

Start crop Point

End crop Point

0 -

10 -

0

250

500

750

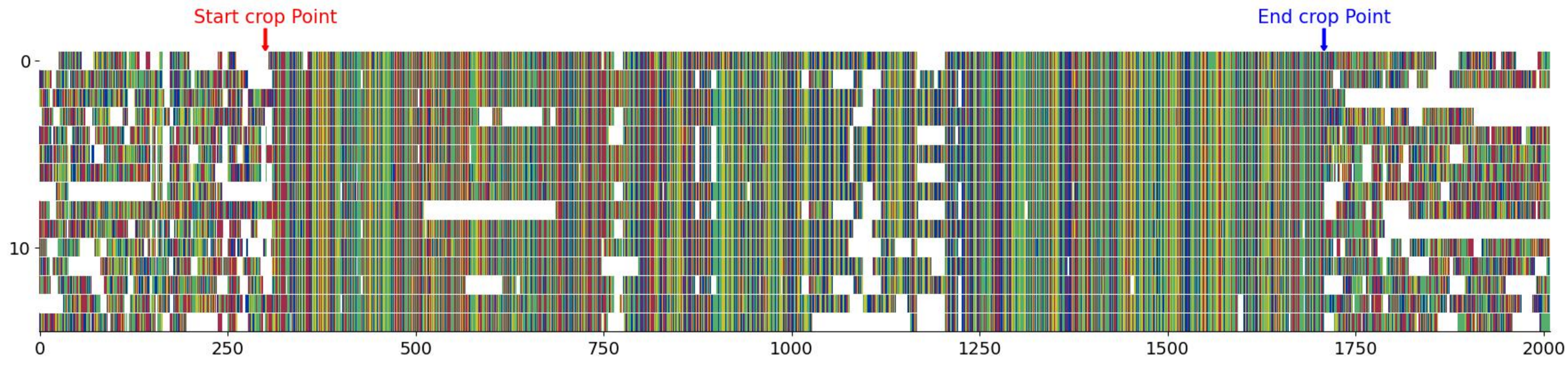
1000

1250

1500

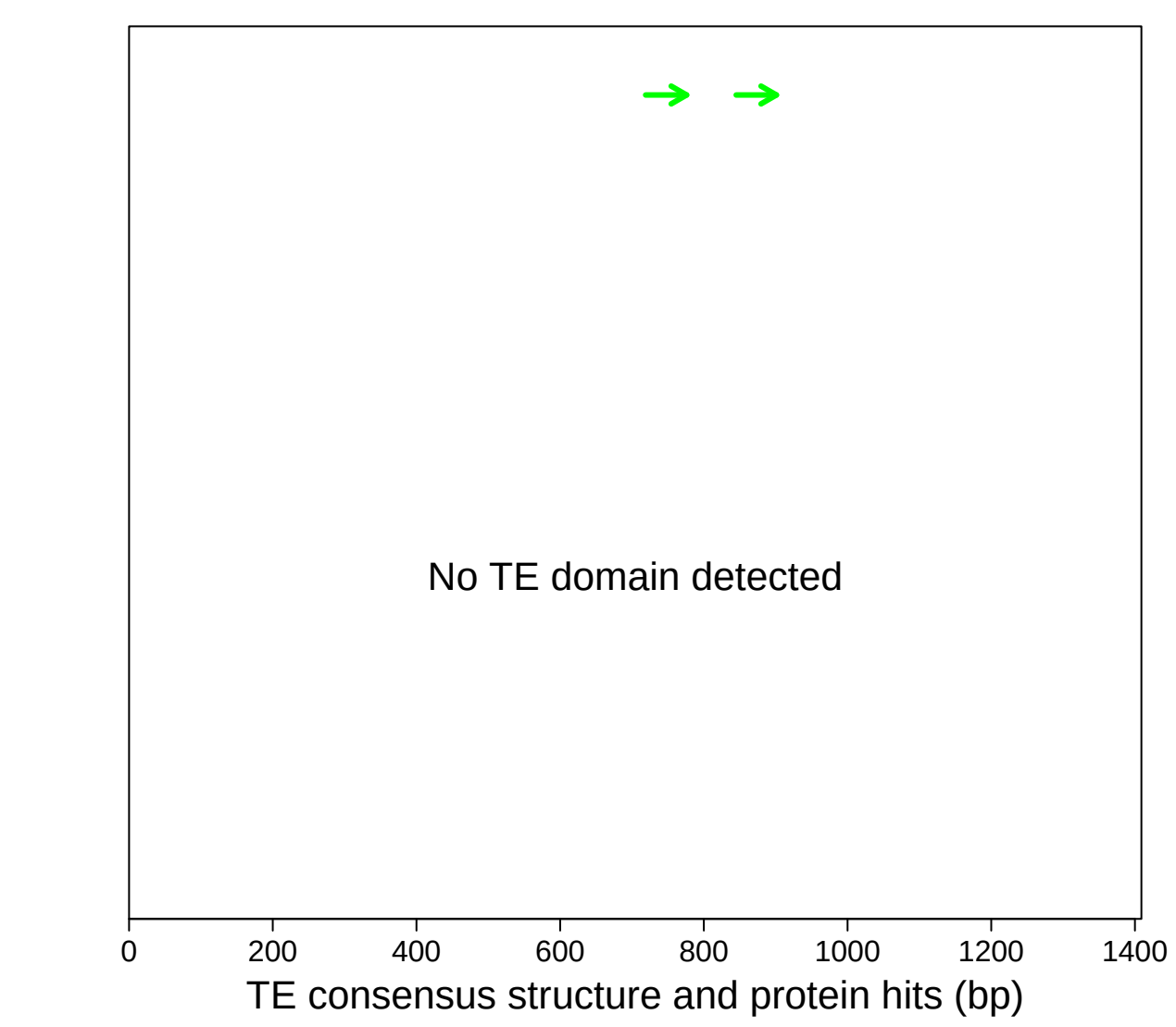
