

Start crop Point

End crop Point

0

10

0

500

1000

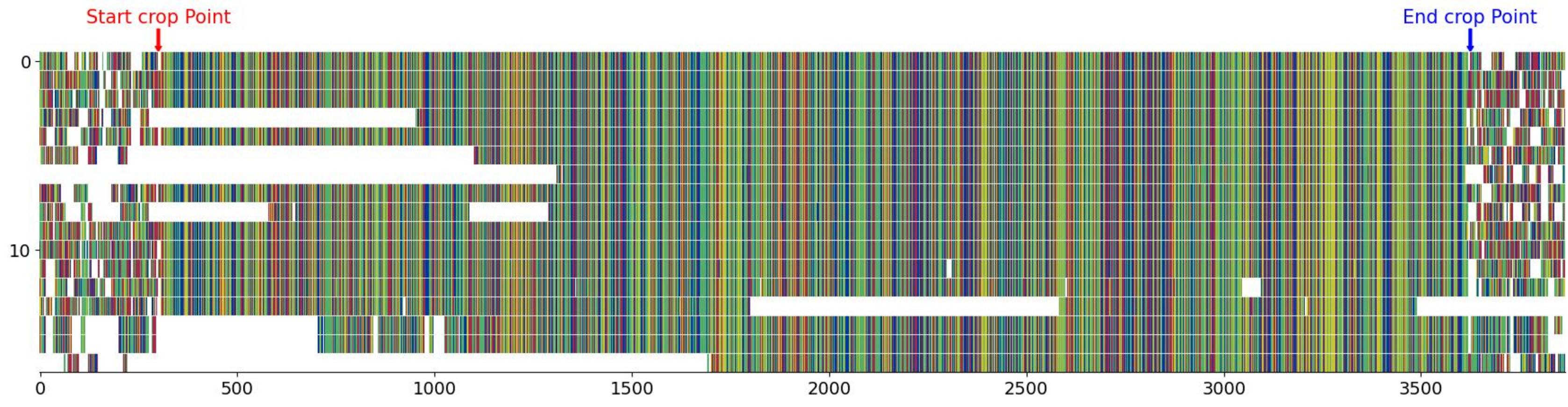
