

Start crop Point

End crop Point

0

10

0

500

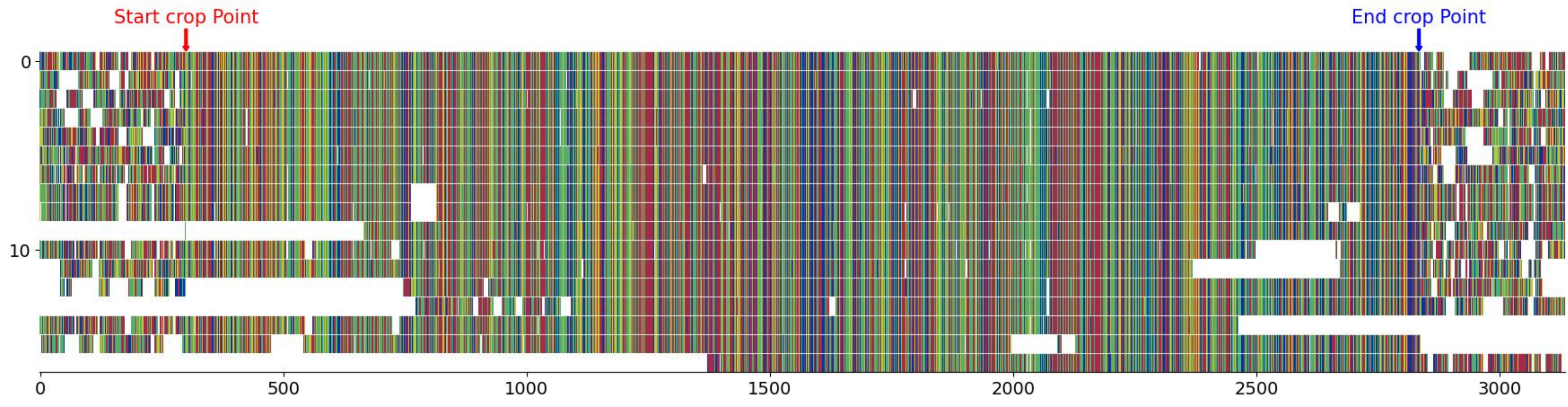