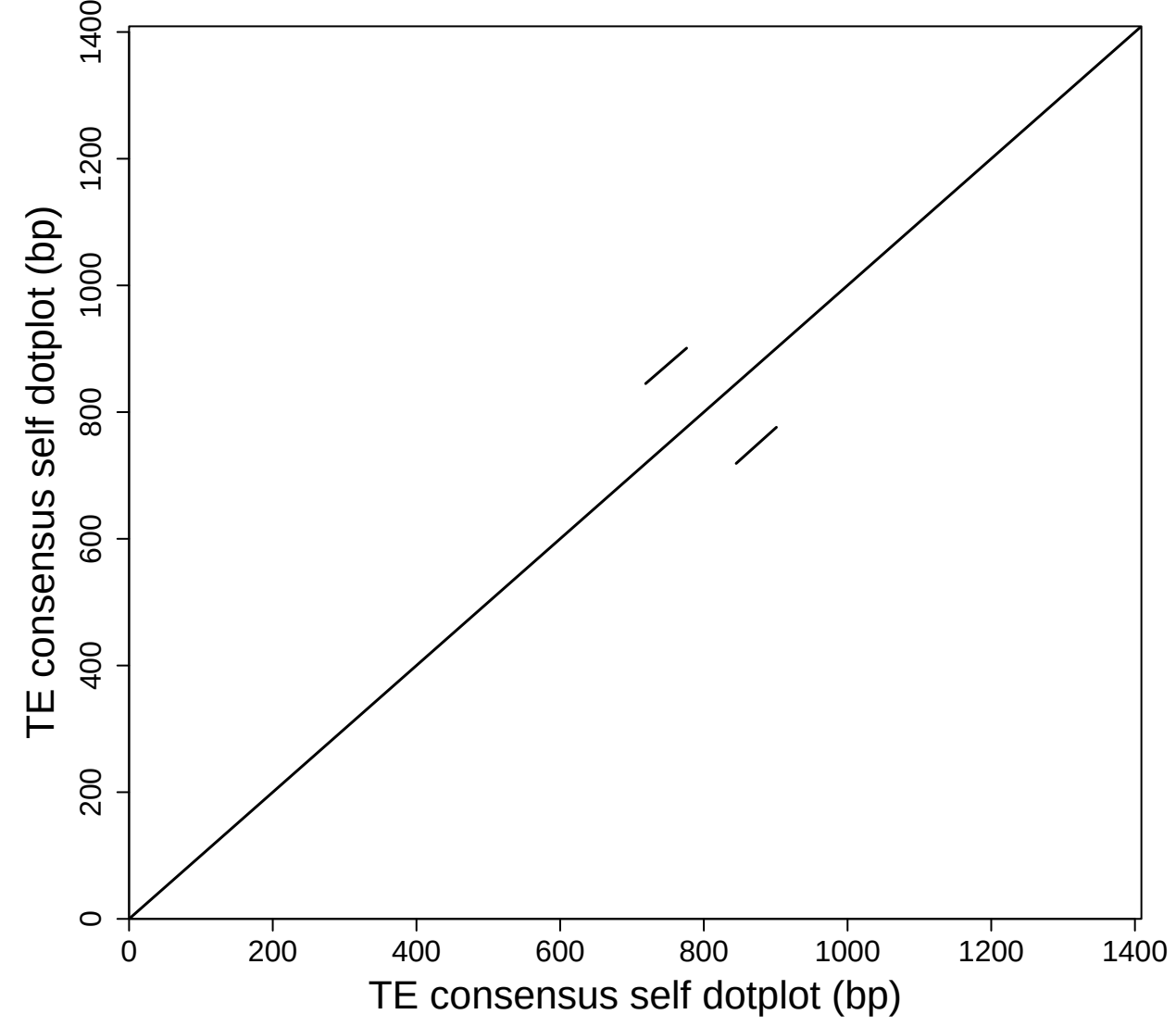
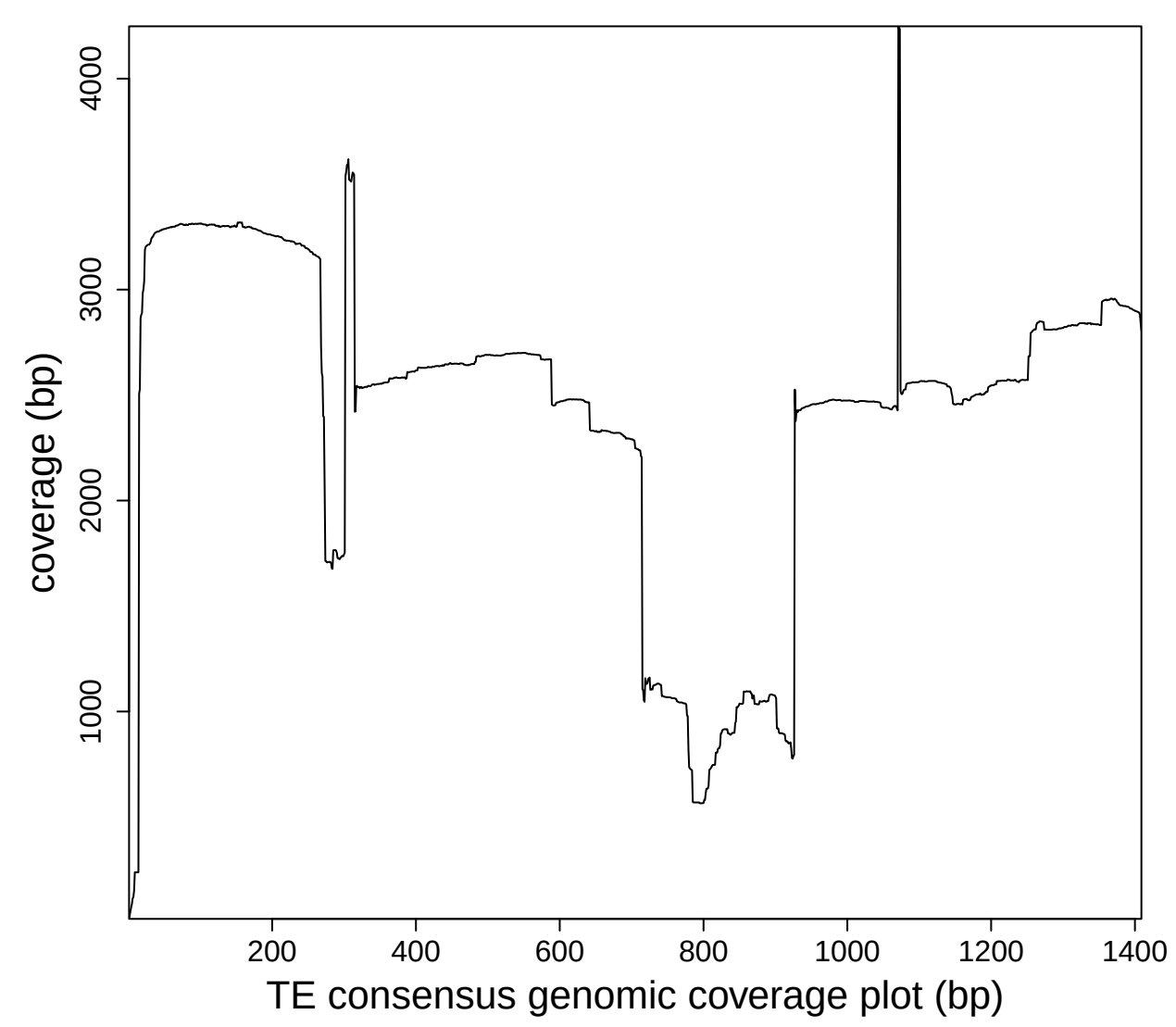
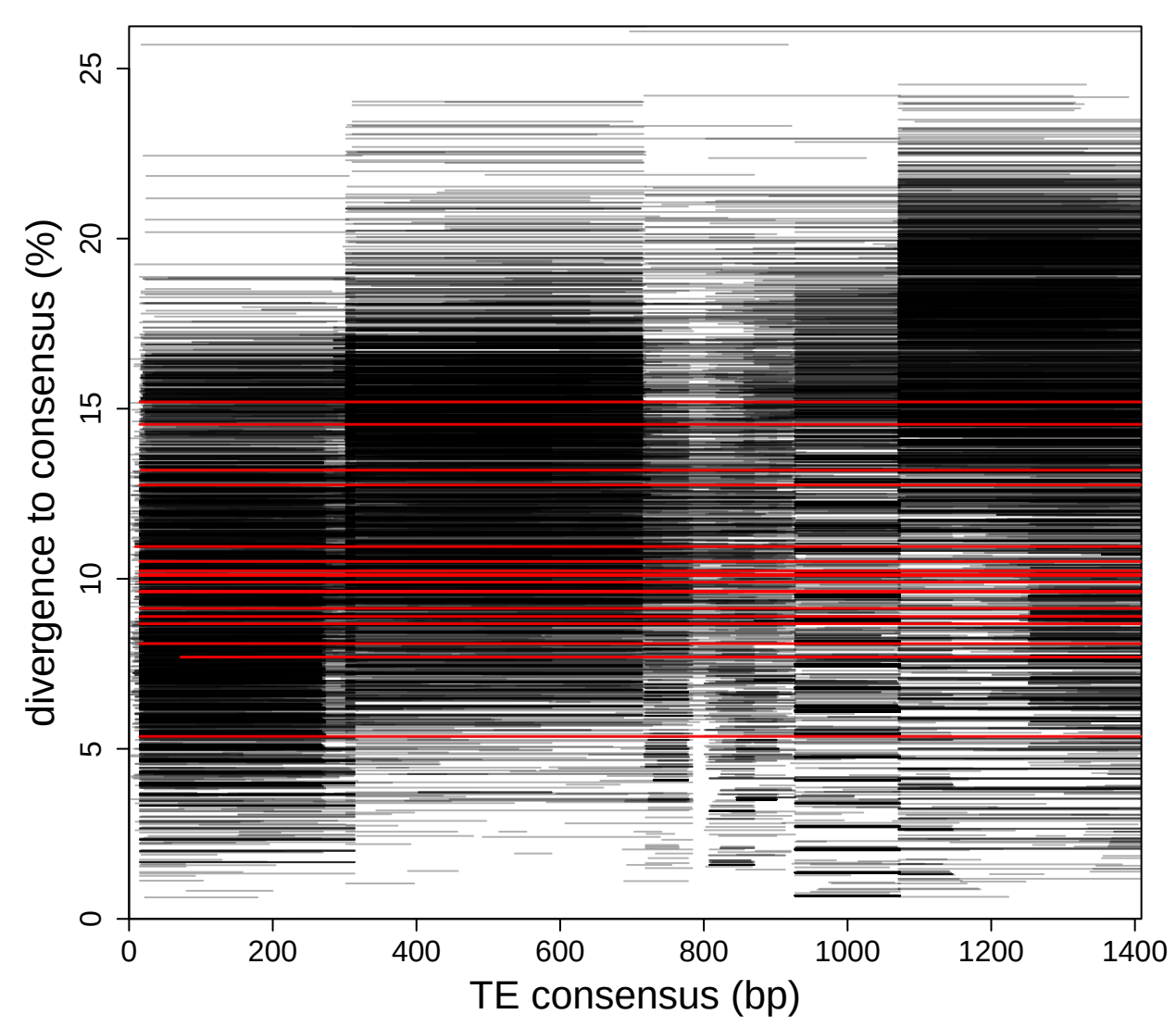
1750