1500

2000

2500

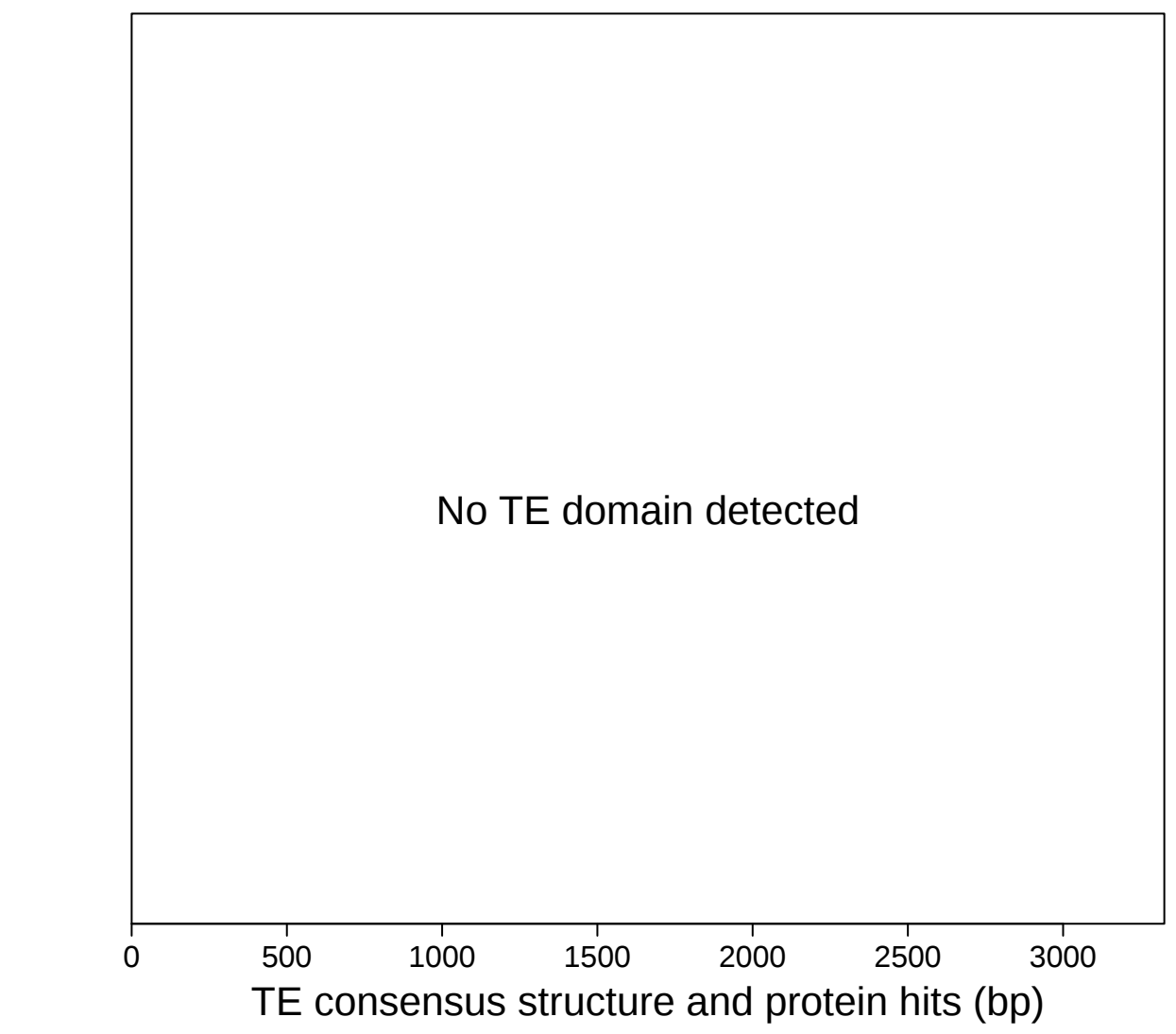
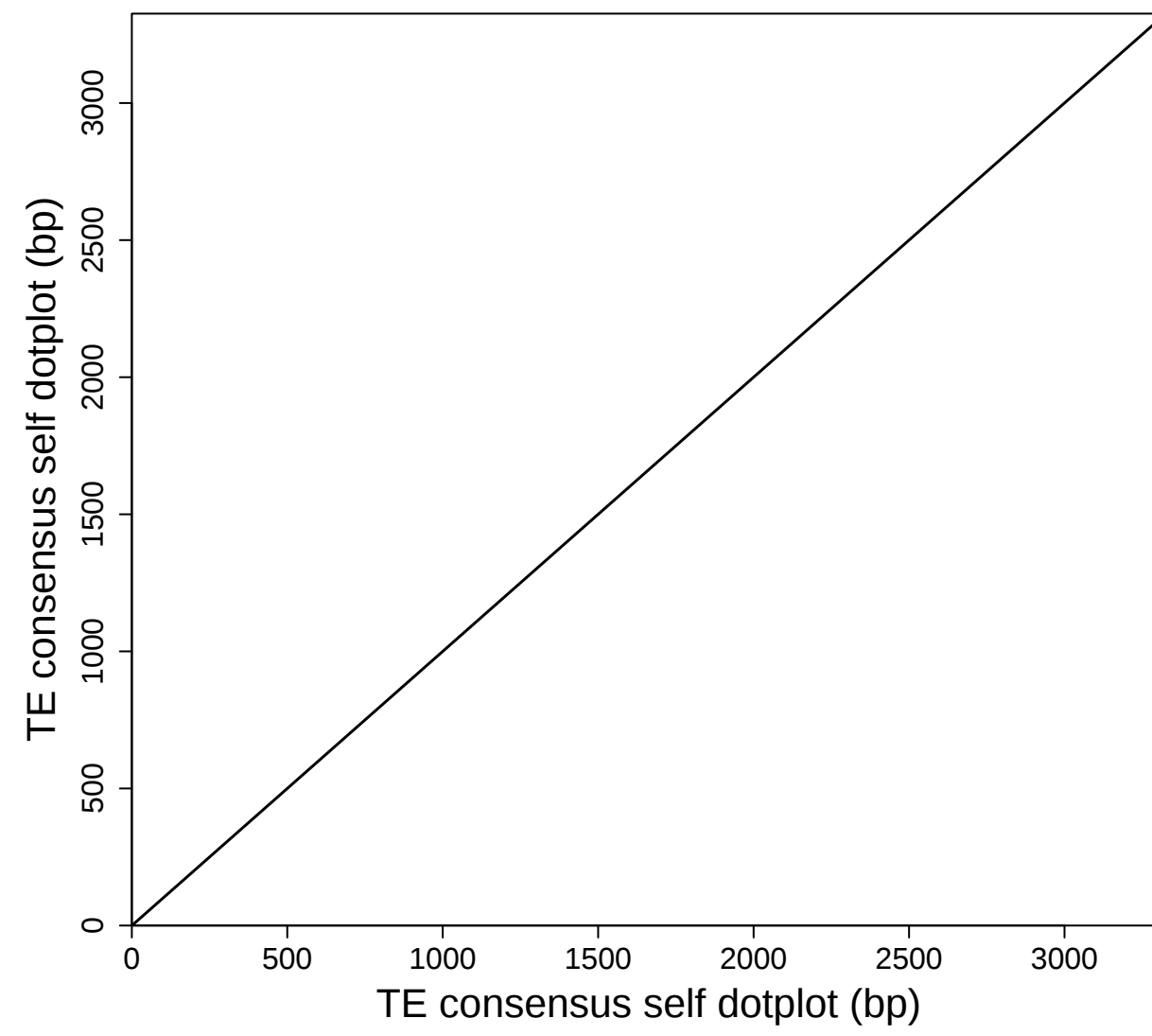
