

Start crop Point

End crop Point

0

10

0

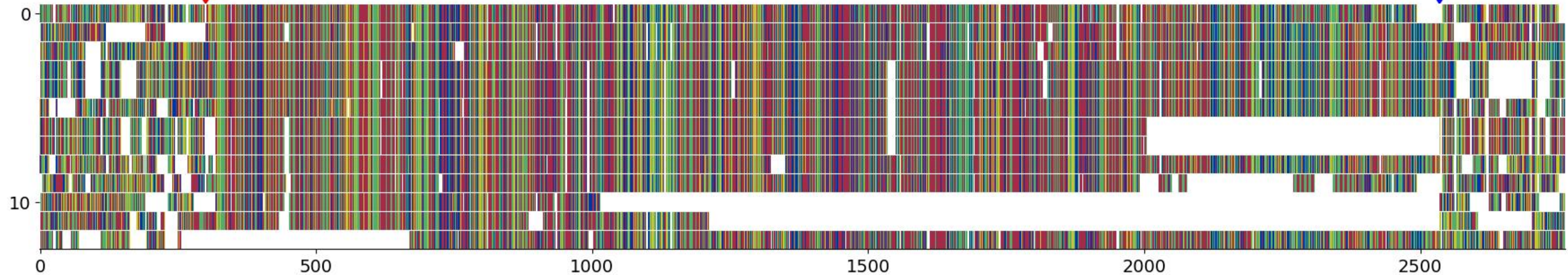
500

1000

1500

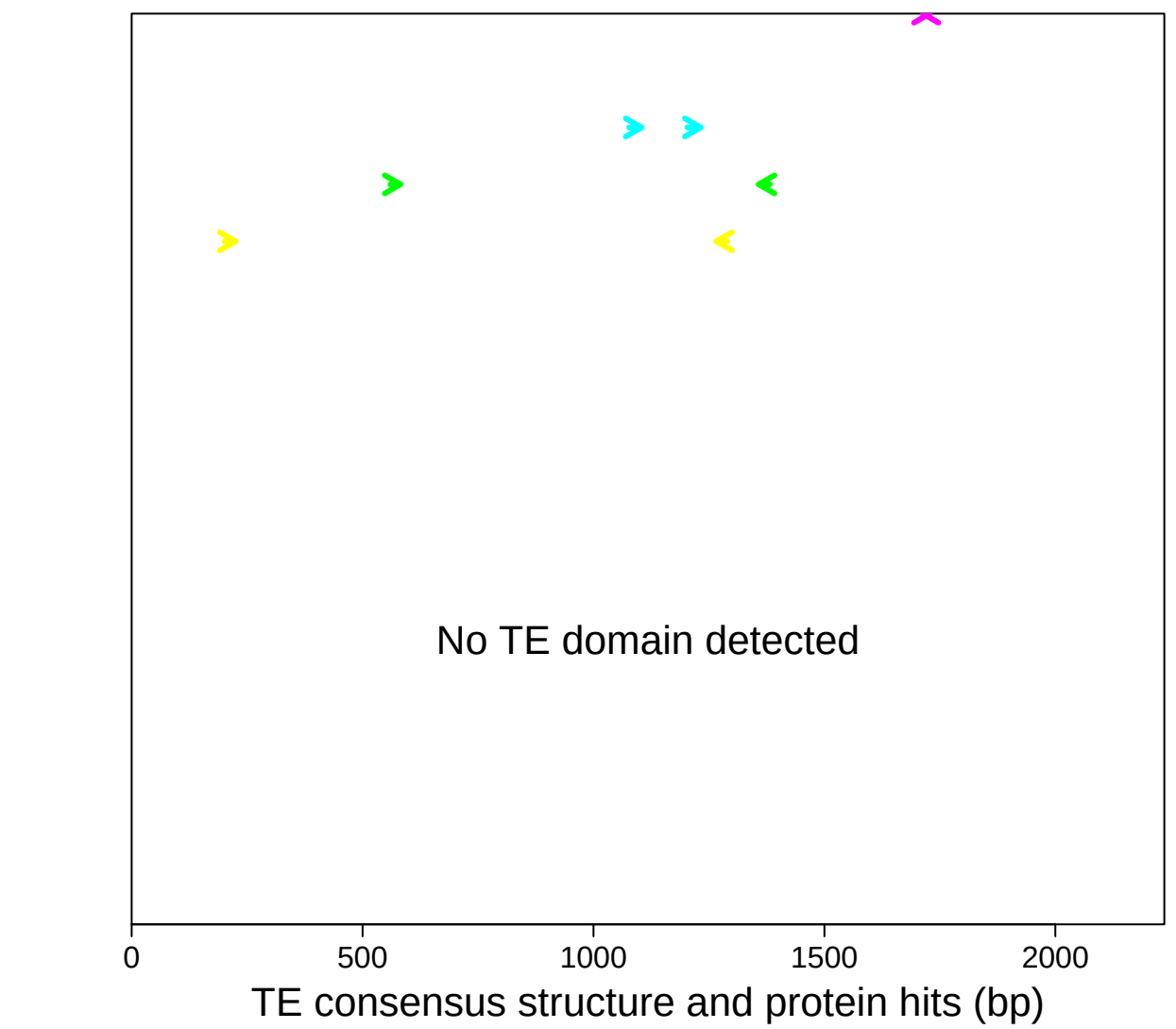
