

Start crop Point

End crop Point

0 -

10 -

0

500

1000

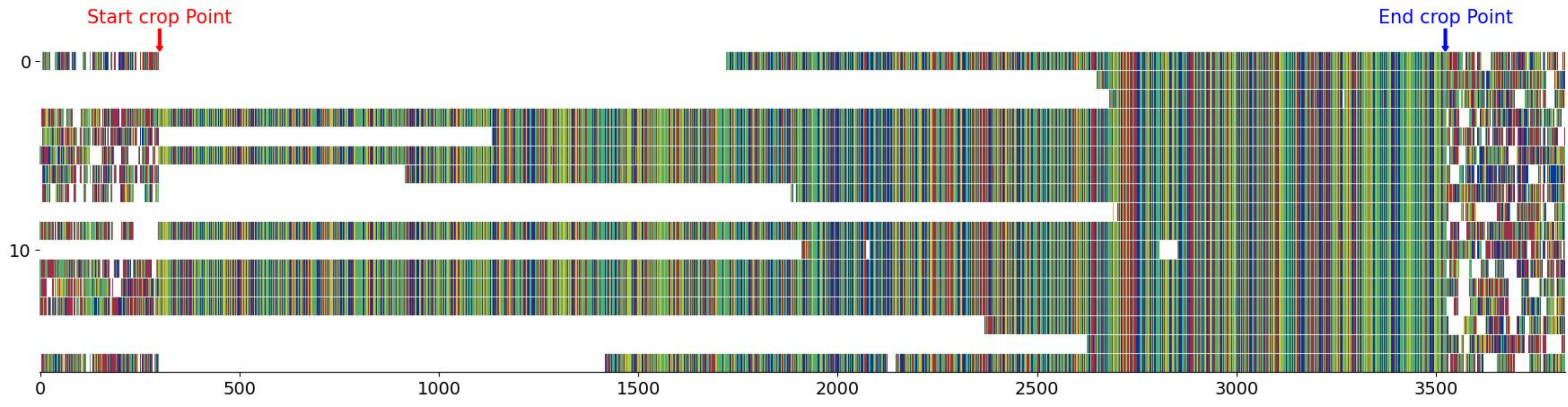
1500

2000

2500

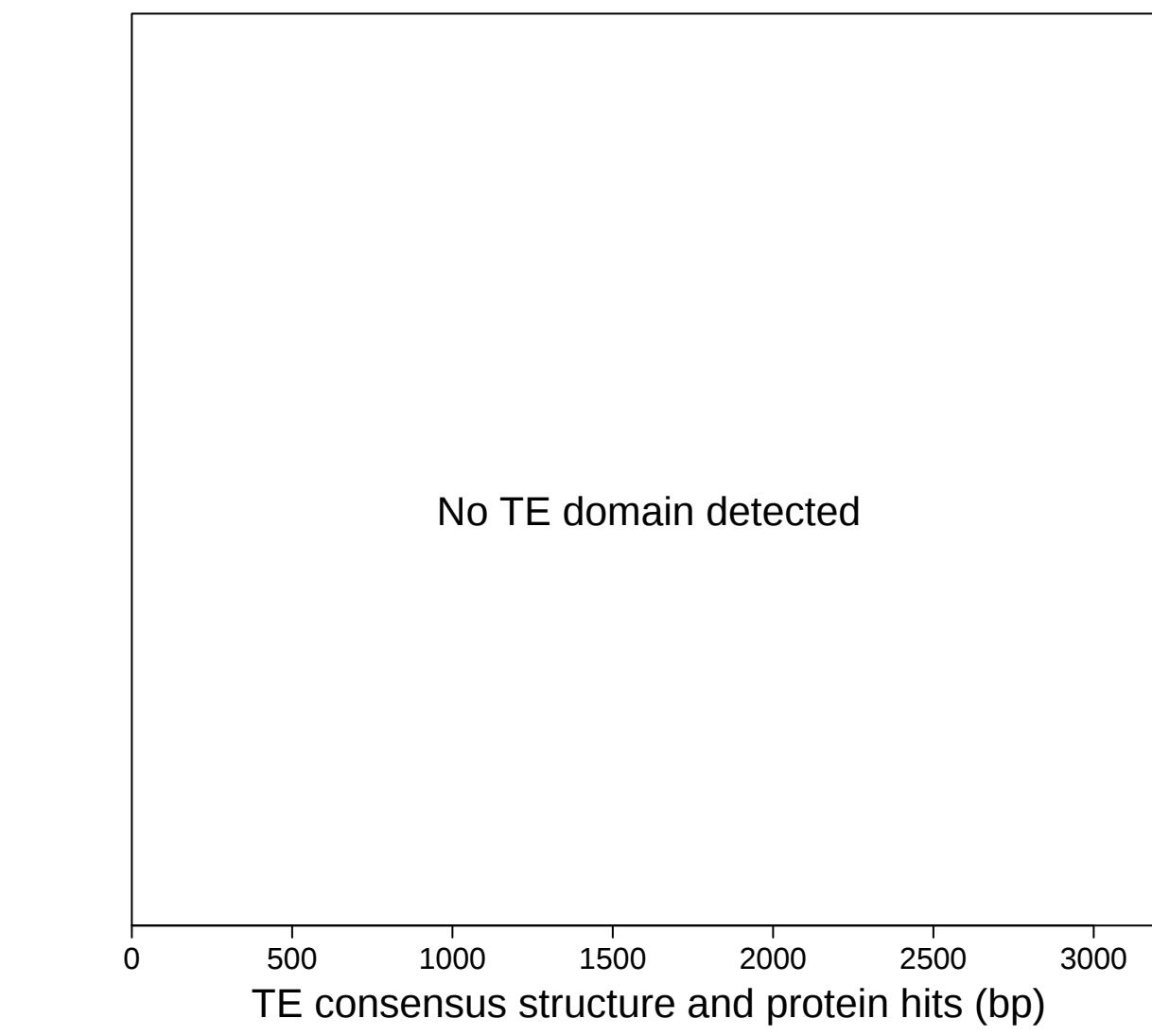