1000

1500

2000

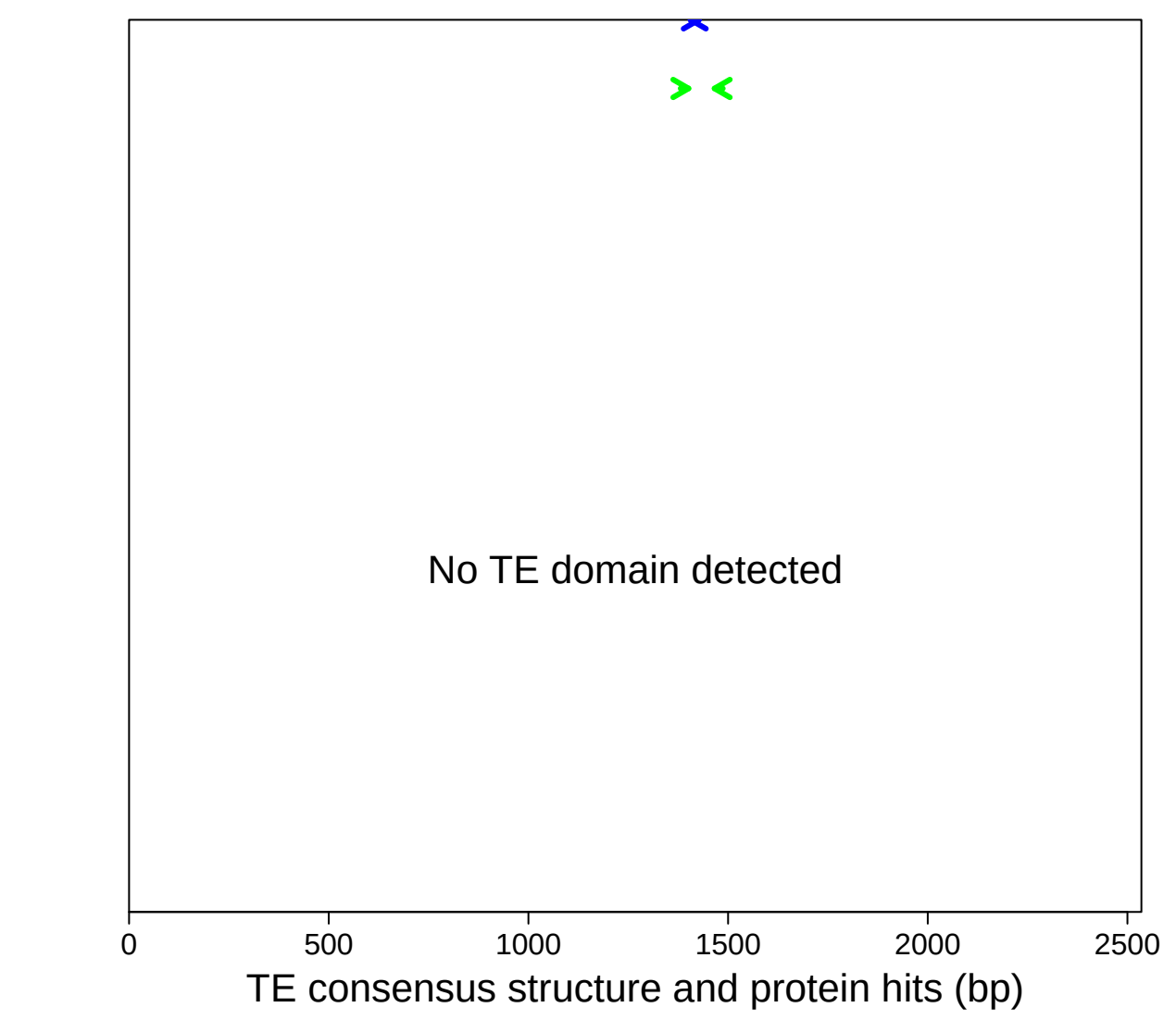
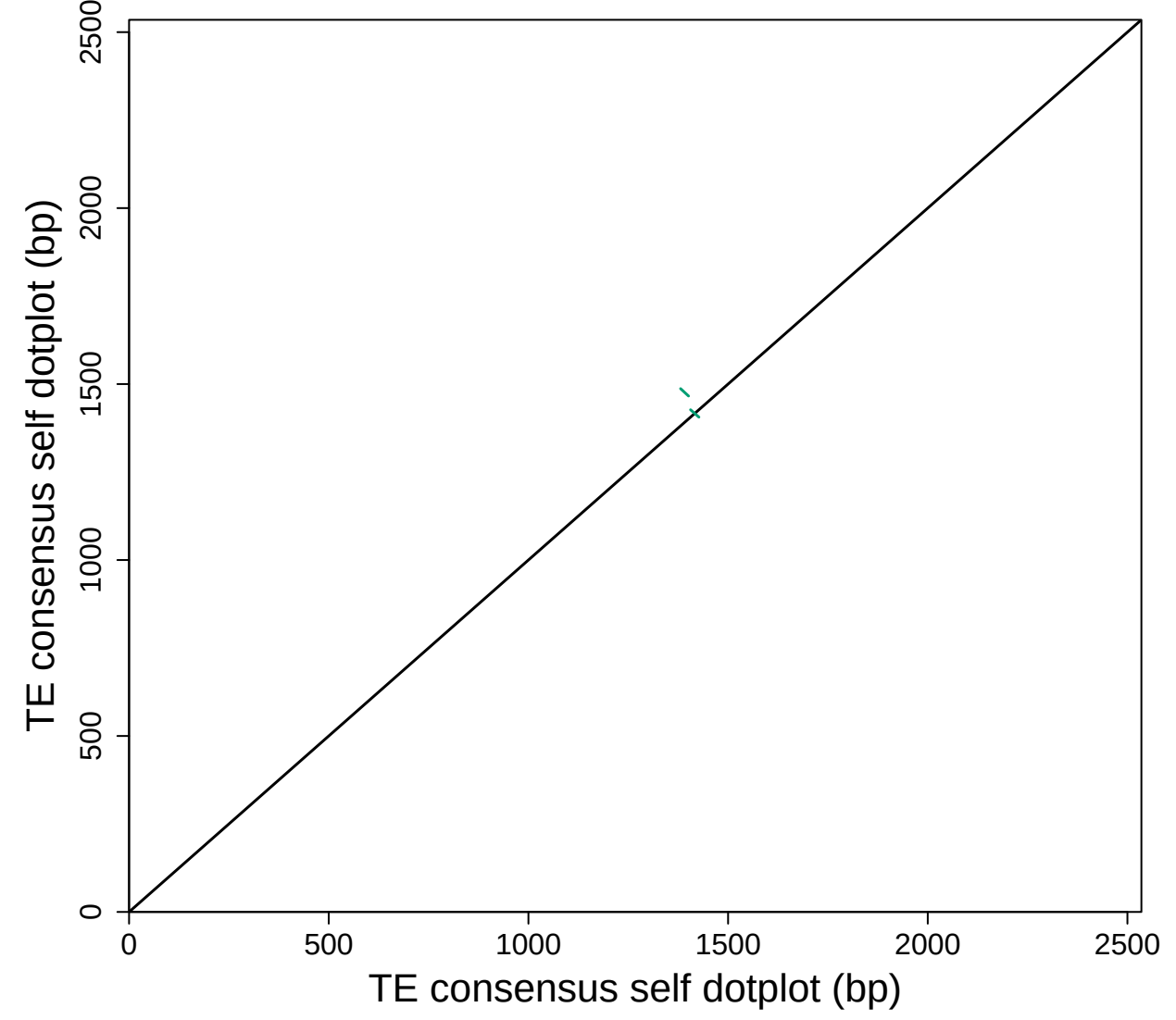
