

Start crop Point



End crop Point



MSA length = 3528



Start crop Point

End crop Point

0

10

0

500

1000

1500

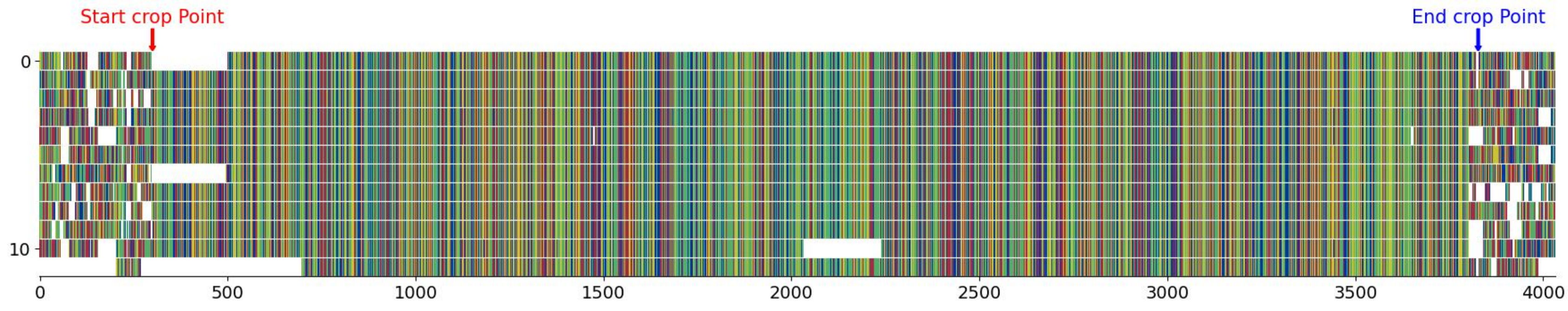
2000

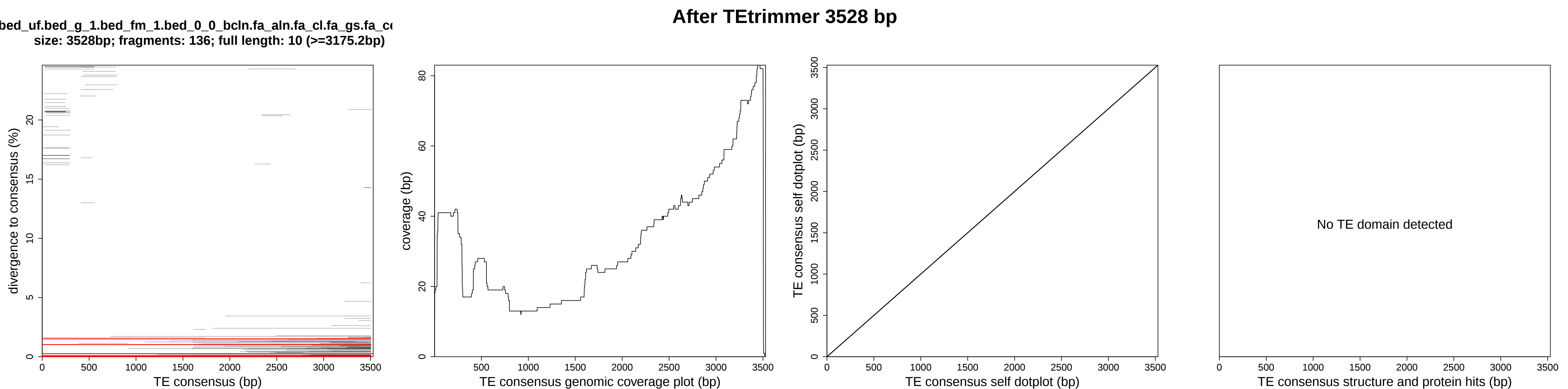
2500

3000

3500

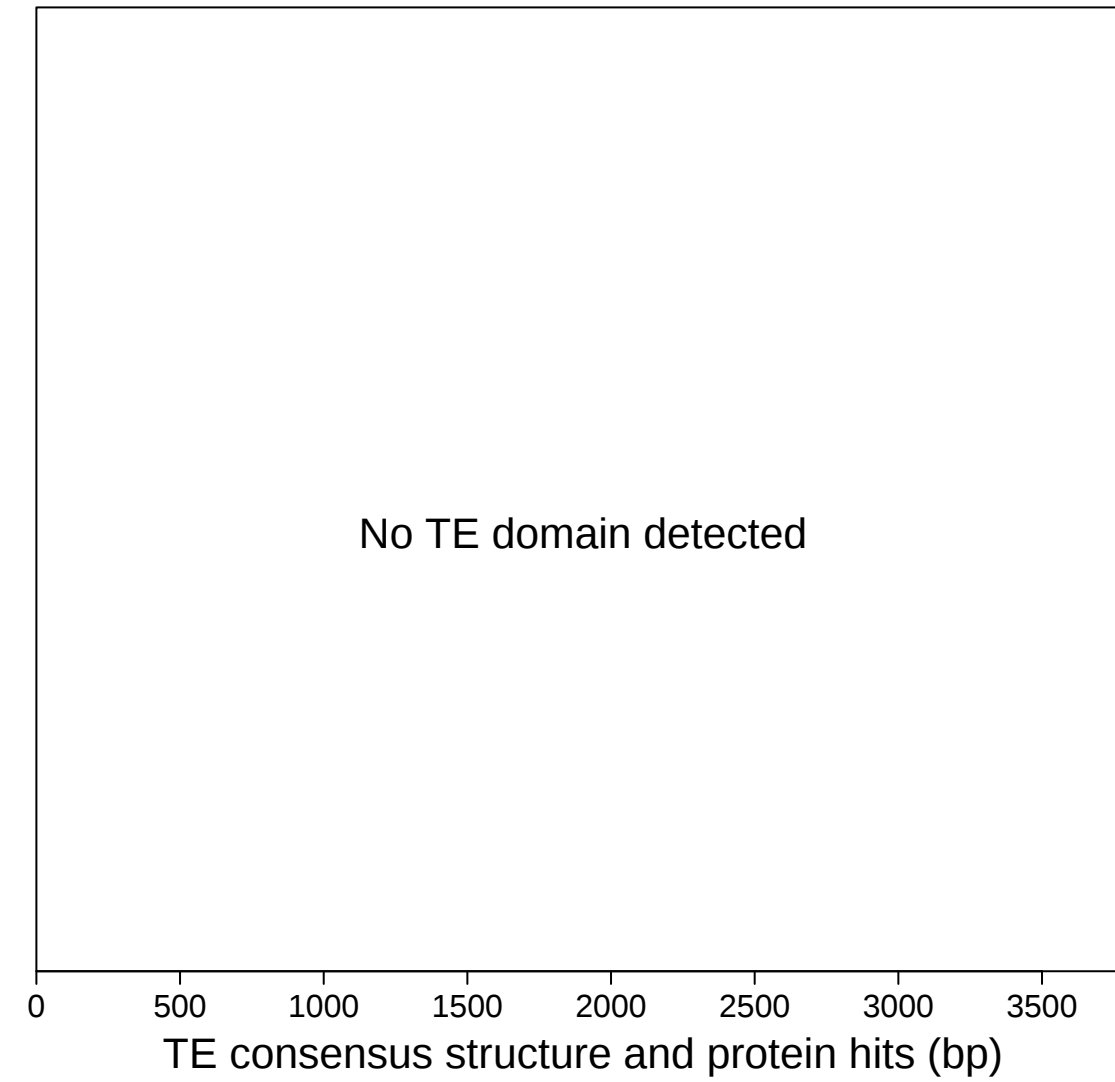
