

Start crop Point

End crop Point

MSA length = 2193



Start crop Point

End crop Point

0

10

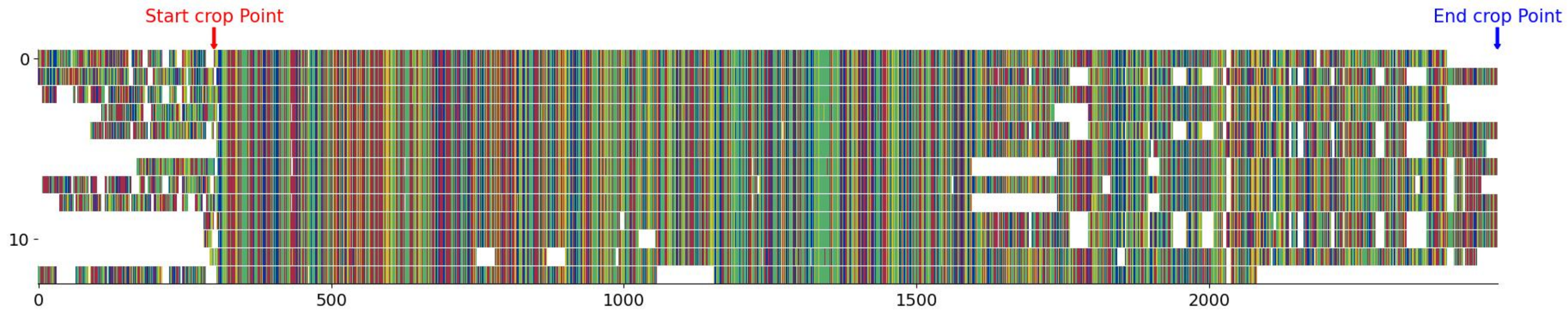
0

500

1000

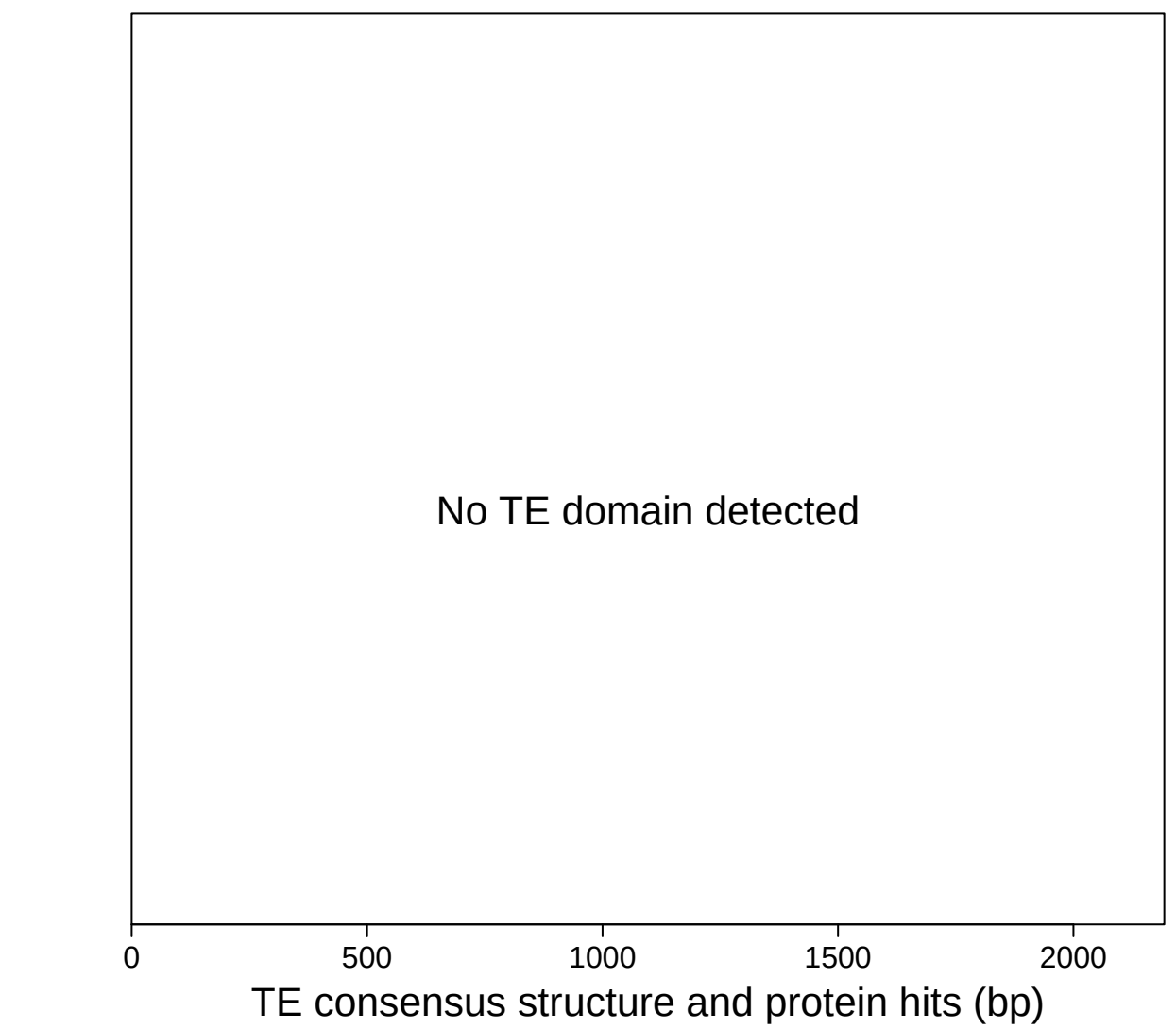
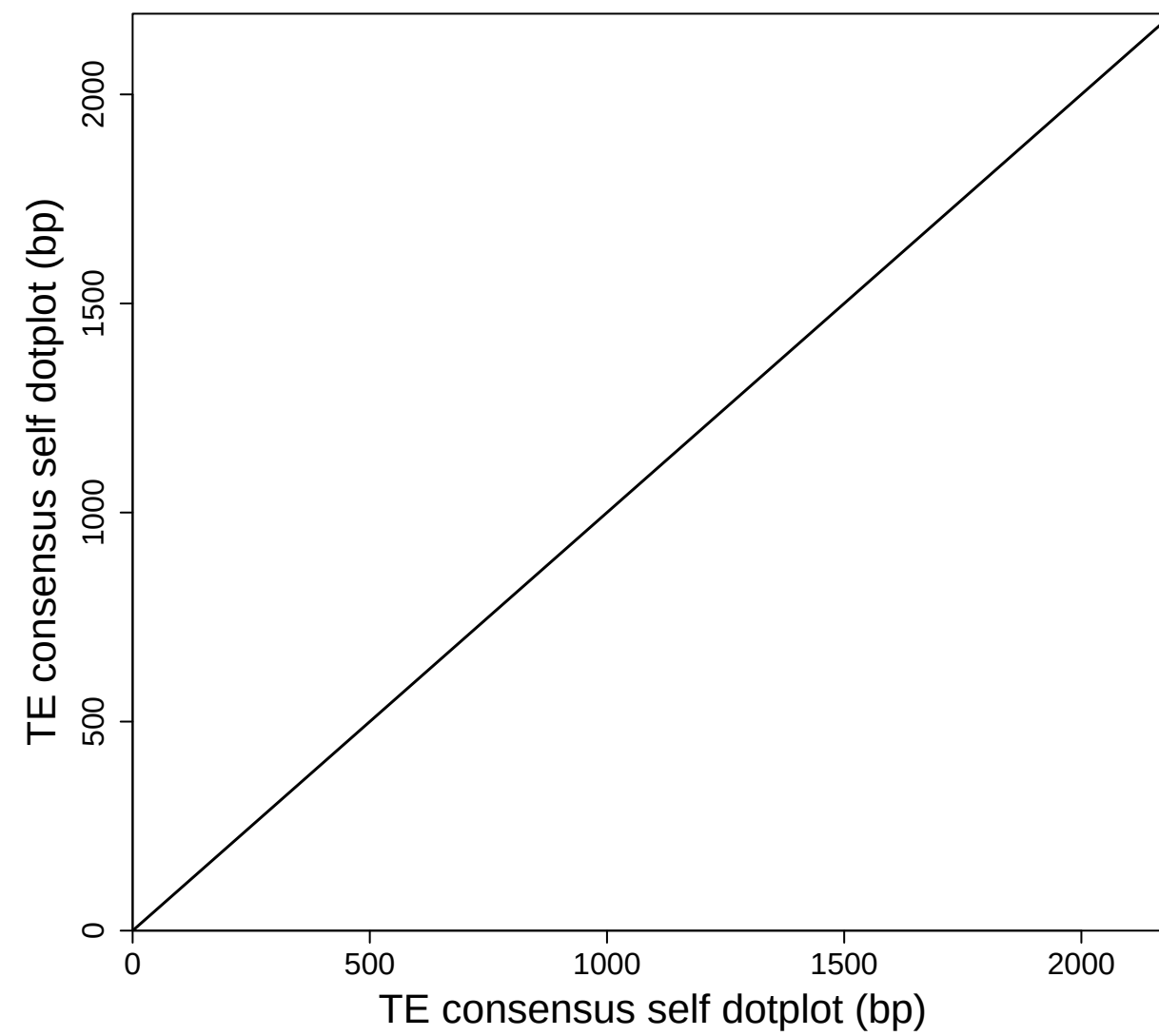
