

Start crop Point

End crop Point

MSA length = 1893



Start crop Point

End crop Point

0

10

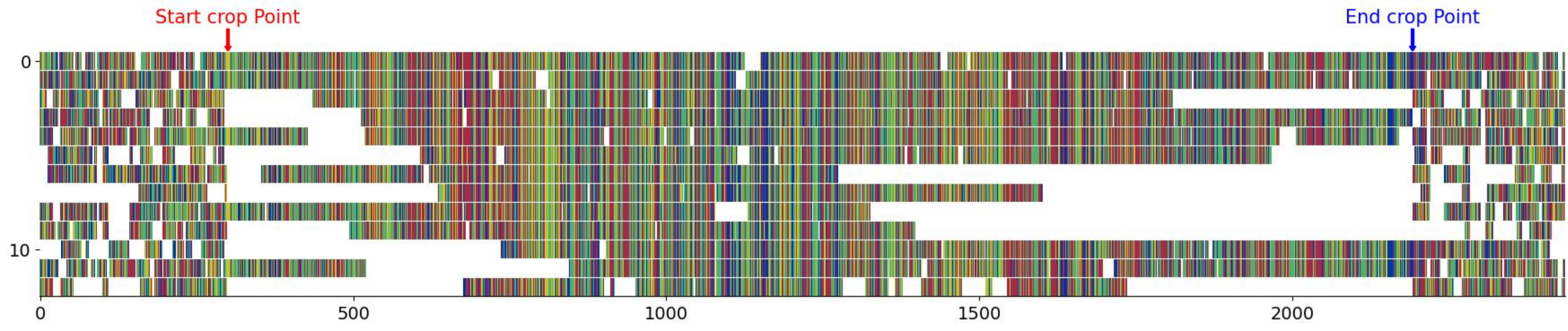
0

500

1000

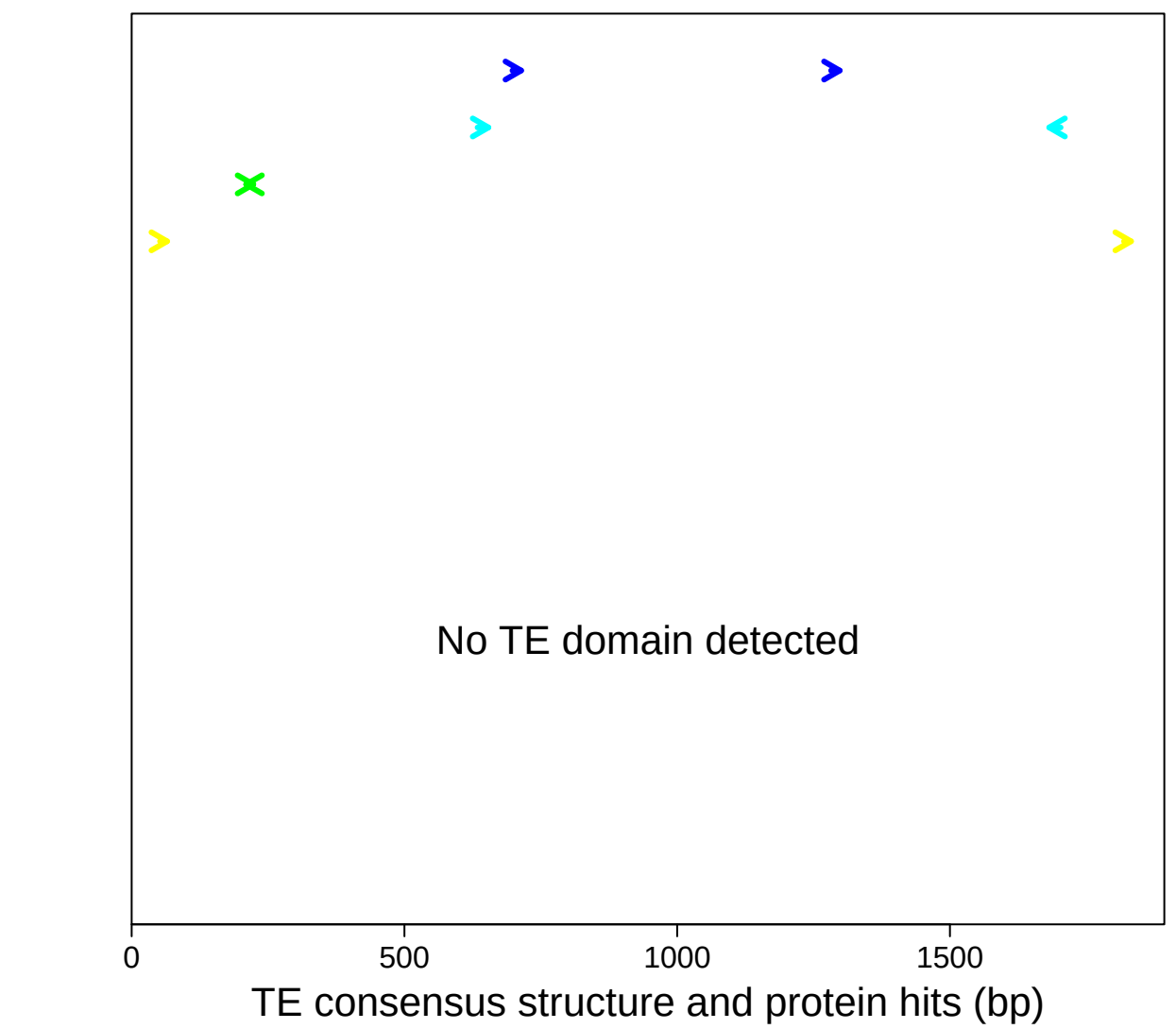
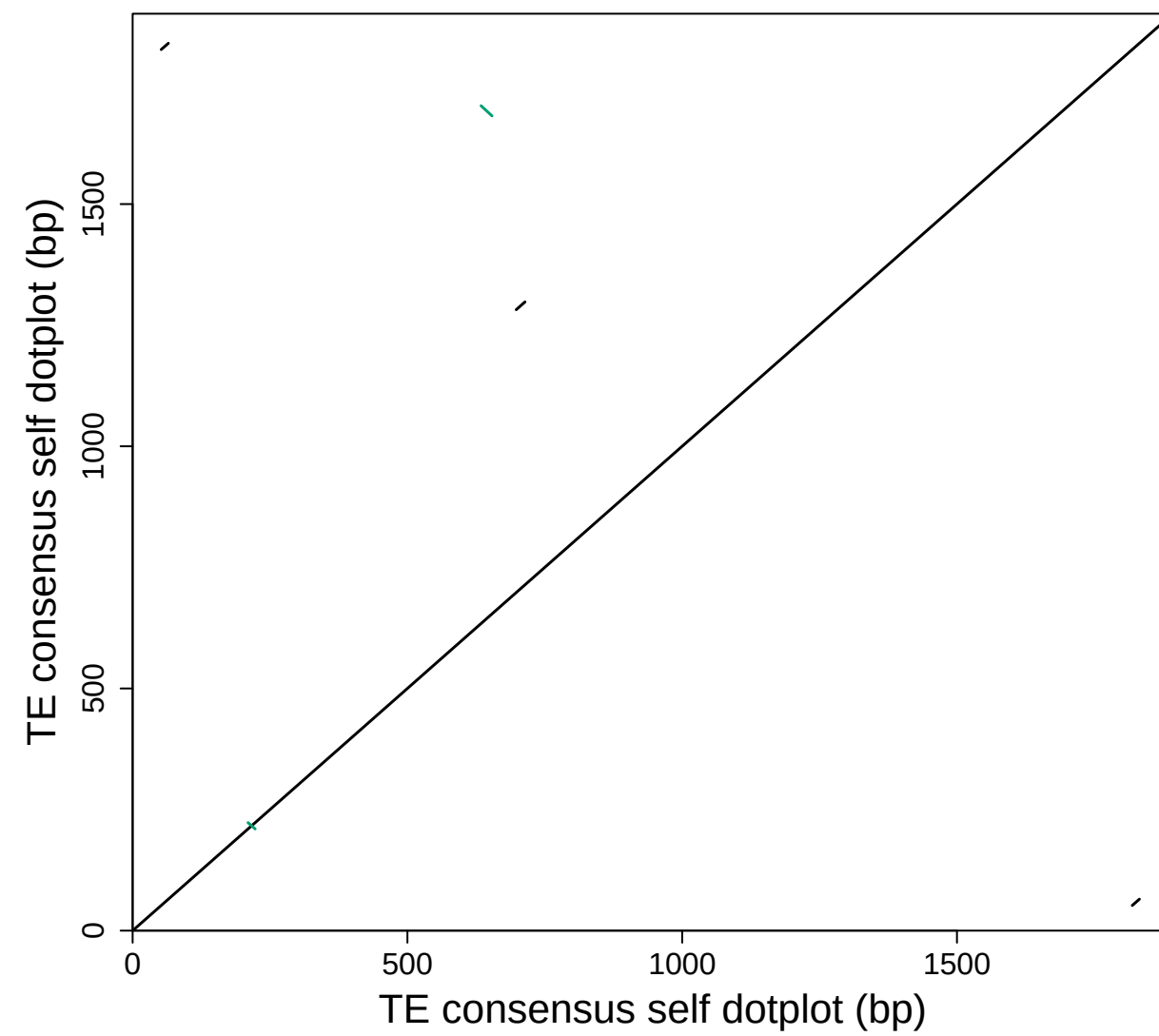
