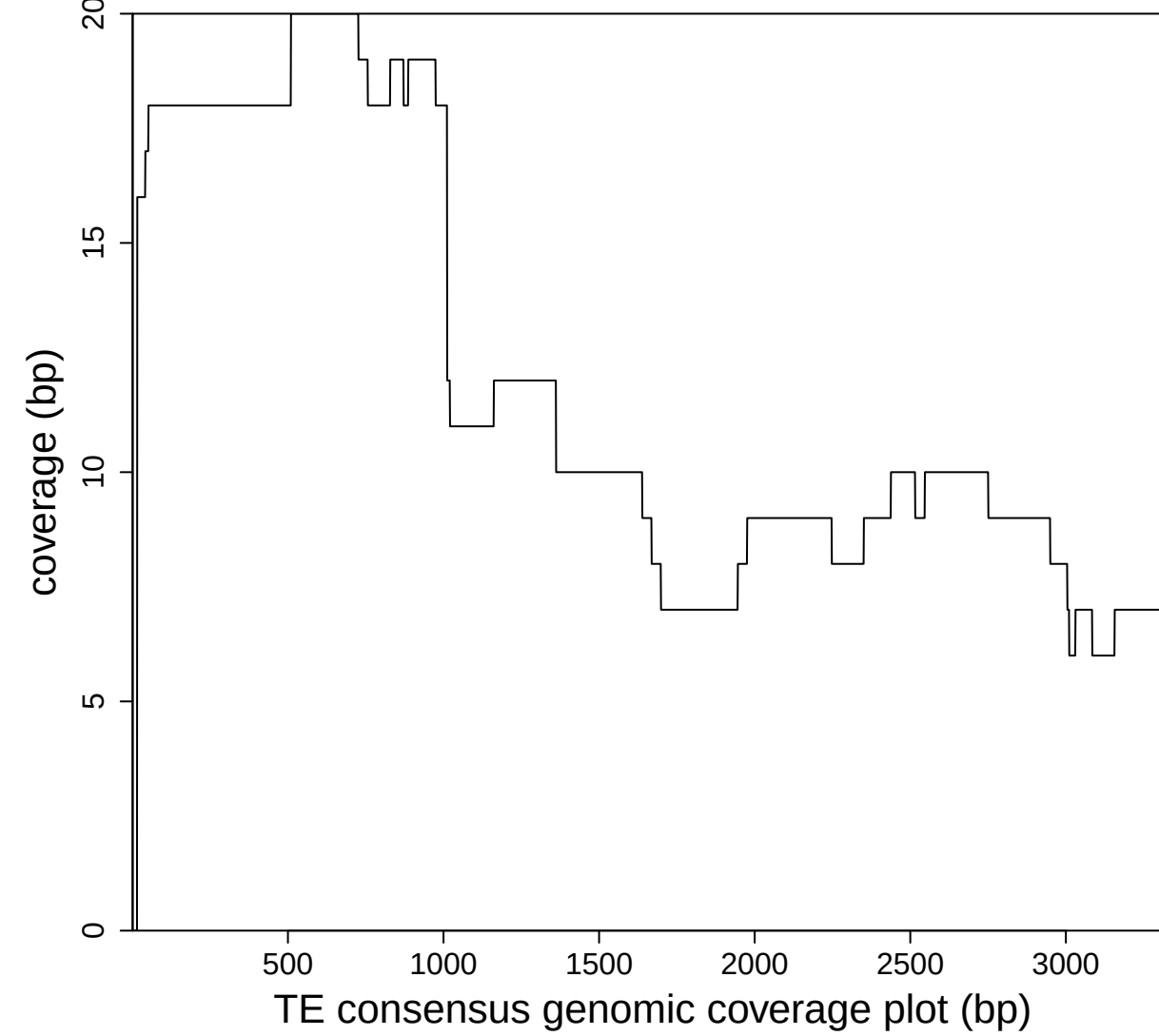
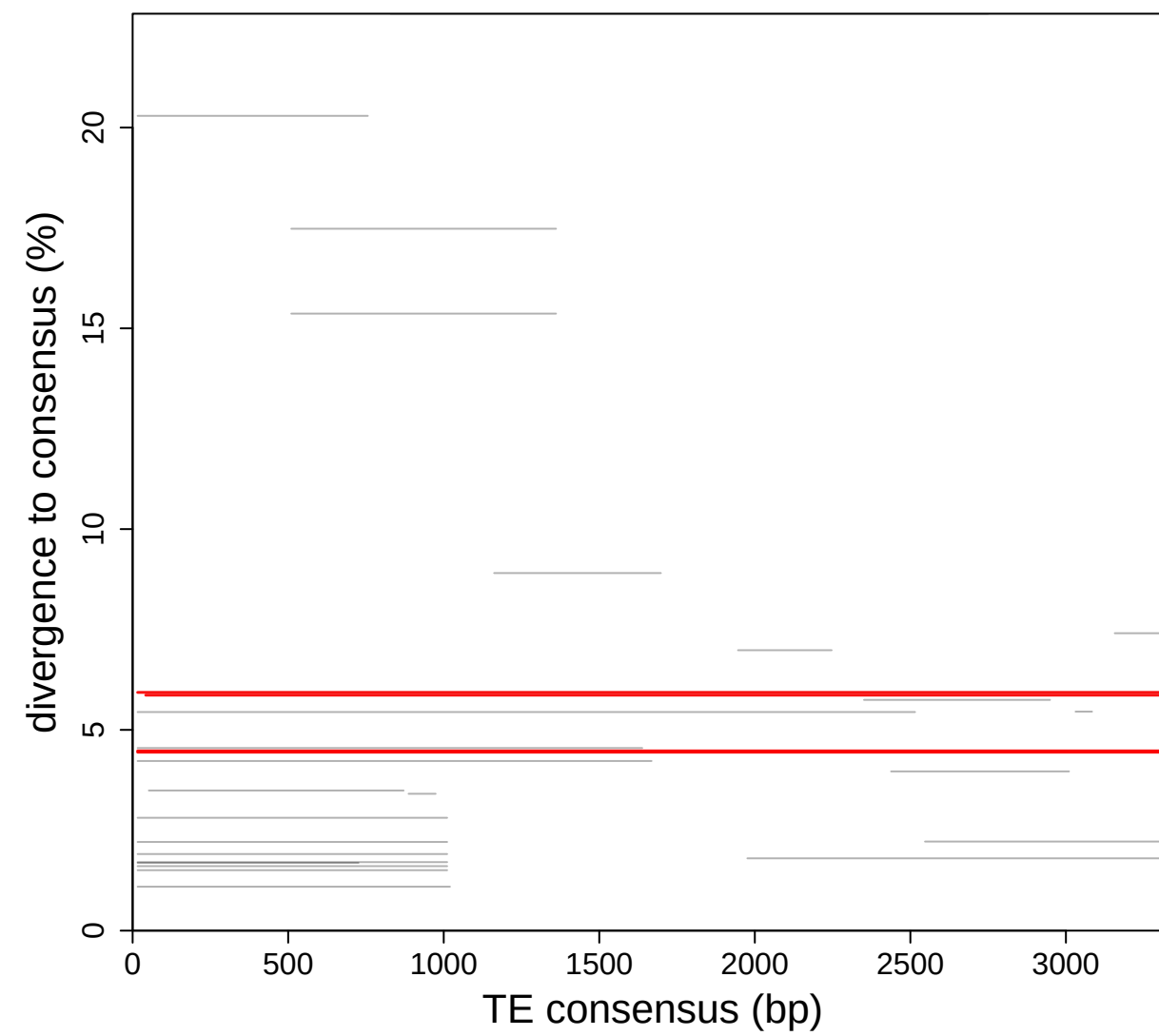
3000

3500



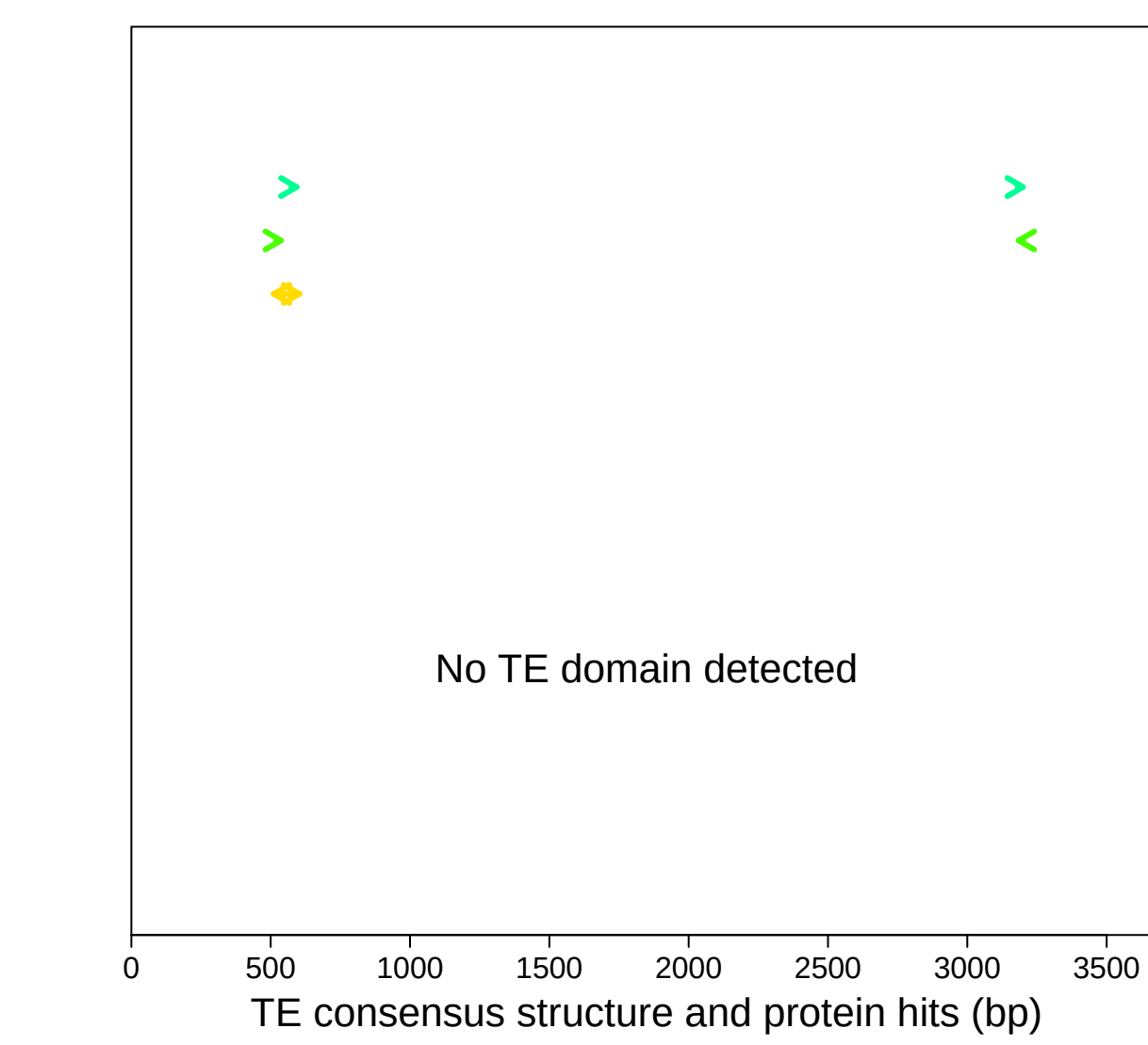