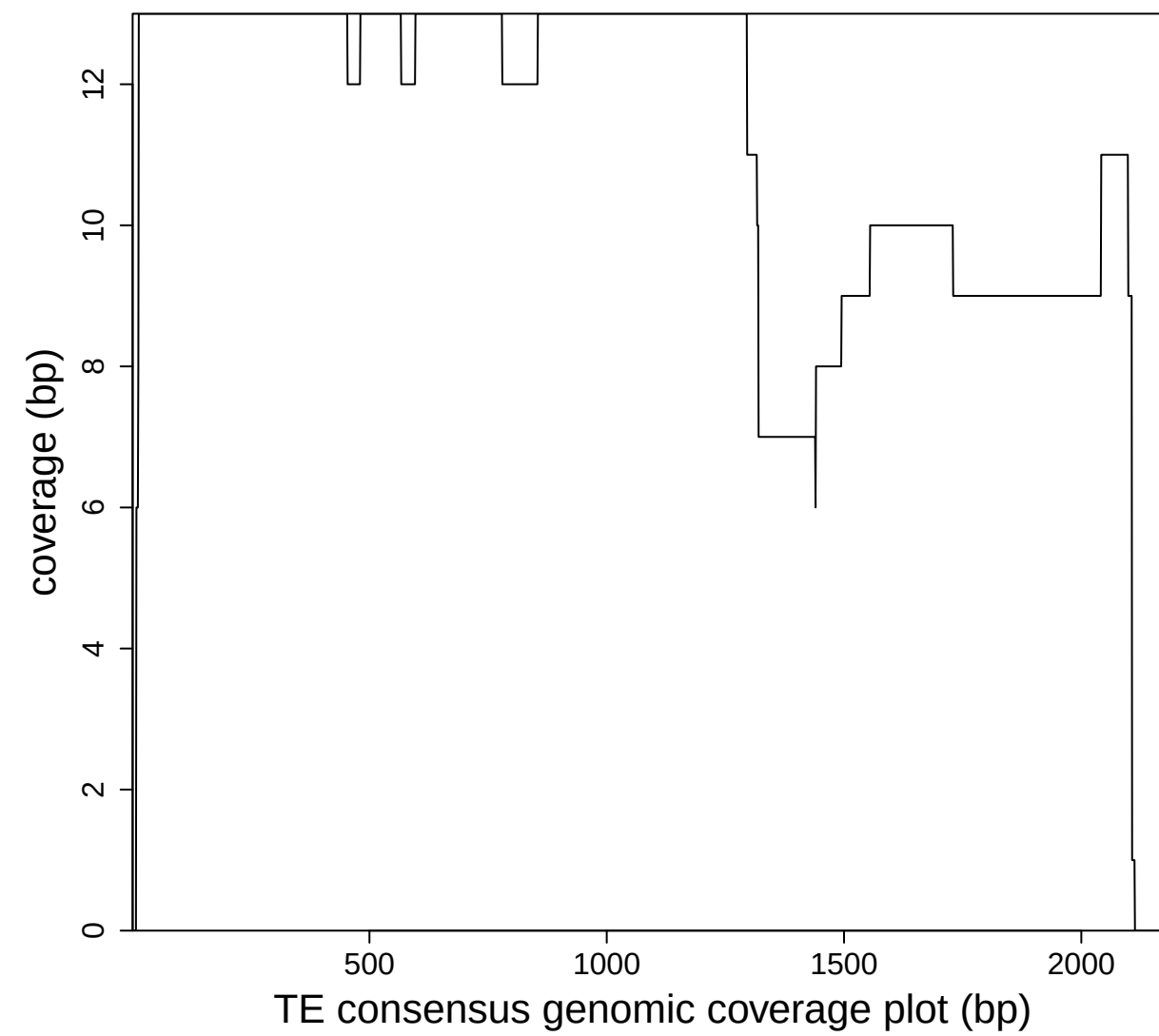
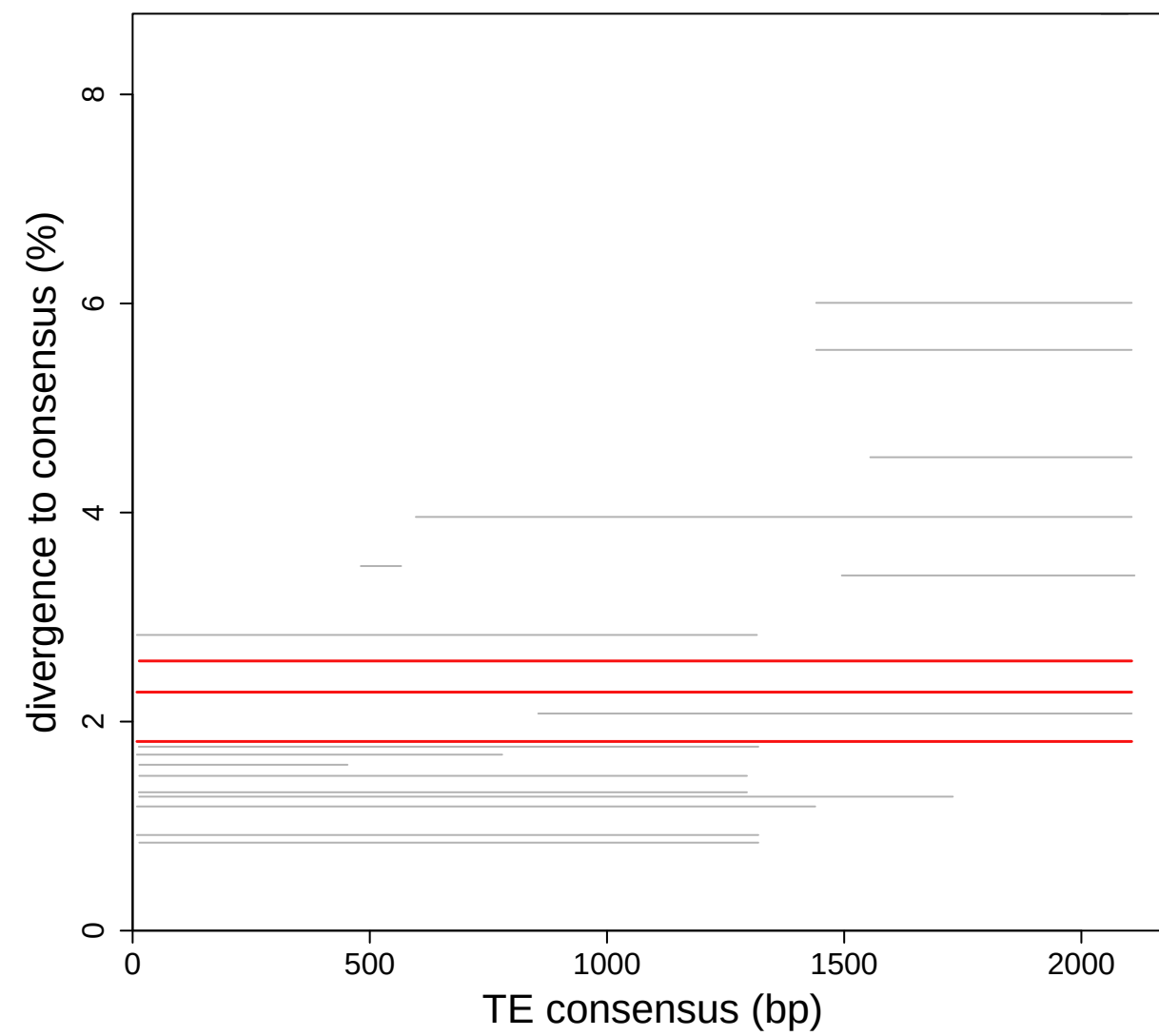
1500