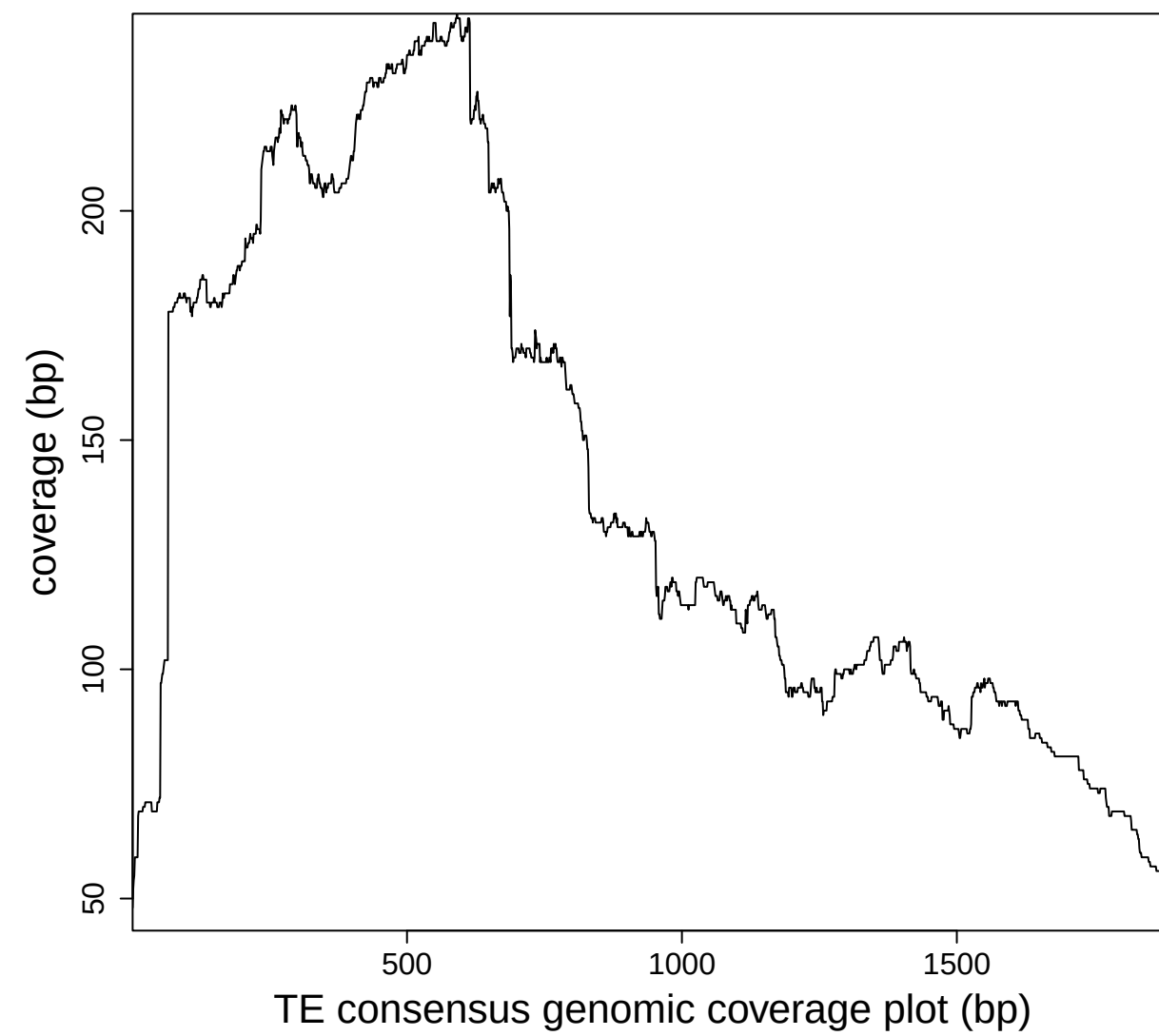
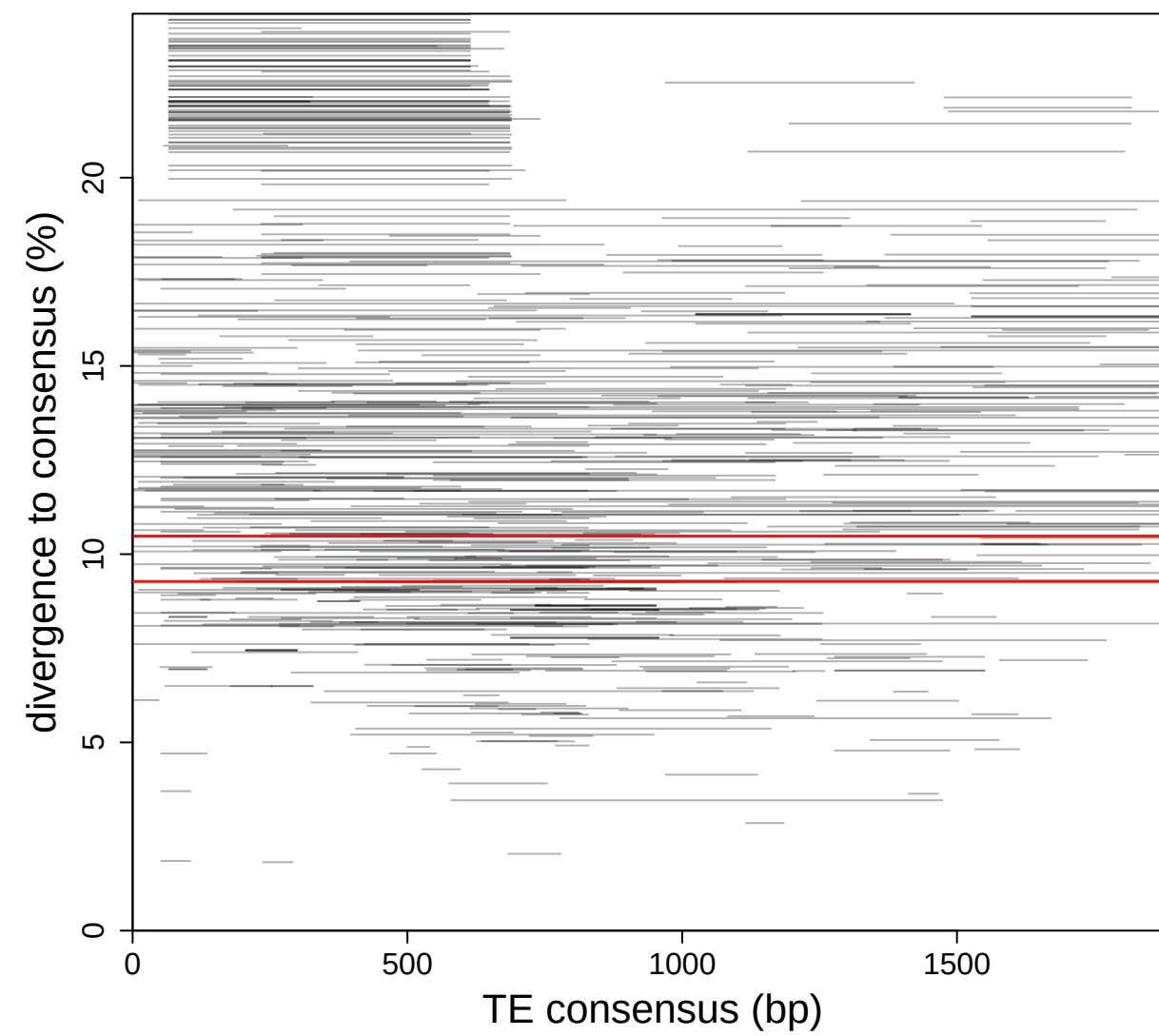
1500