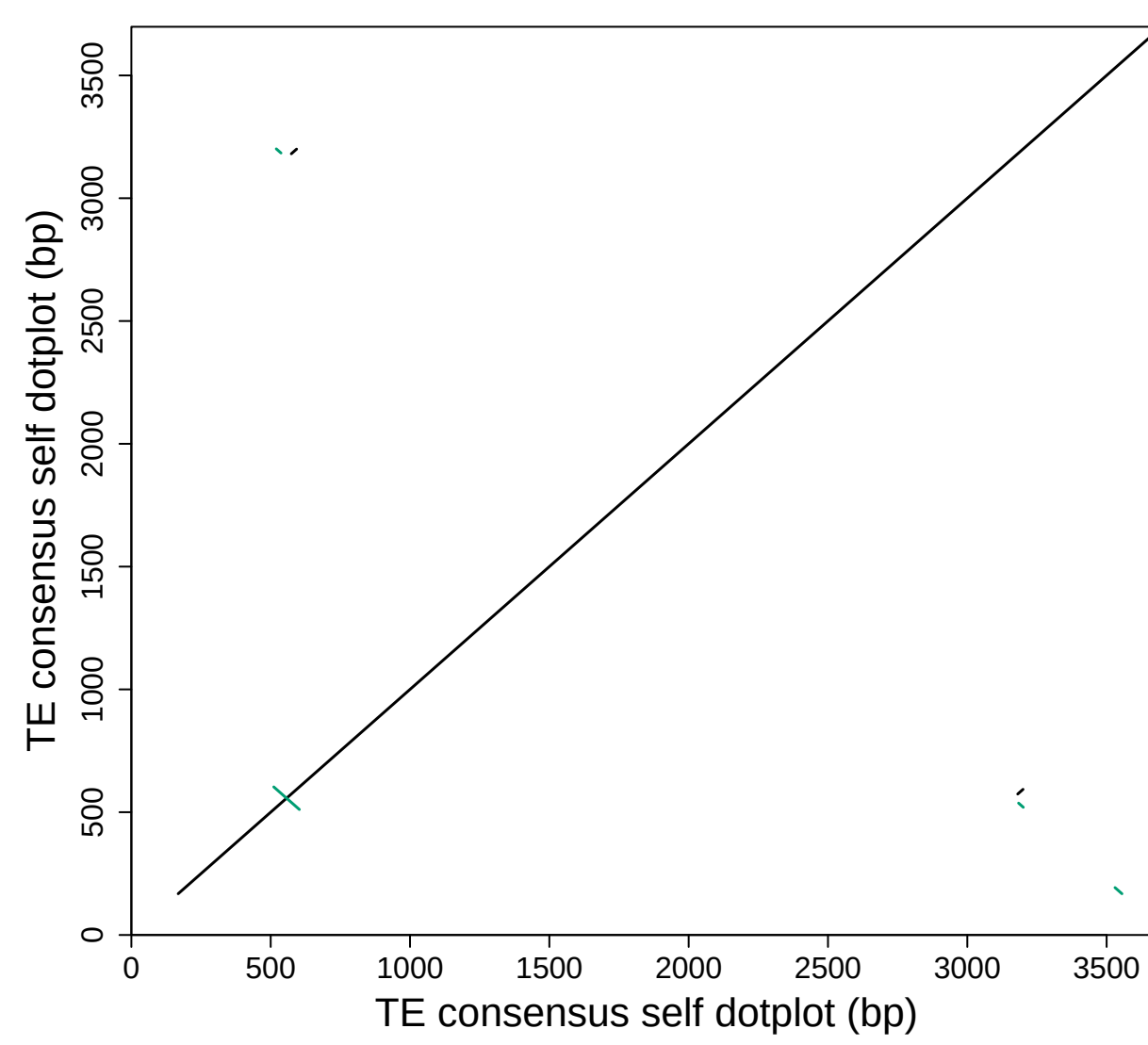
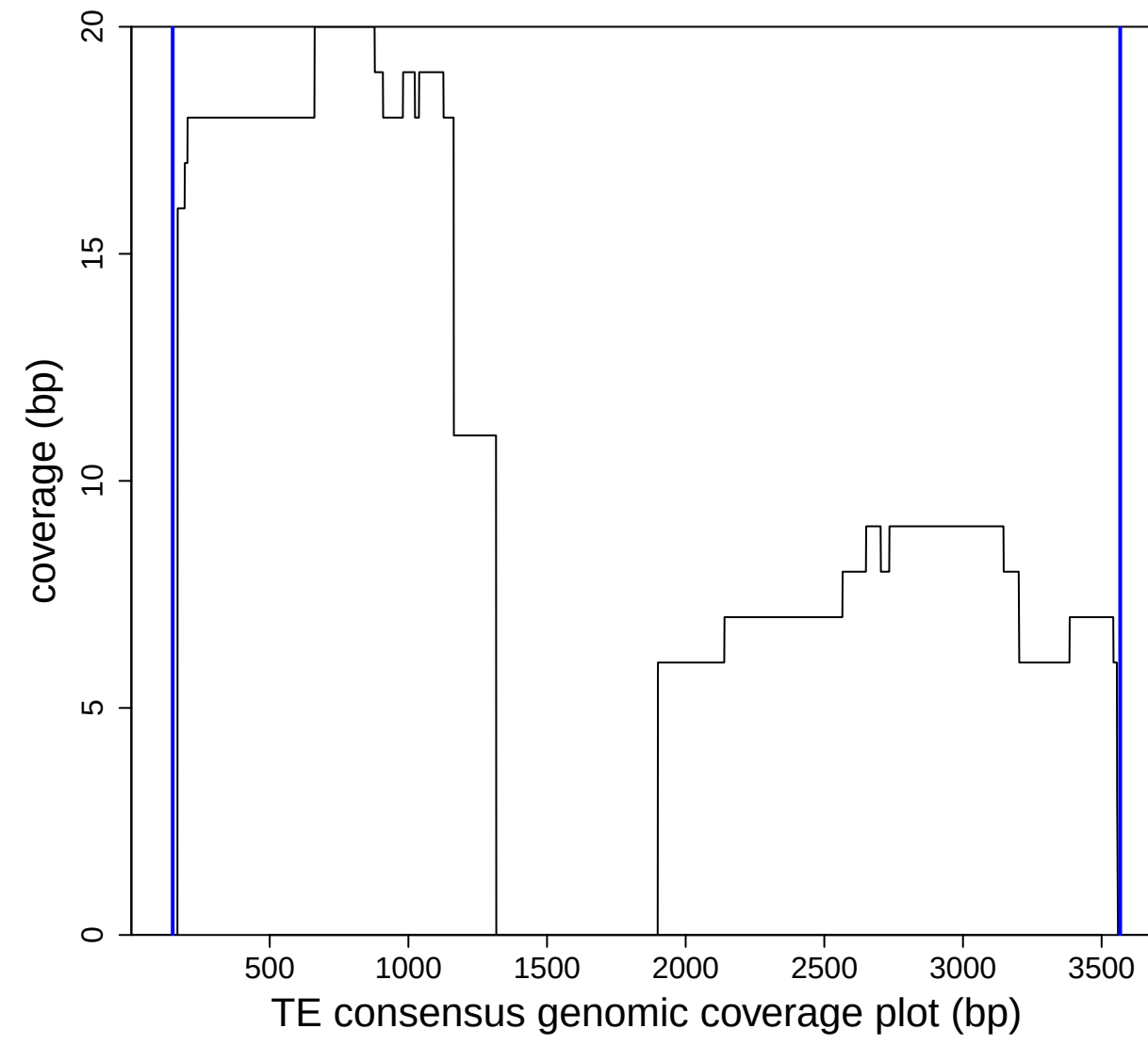