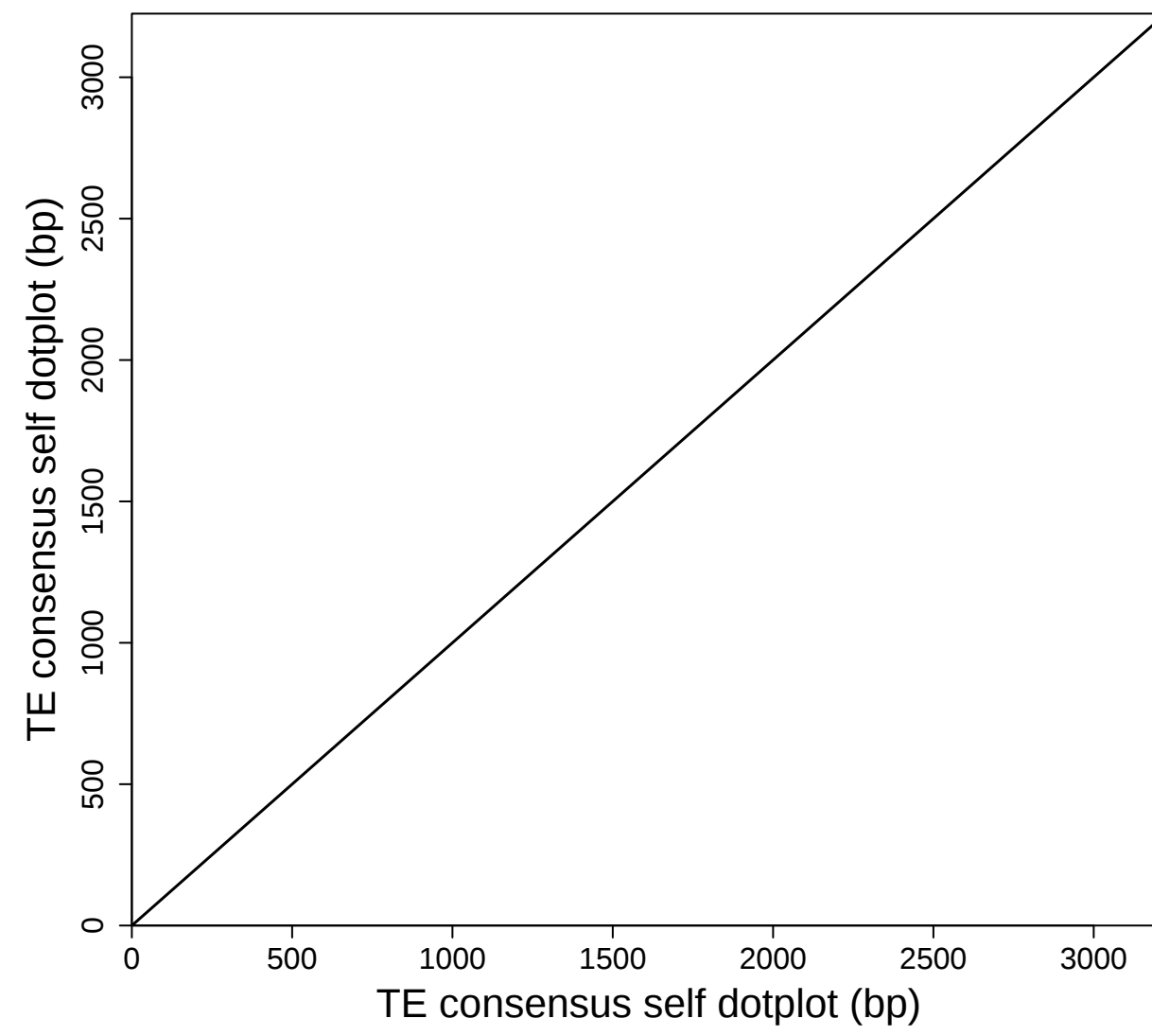