2000



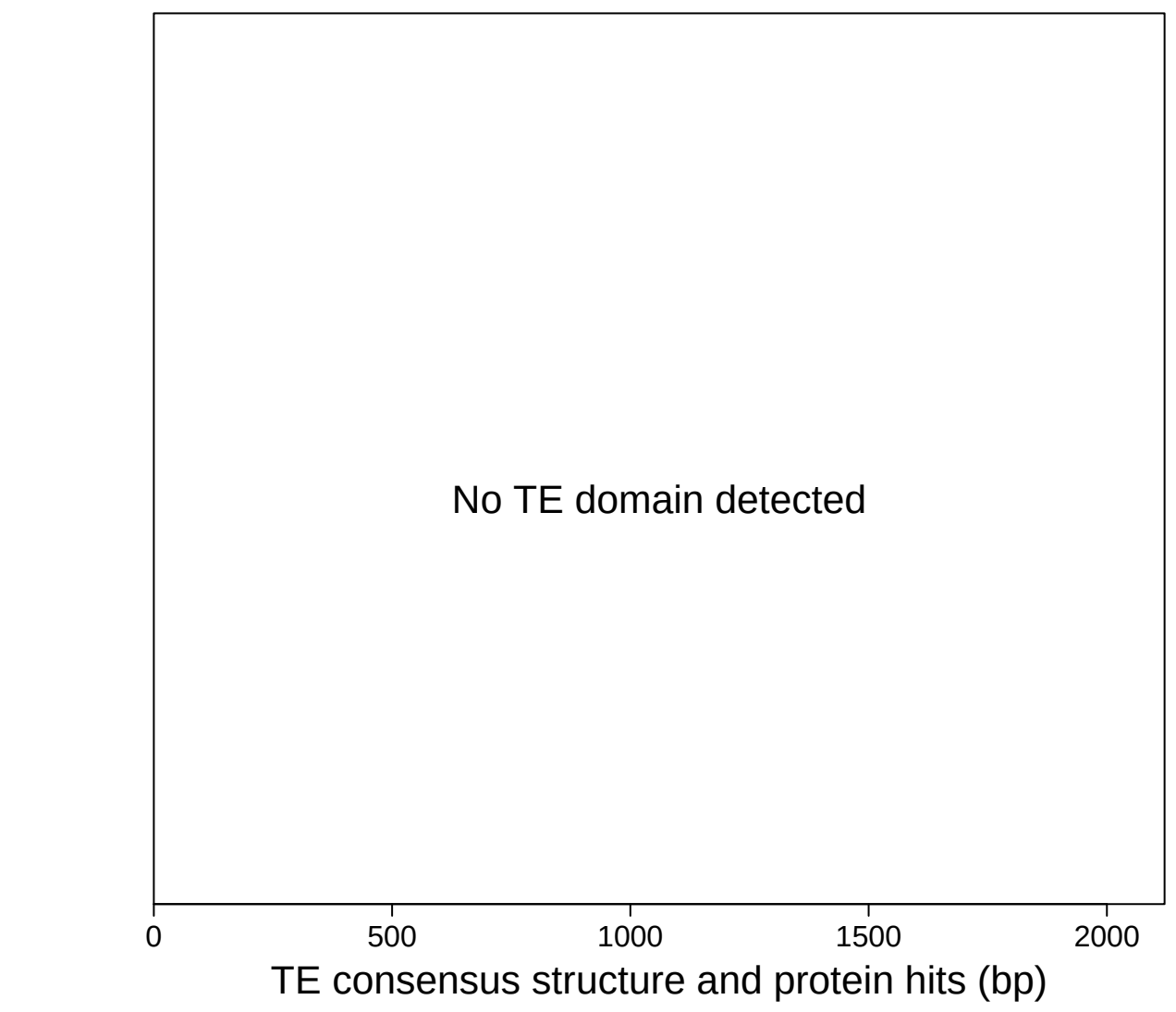
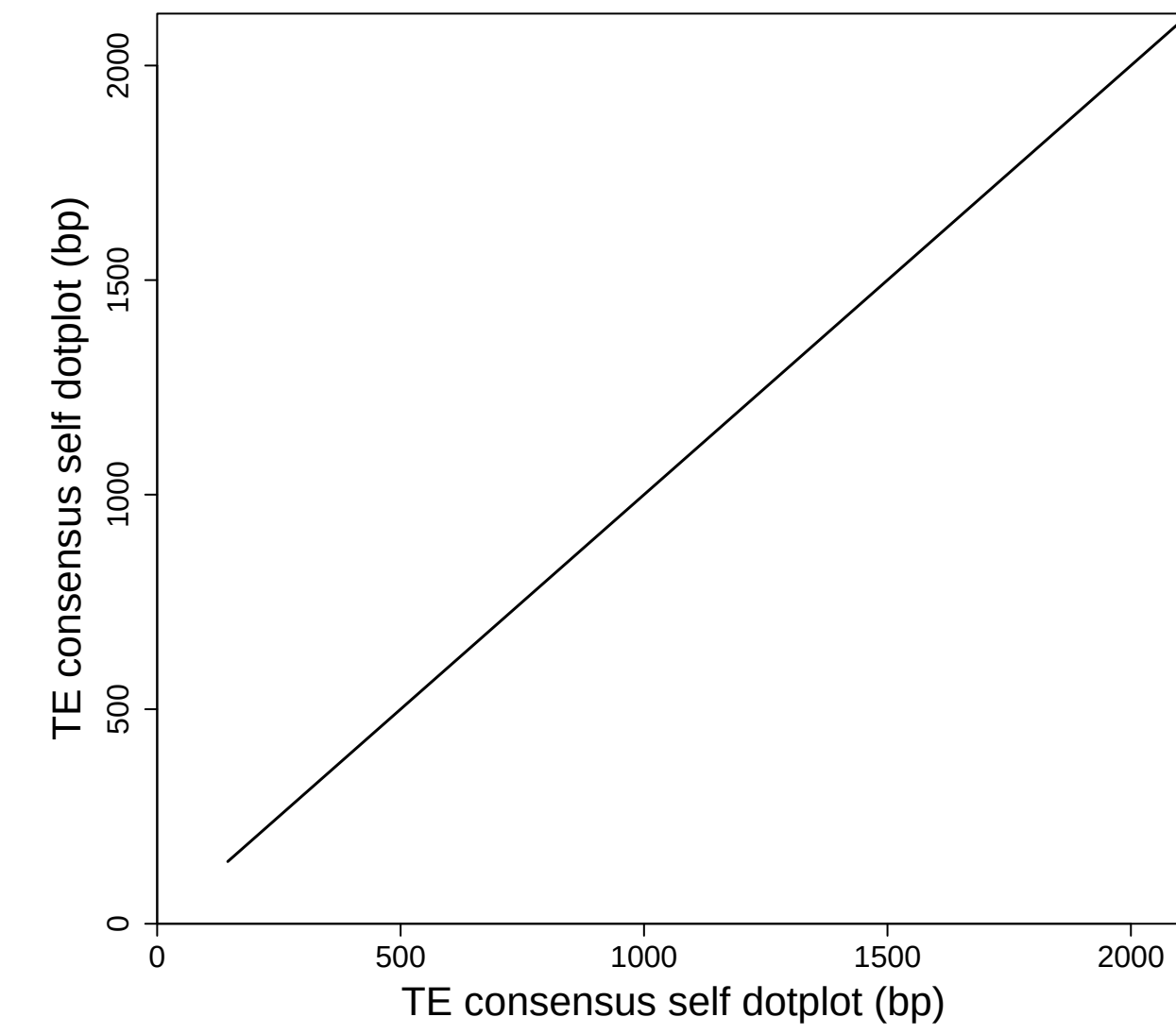
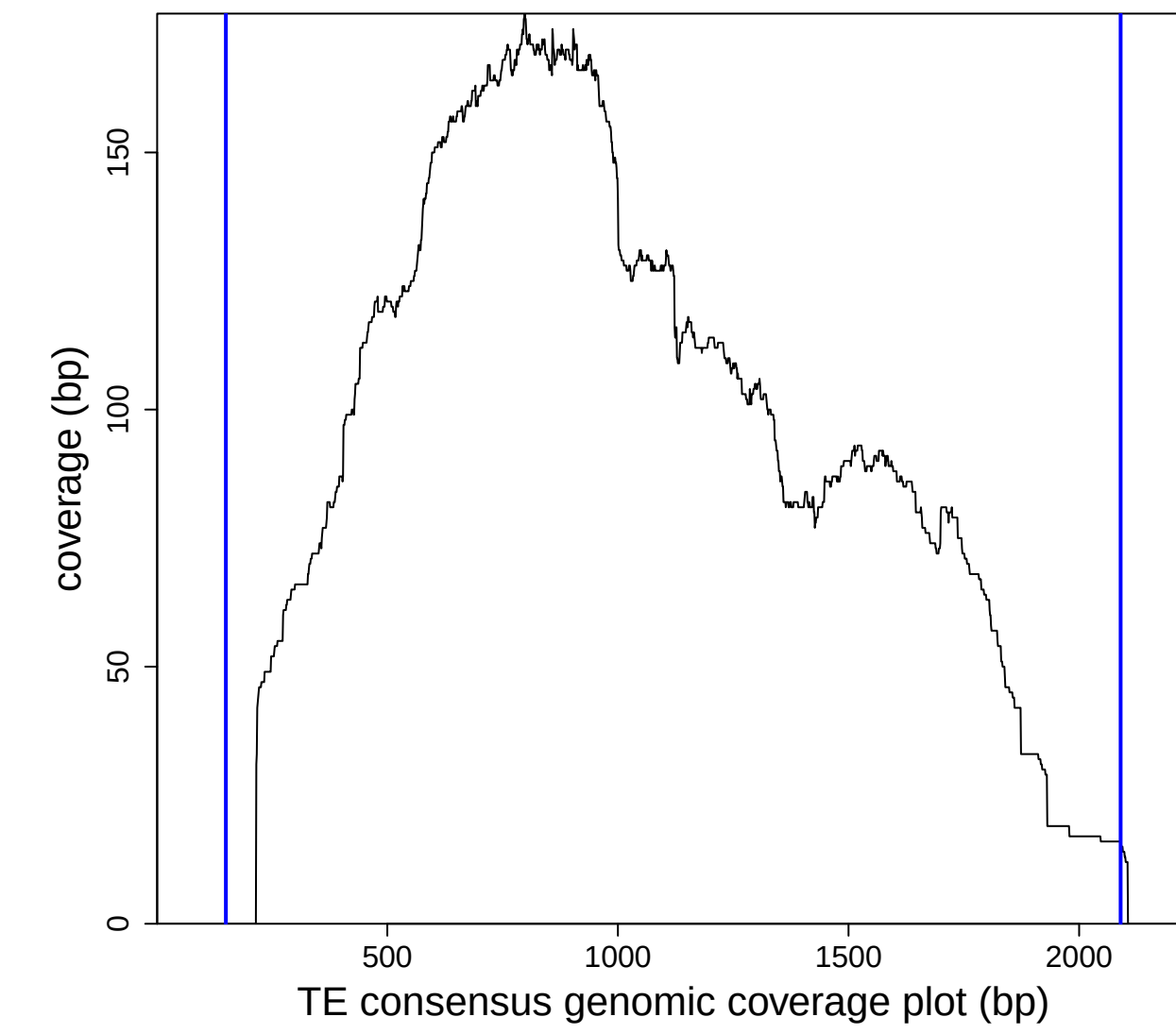
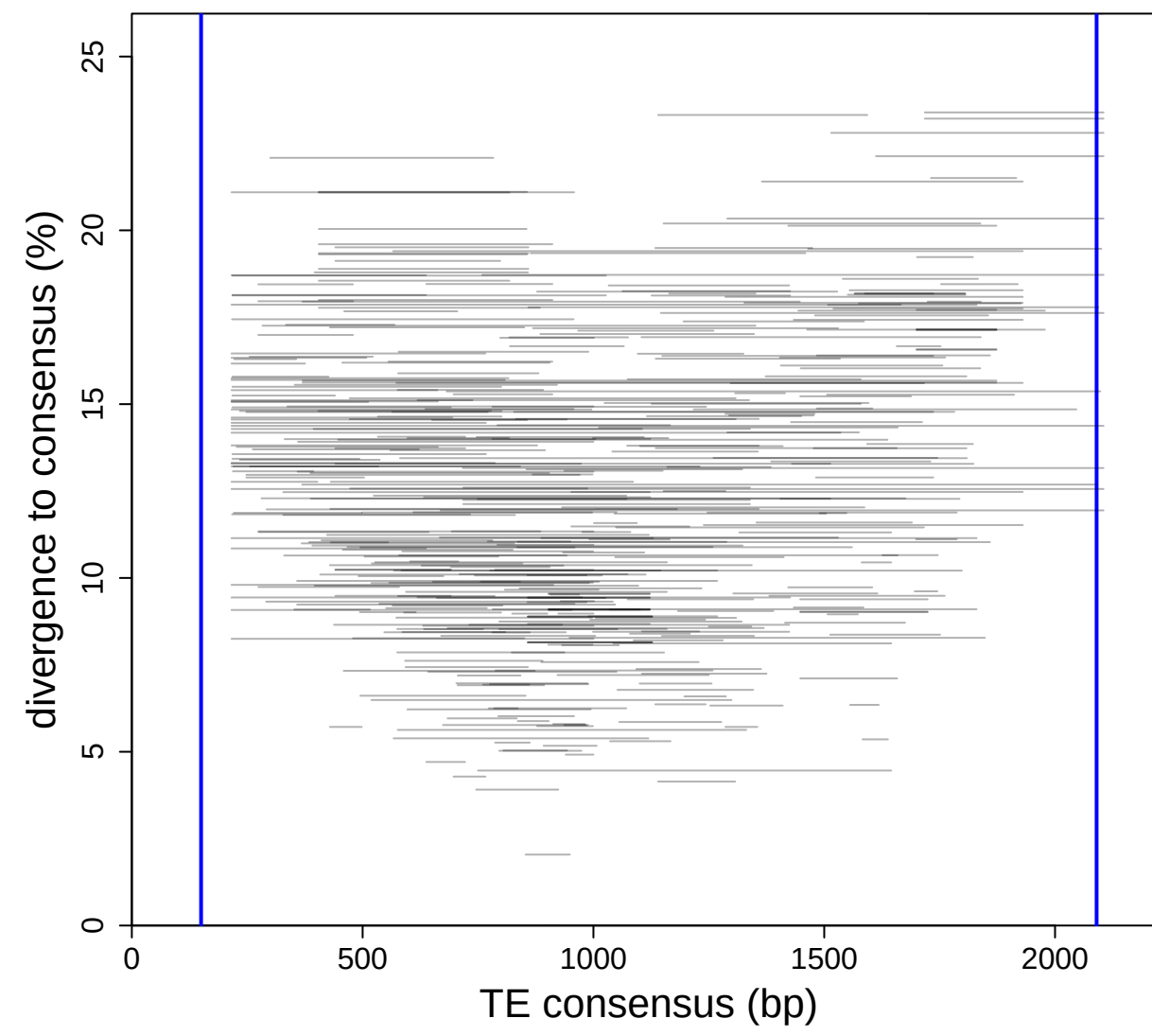
l.b.bed uf.bed g_2.bed fm_1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa
size: 1893bp; fragments: 616; full length: 2 (>=1703.7bp)