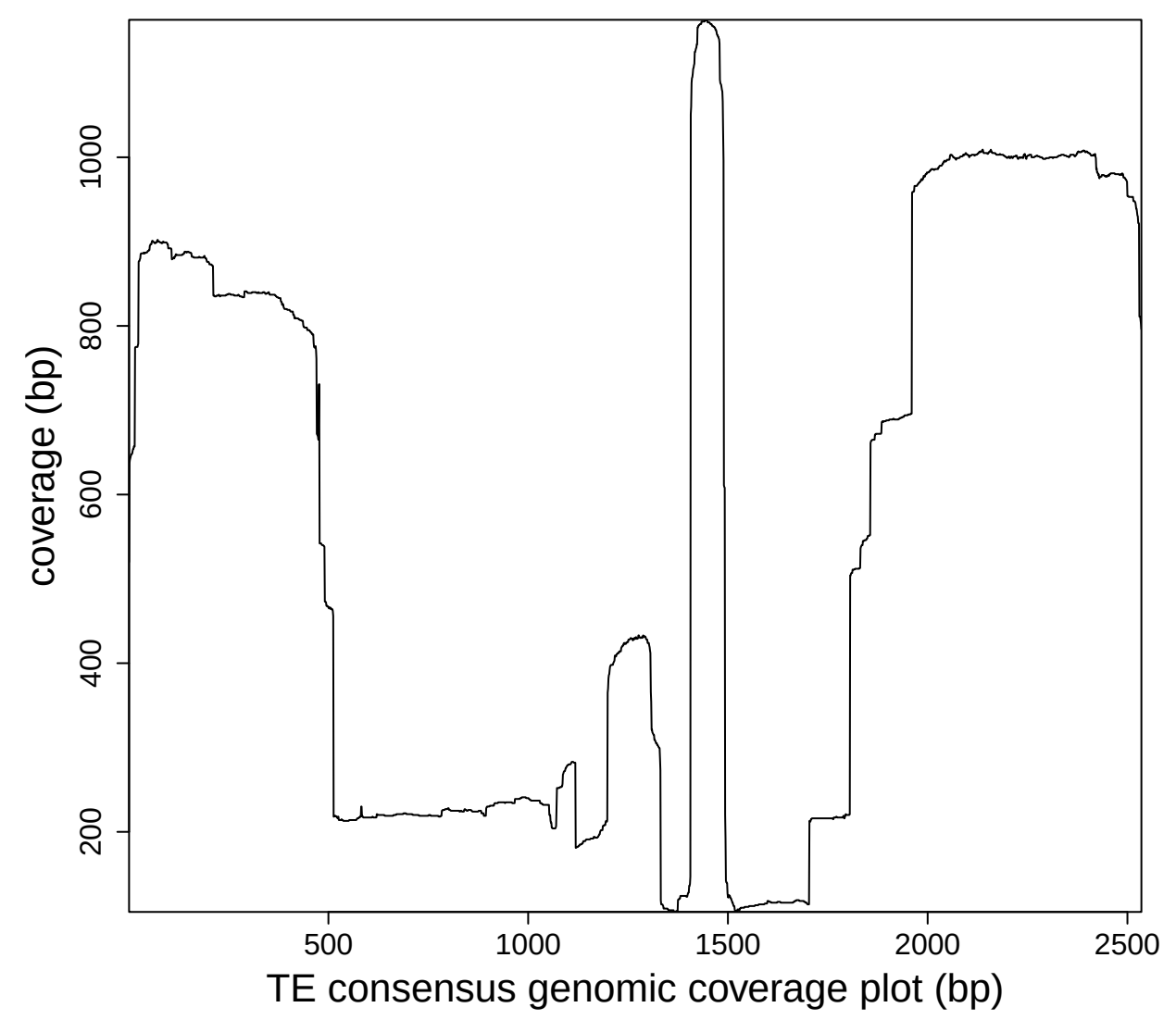
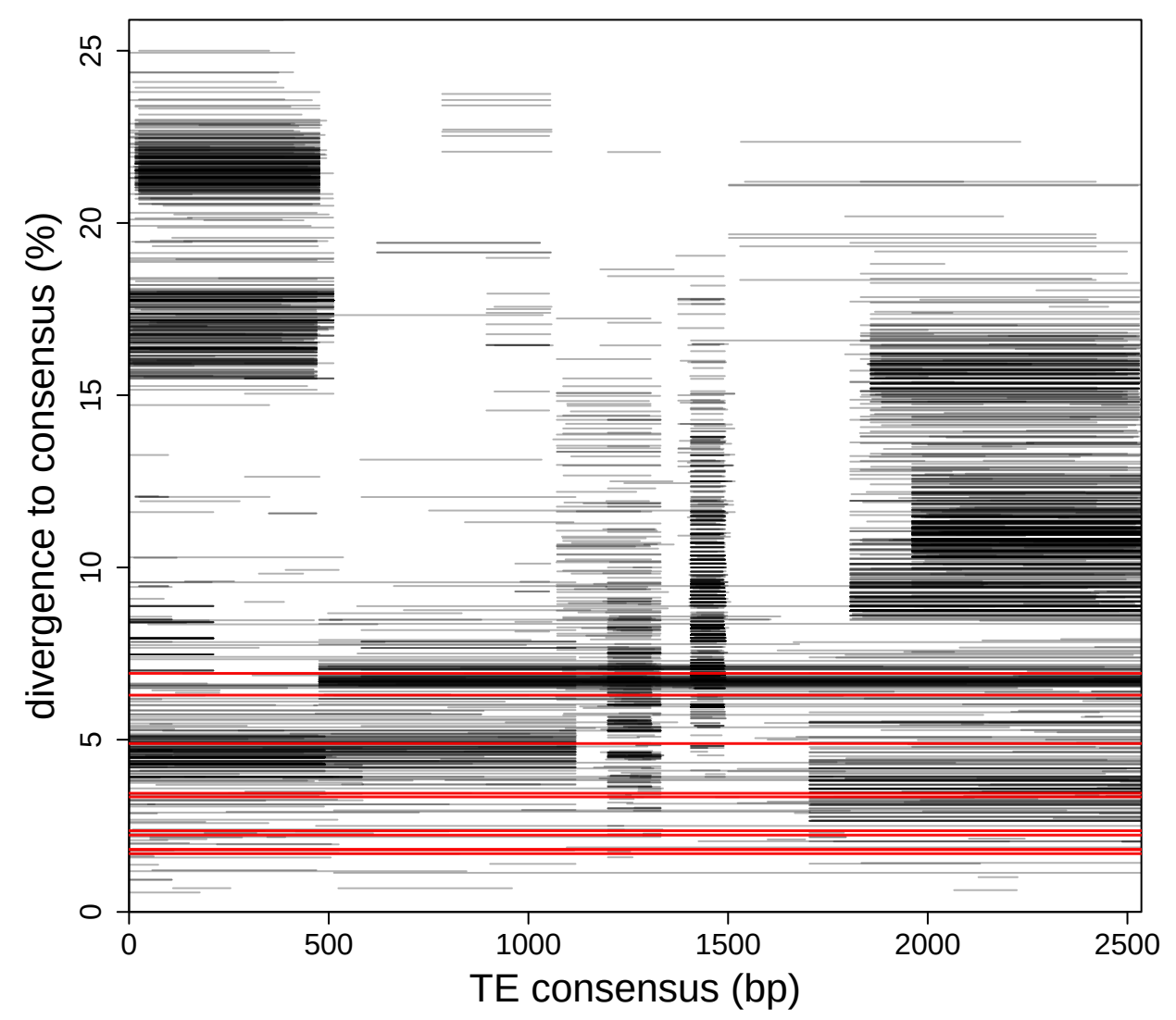
2500