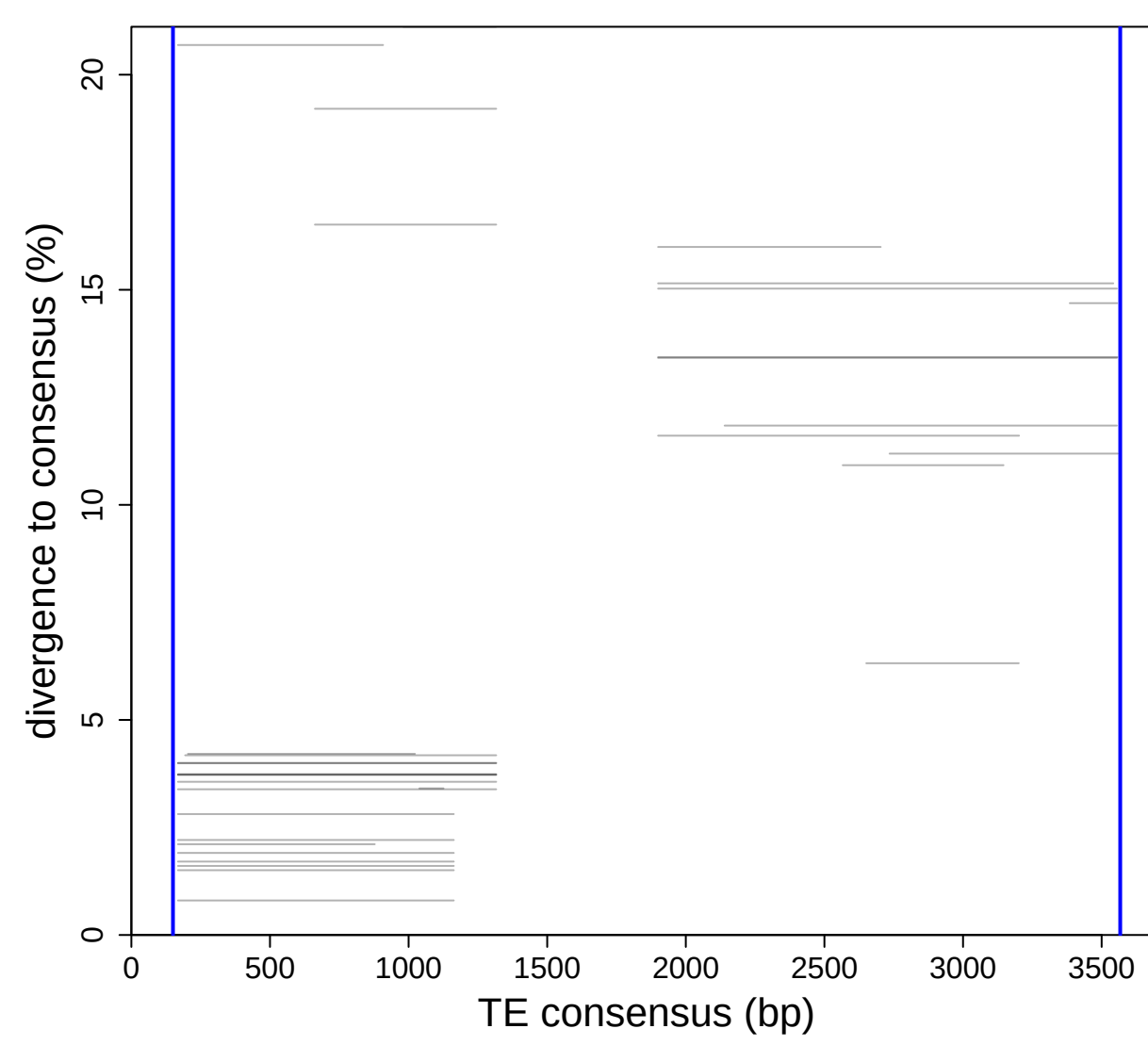
b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 3344bp; fragments: 30; full length: 4 (>=3009.6bp)