After TEtrimmer 1893 bp

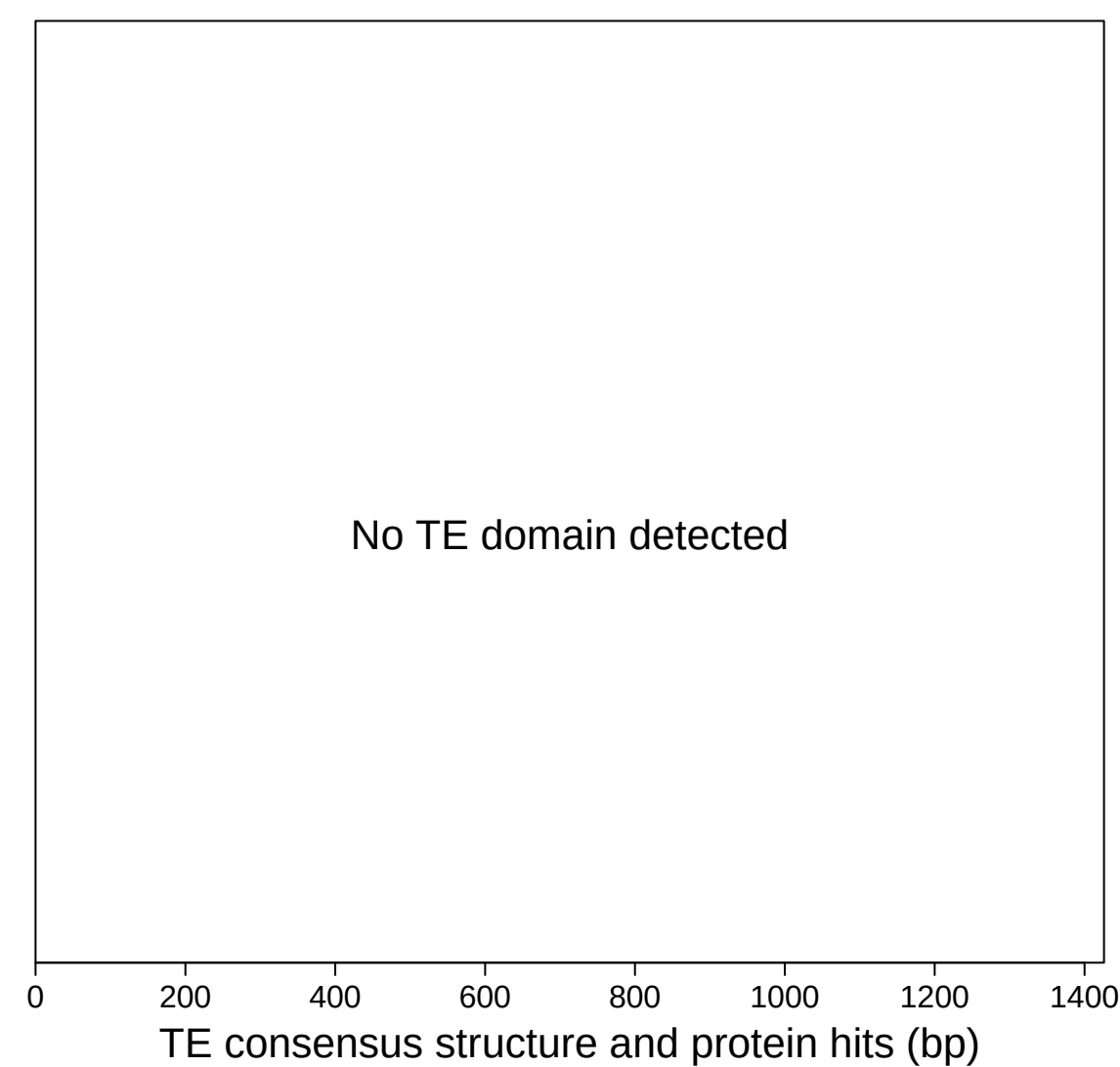
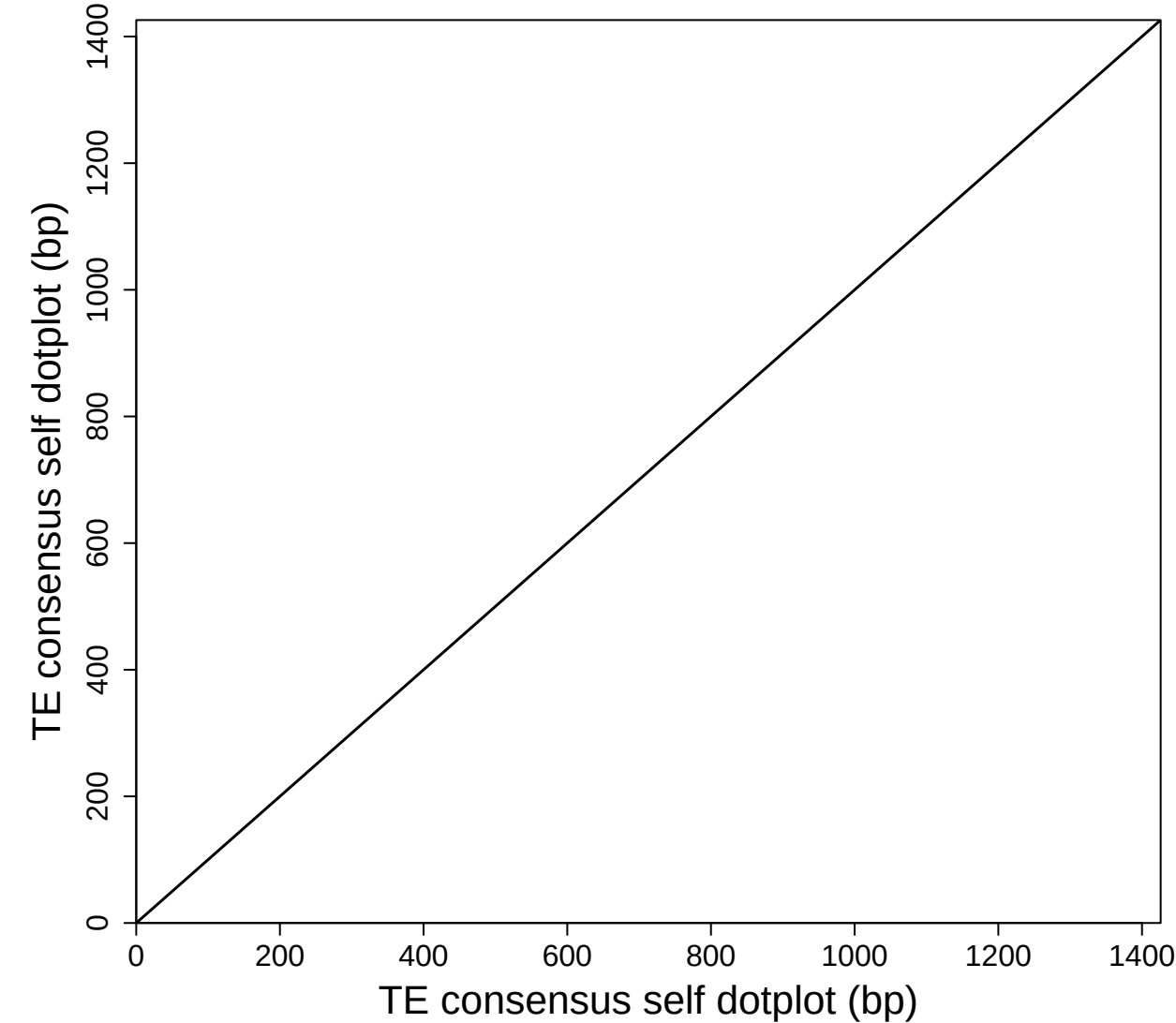