3000



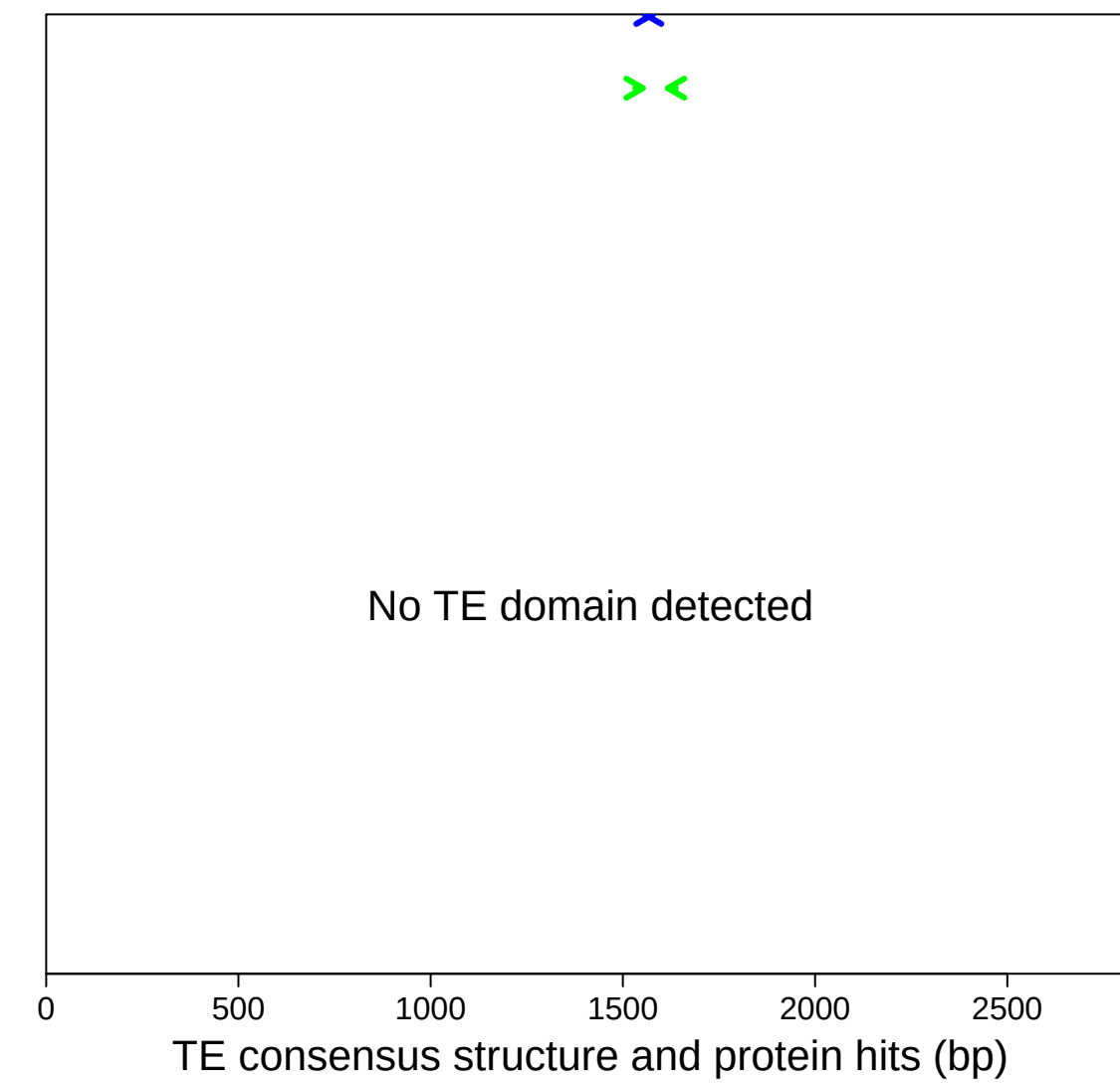
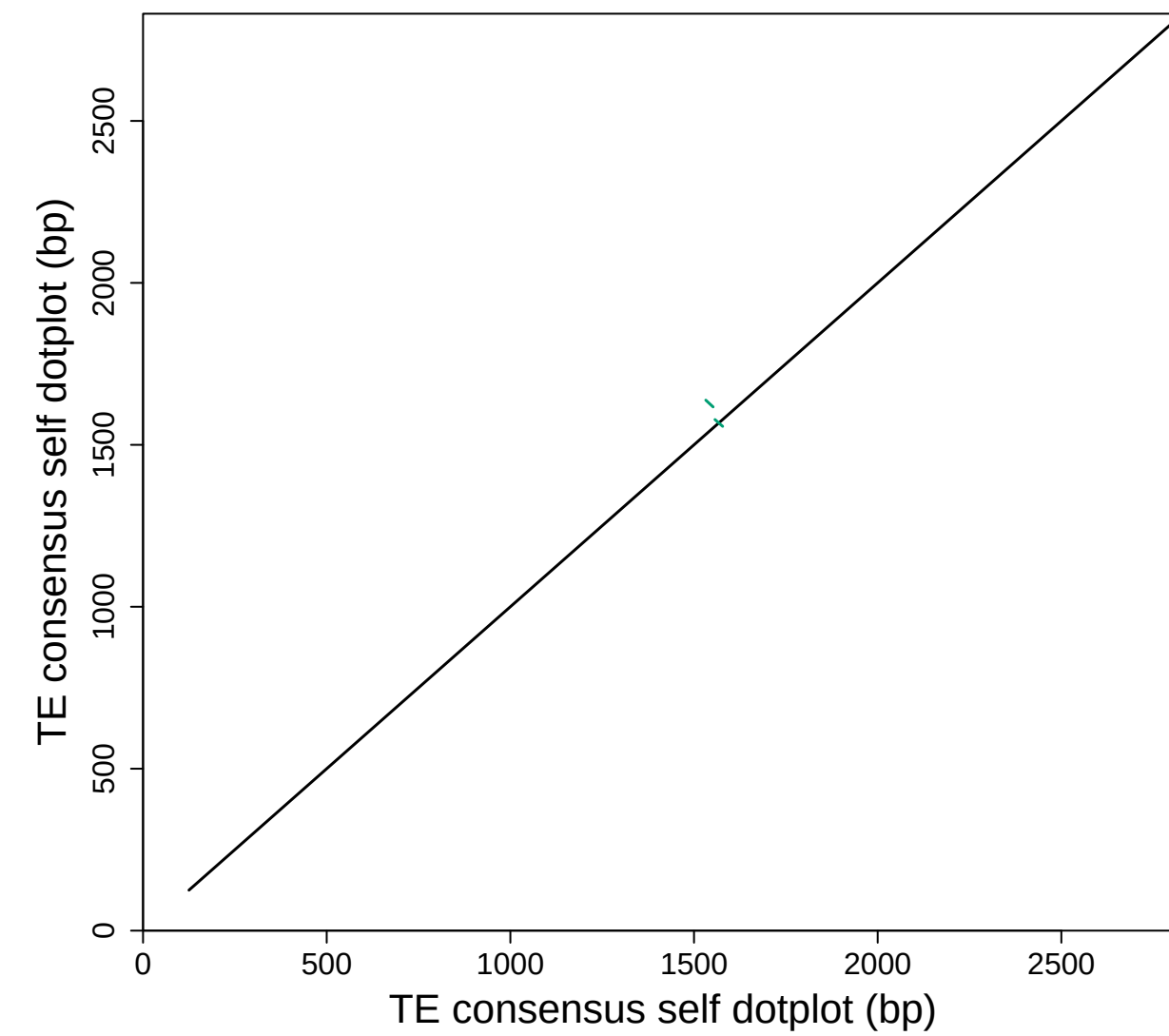
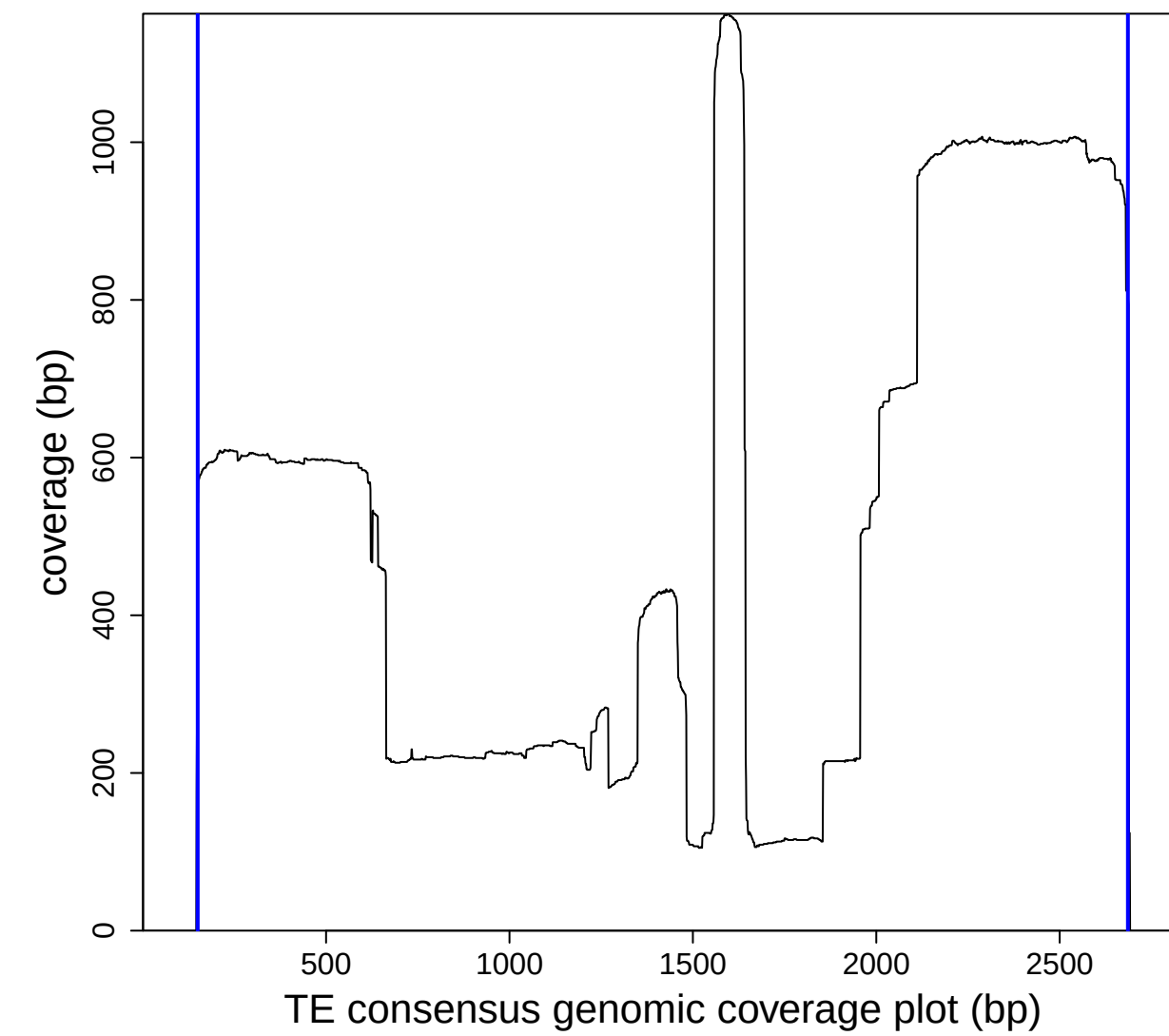