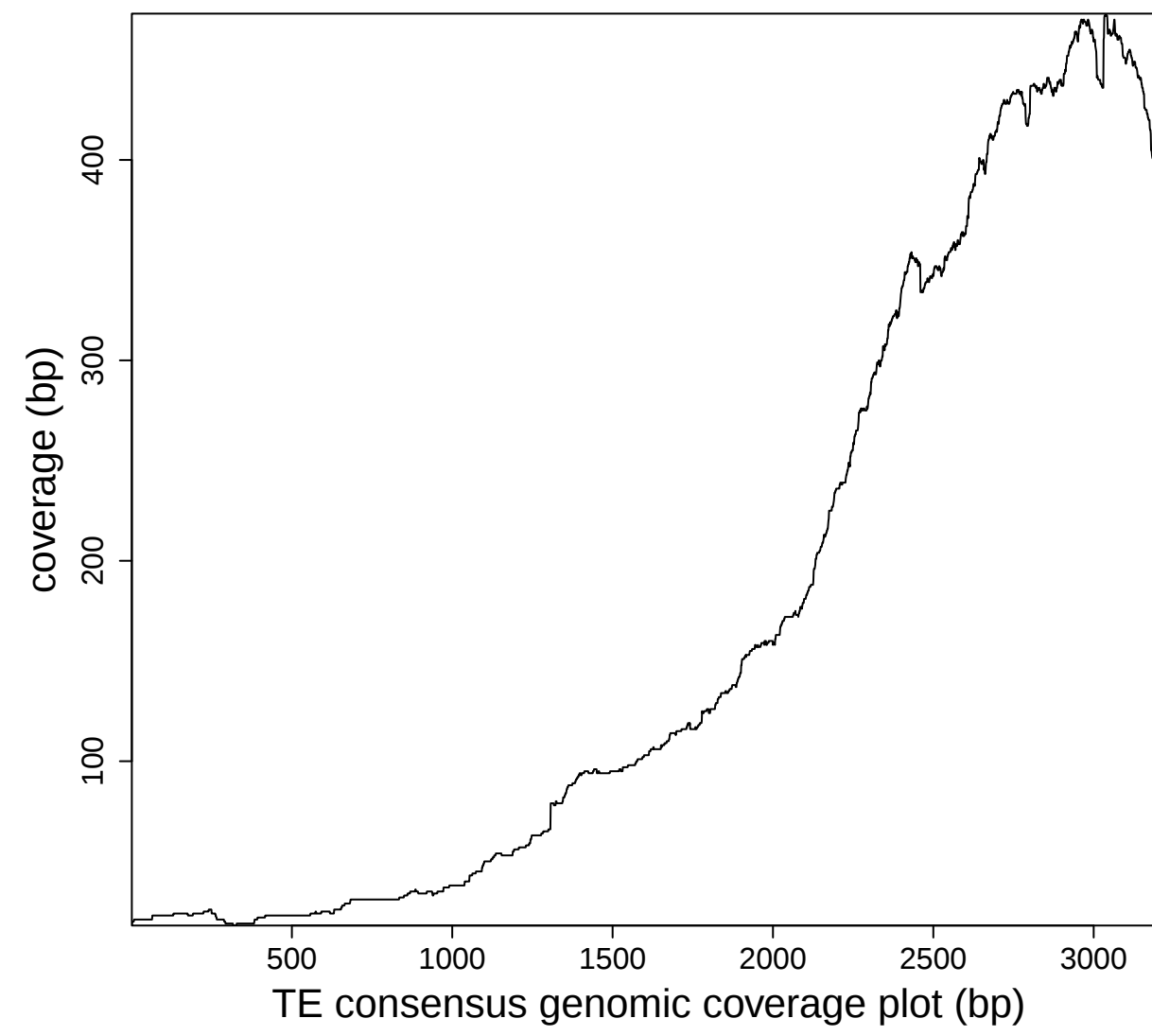
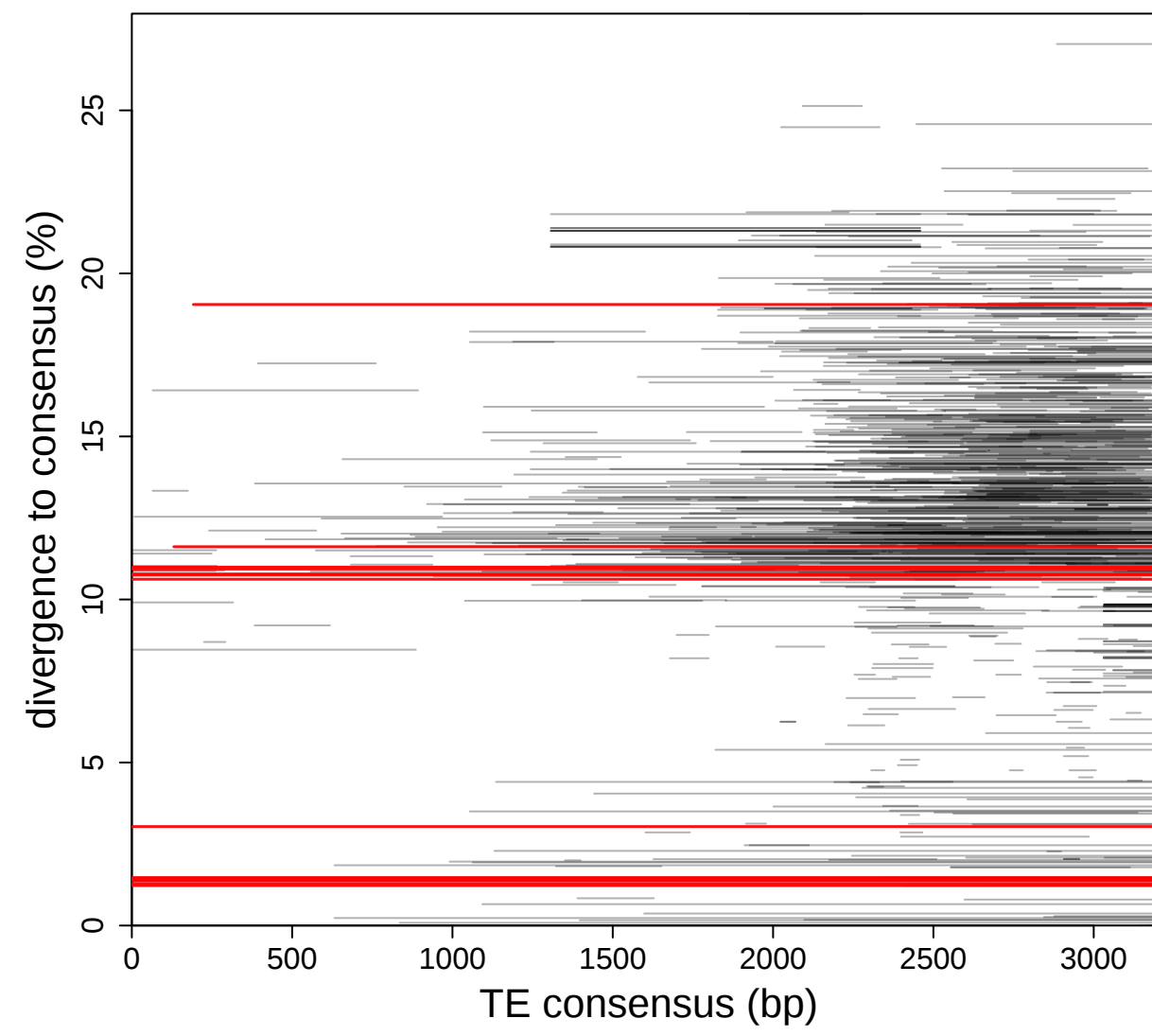
3000