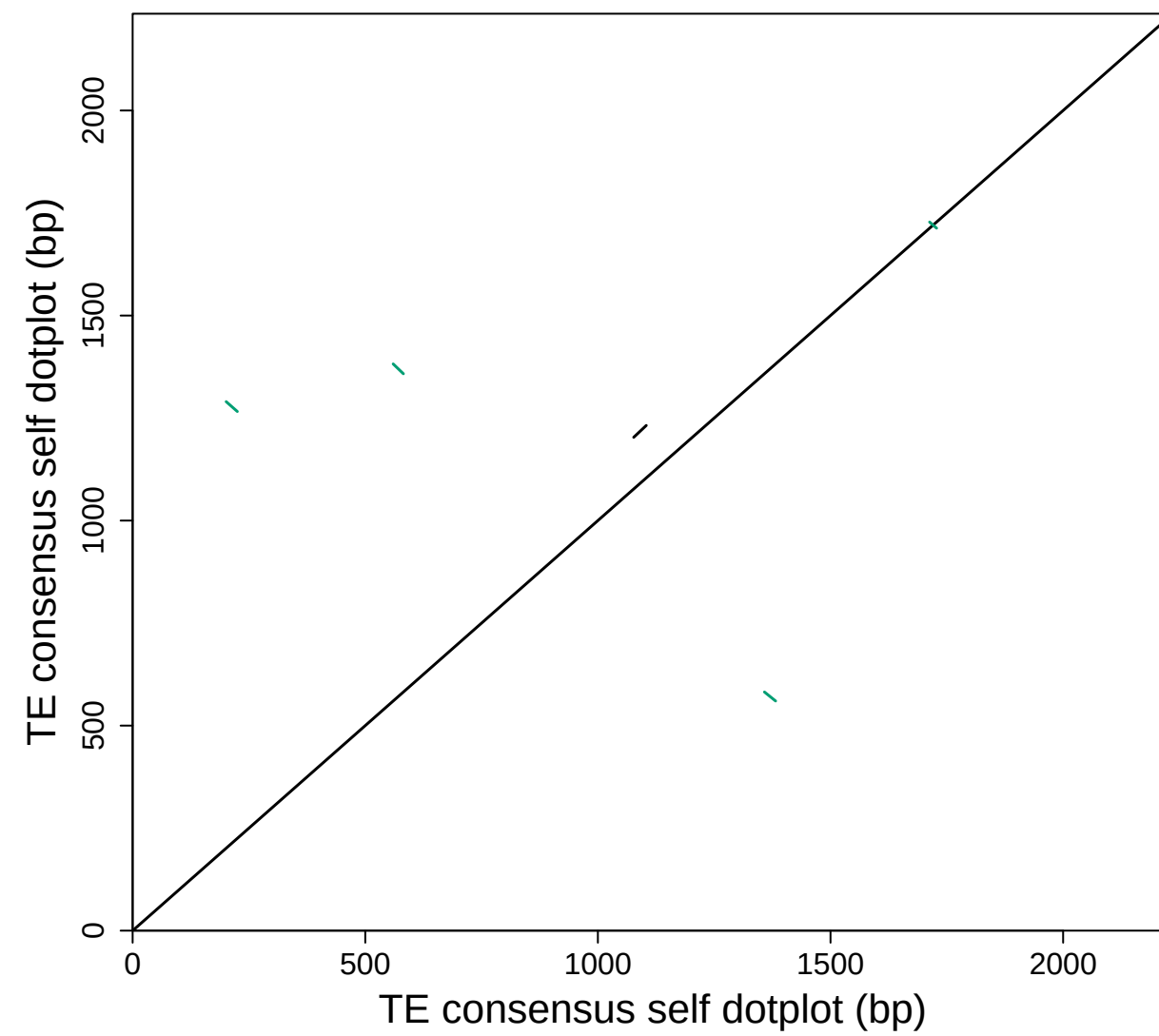
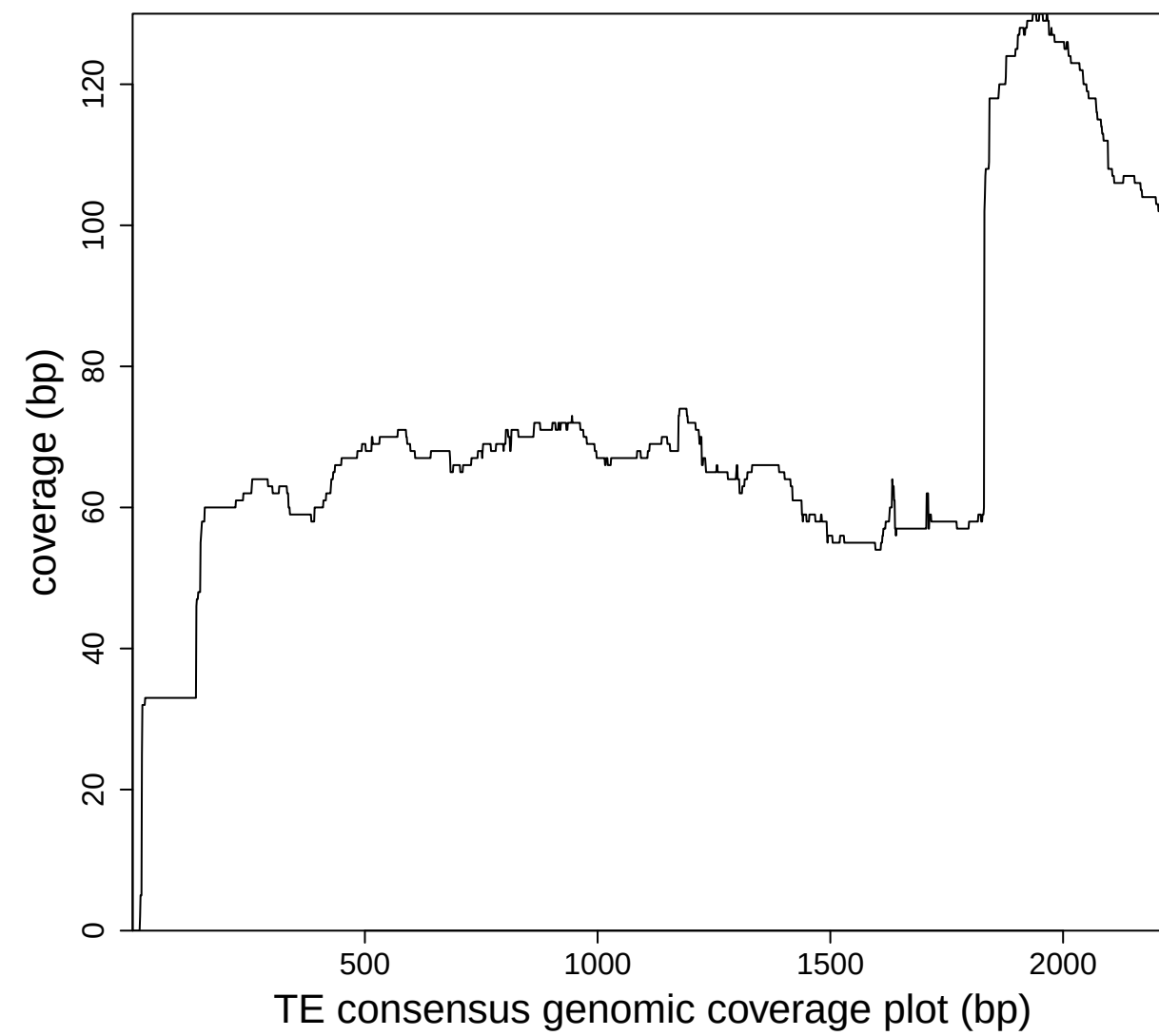
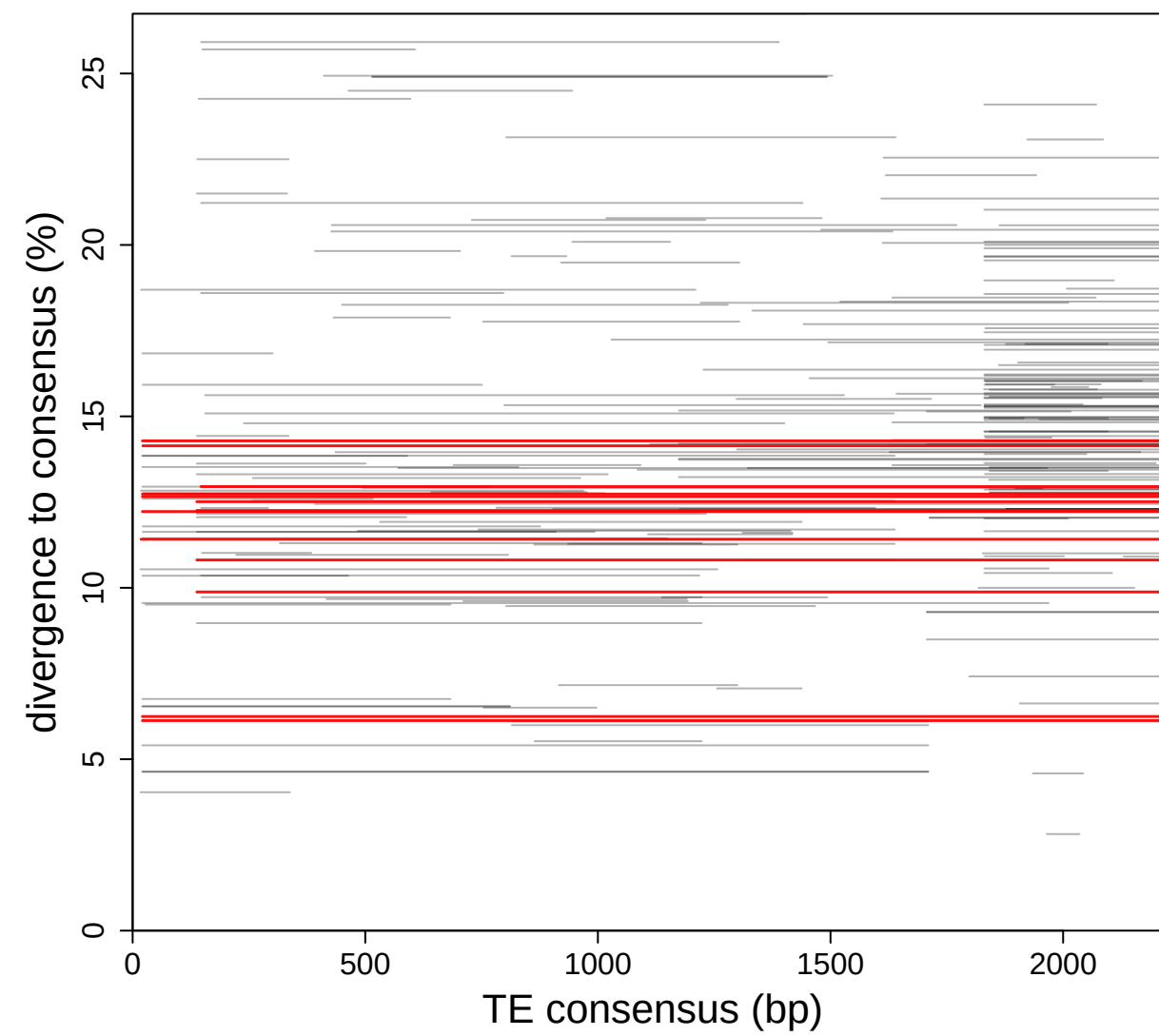
2000