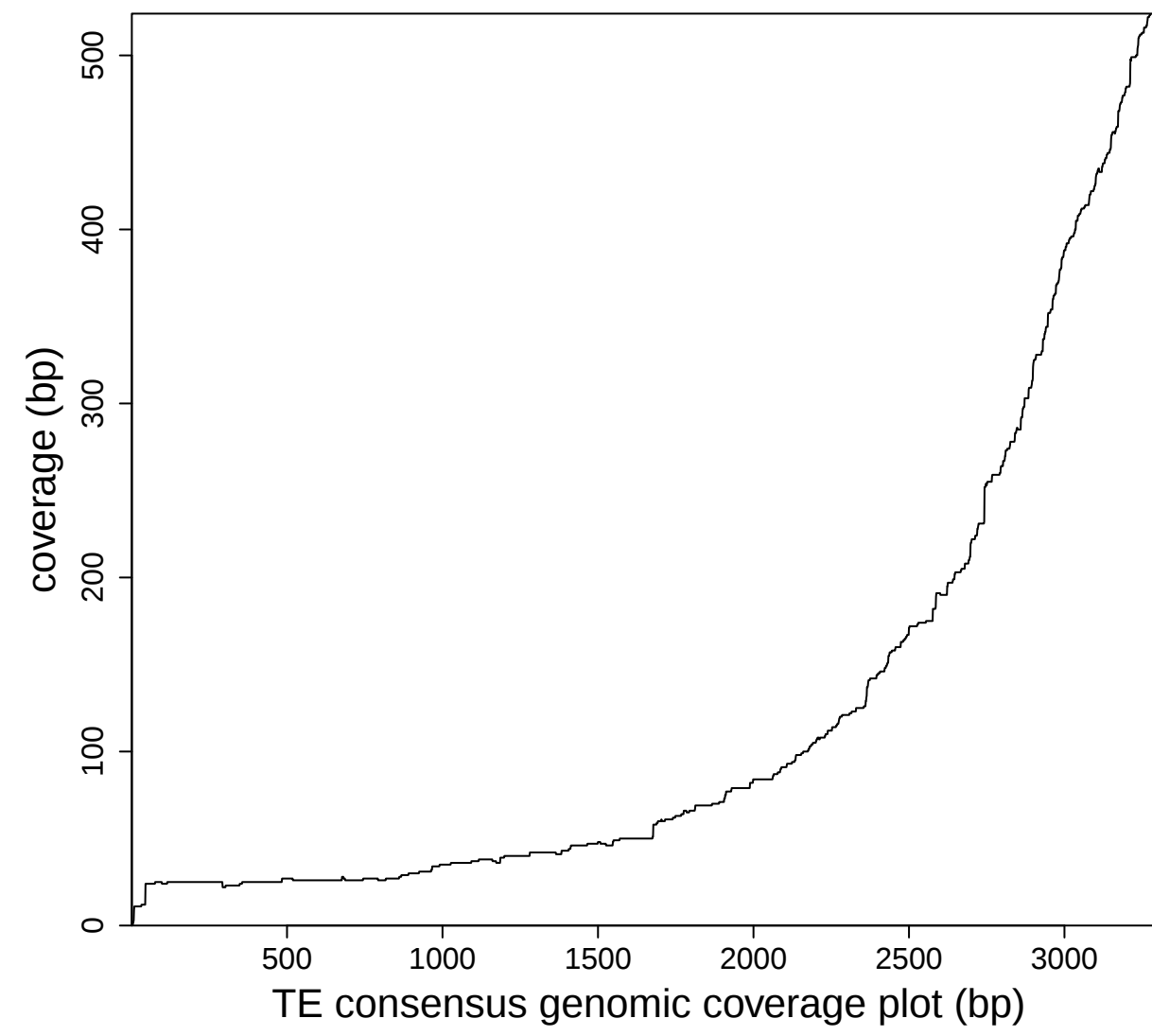
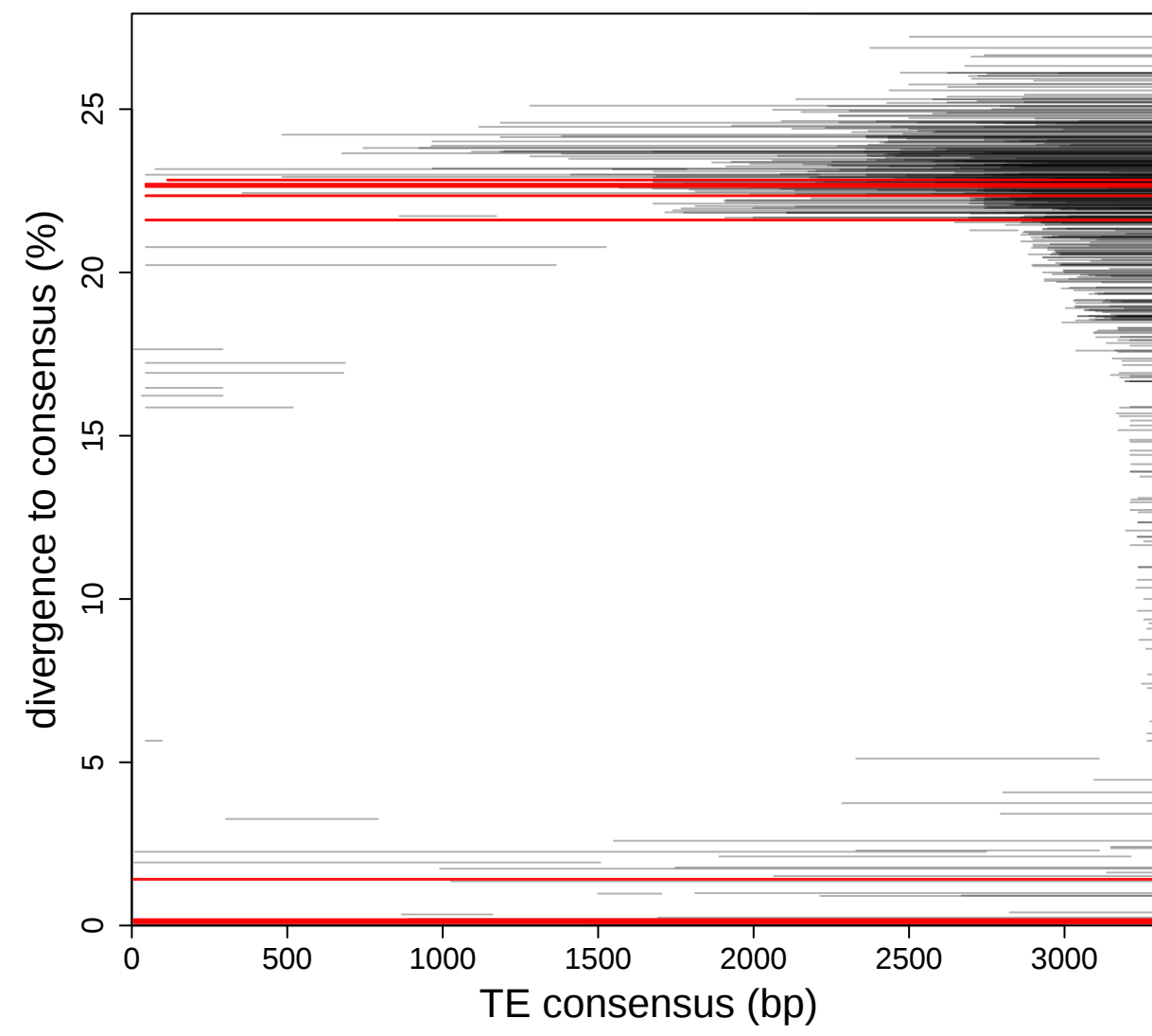
3000