After TEtrimmer 3344 bp

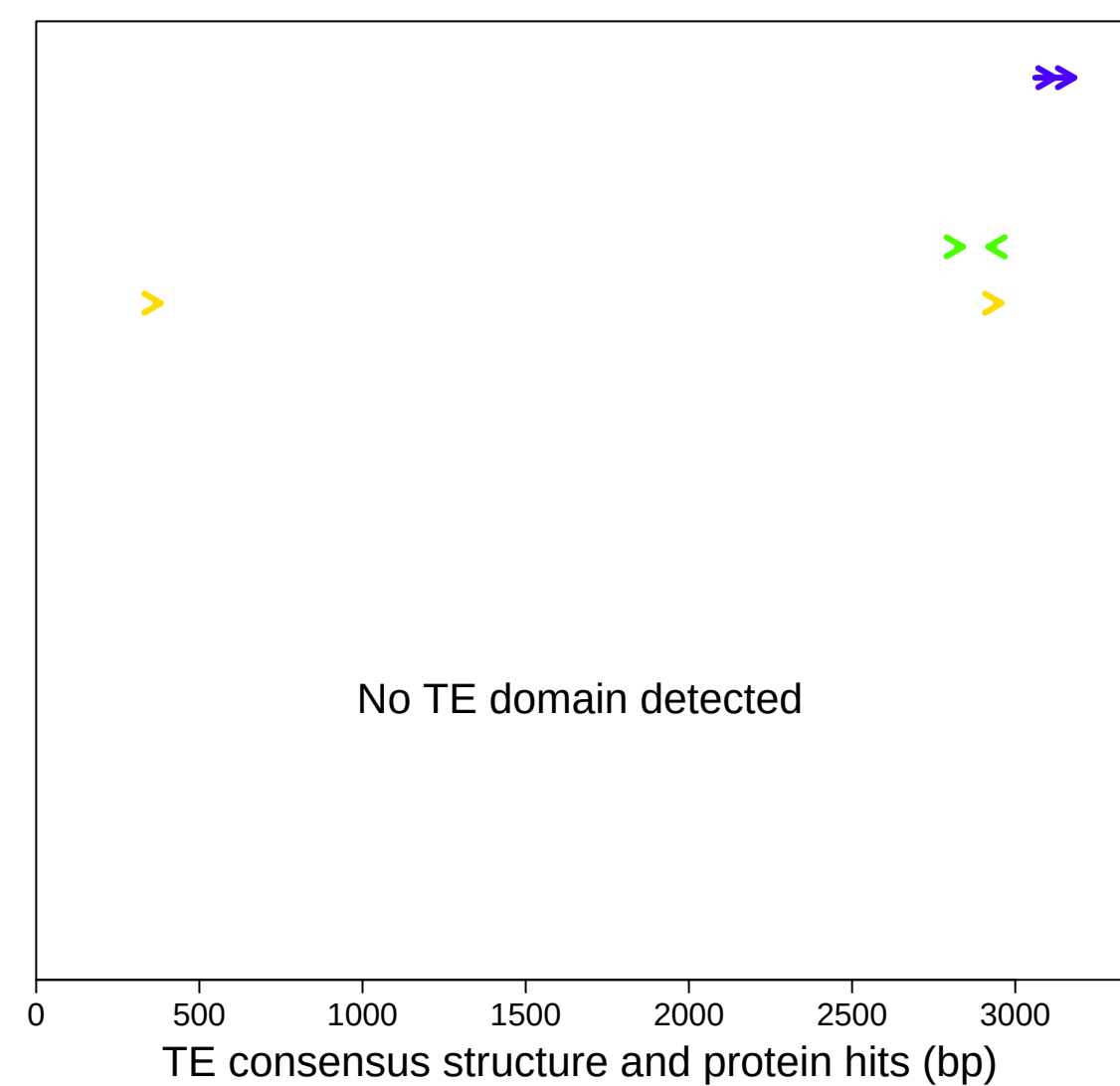
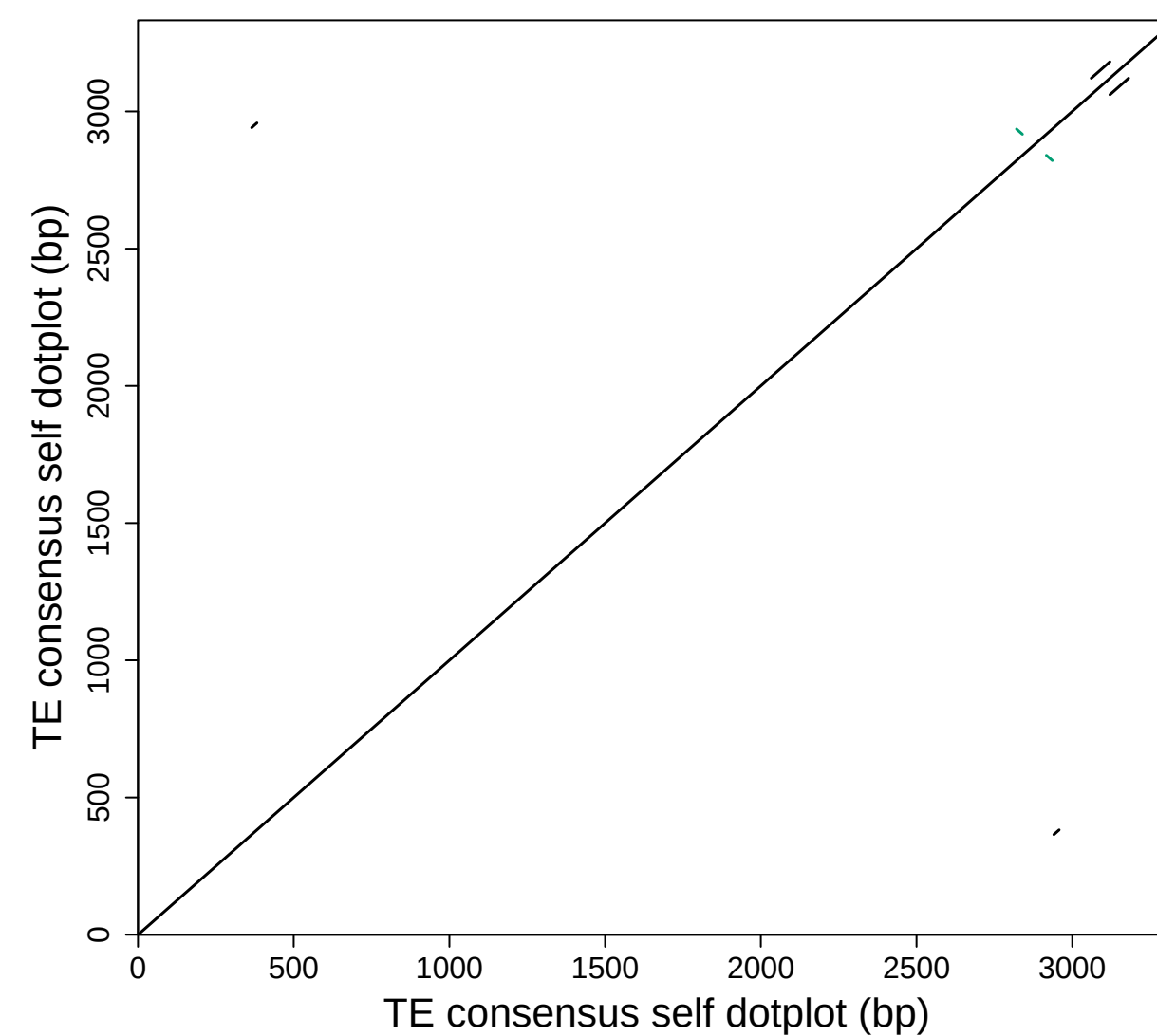