2000



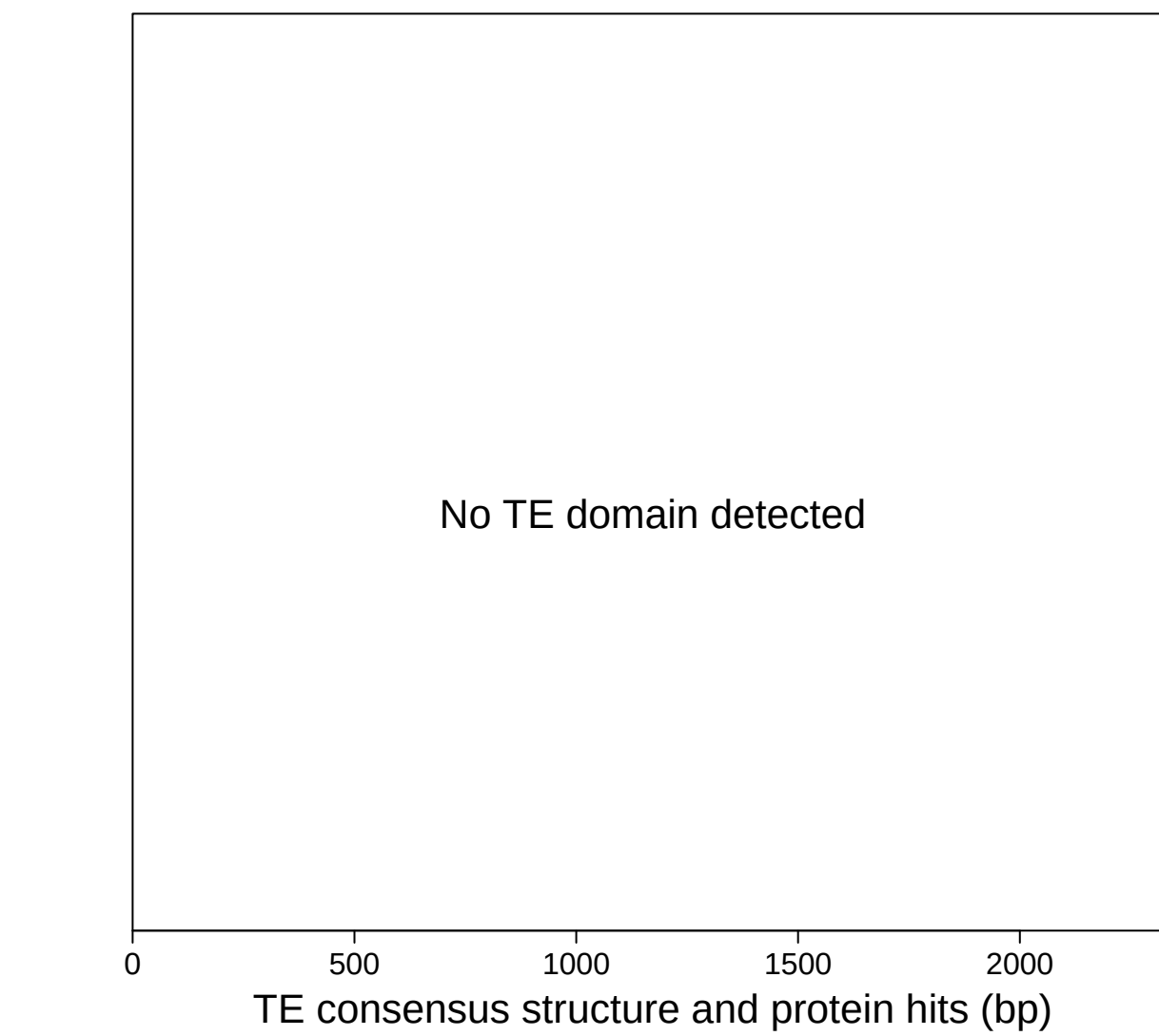
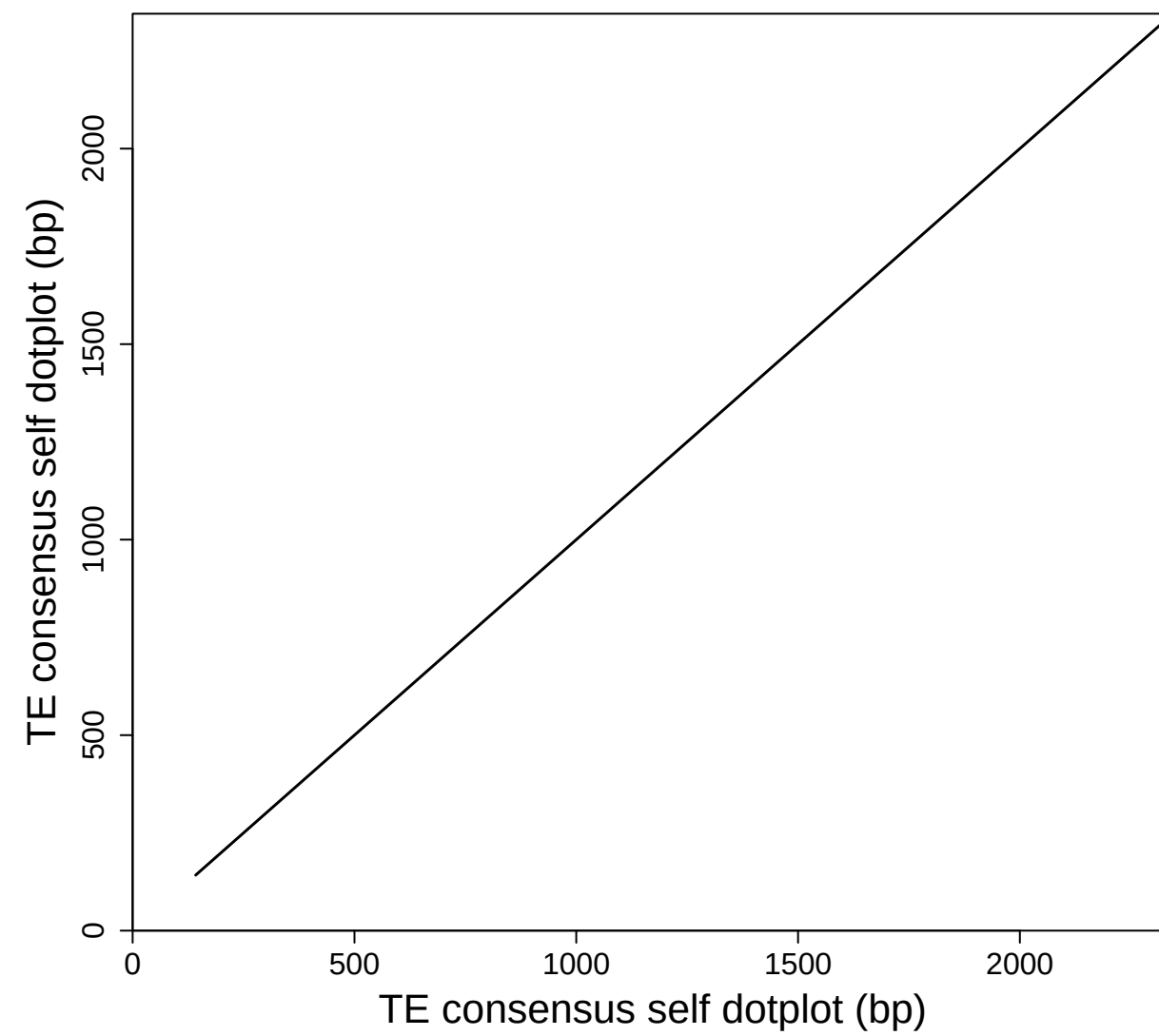
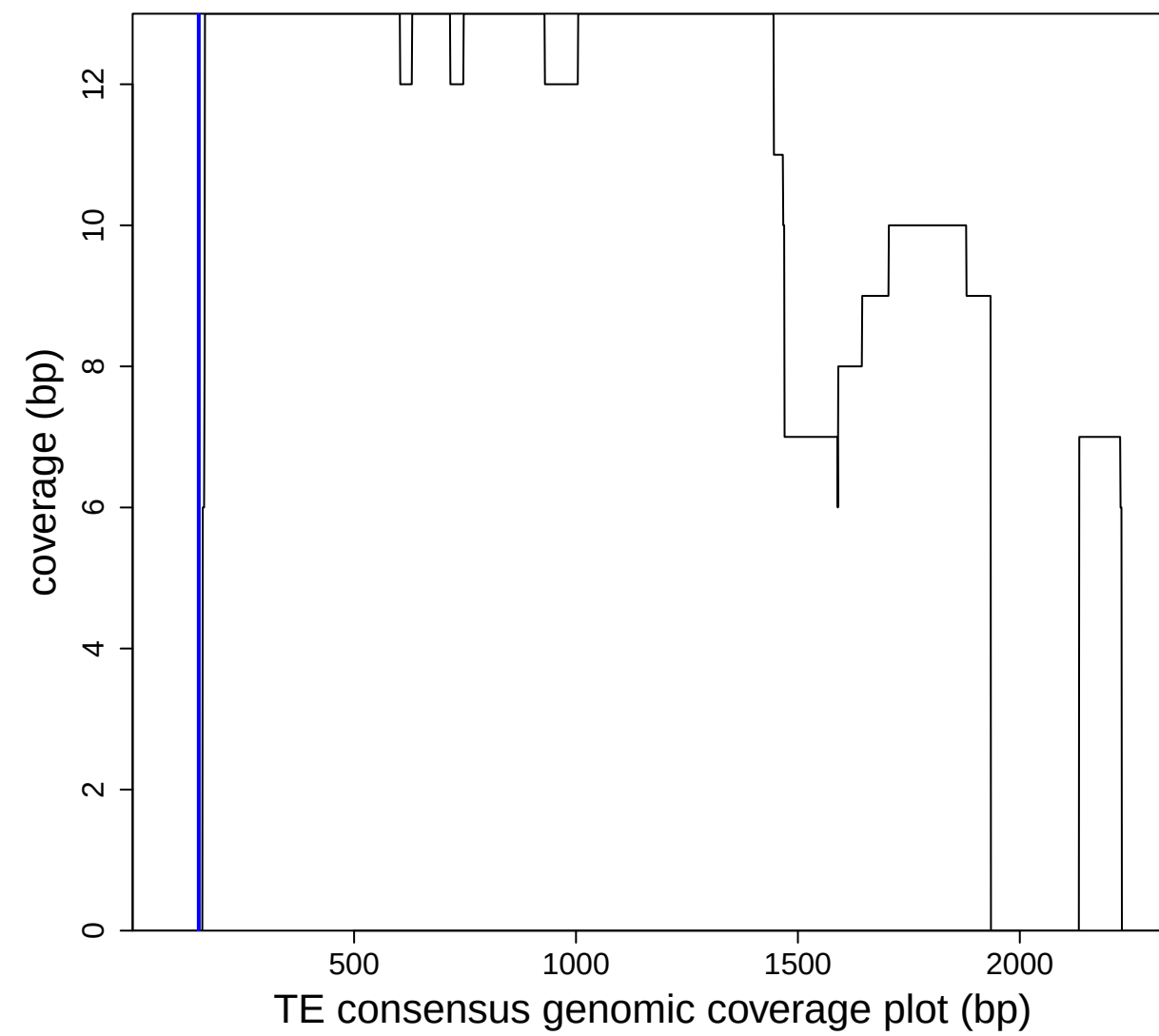