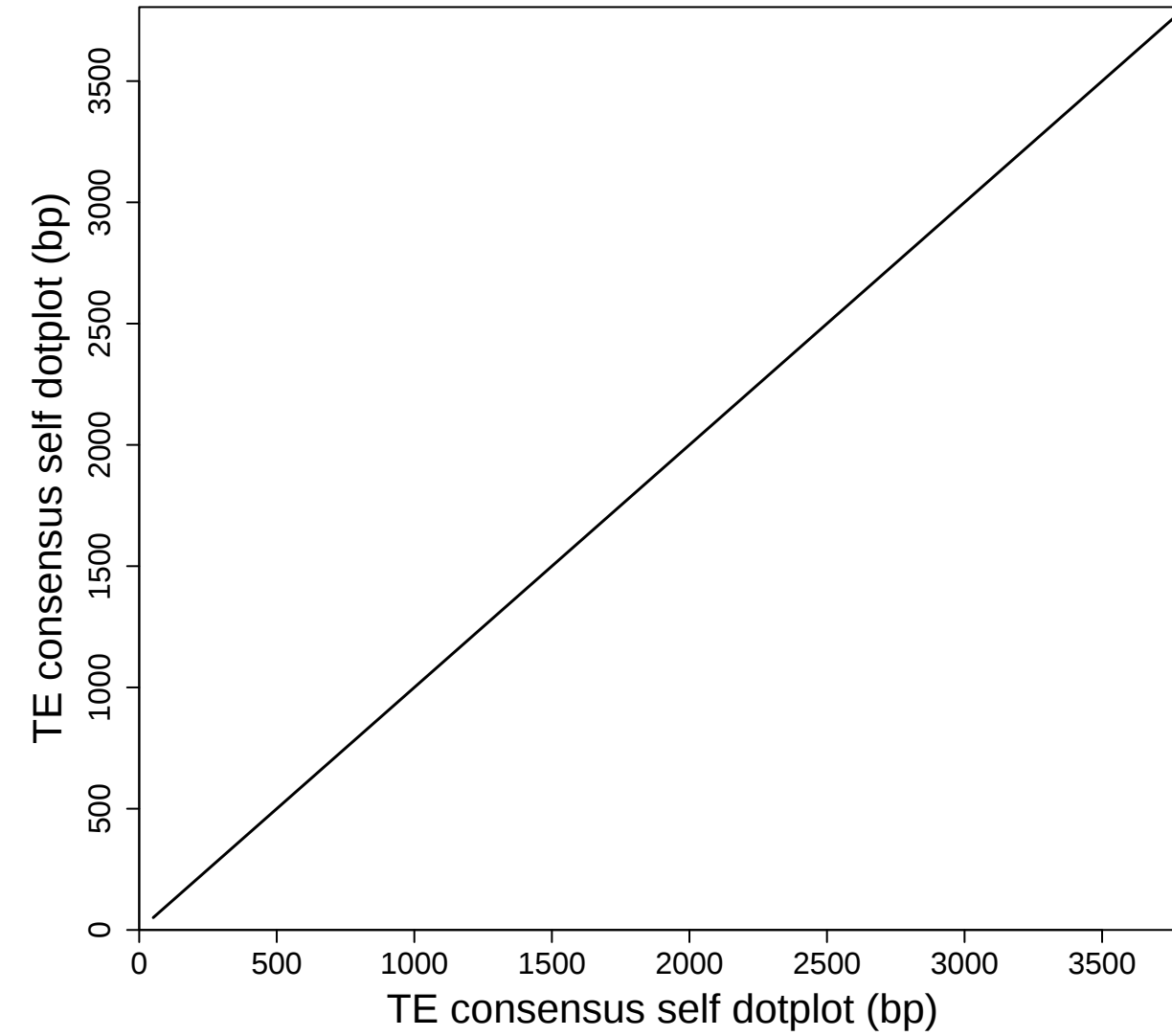
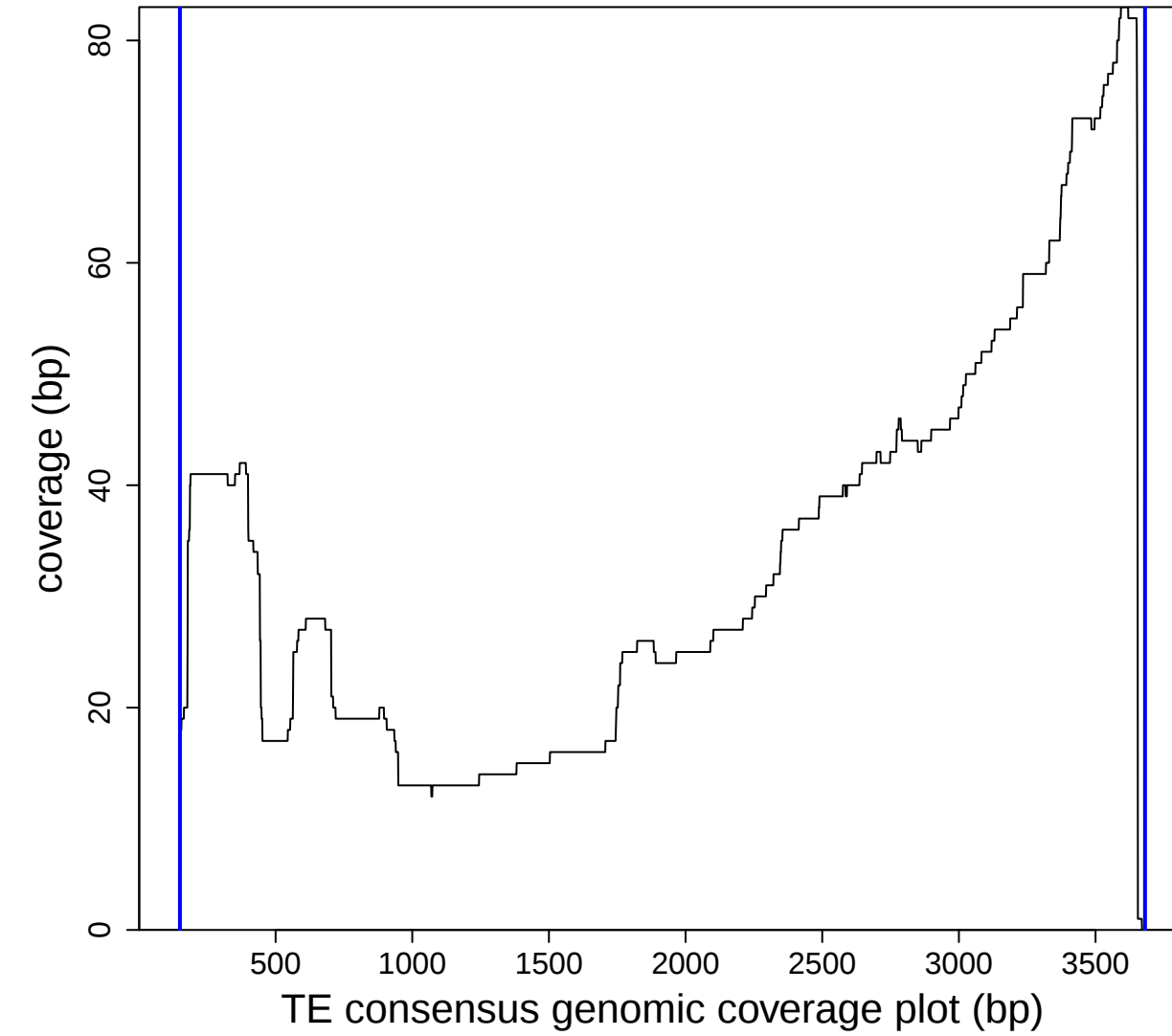
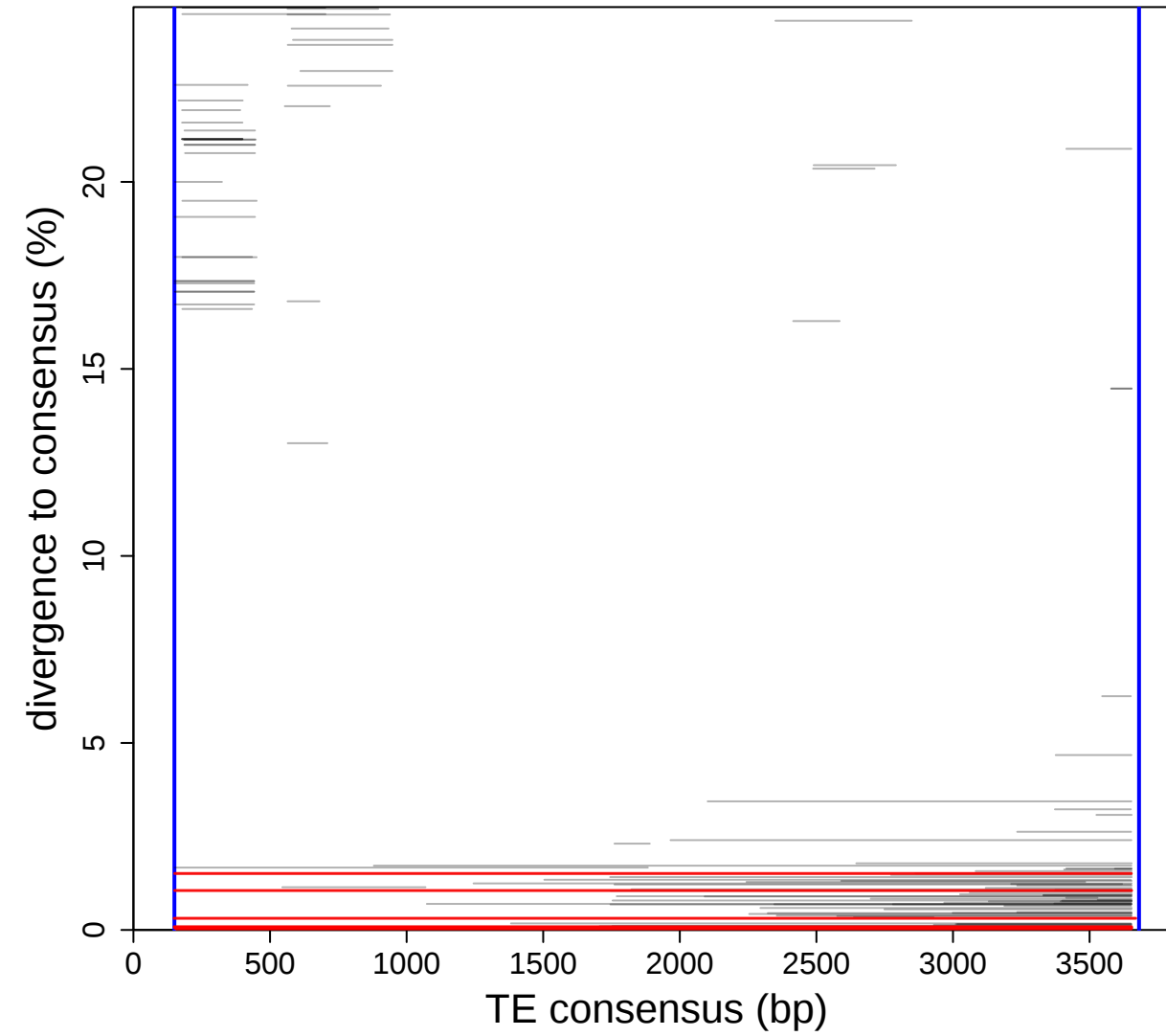
4000