3500



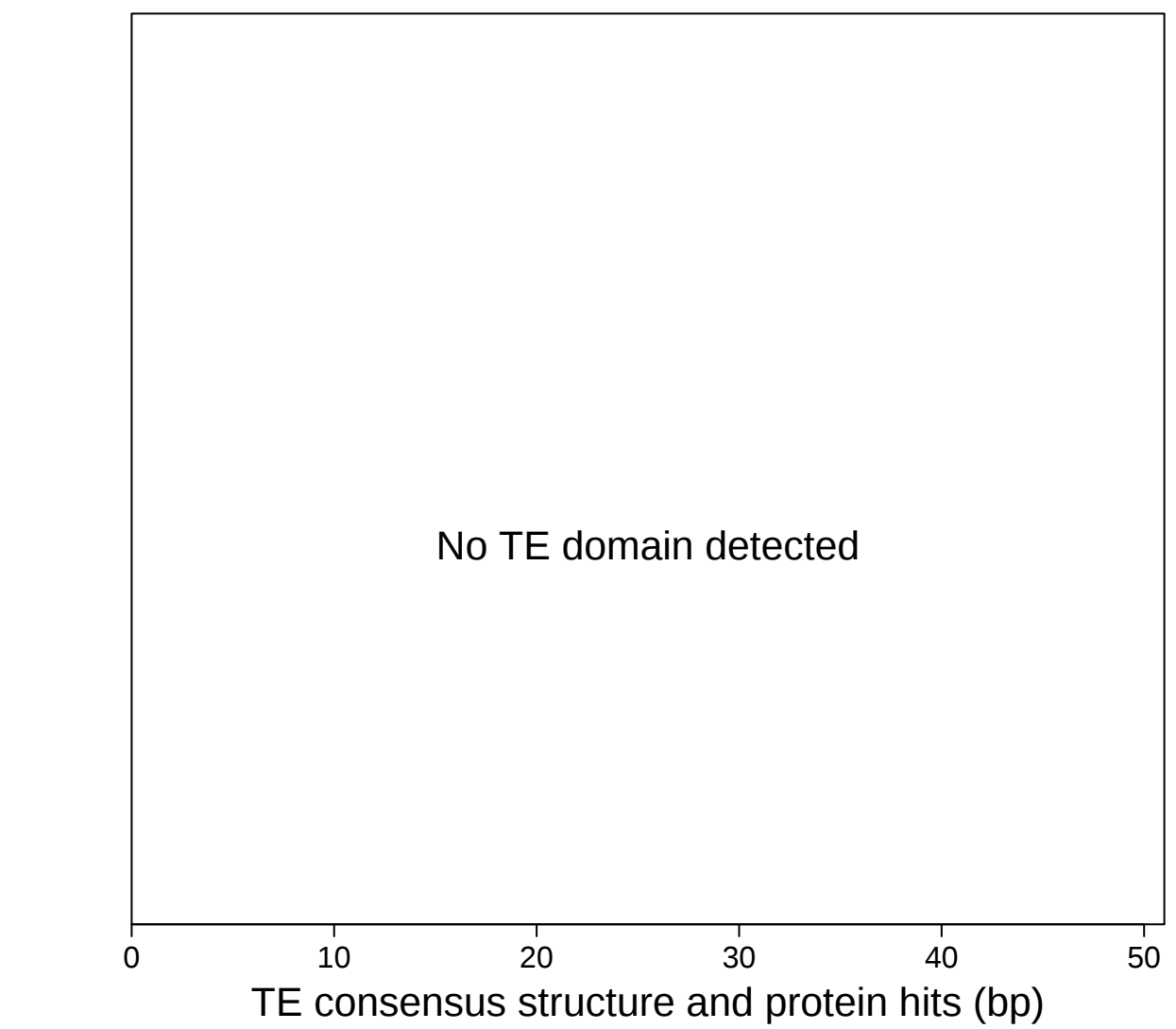
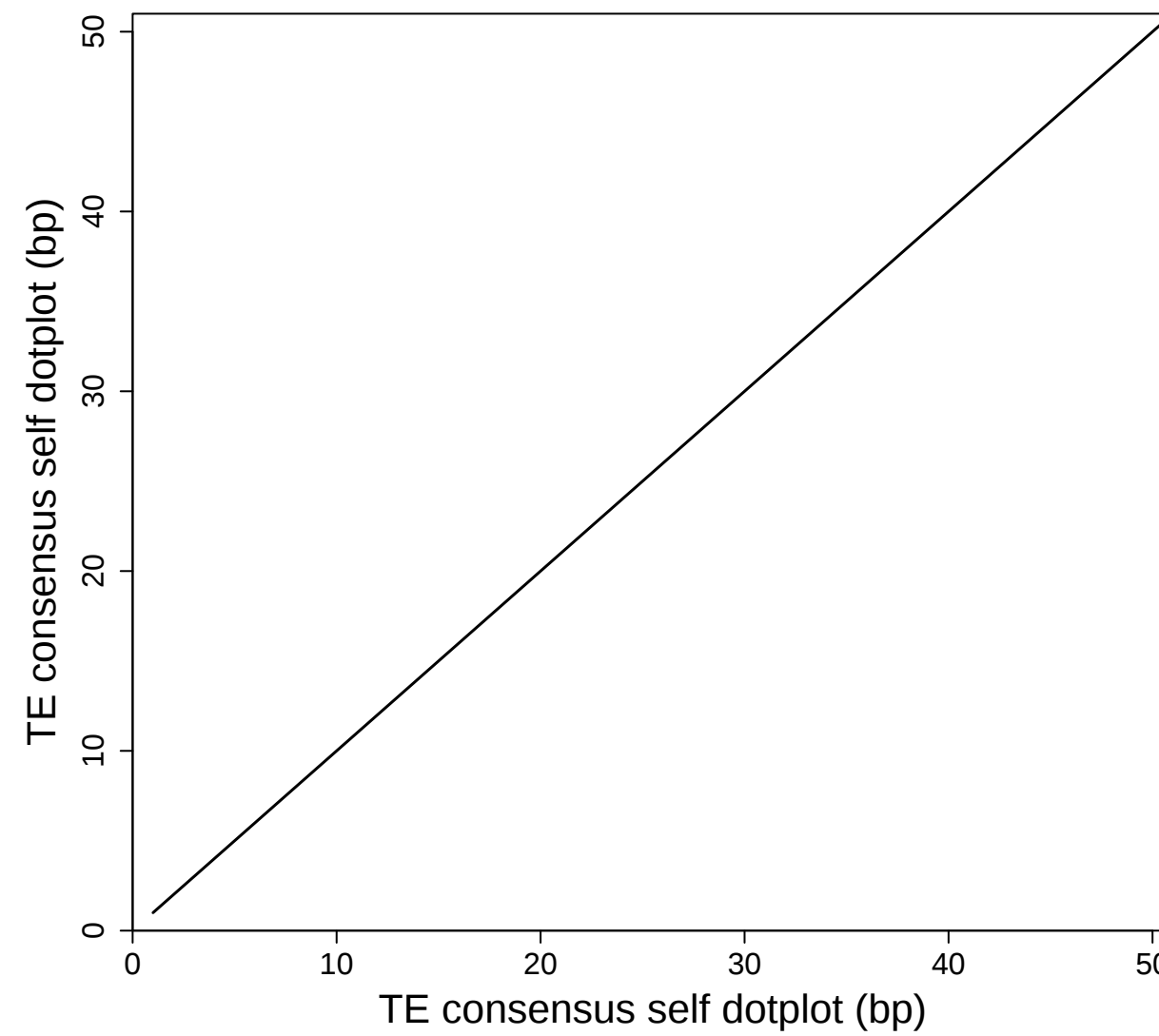
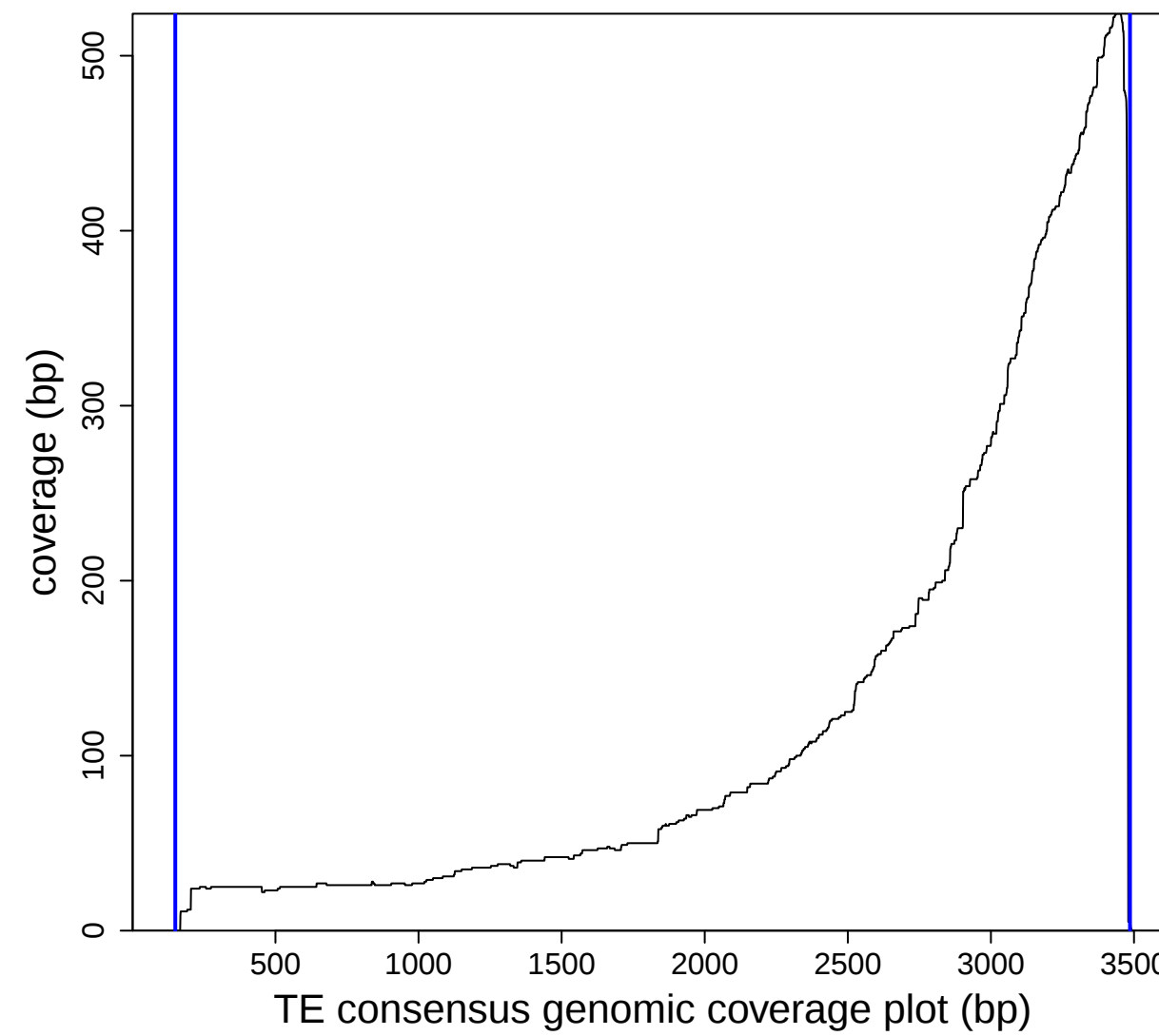