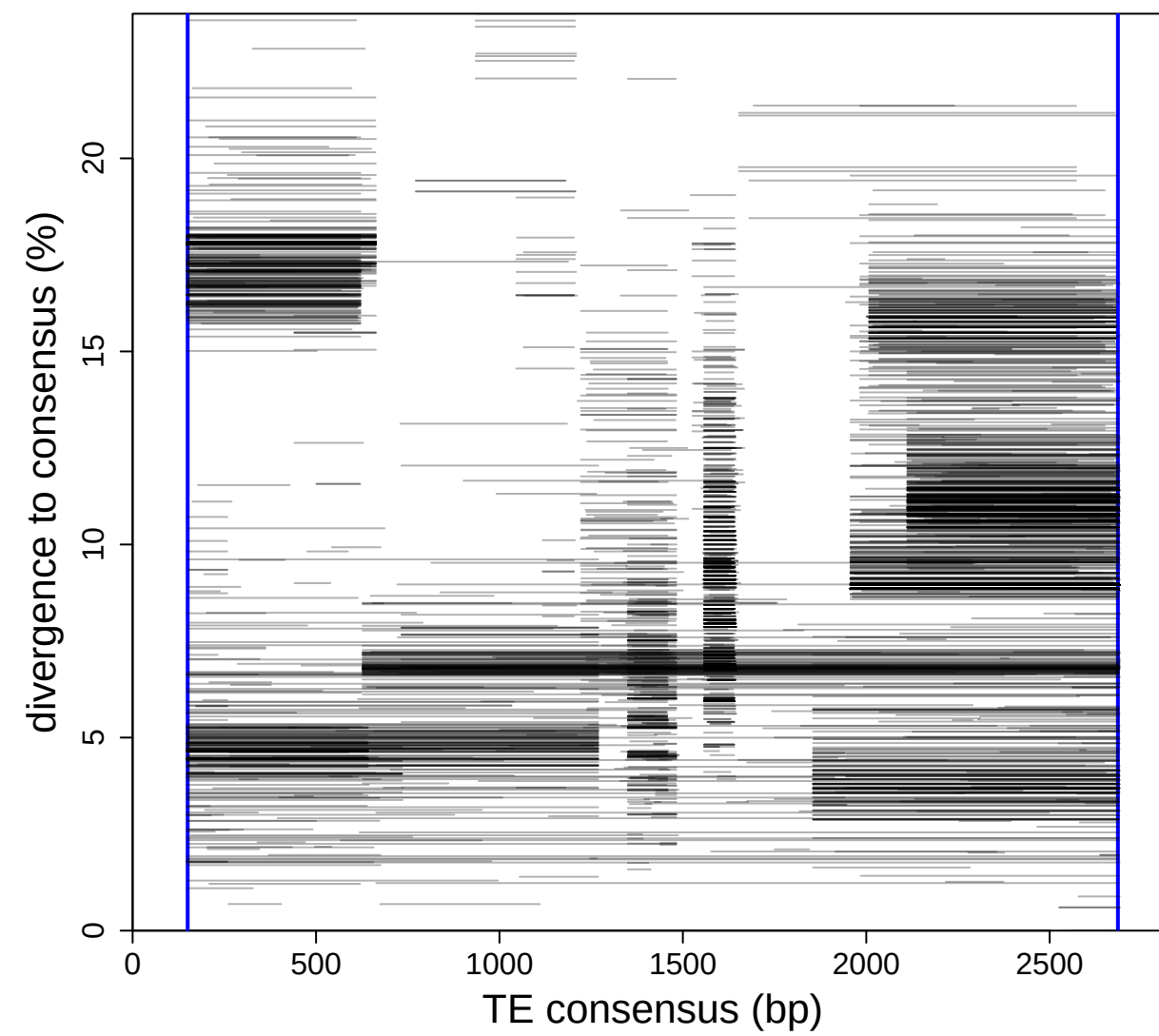
b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 2535bp; fragments: 3559; full length: 10 (≥ 2281.5 bp)