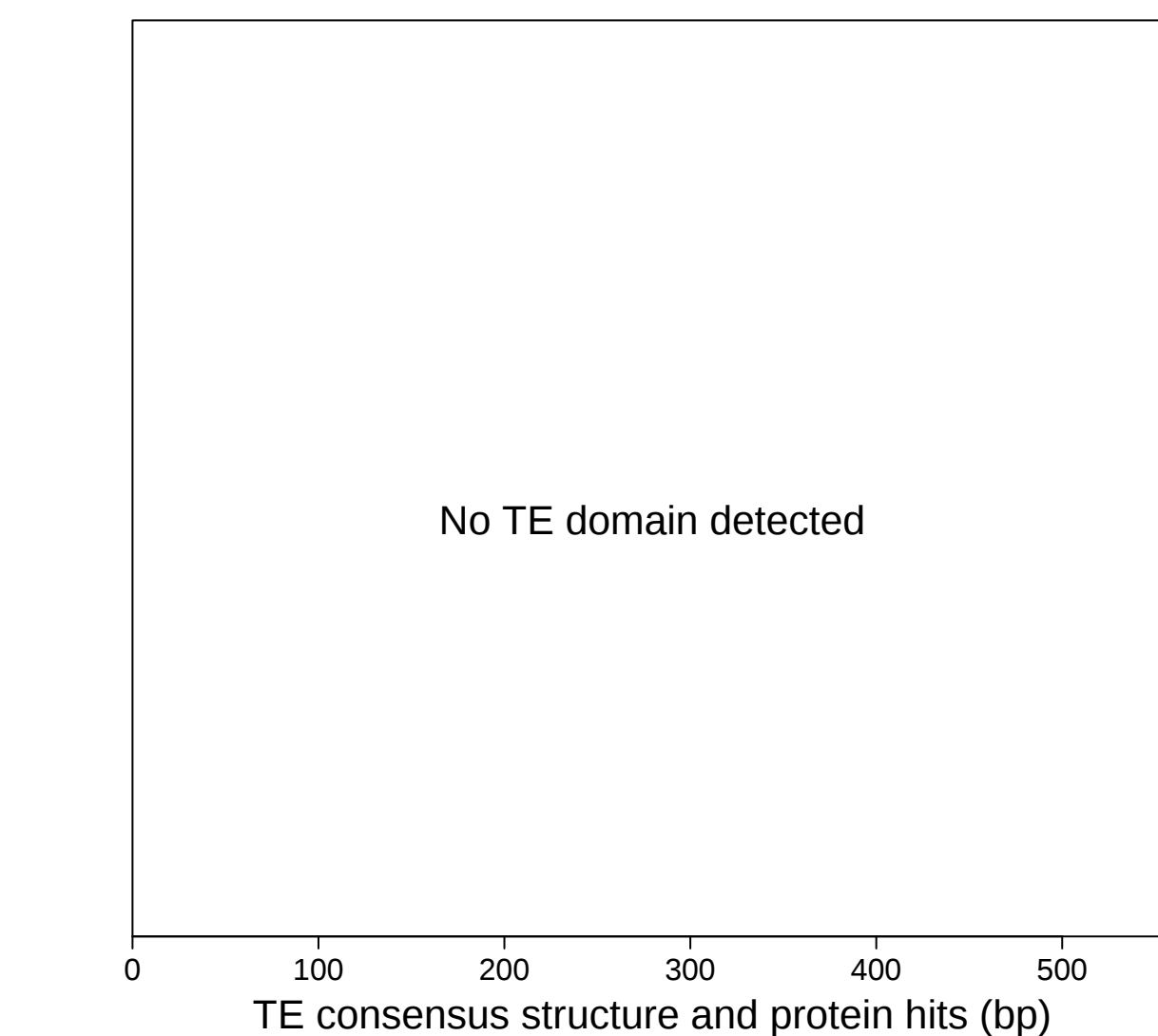