2500



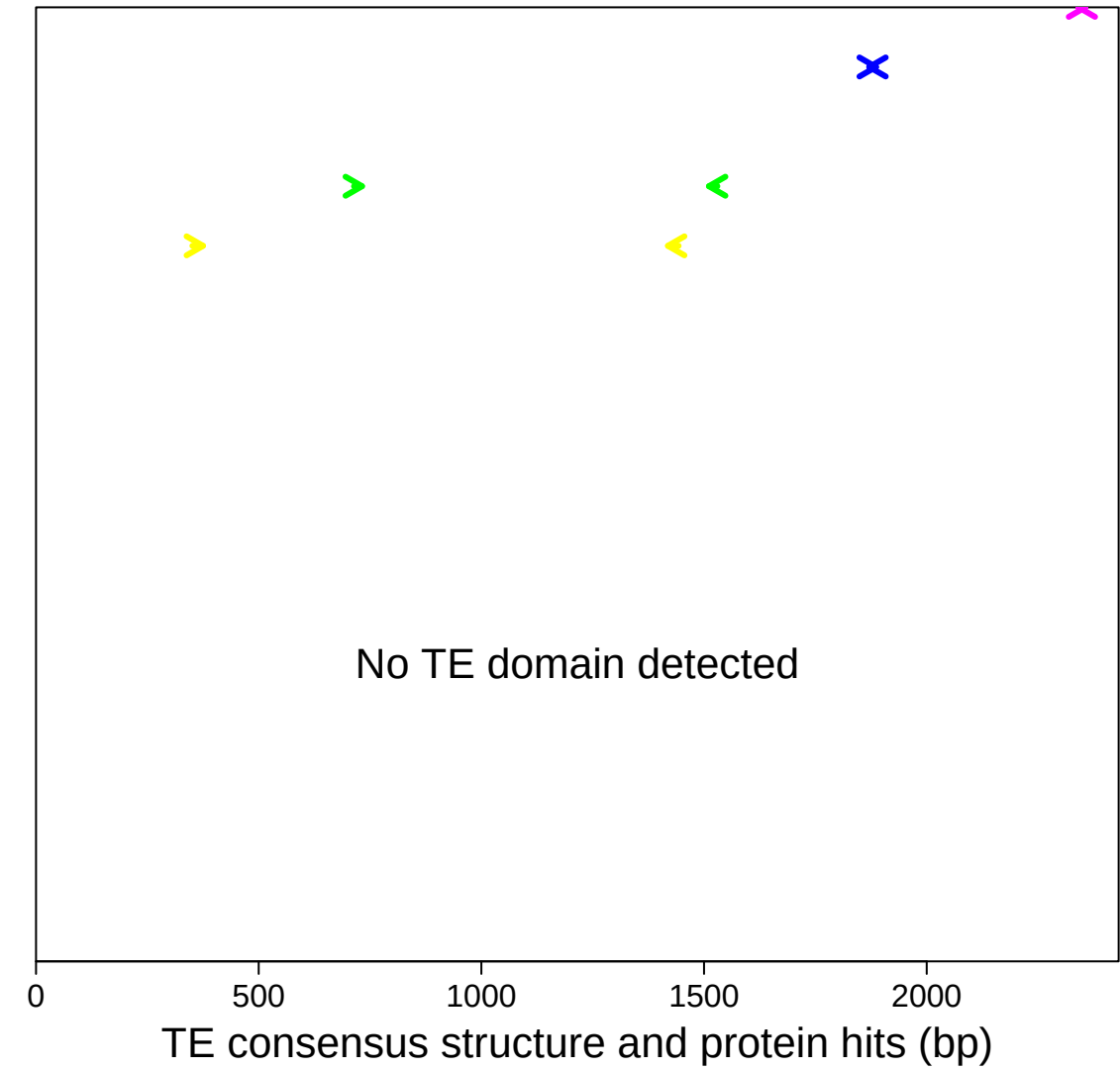
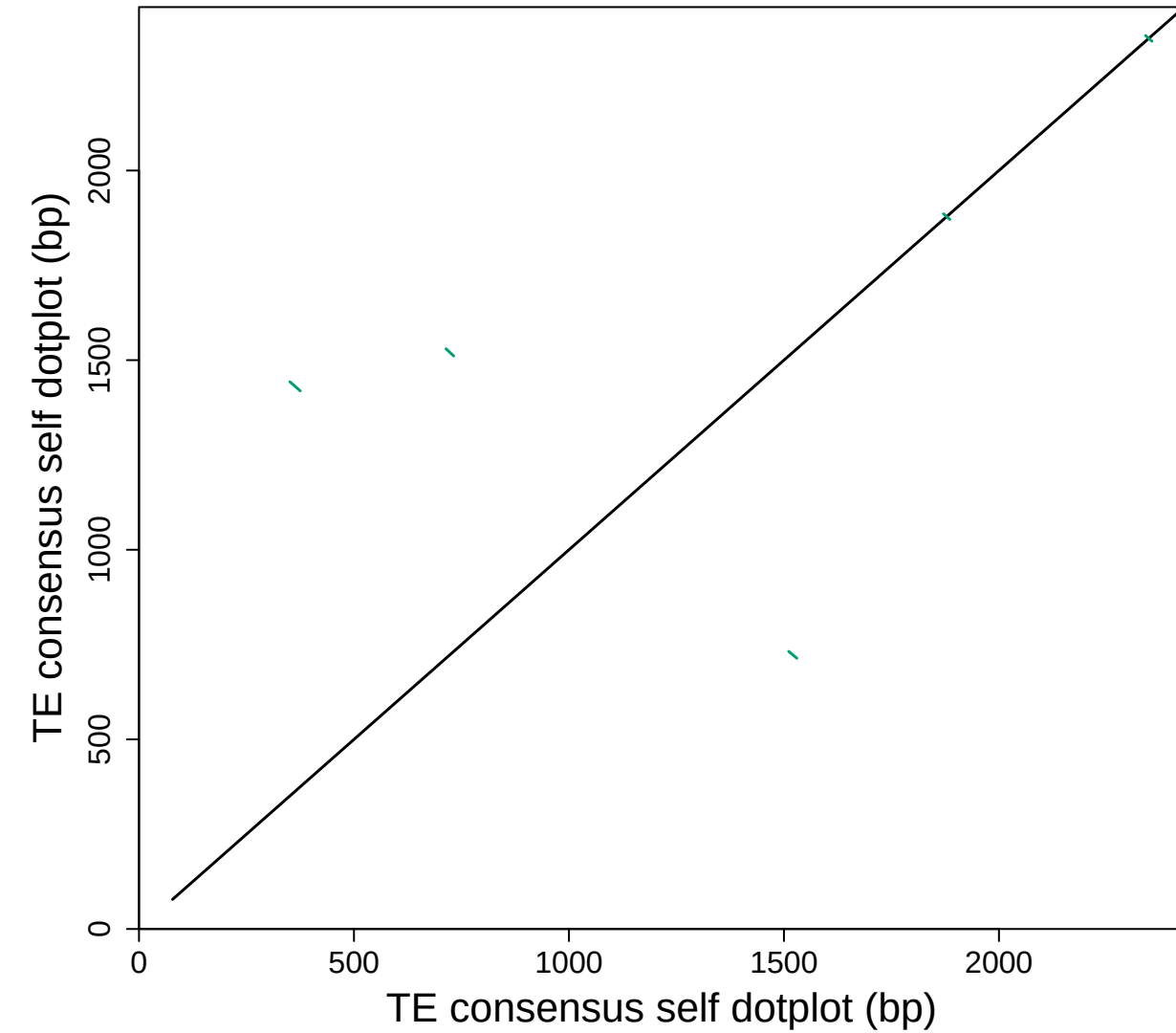
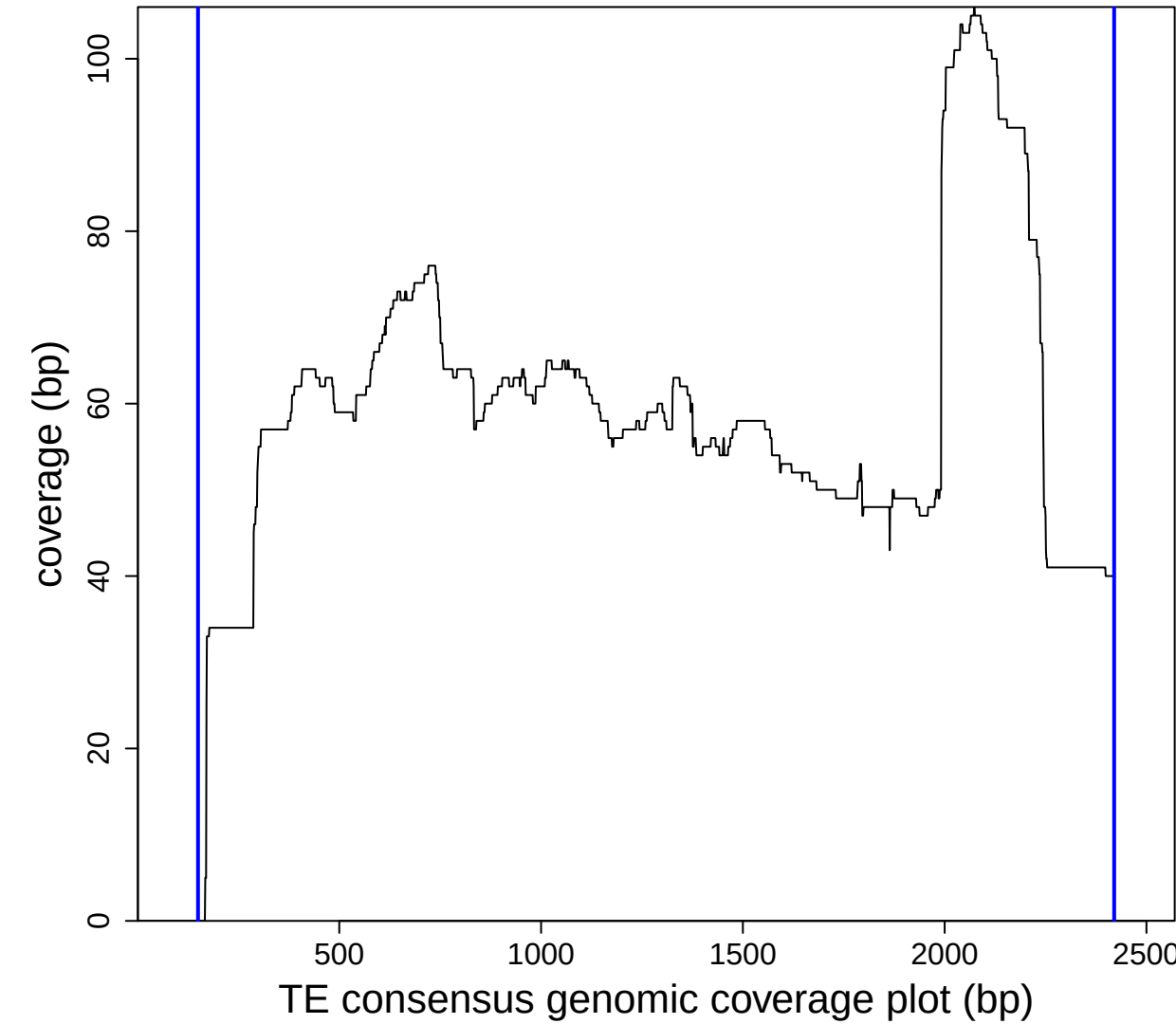