2000



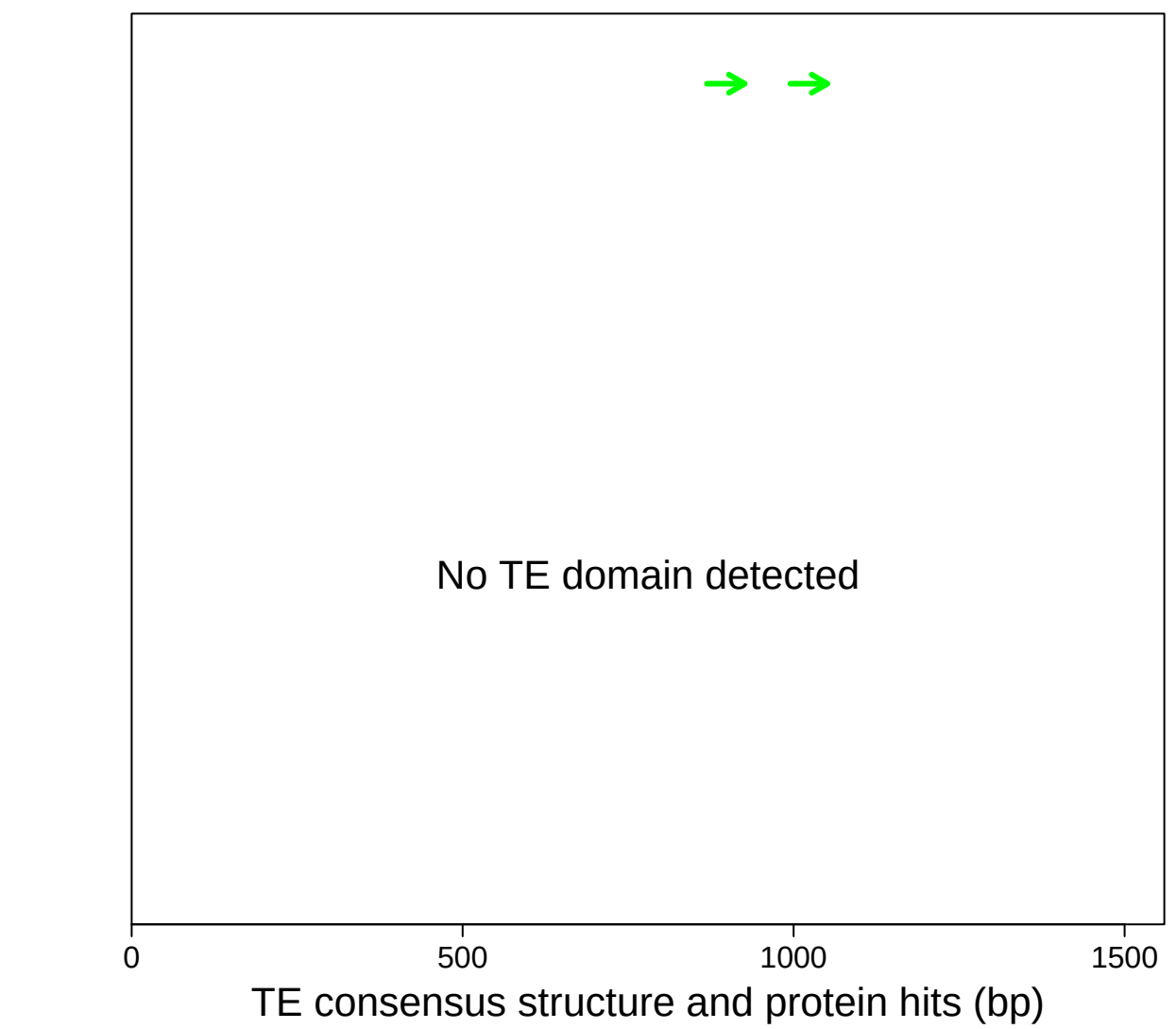
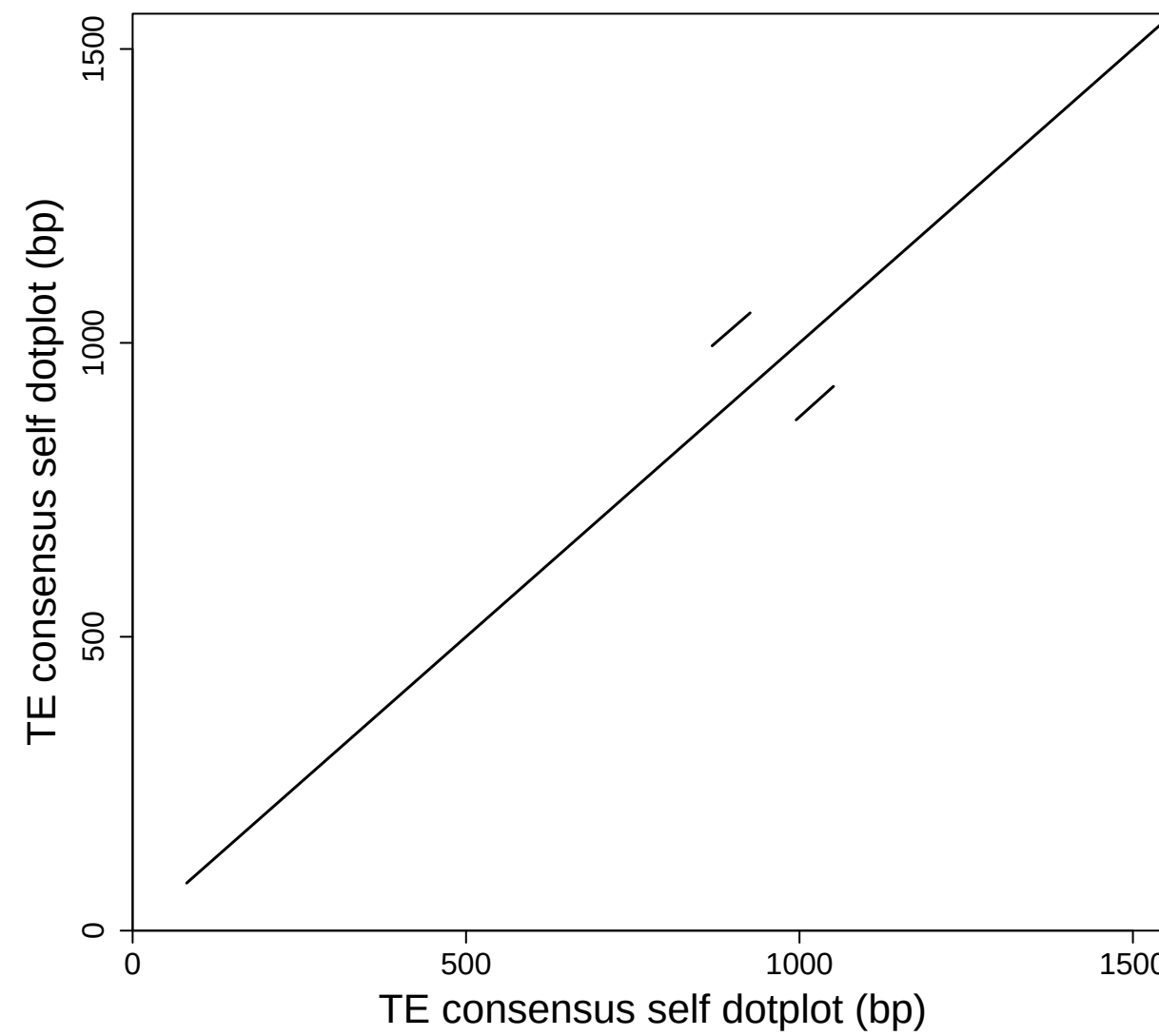
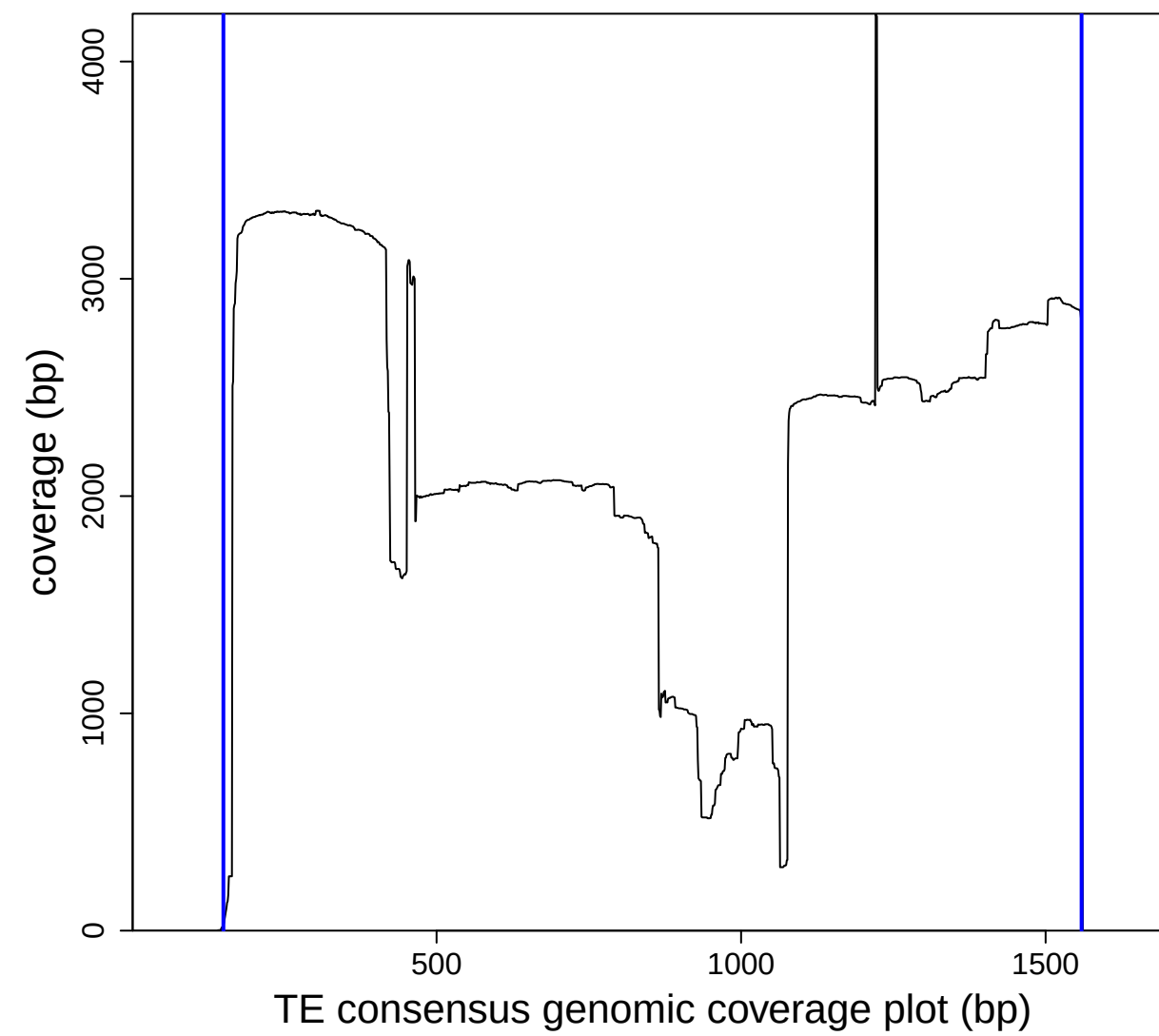