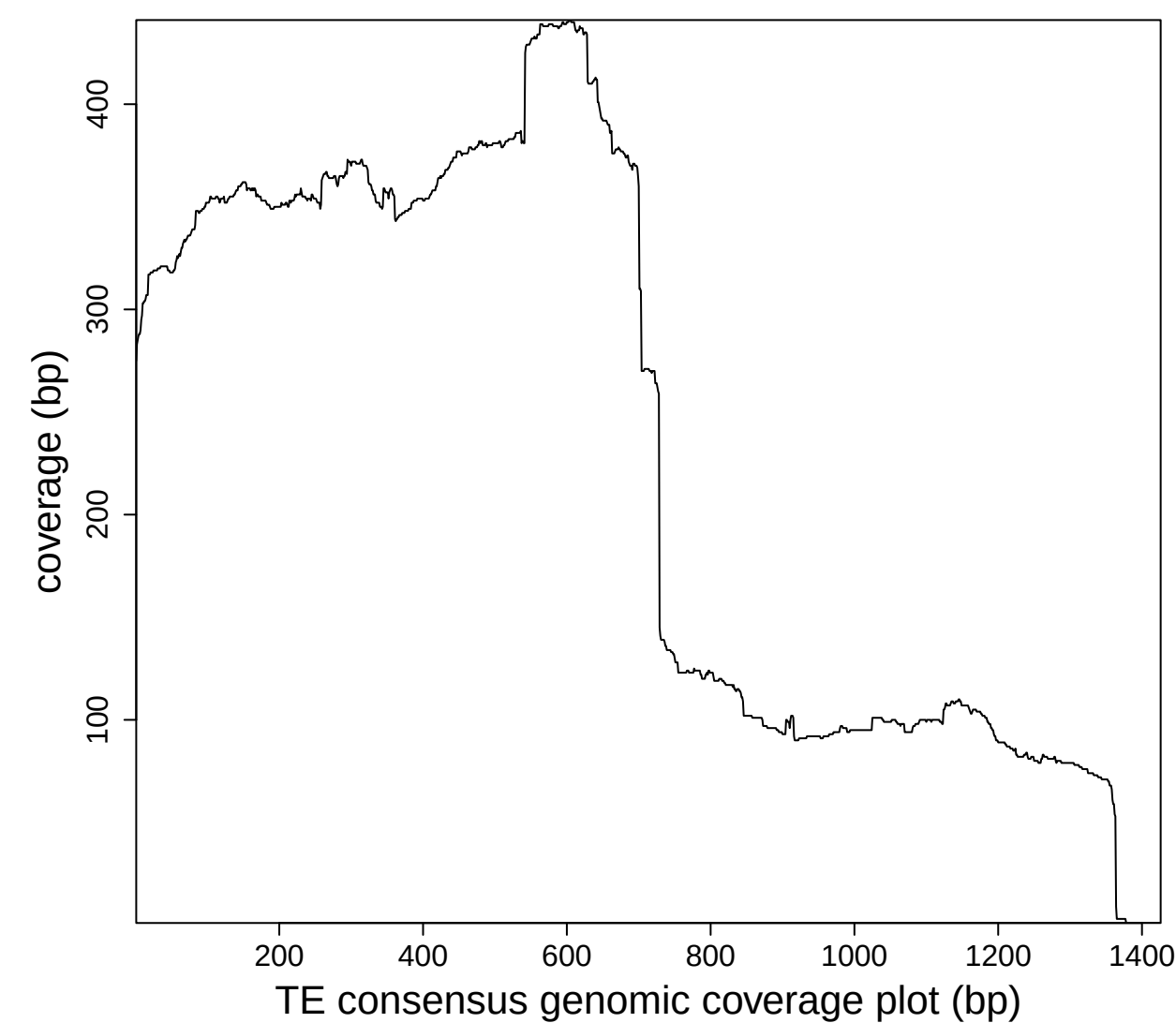
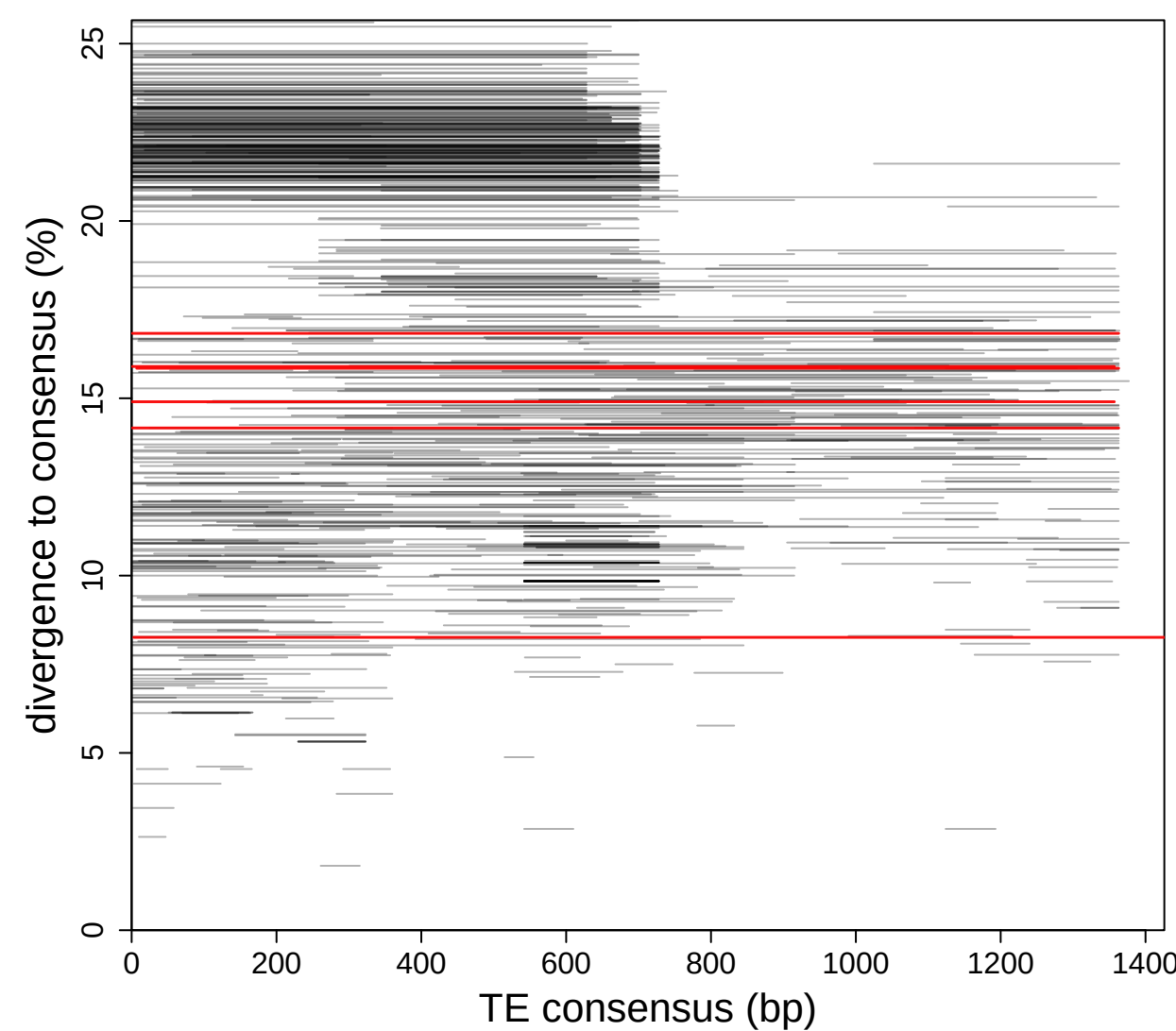