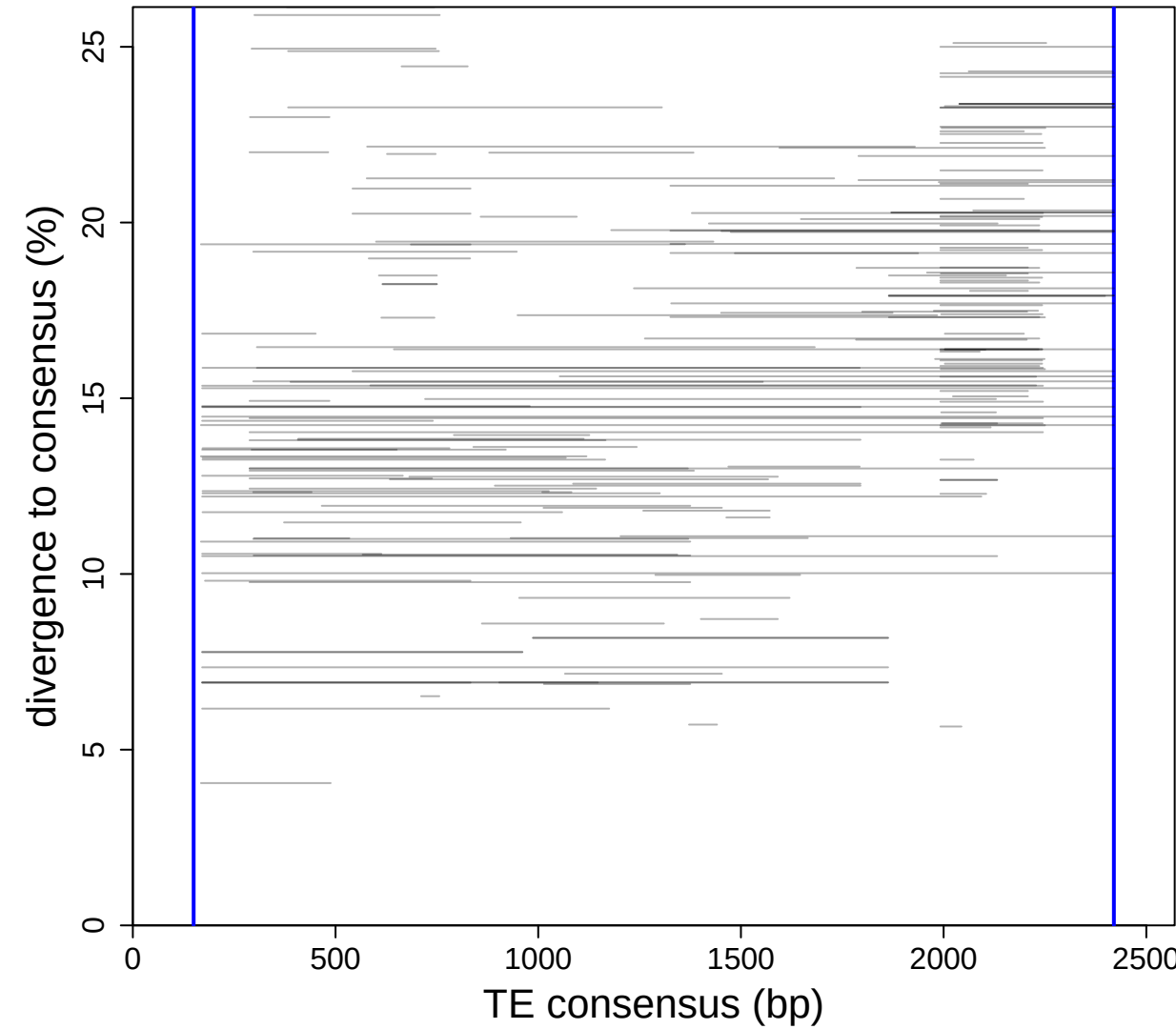
ta.b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.f
size: 2236bp; fragments: 226; full length: 13 (>=2012.4bp)