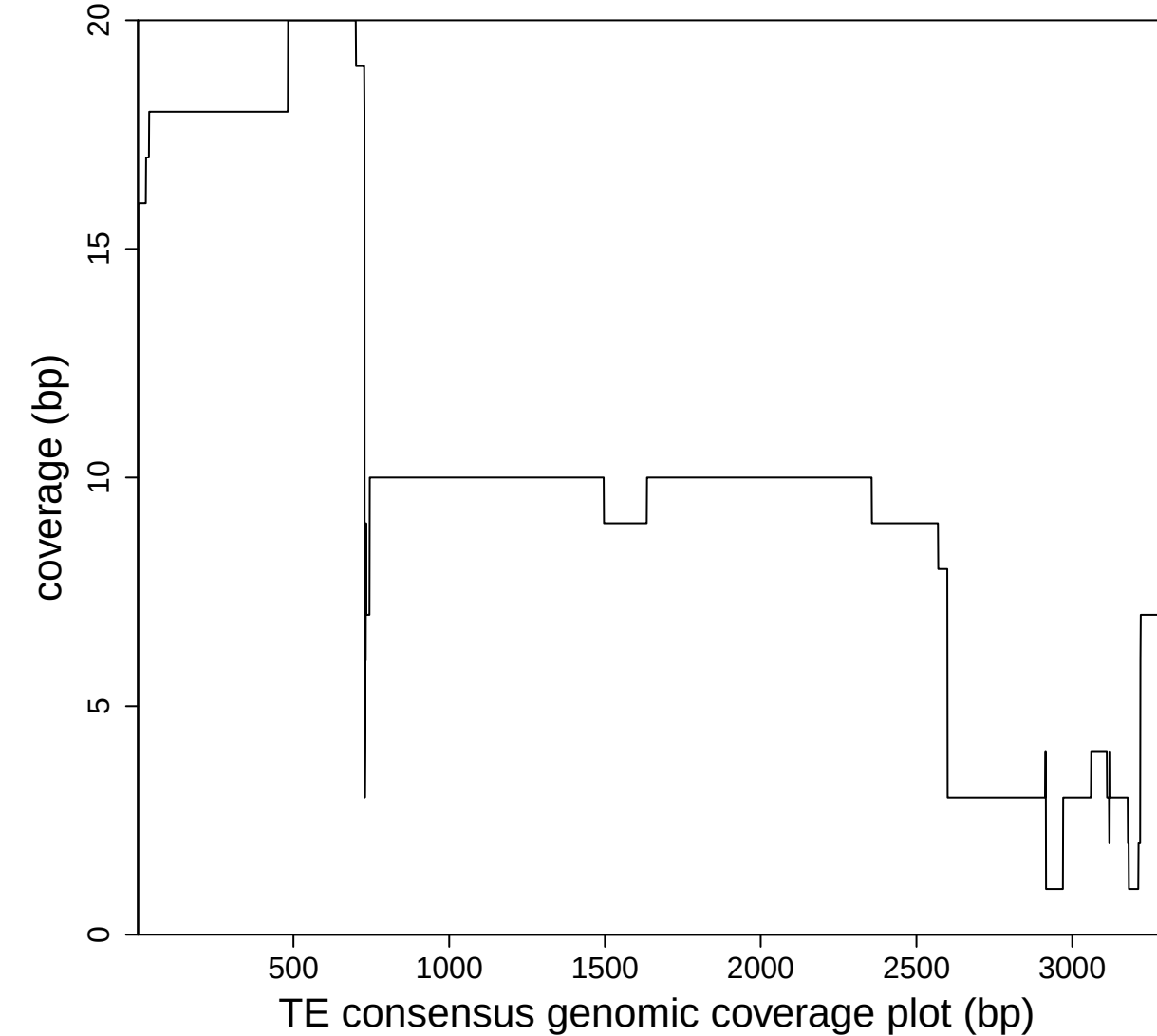
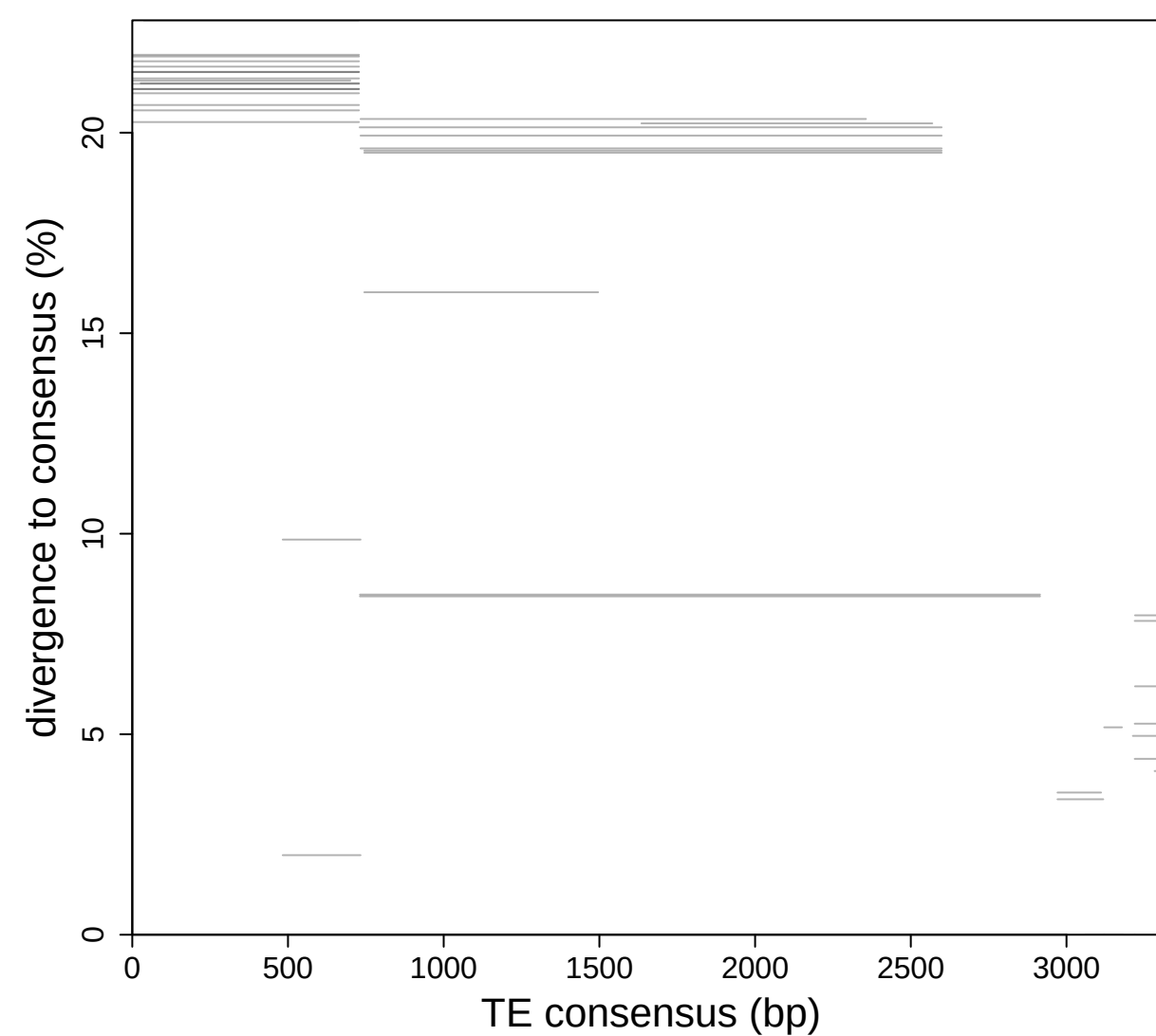


.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 