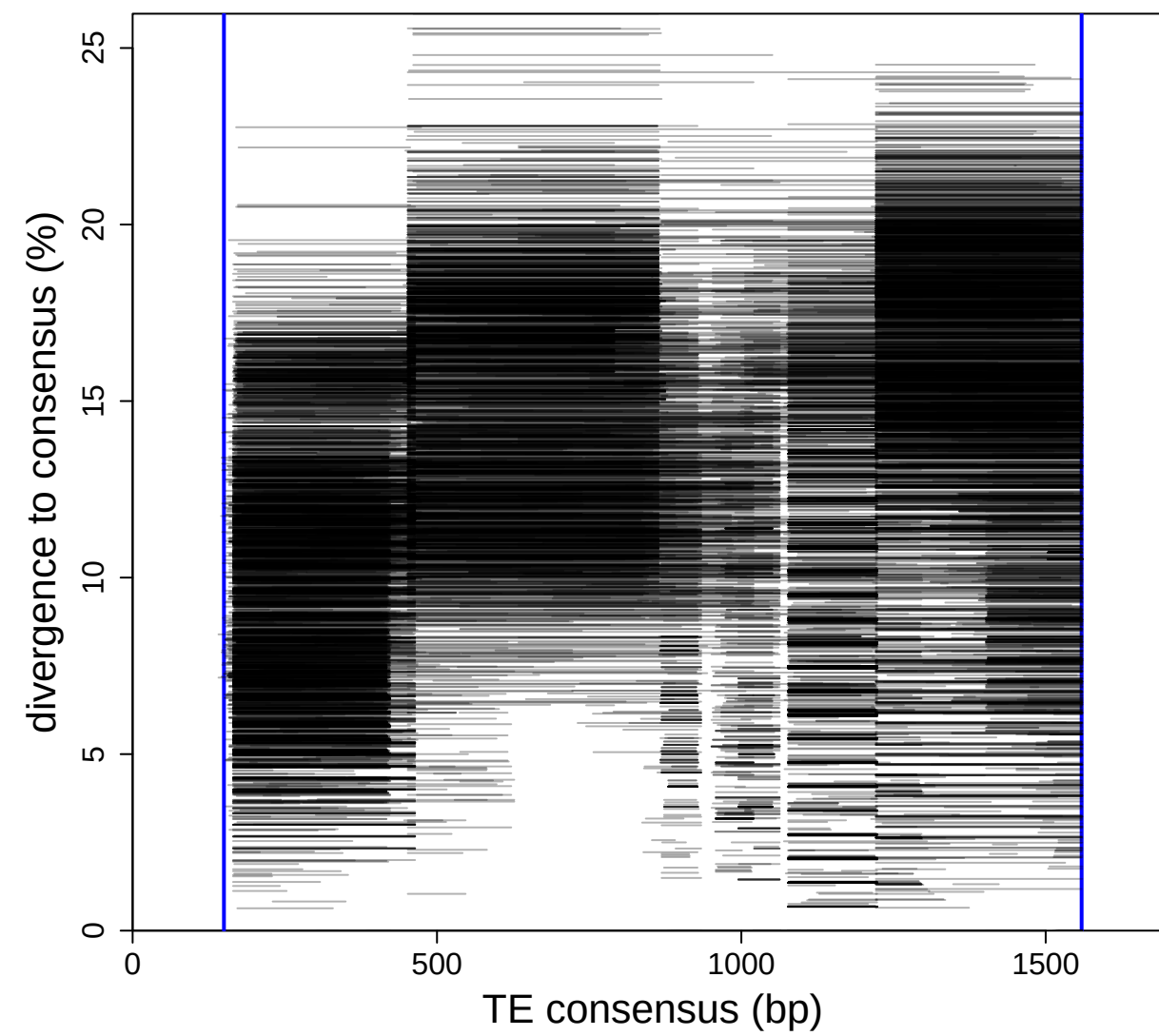
b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 1409bp; fragments: 11882; full length: 18 (≥ 1268.1 bp)