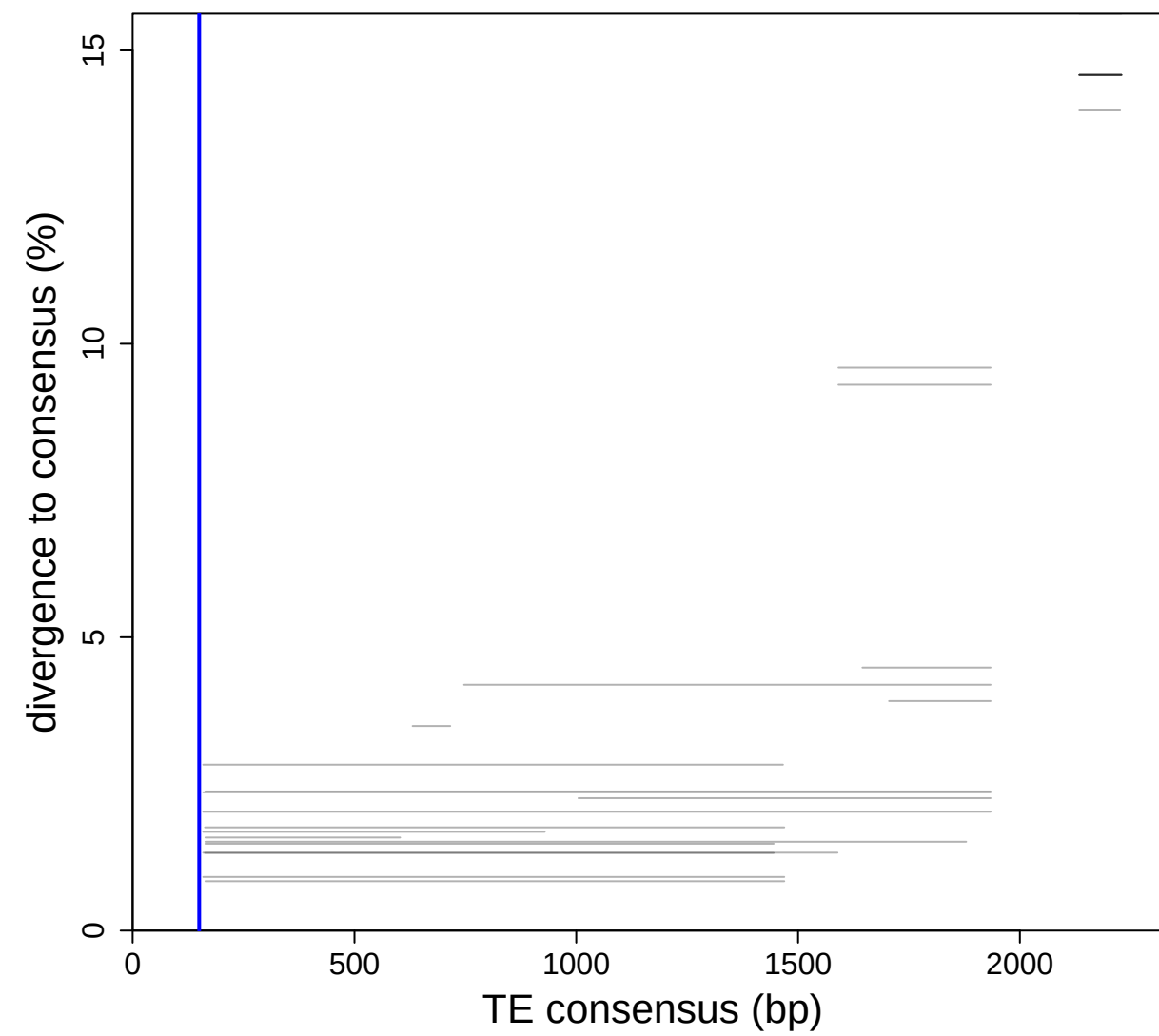
a.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 2193bp; fragments: 22; full length: 3 (>=1973.7bp)