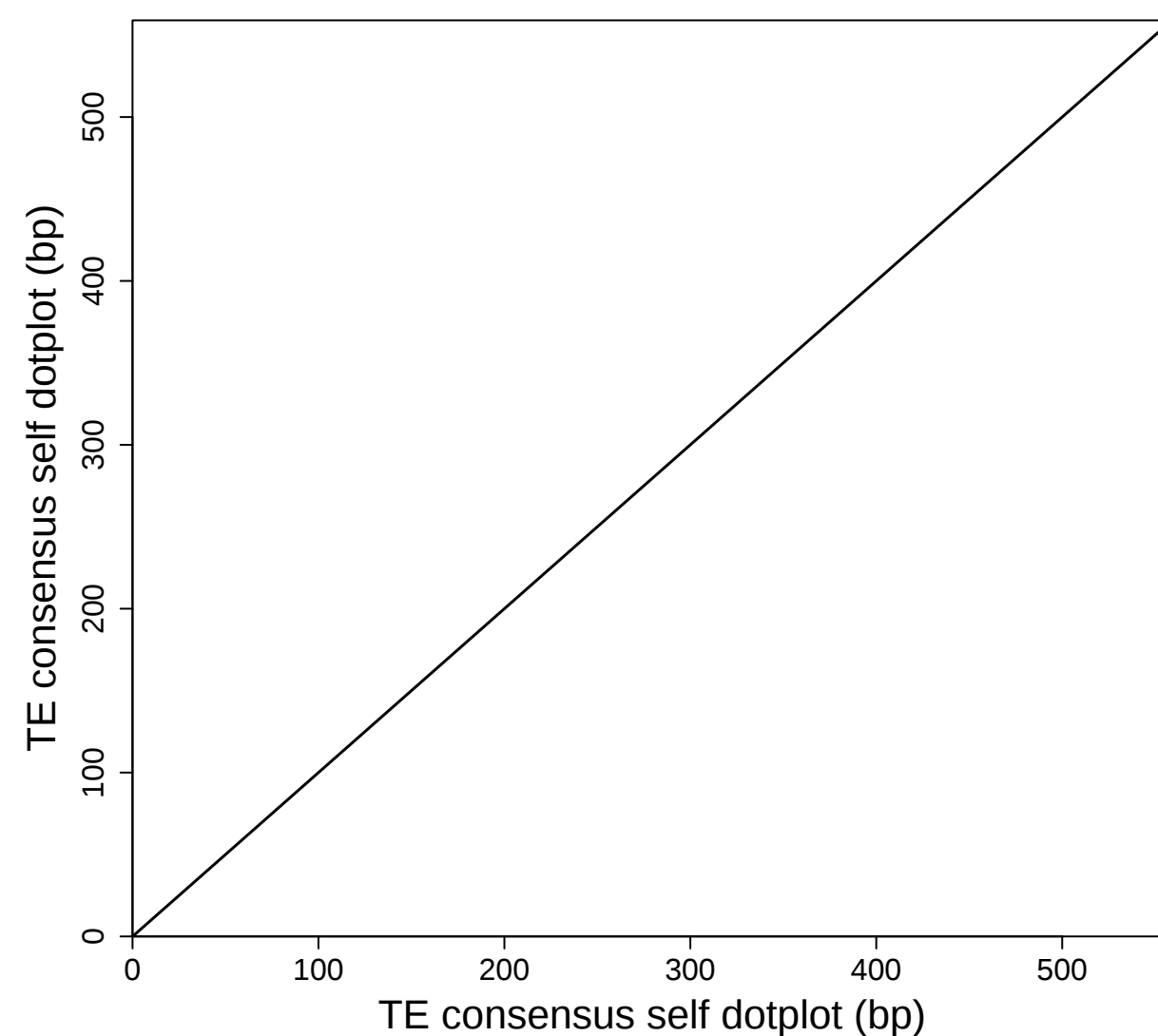
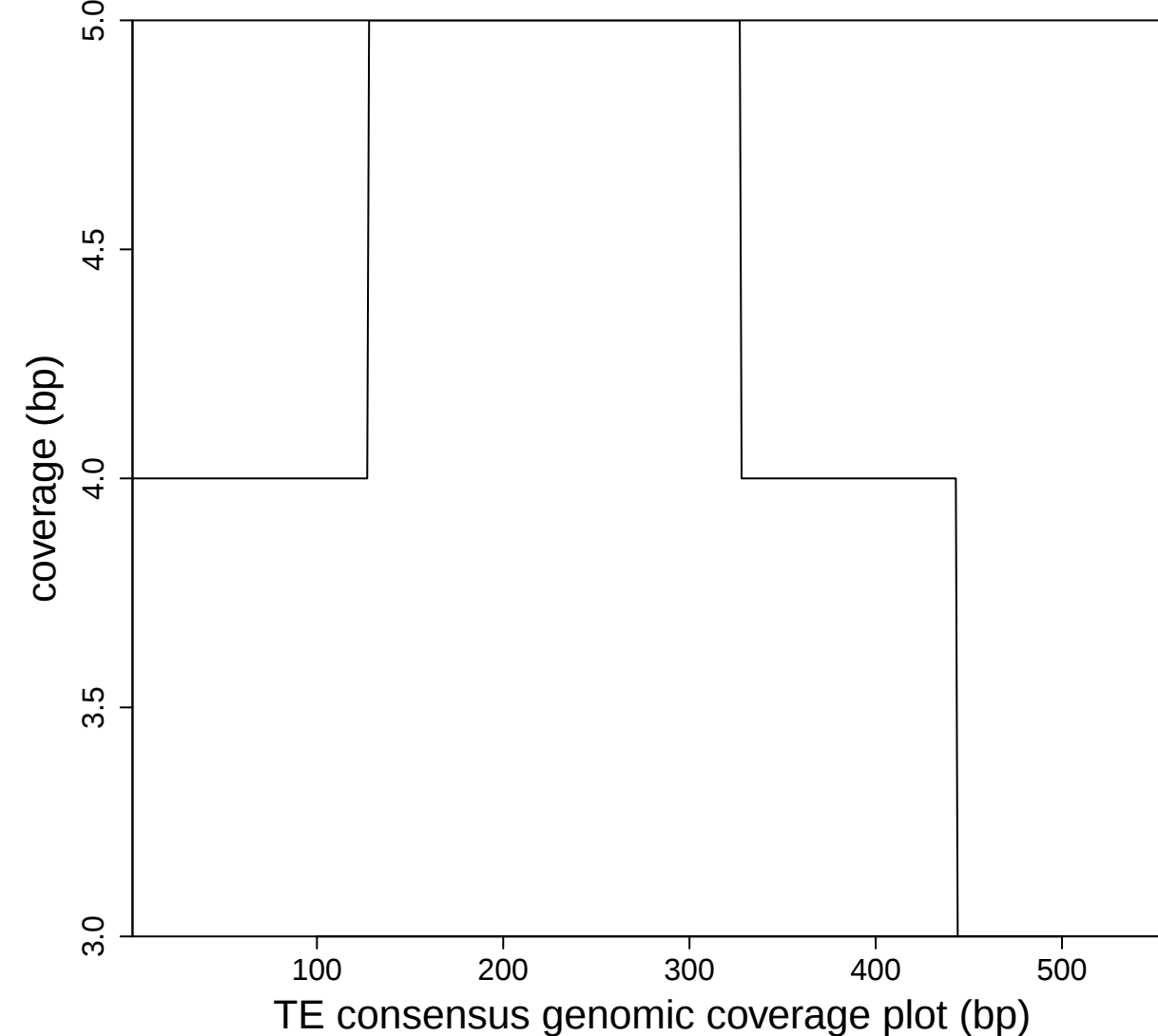