After TEtrimmer 2535 bp



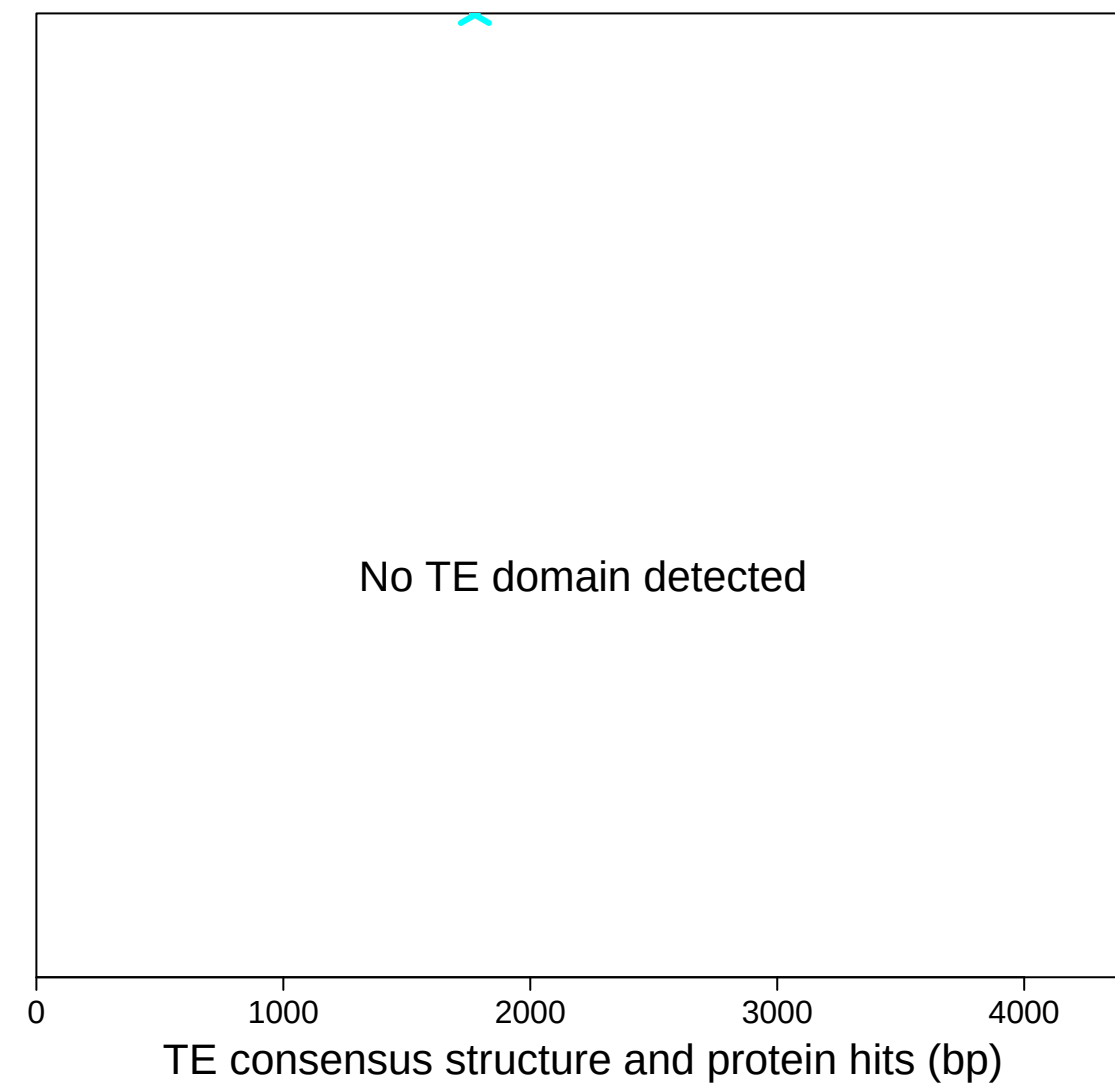
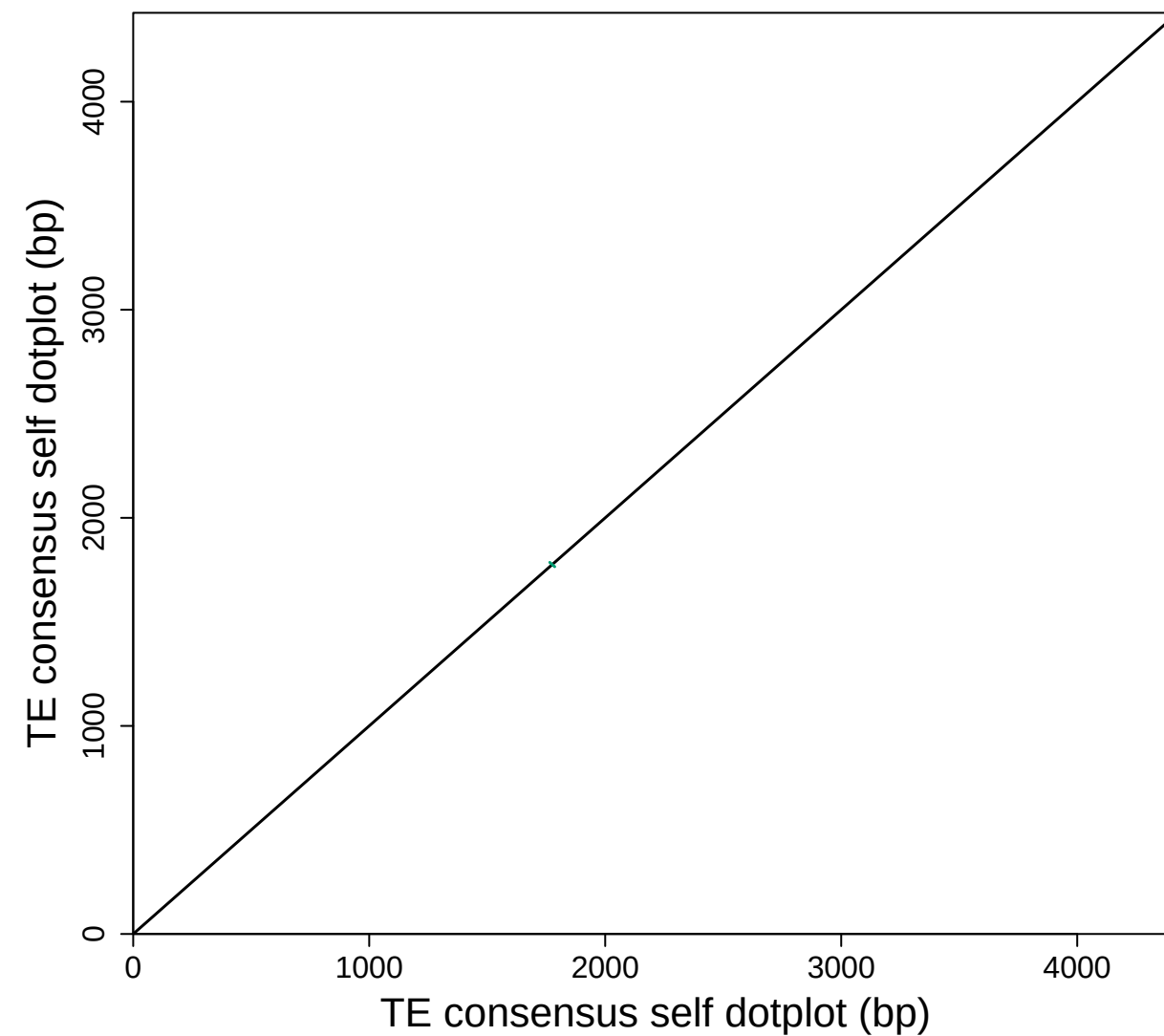
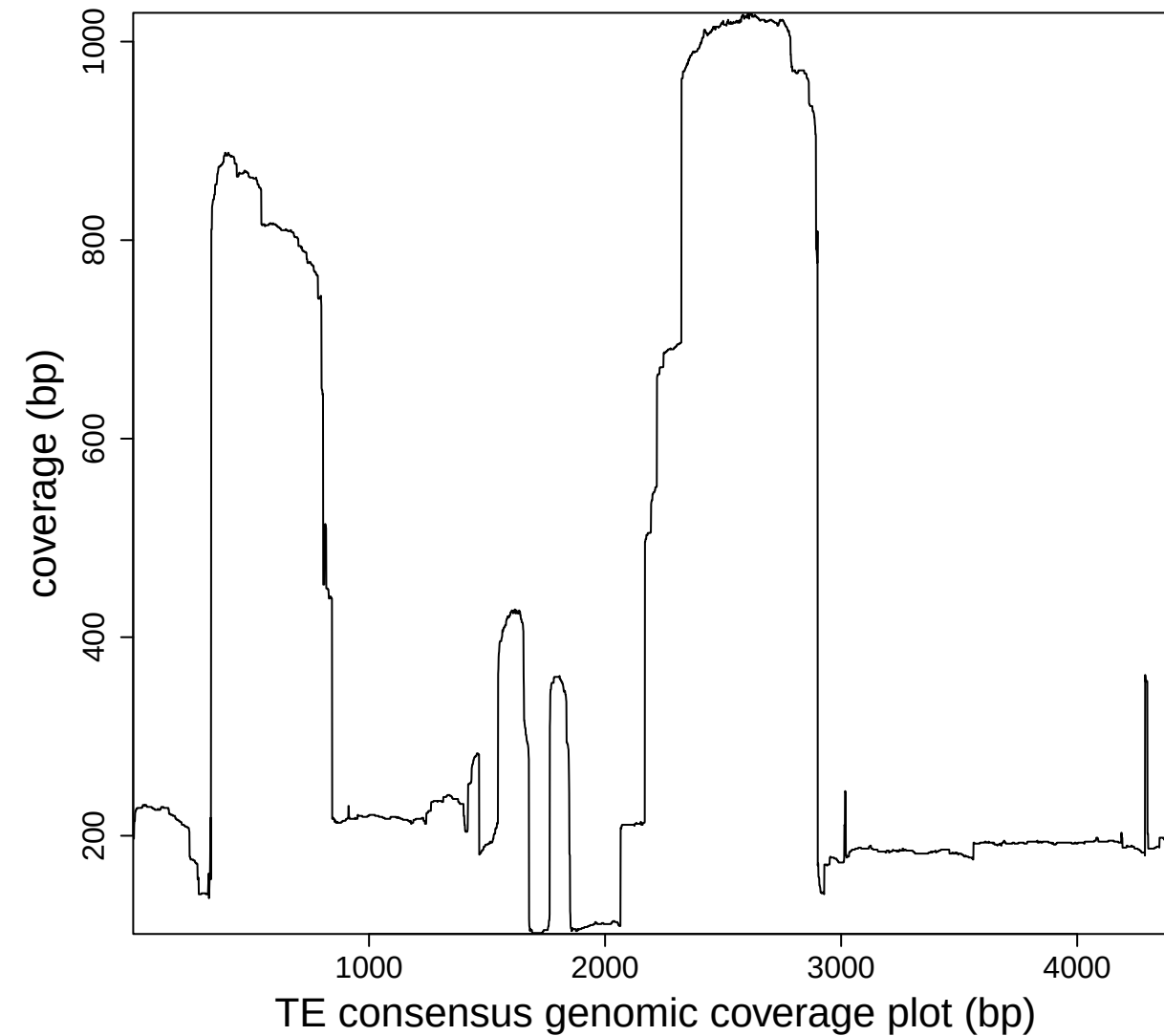
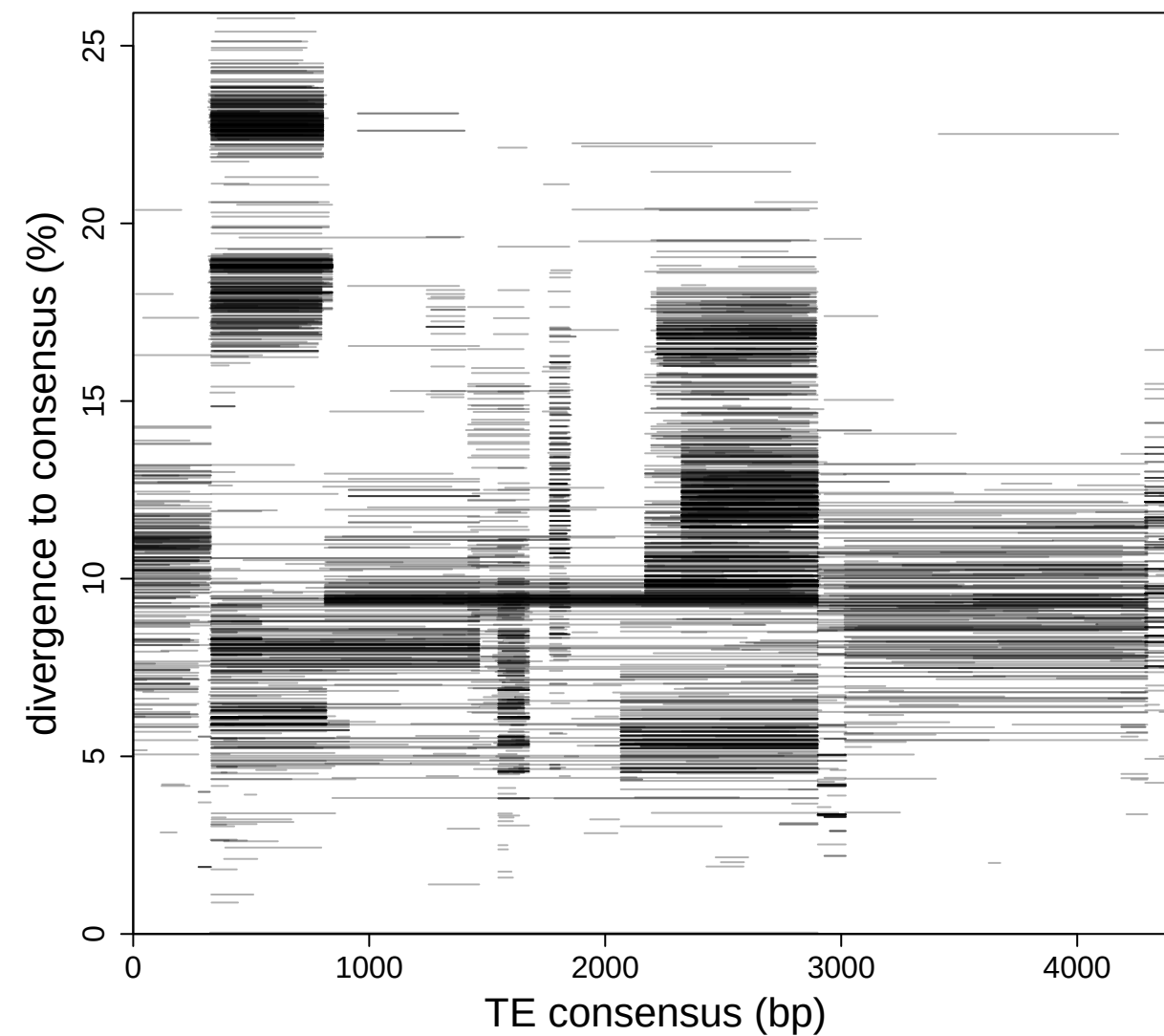
.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 2836bp; fragments: 3265; full length: 0 (>=2552.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