After TEtrimmer 2193 bp



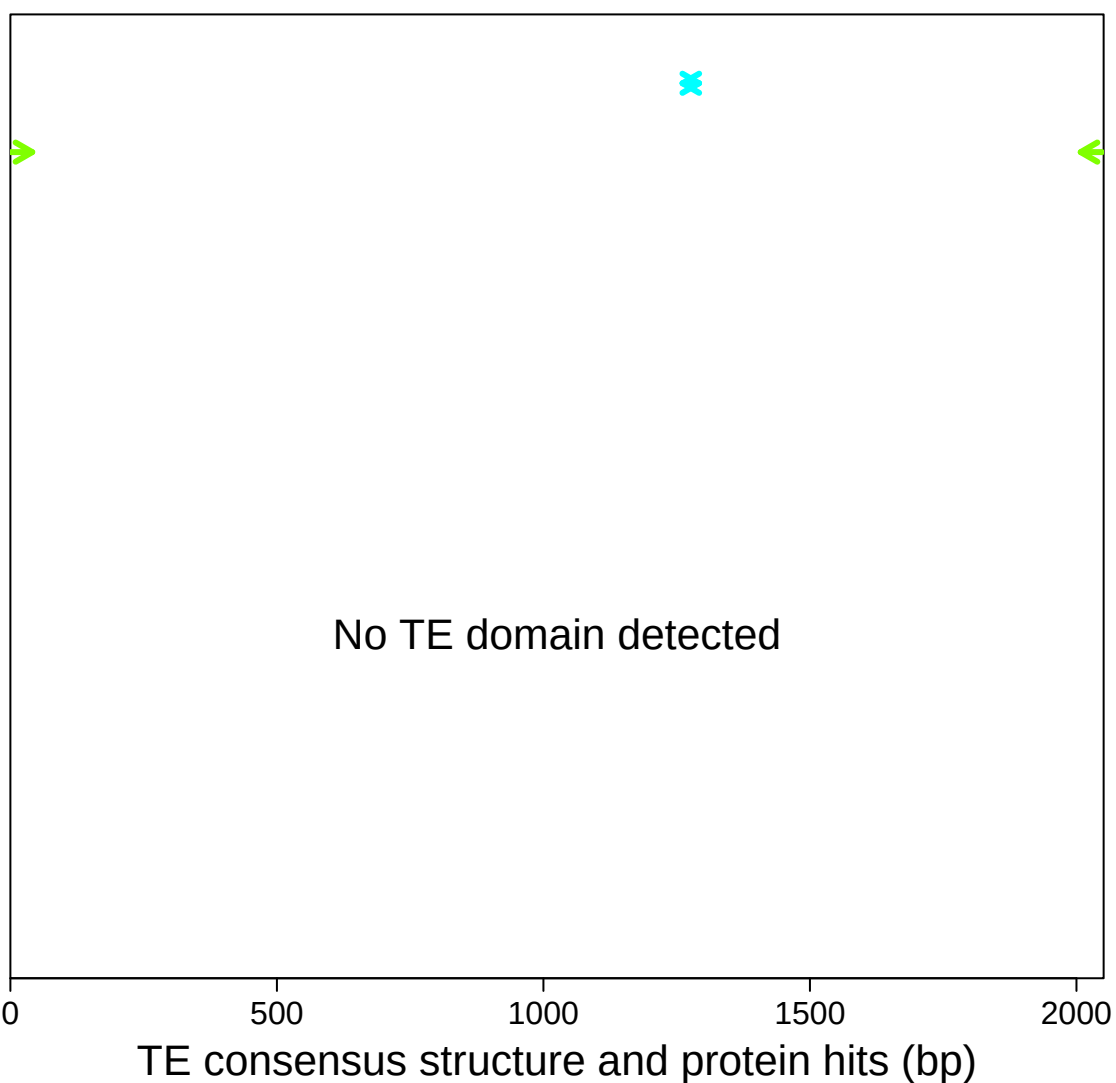
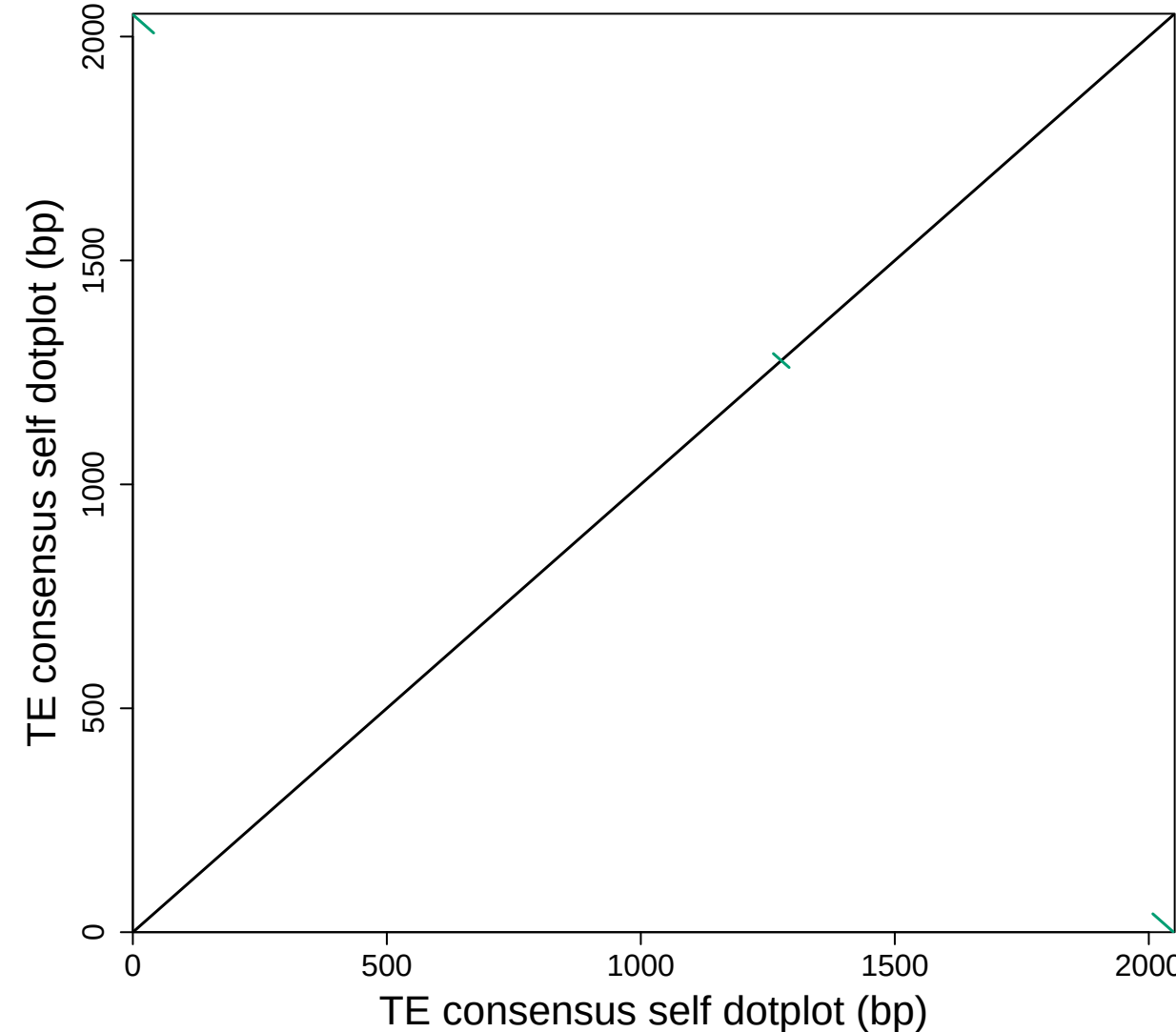
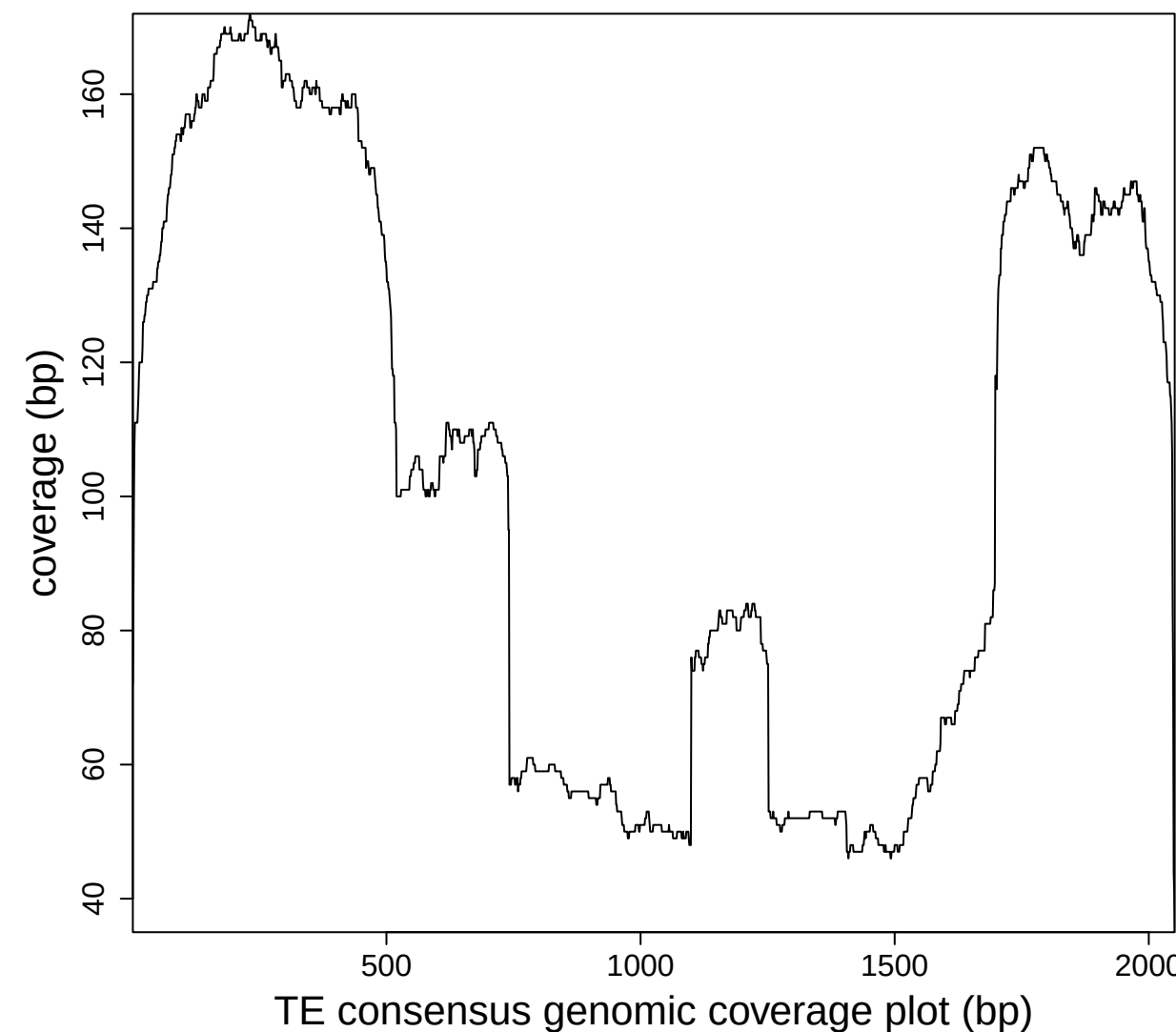
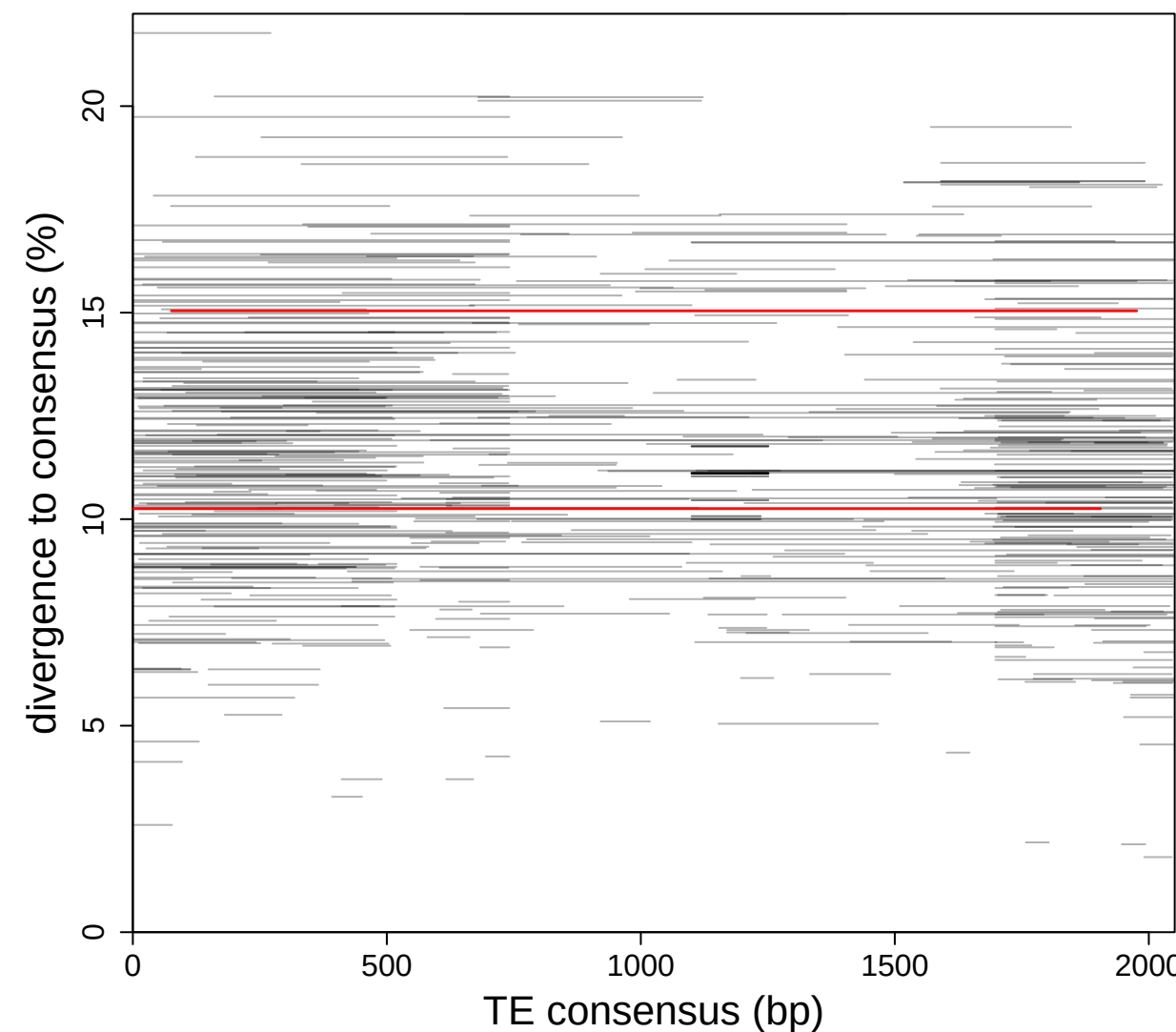
5.fasta.b.bed uf.bed g_1.bed fm_1.bed 0_0_bcln.fa aln.fa cl.f
size: 2345bp; fragments: 27; full length: 0 (>=2110.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