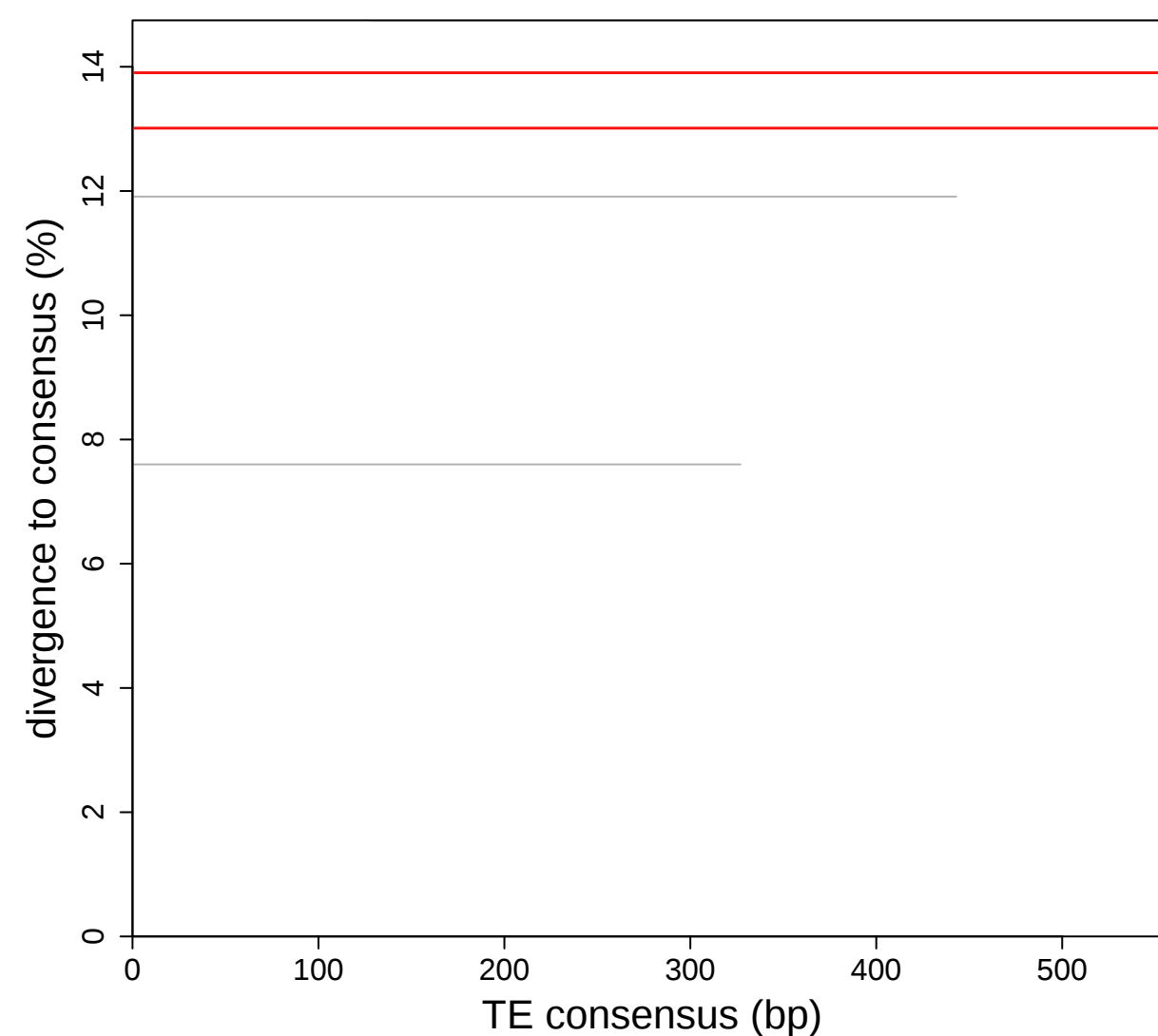