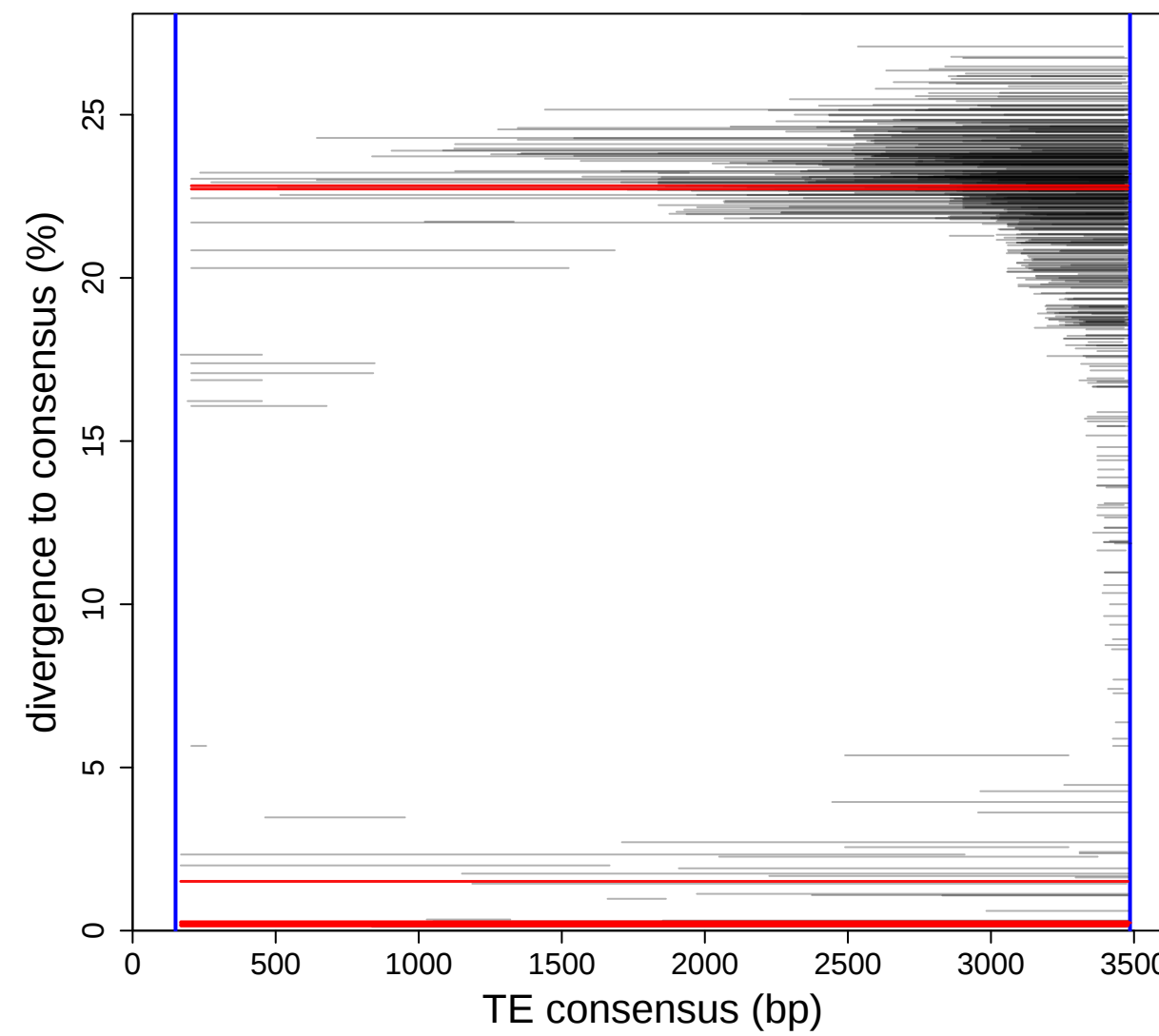
bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 3326bp; fragments: 547; full length: 13 (≥ 2993.4 bp)