After TEtrimmer 1409 bp

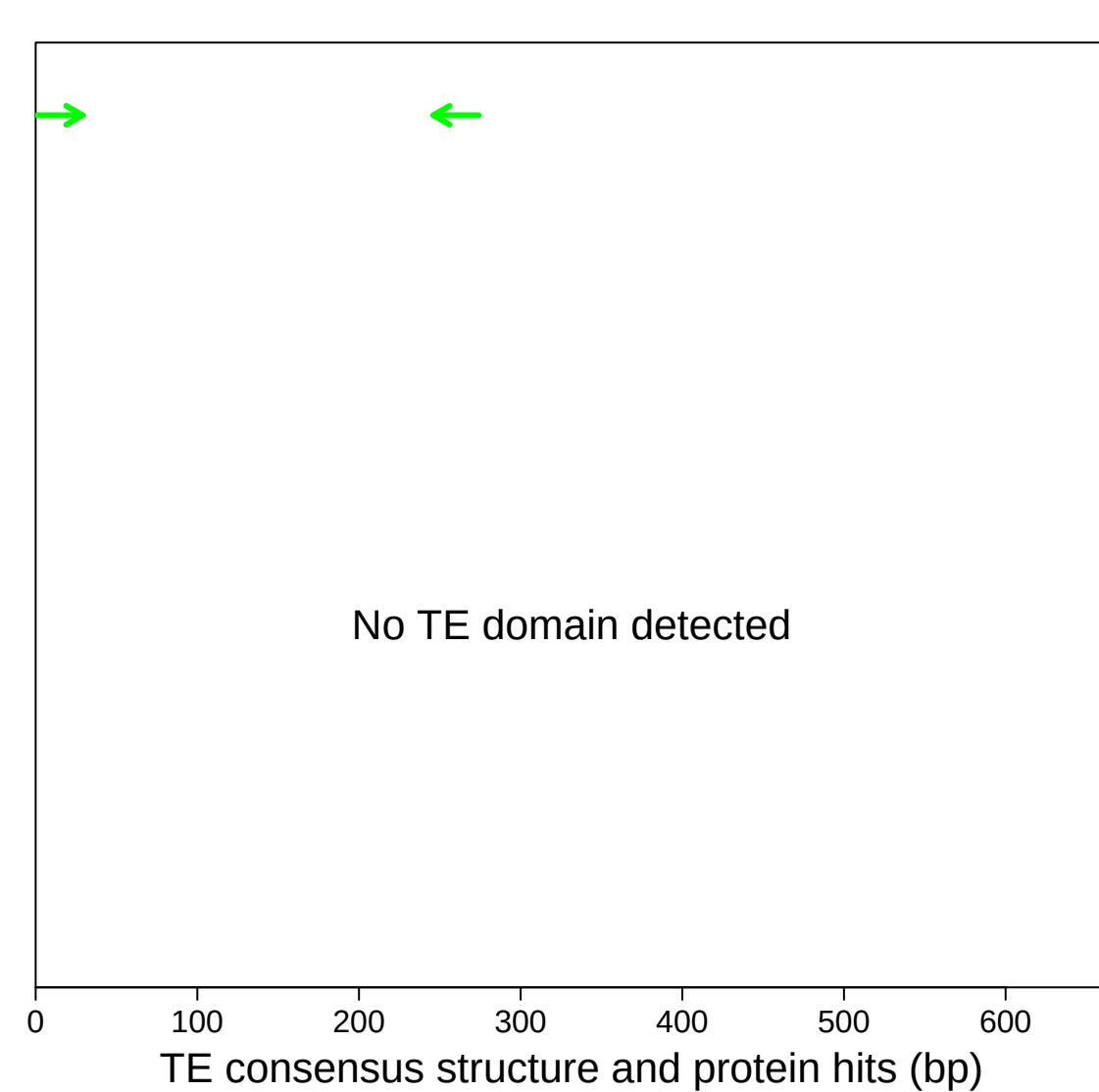
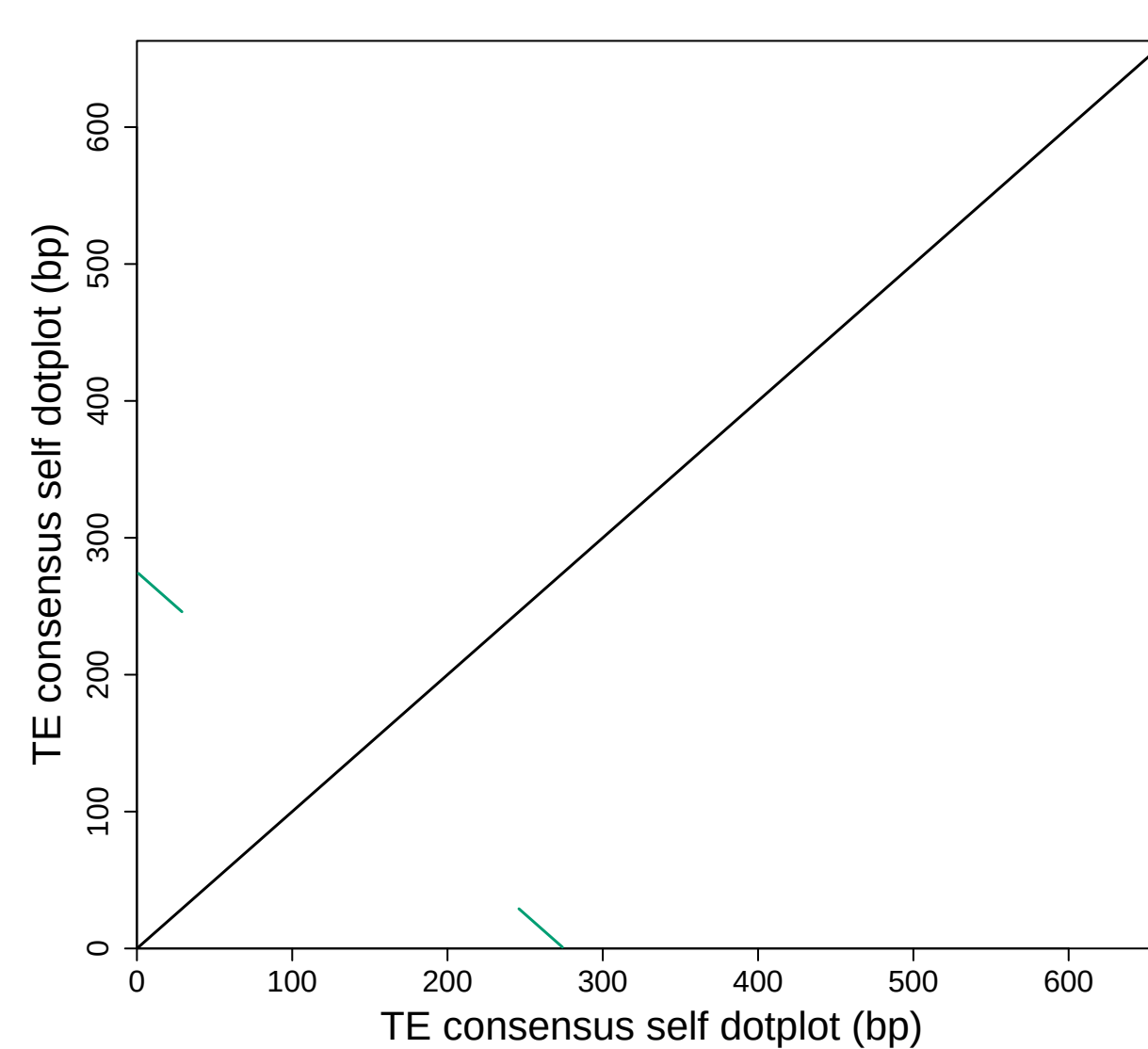