2.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcIn.fa_aln.fa_cl.f
size: 3831bp; fragments: 136; full length: 8 (>=3447.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

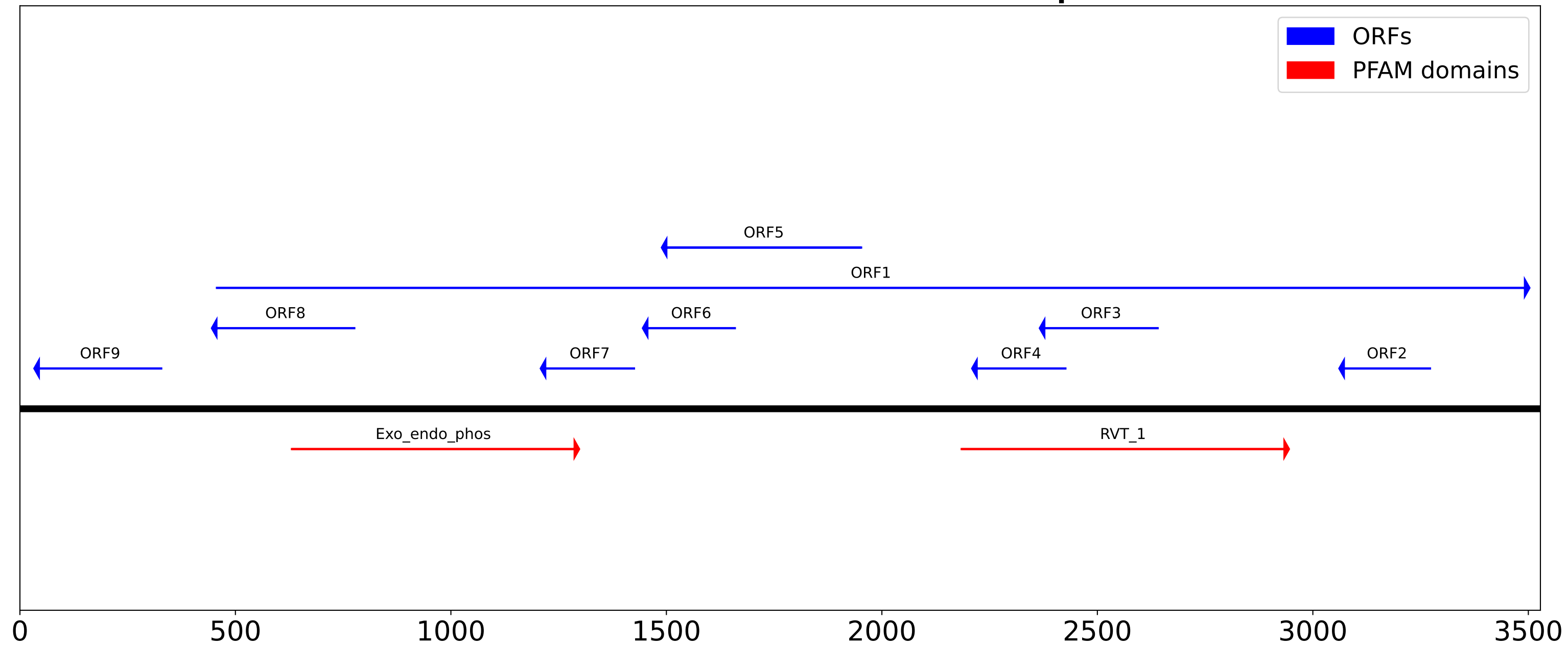


TE: rnd_5_family_1332
size: 559bp; fragments: 5; full length: 2 (≥ 503.1 bp)