After TEtrimmer 3326 bp

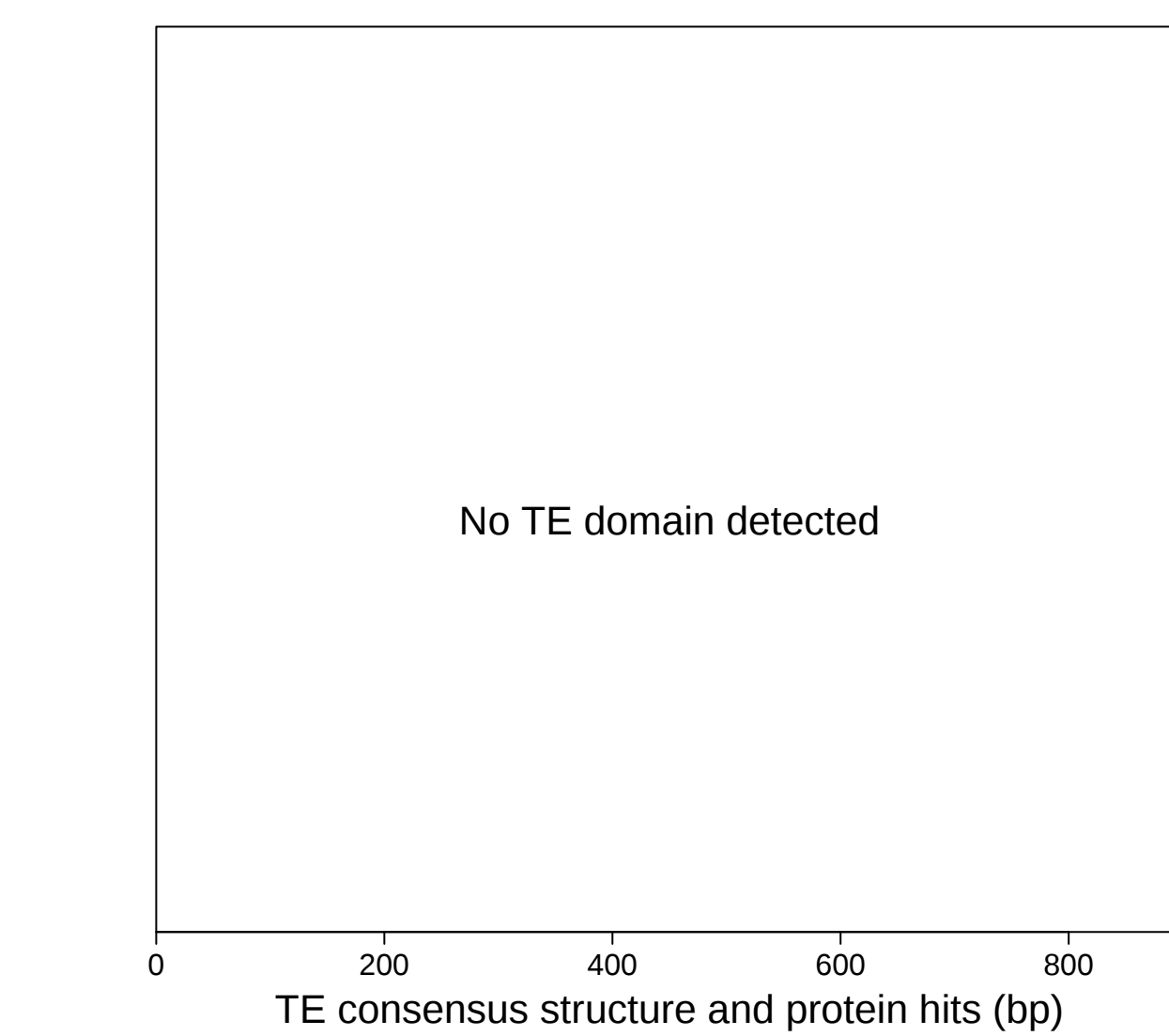
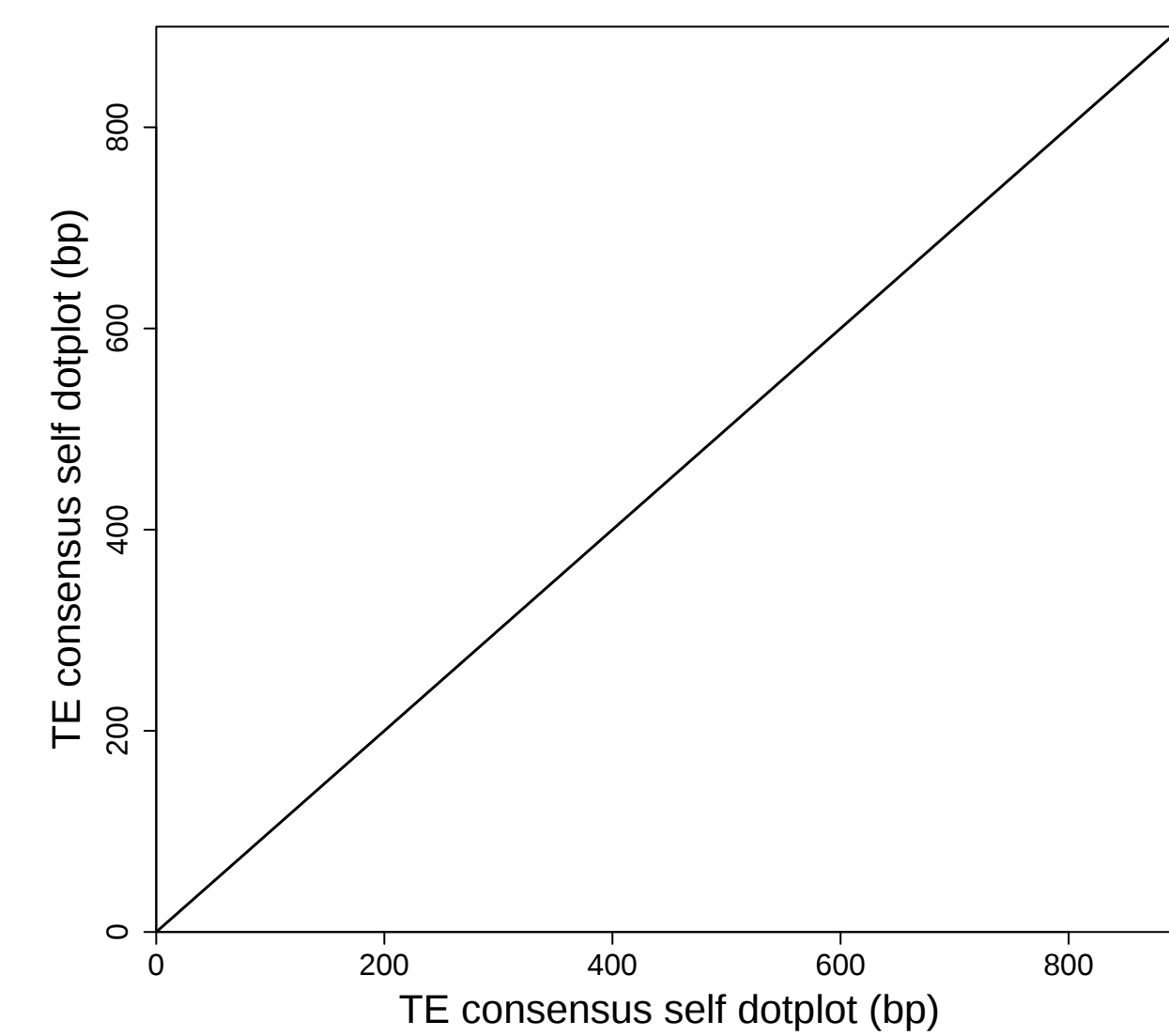