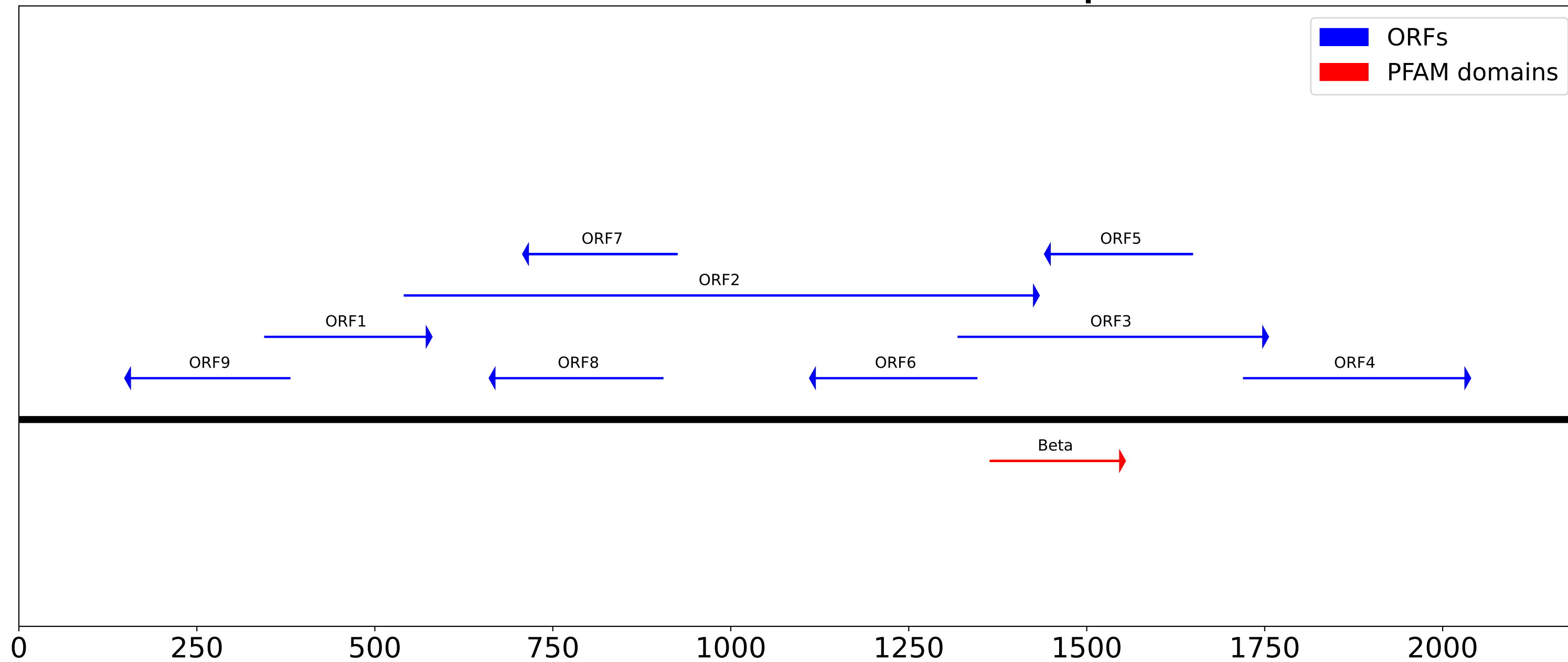


TE: rnd_5_family_1135
size: 2051bp; fragments: 506; full length: 2 (≥ 1845.9 bp)

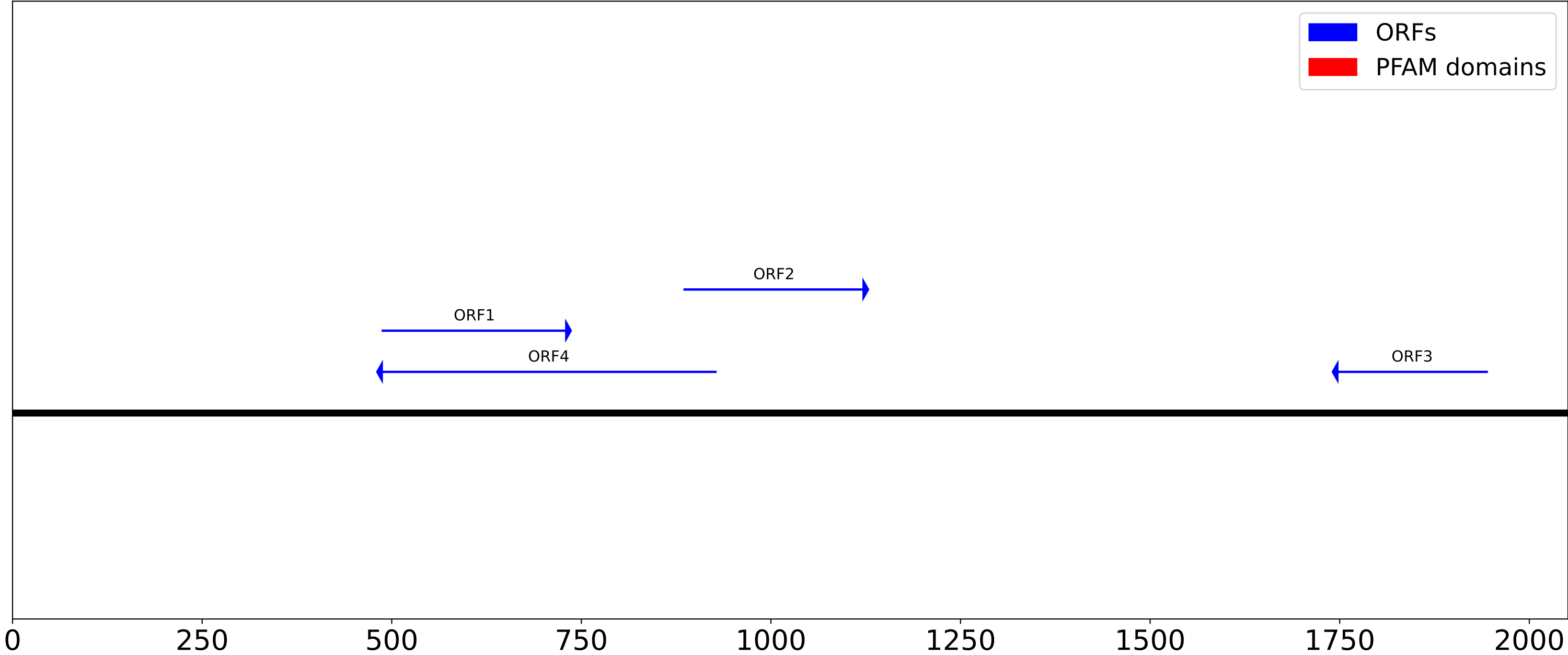
Before TEtrimmer 2051 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