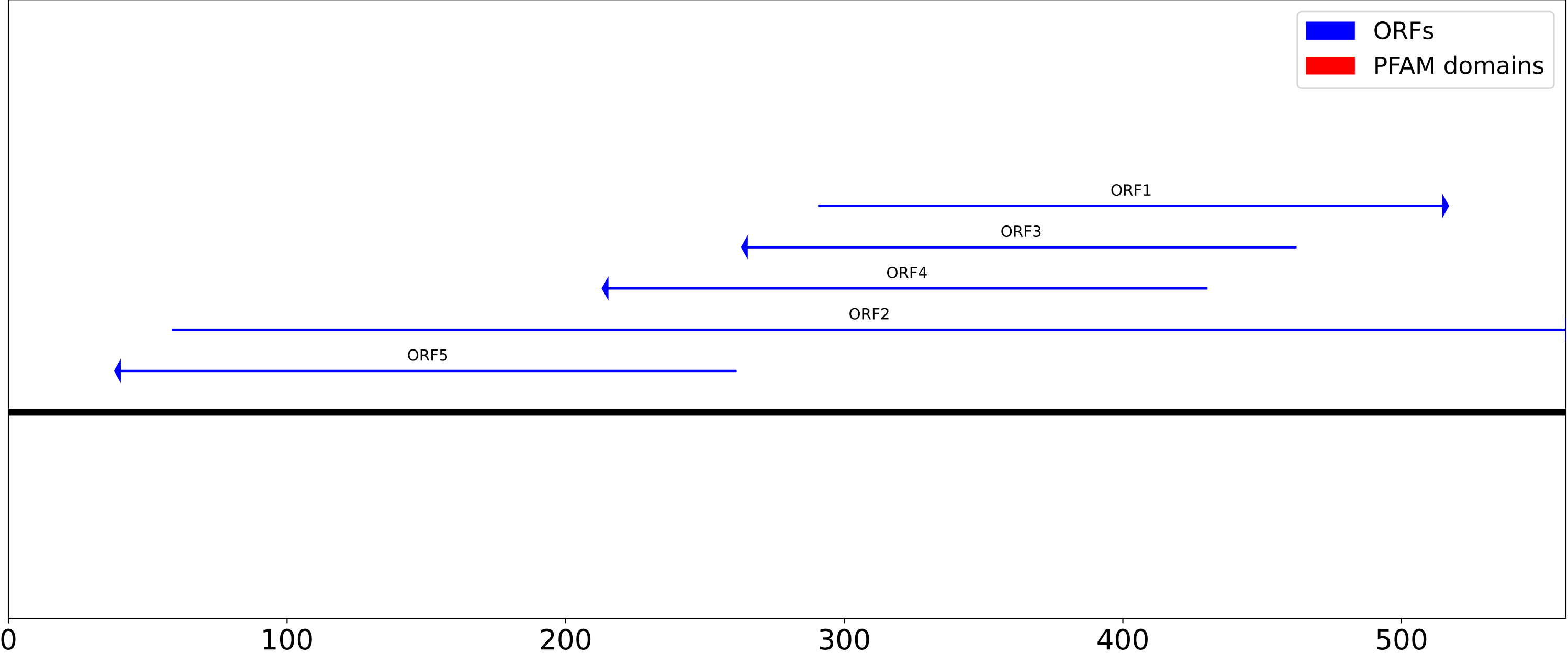


Before TEtrimmer 559 bp

After TEtrimmer ORF and PFAM domain plot



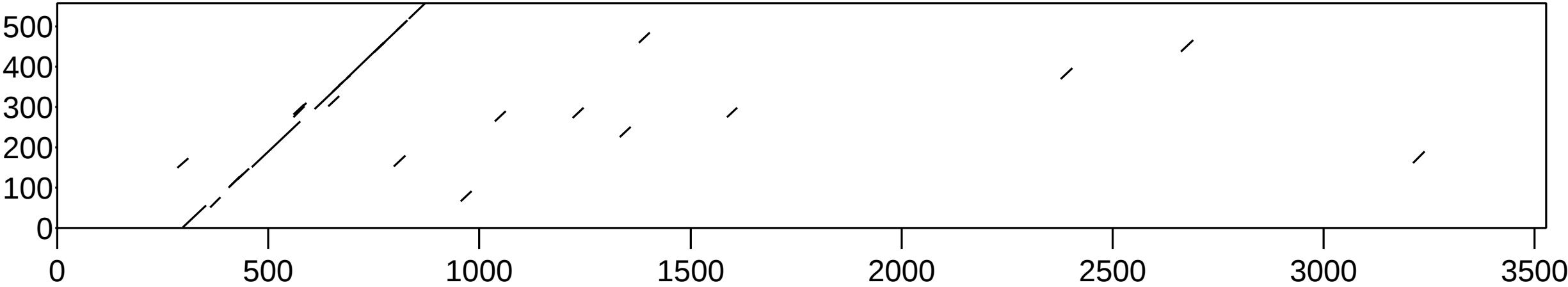
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)