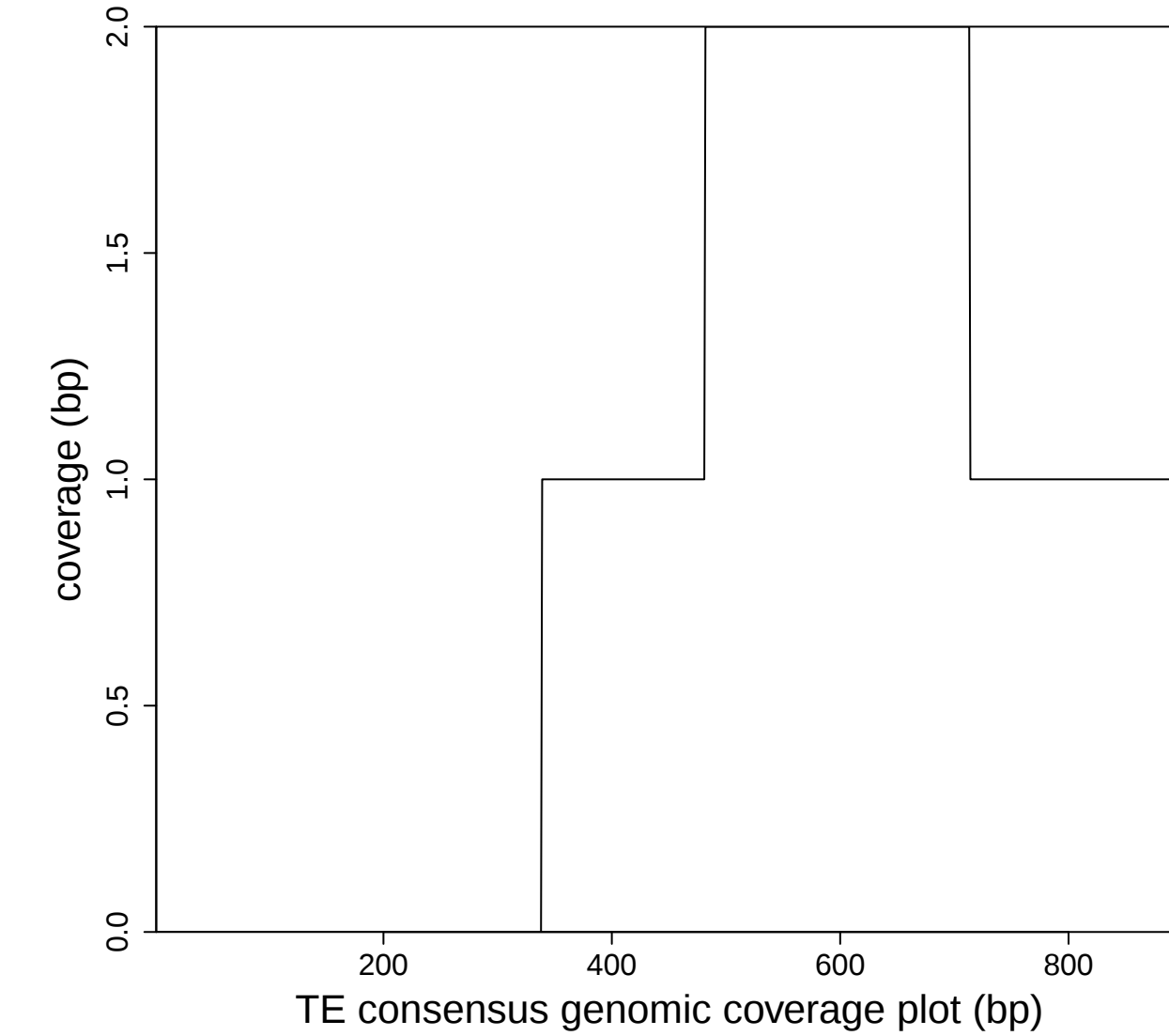
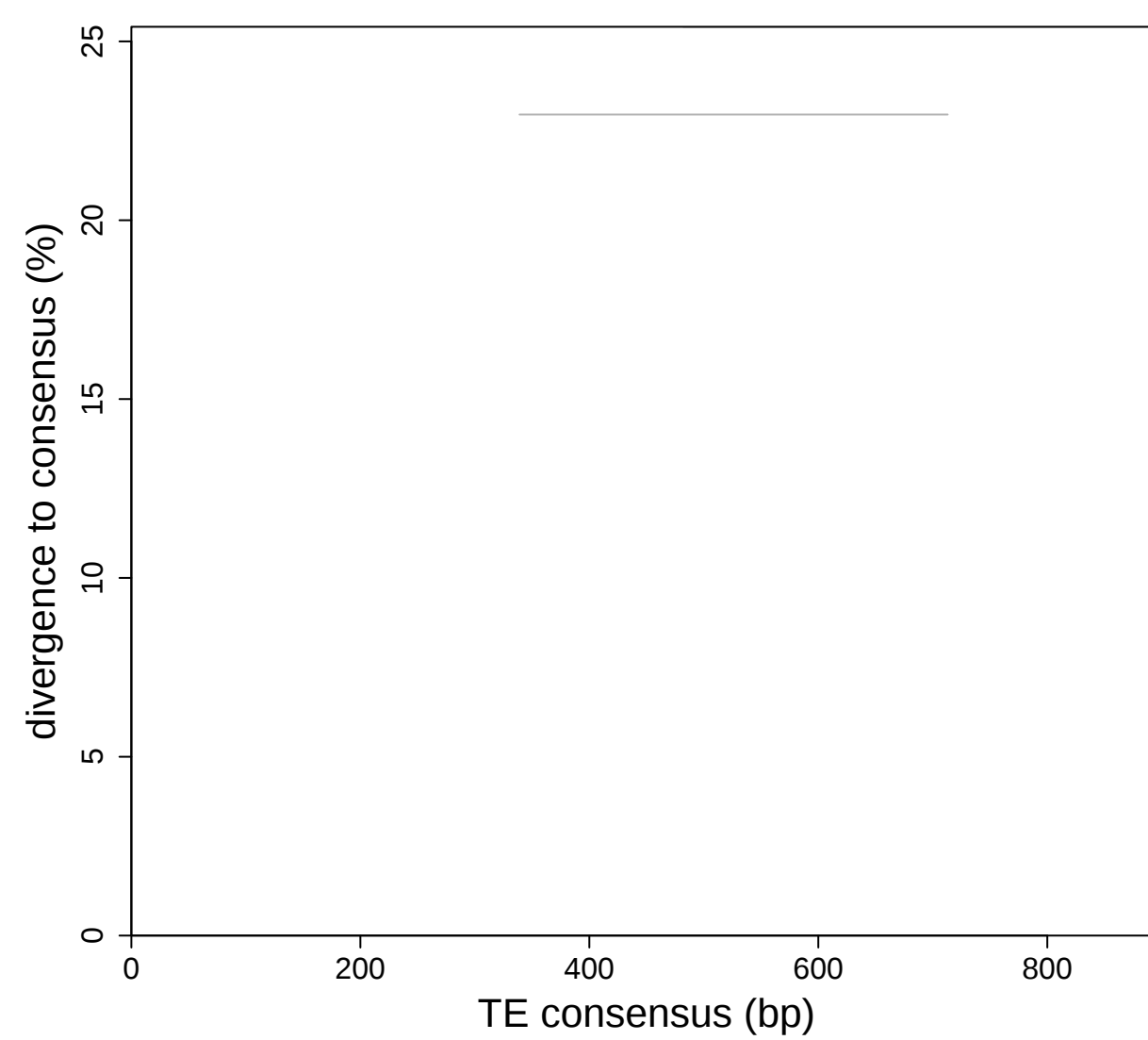