After TEtrimmer 2236 bp



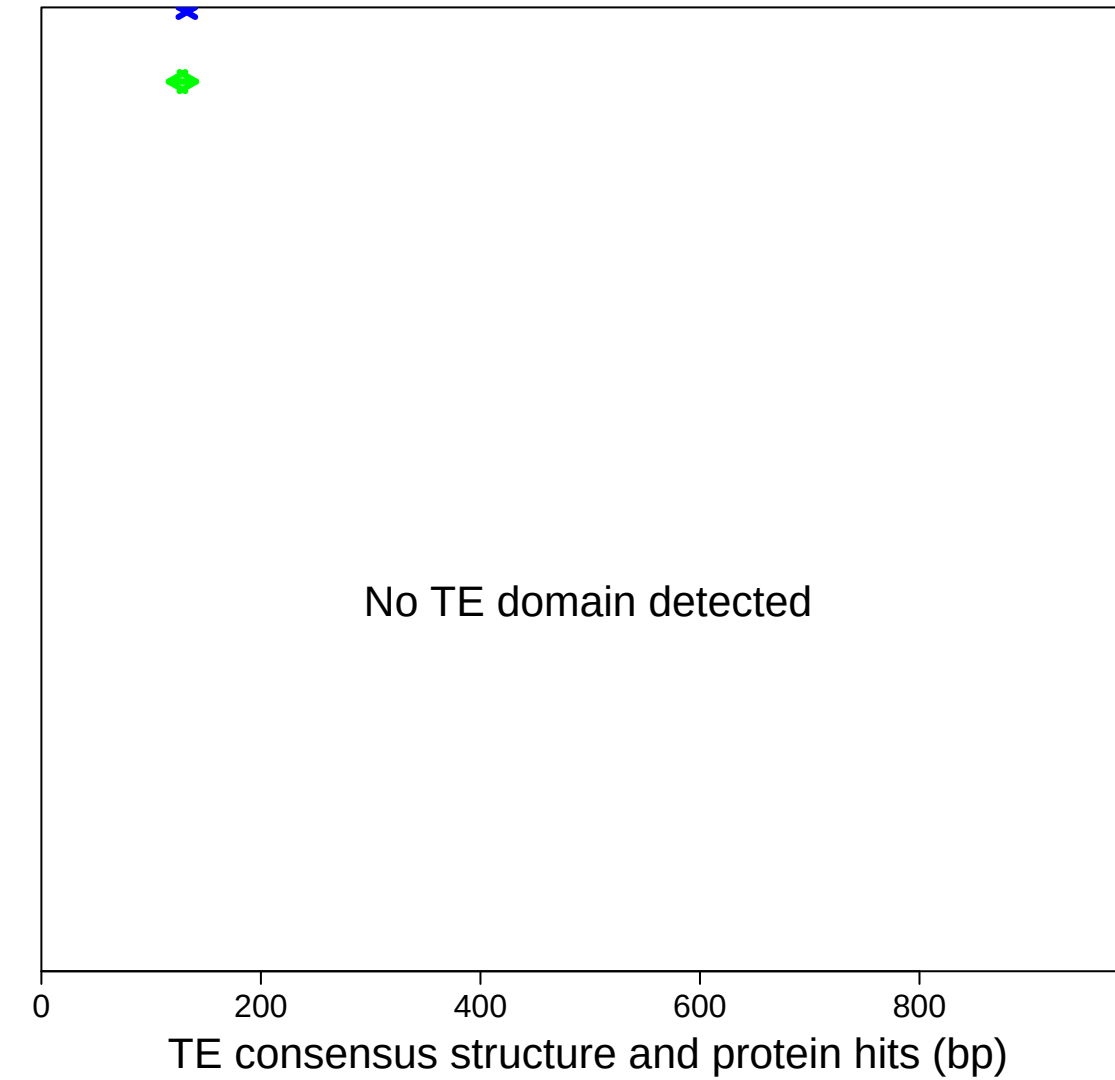
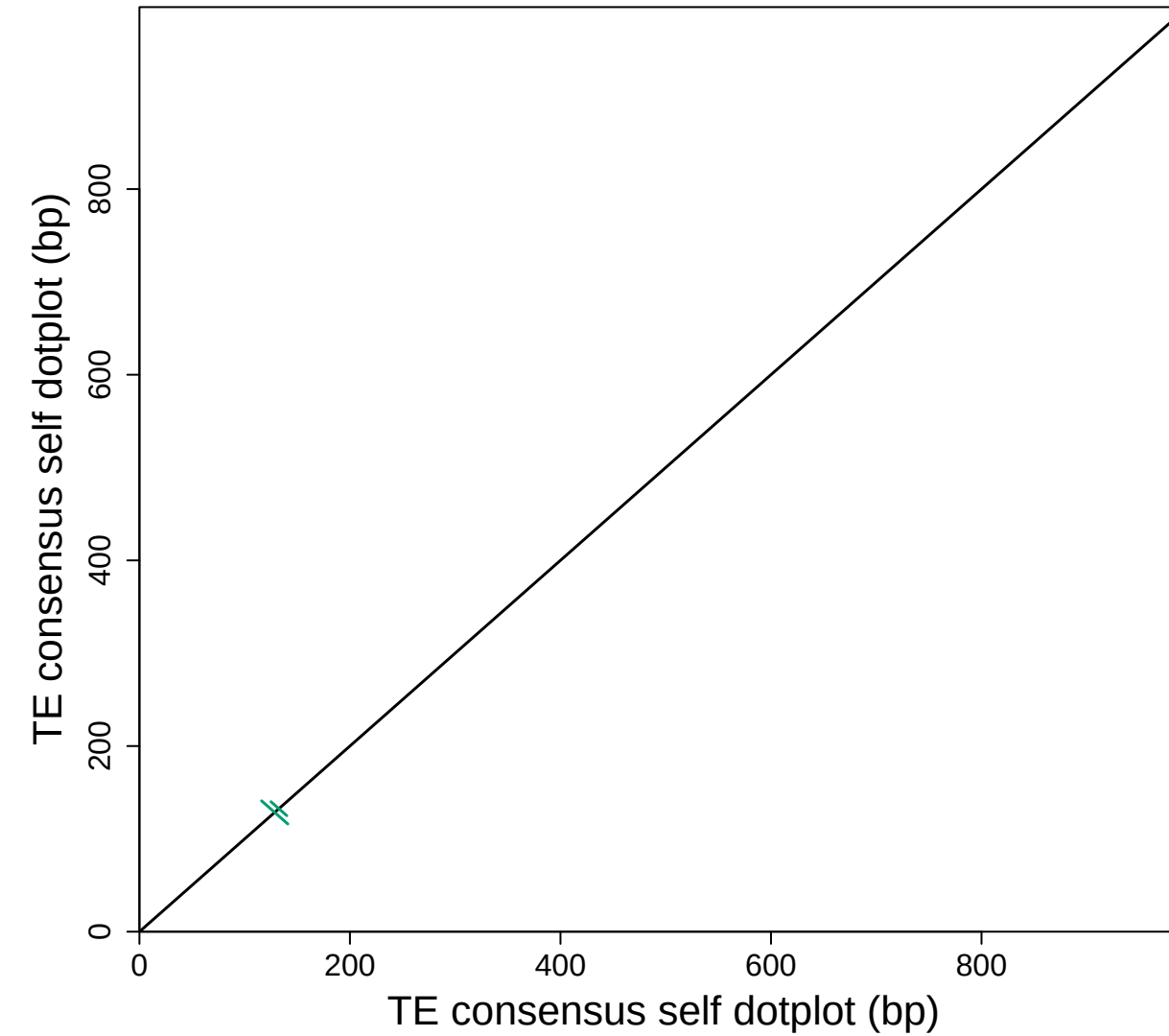
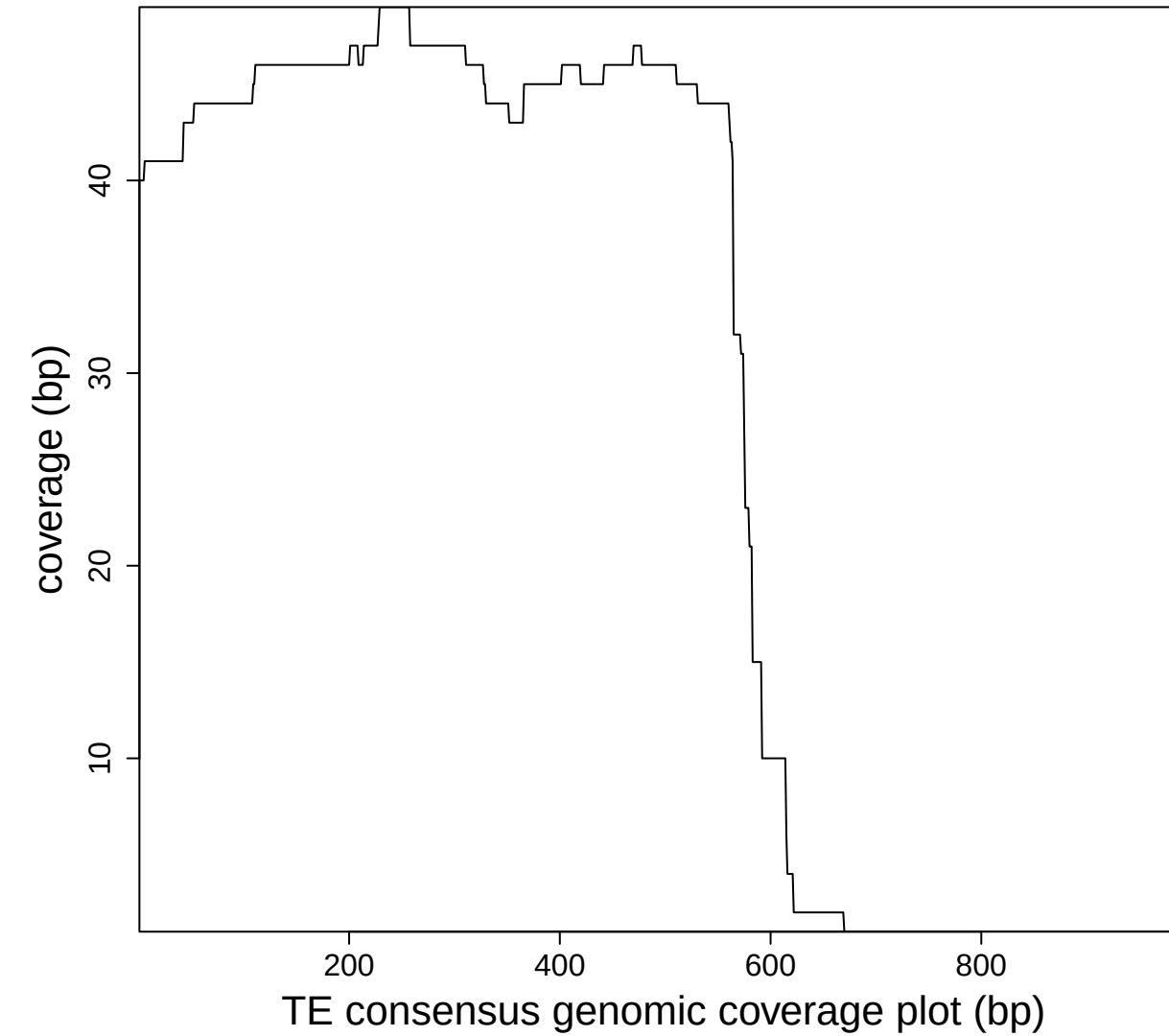