3.fasta.b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa
size: 2240bp; fragments: 427; full length: 0 (>=2016bp)

After TEtrimmer Extended plot Blue lines are boundaries

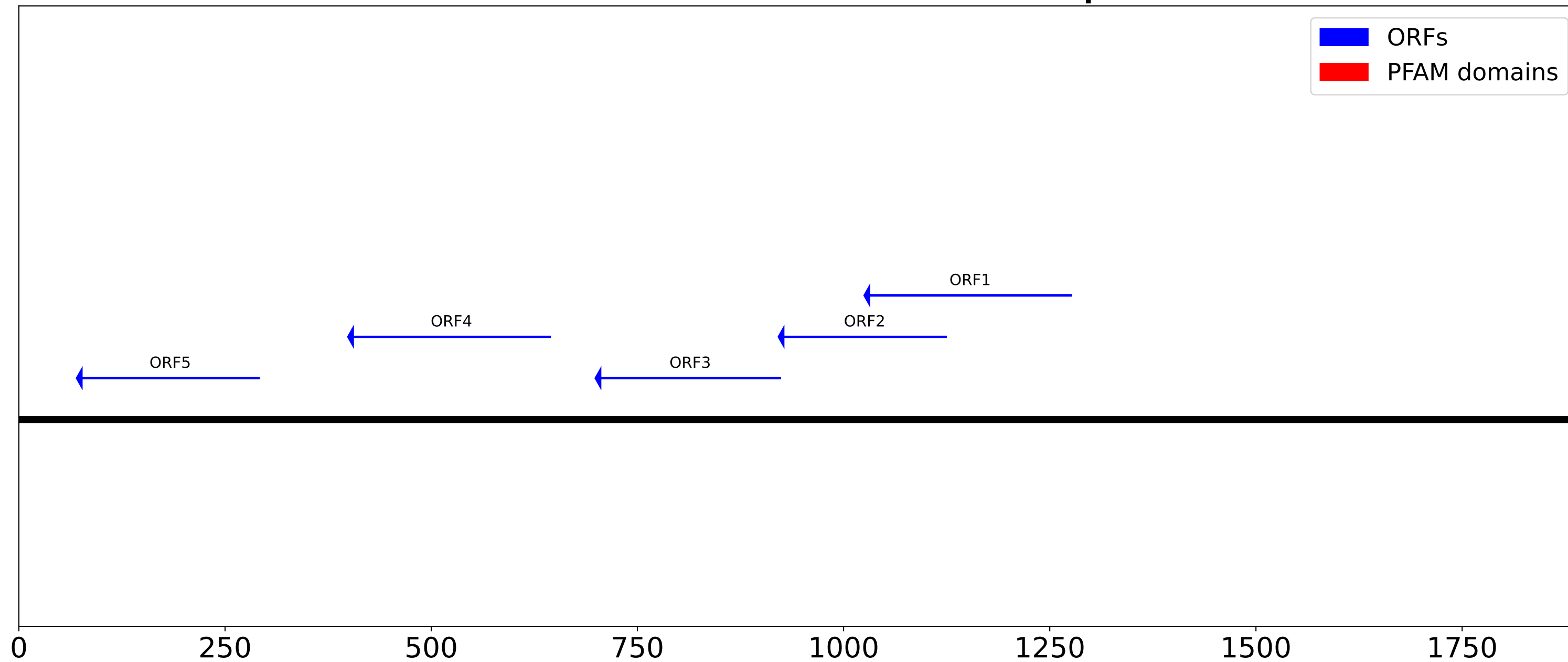


TE: rnd_4_family_273
size: 1426bp; fragments: 720; full length: 6 (≥ 1283.4 bp)

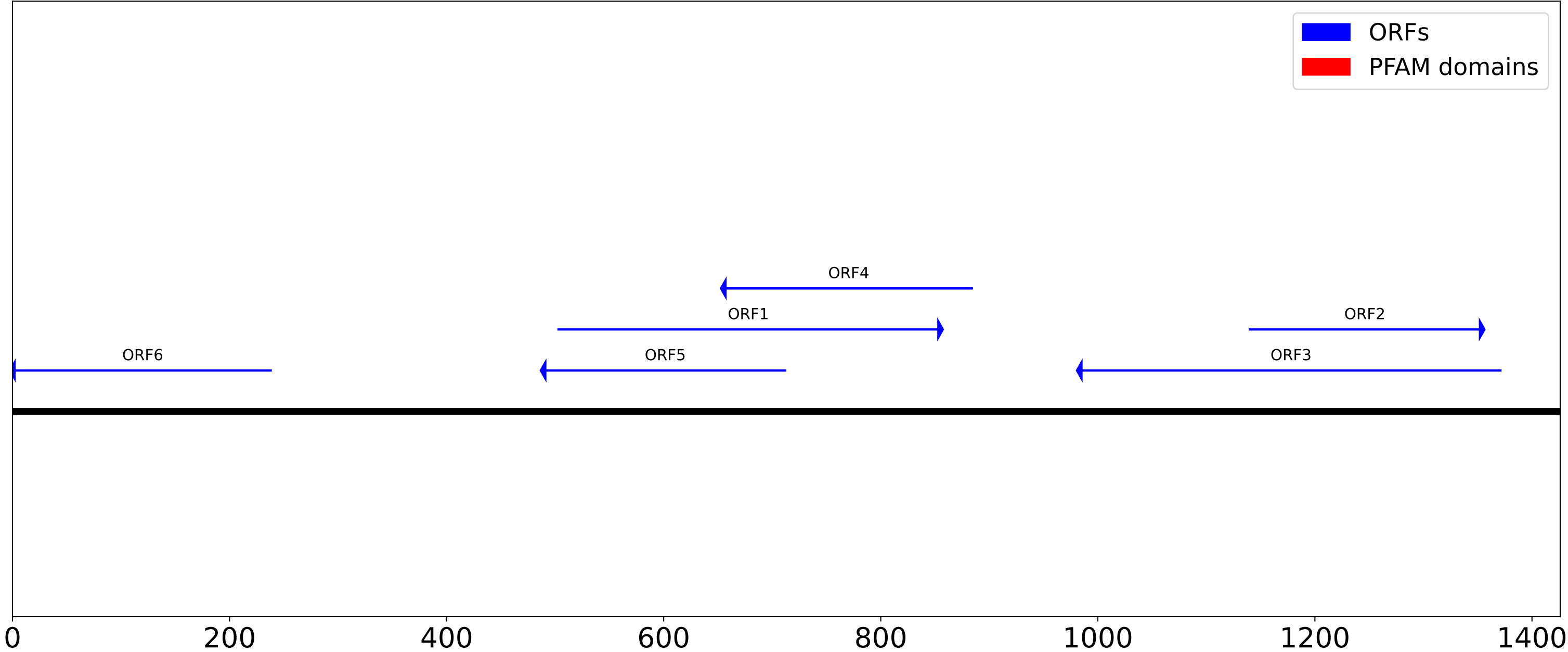


Before TETrimmer 1426 bp

After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