3717bp; fragments: 33; full length: 0 (>=3345.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

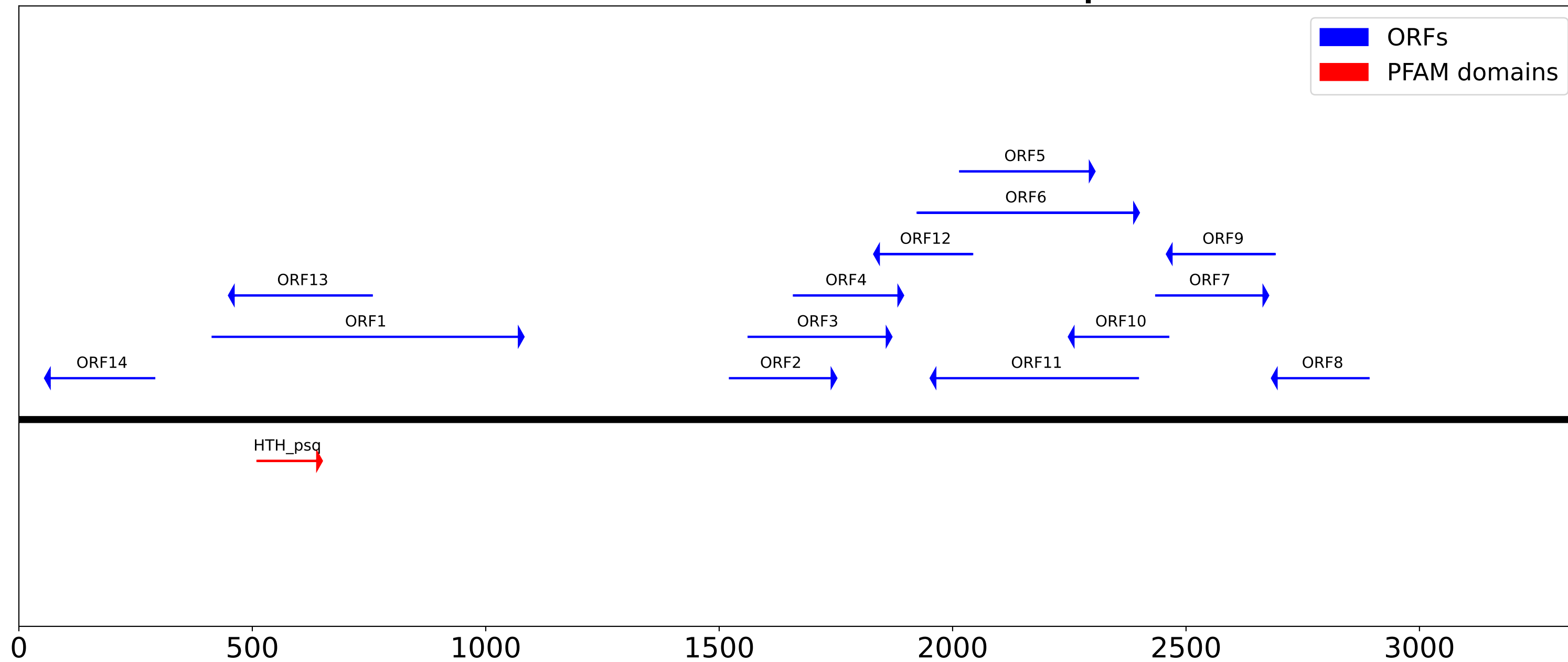


TE: ltr_1_family_796
size: 3332bp; fragments: 44; full length: 0 (≥ 2998.8 bp)