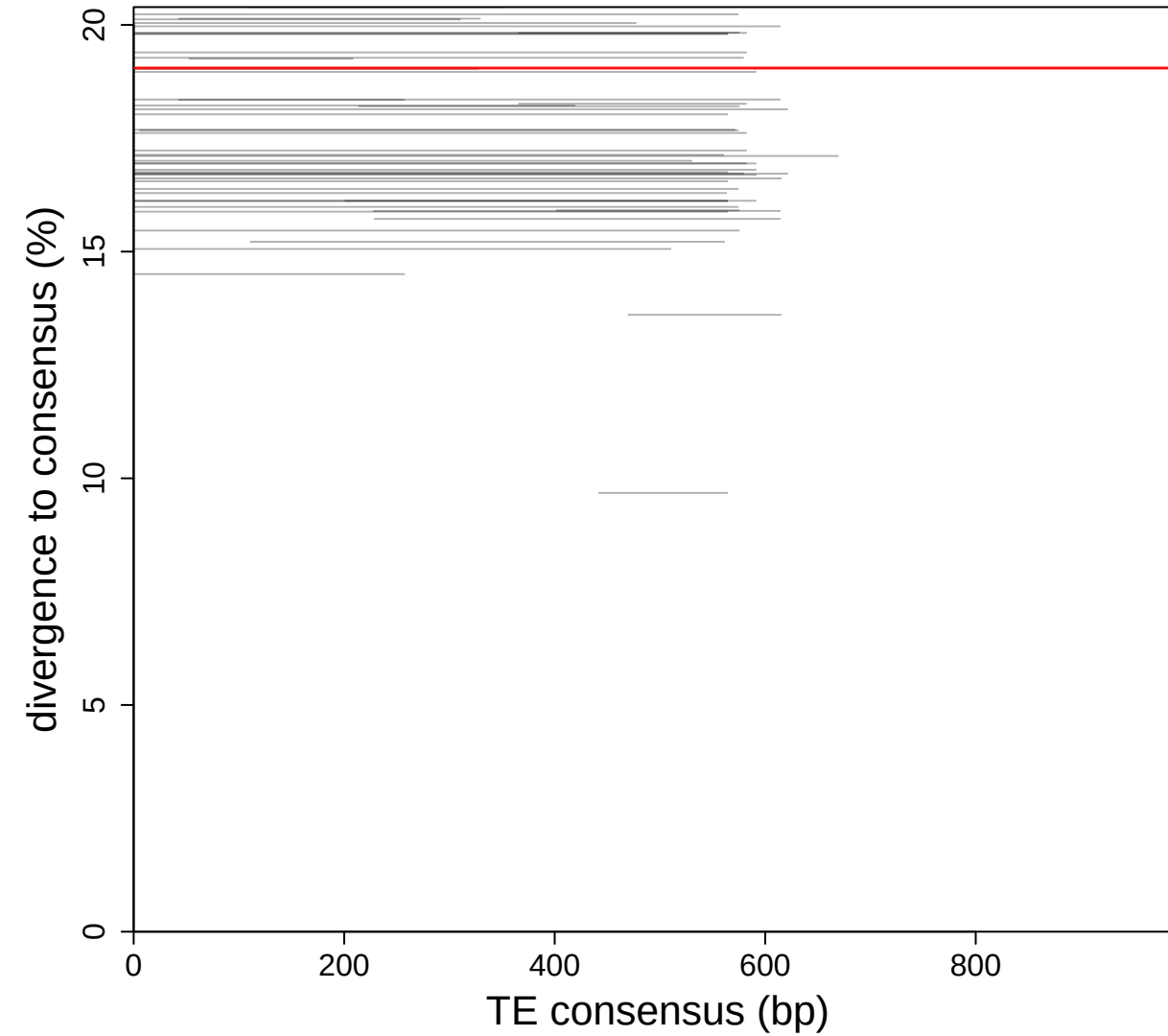
26.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.1
size: 2570bp; fragments: 201; full length: 0 (>=2313bp)