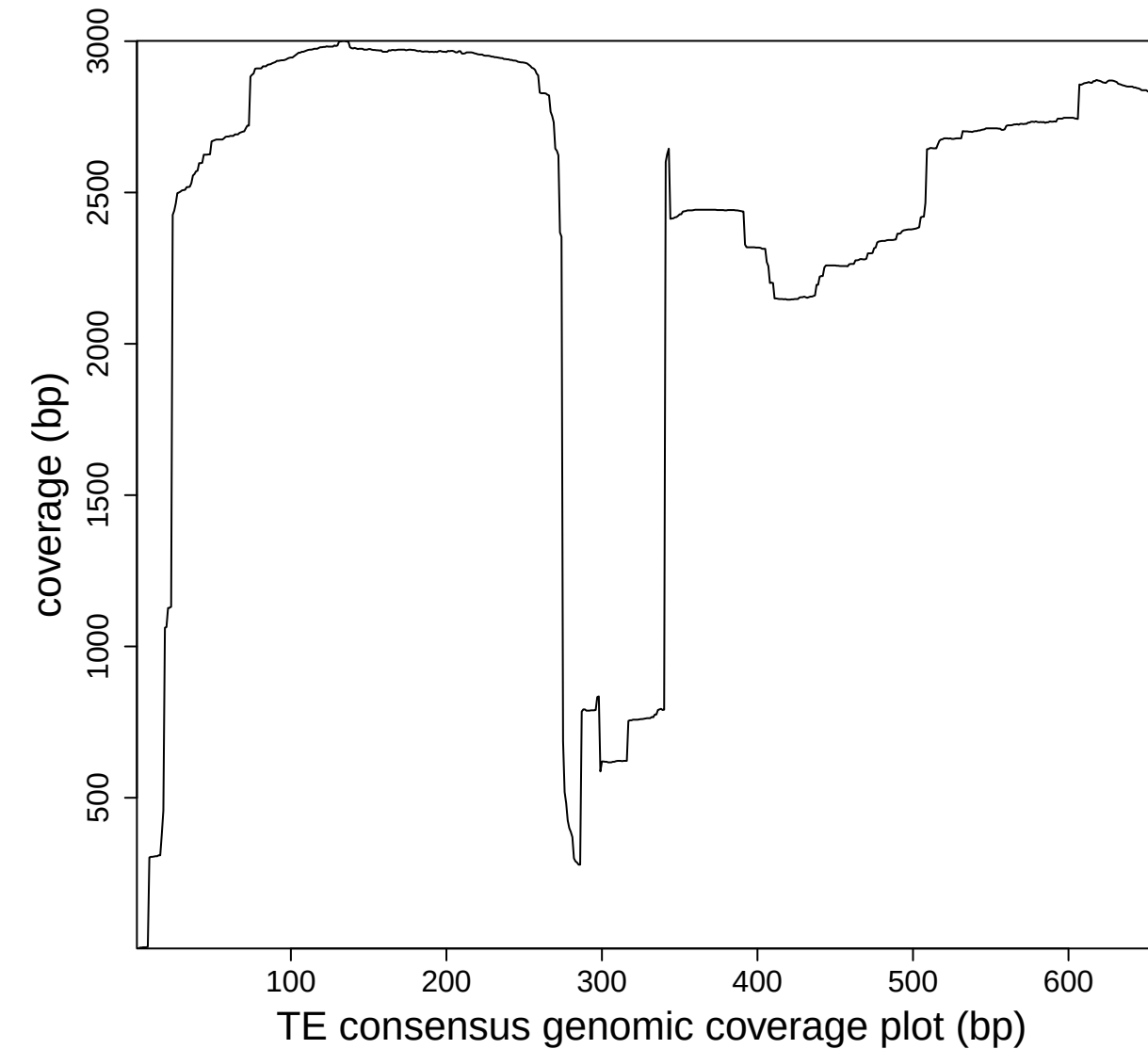
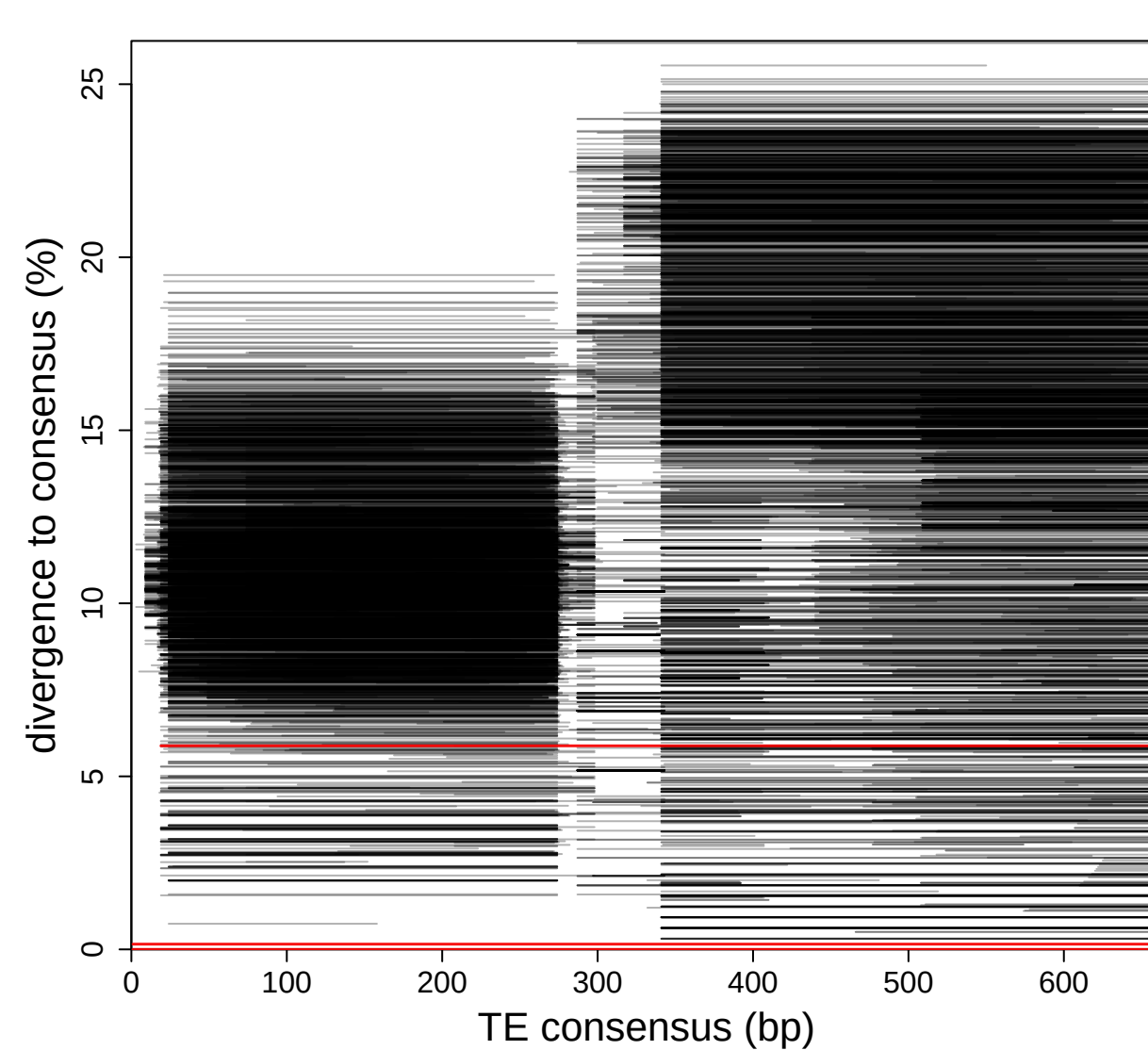