2.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 3636bp; fragments: 547; full length: 10 (>=3272.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

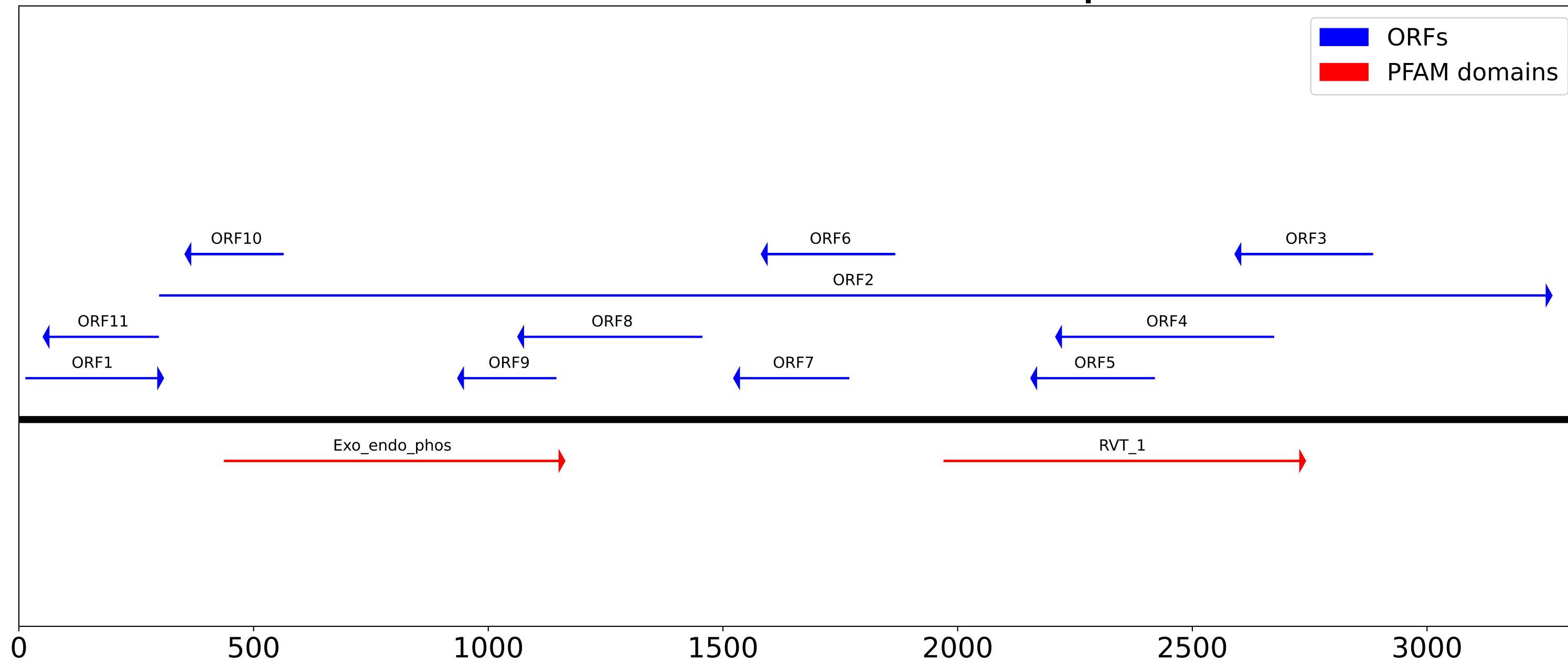


TE: rnd_4_family_1612
size: 900bp; fragments: 2; full length: 0 (≥ 810 bp)

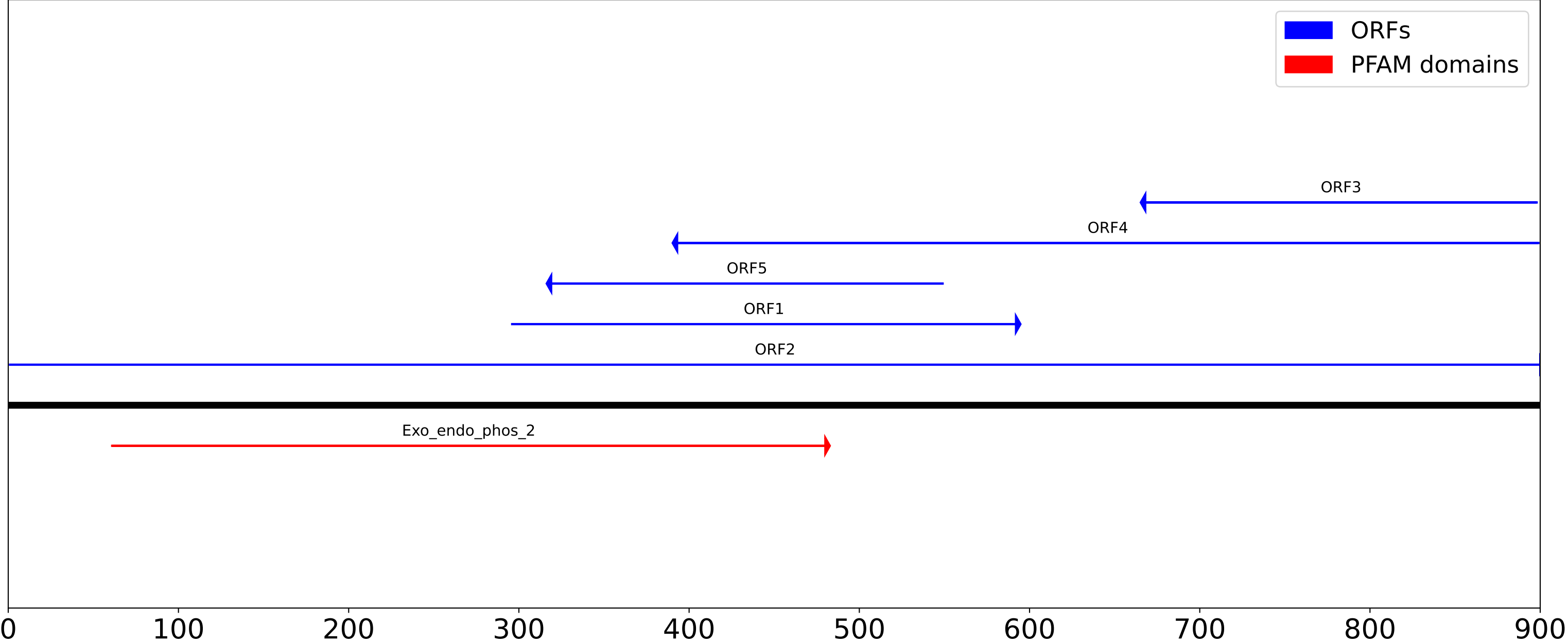


Before TEtrimmer 900 bp

After TEtrimmer ORF and PFAM domain plot



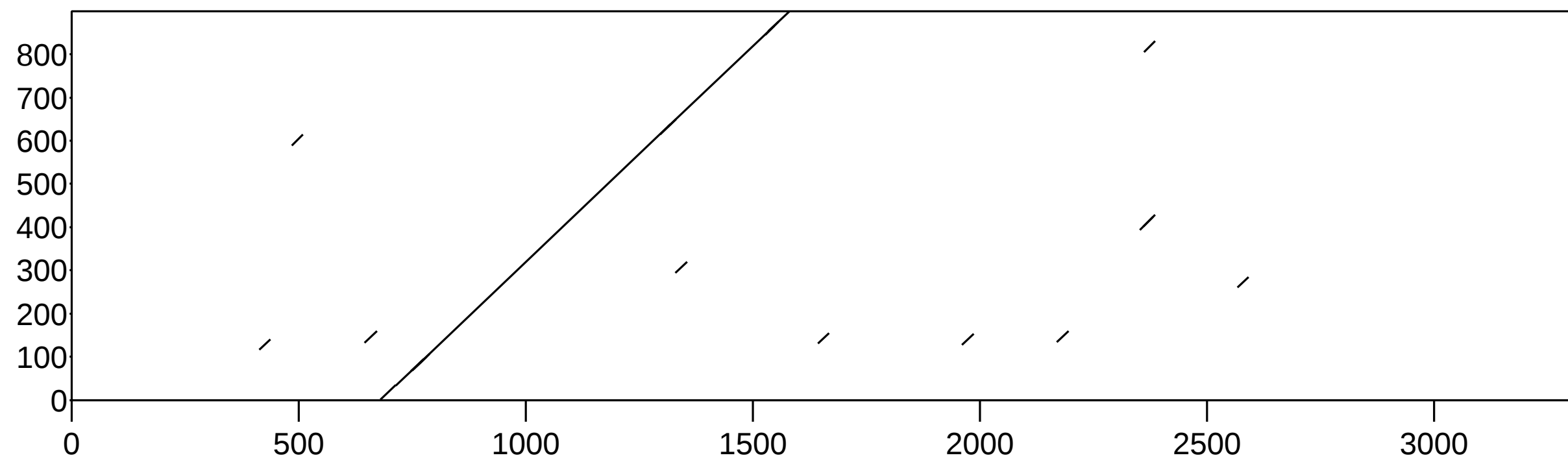
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)