After TEtrimmer Extended plot Blue lines are boundaries

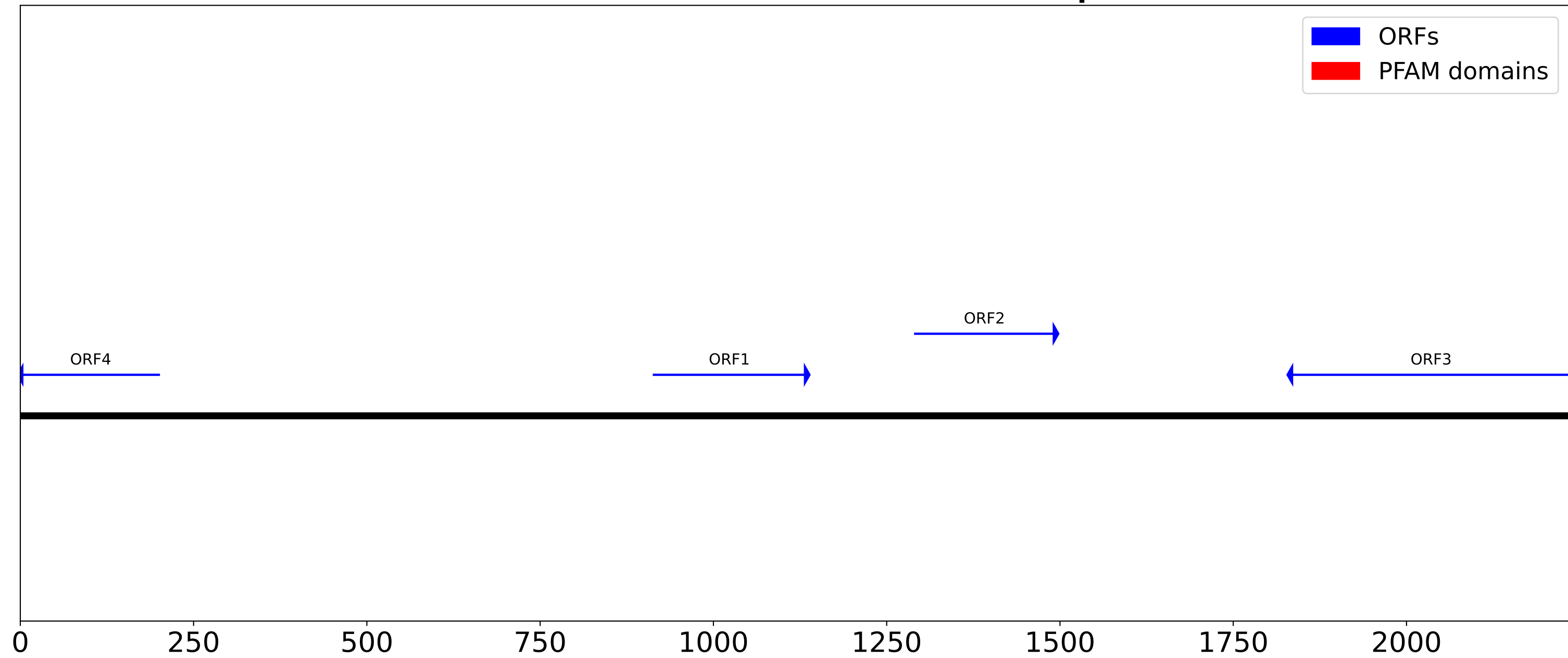


Before TEtrimmer 996 bp

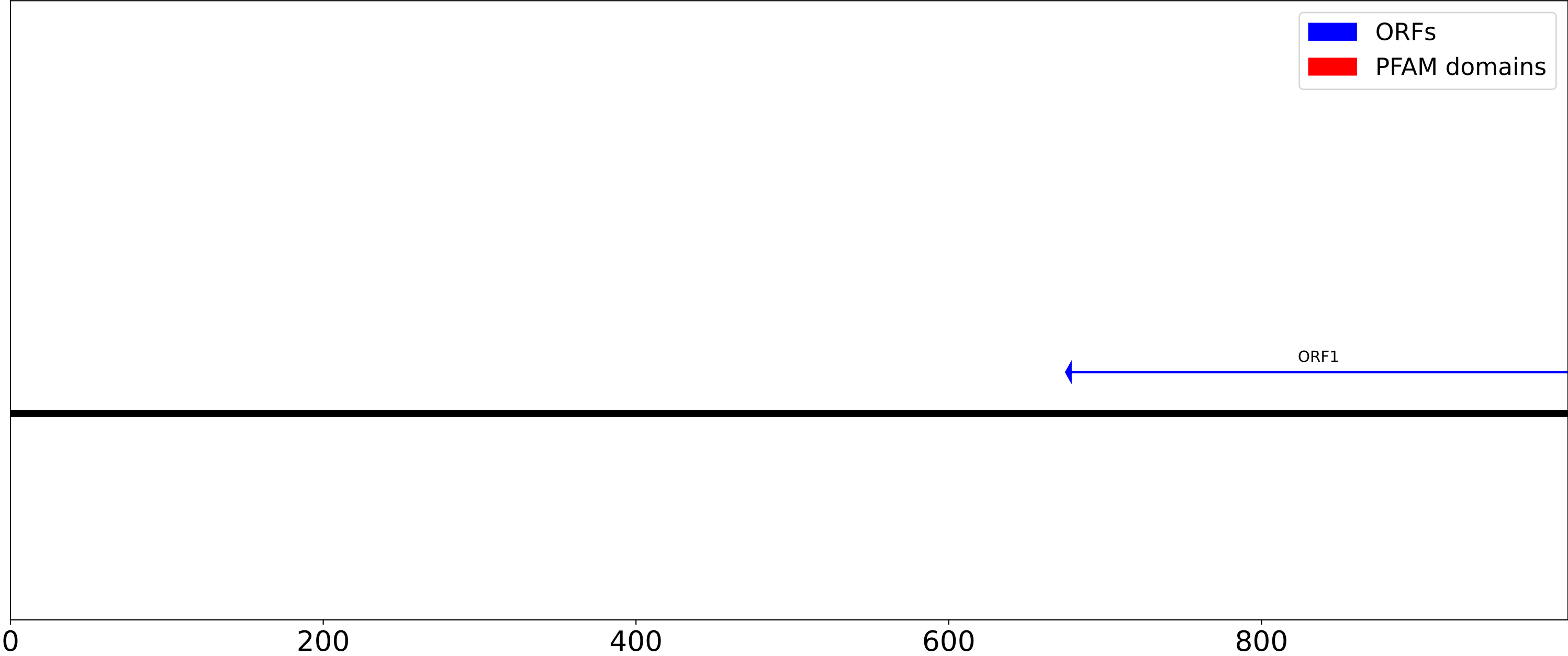
TE: rnd_5_family_17126