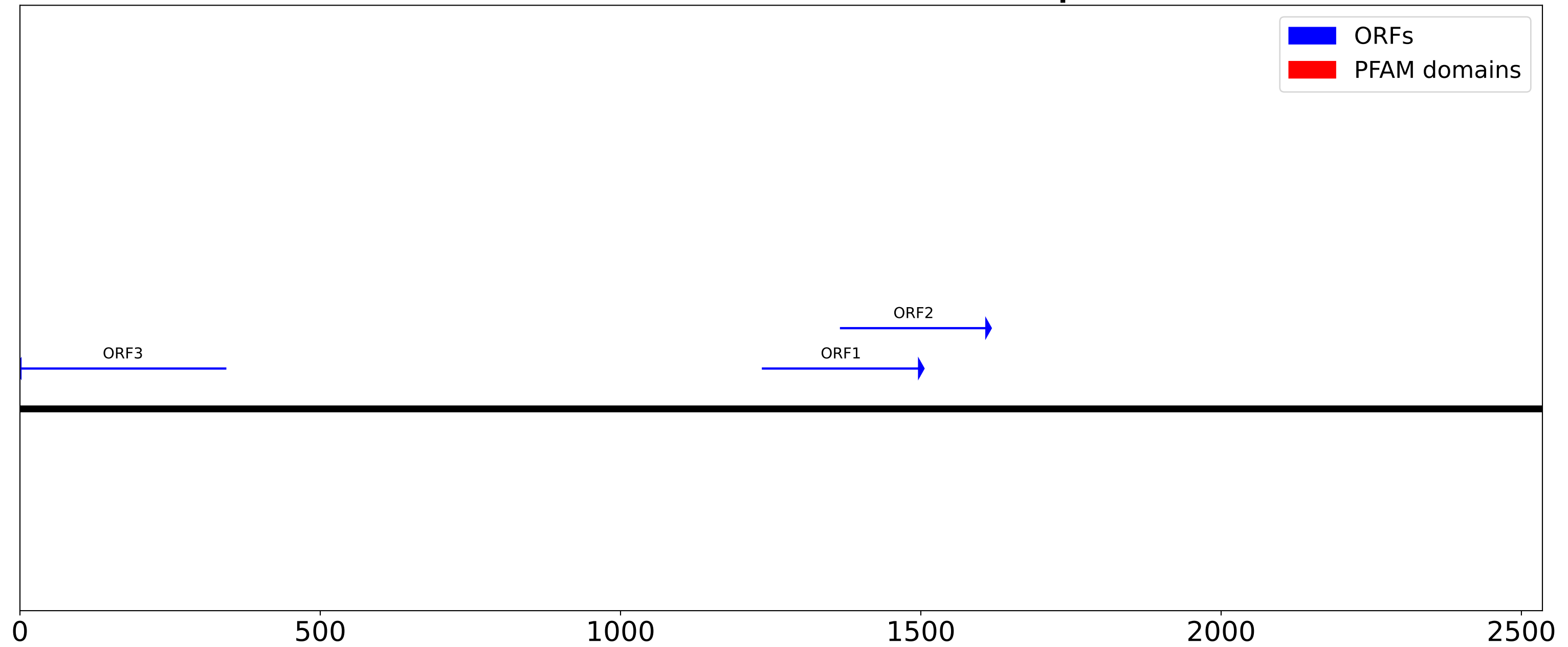


TE: ltr_1_family_987
size: 4427bp; fragments: 3511; full length: 0 (≥ 3984.3 bp)

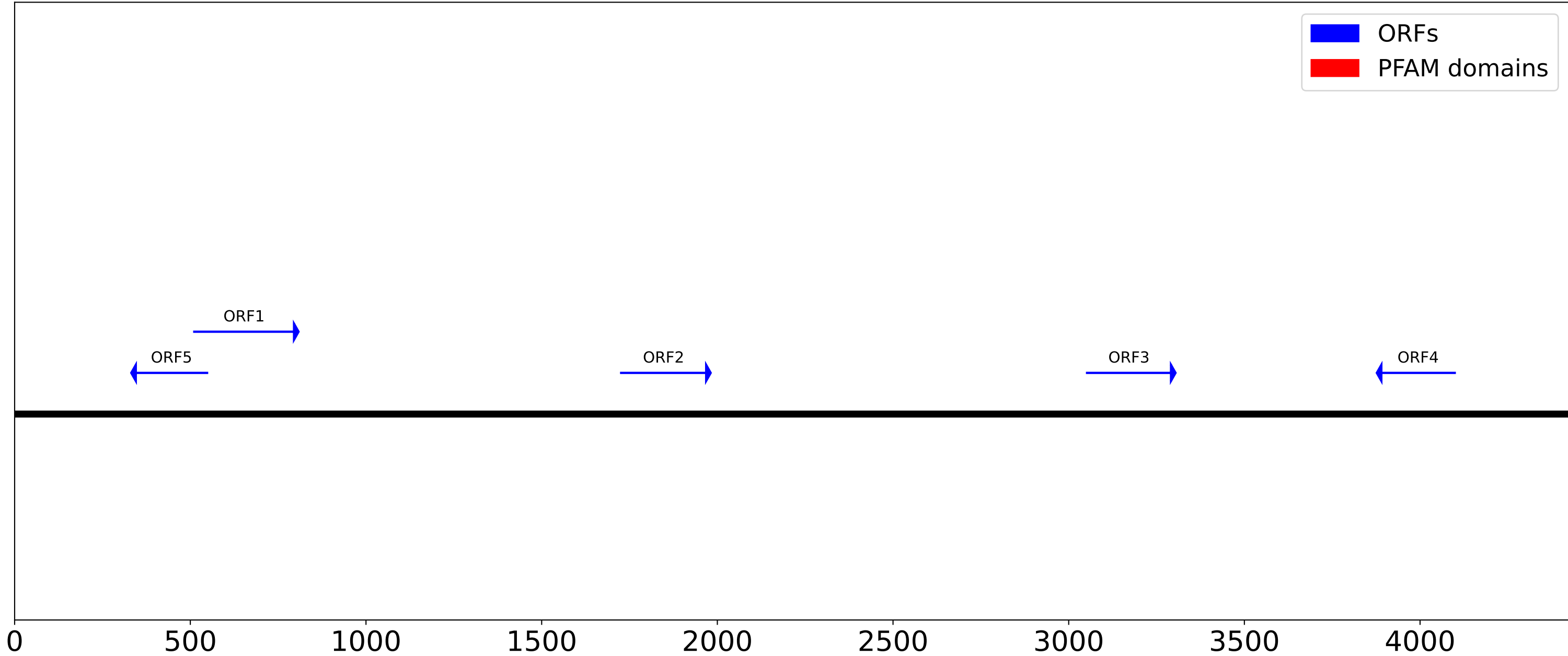
Before TEtrimmer 4427 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