3.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 1709bp; fragments: 11167; full length: 0 (>=1538.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

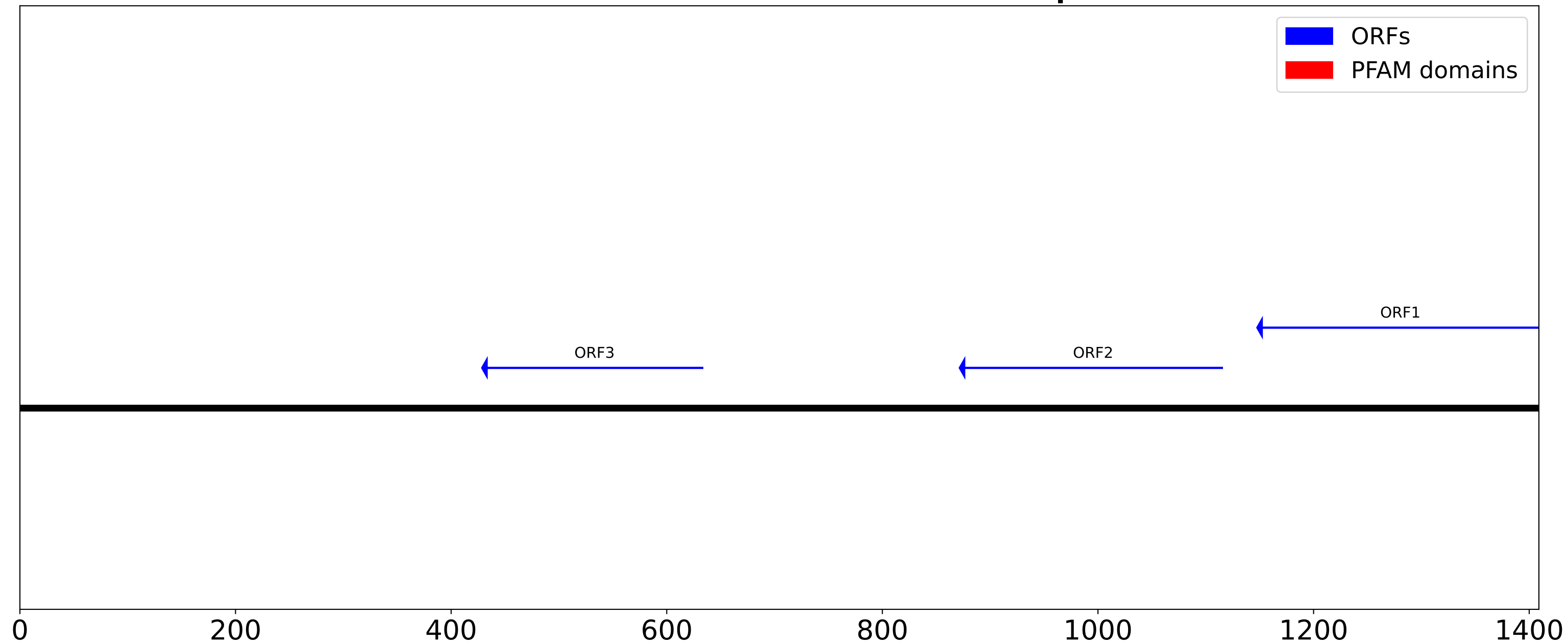


TE: ltr_1_family_257
size: 663bp; fragments: 6624; full length: 3 (≥ 596.7 bp)