size: 996bp; fragments: 55; full length: 1 (≥ 896.4 bp)



After TEtrimmer ORF and PFAM domain plot



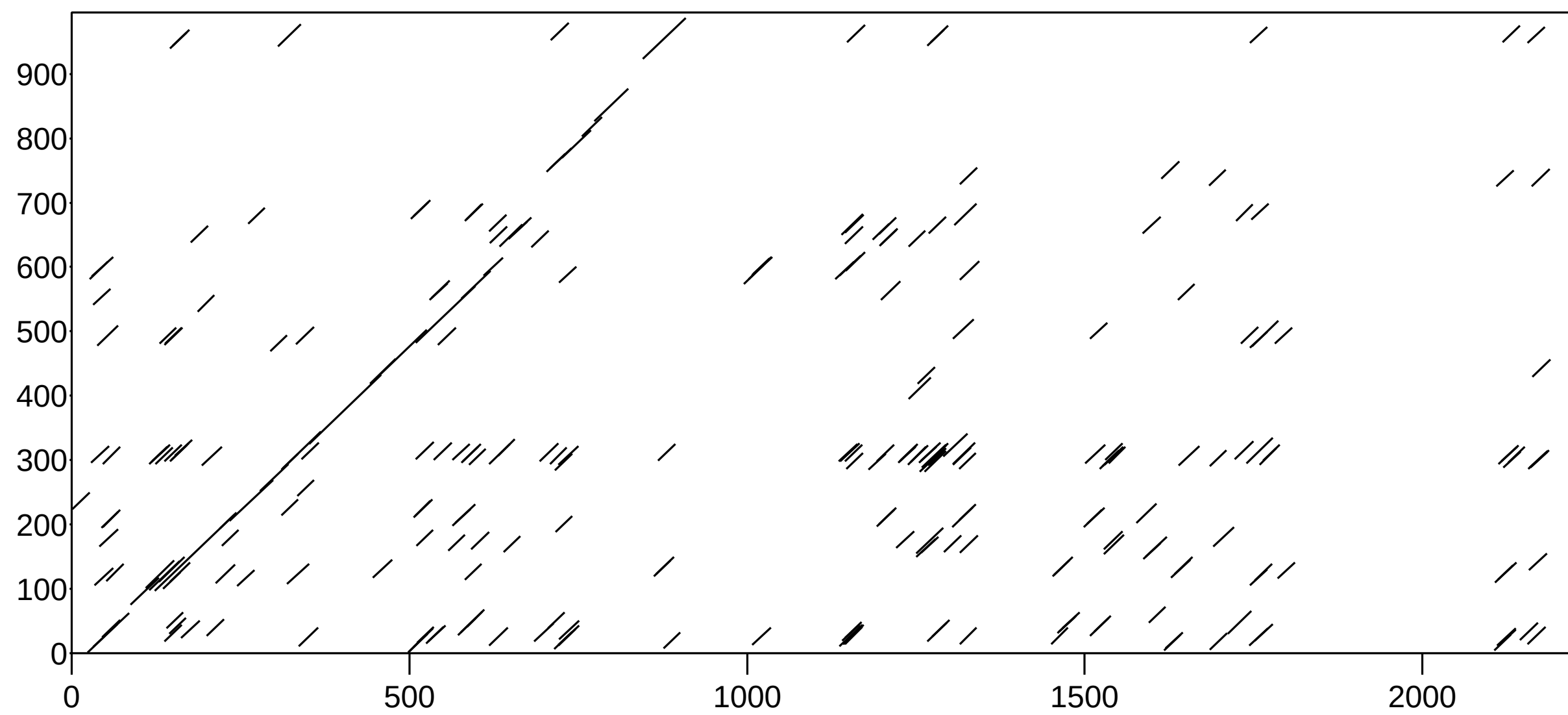
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)