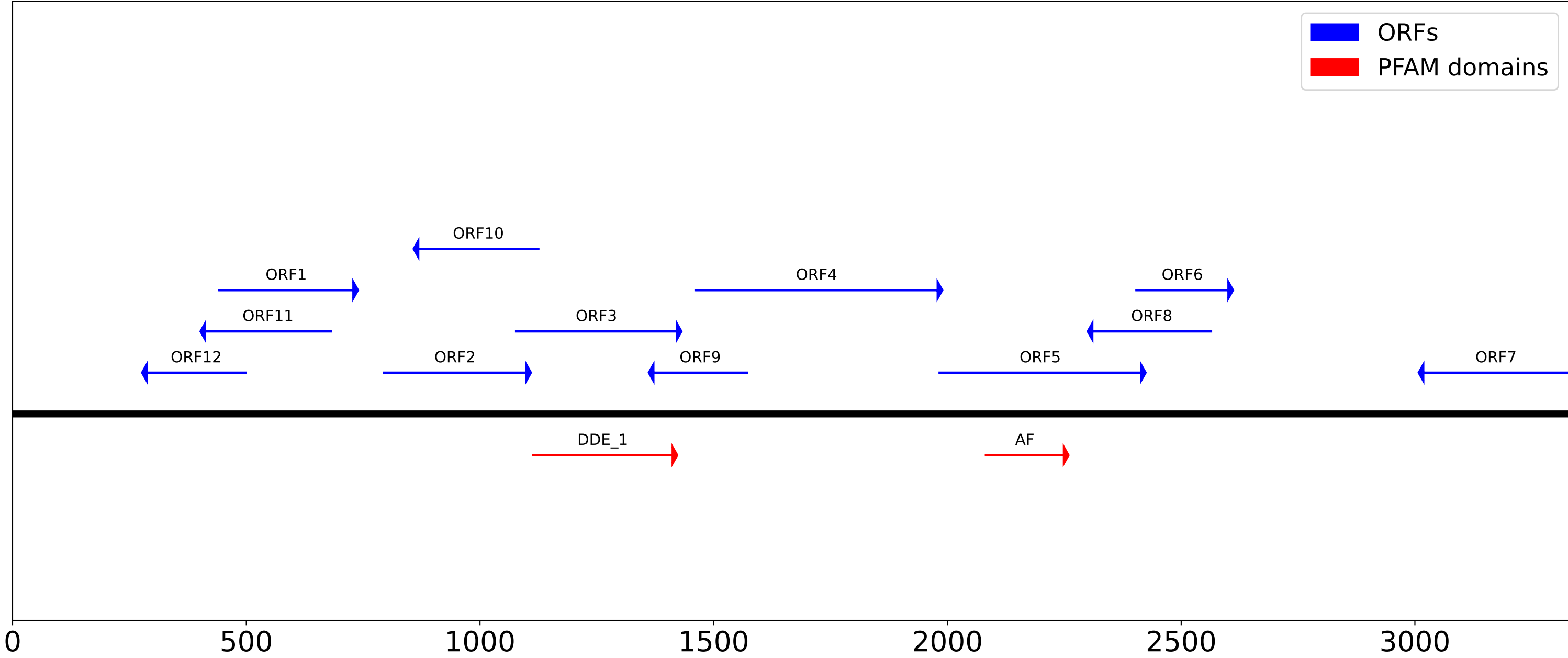


Before TEtrimmer 3332 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