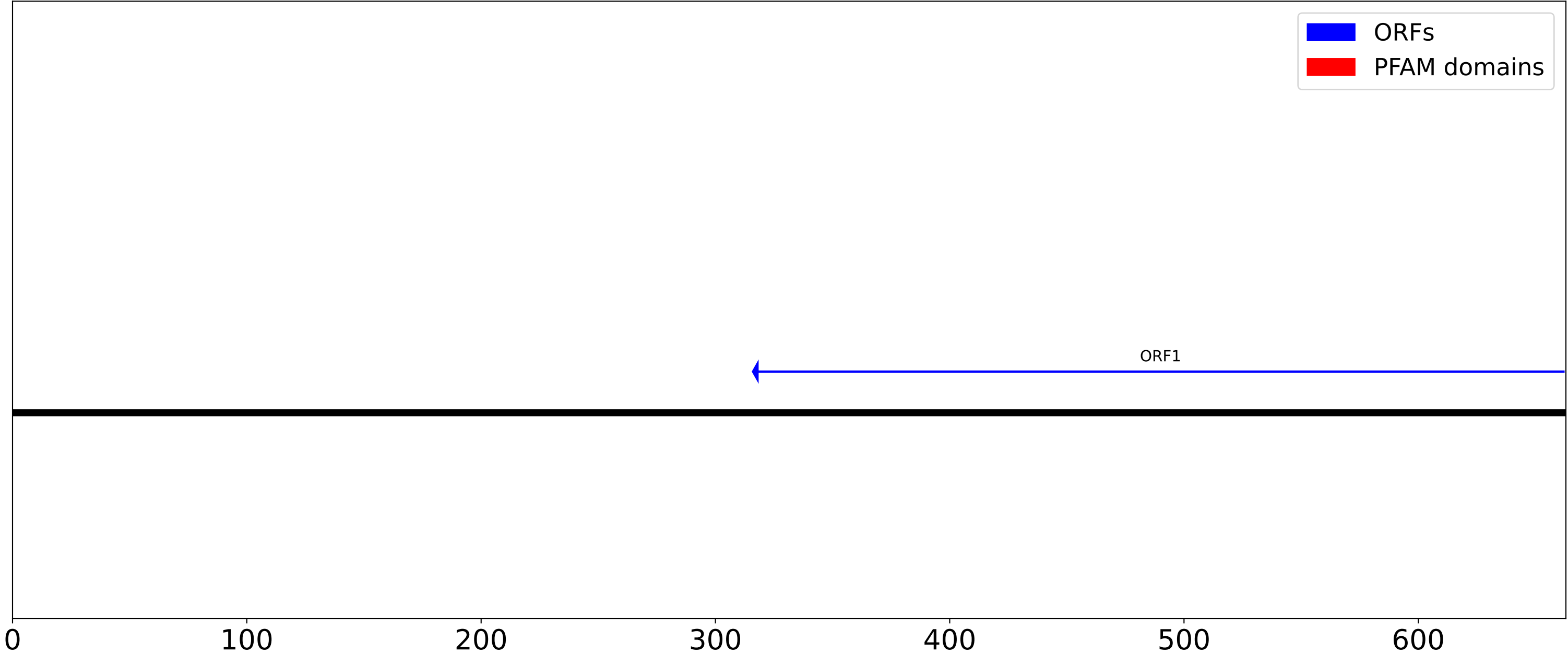


Before TEtrimmer 663 bp

After TEtrimmer ORF and PFAM domain plot



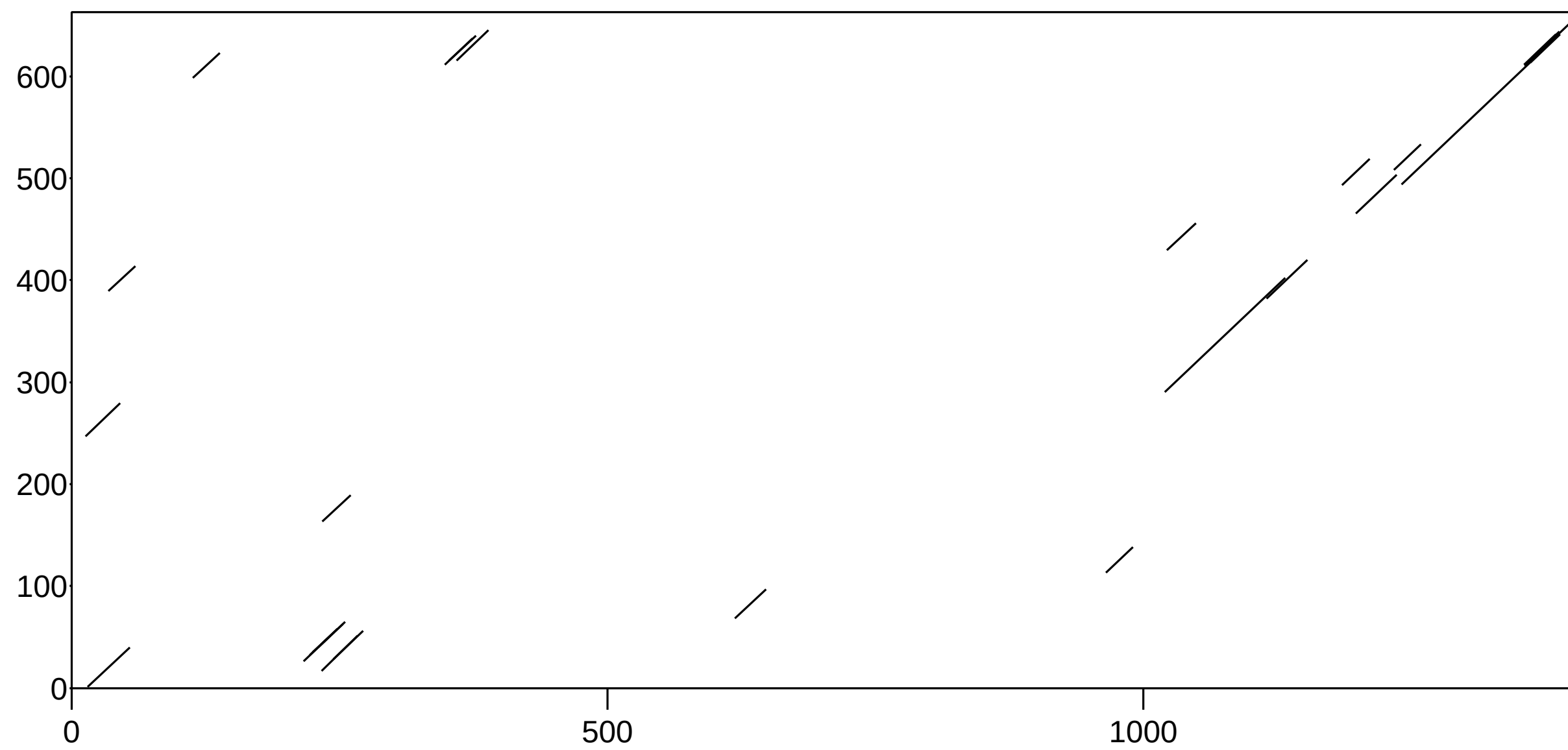
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)