3500



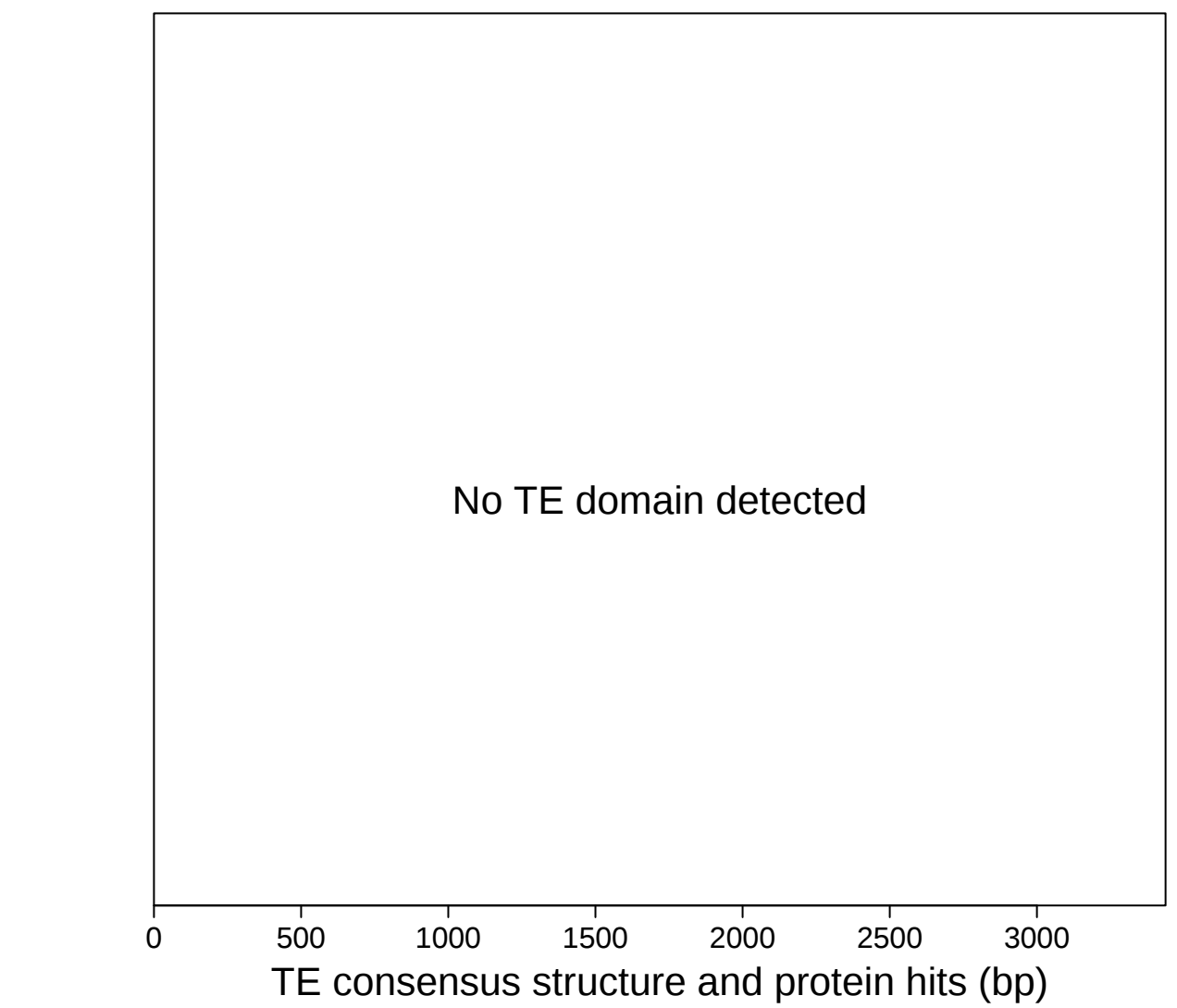
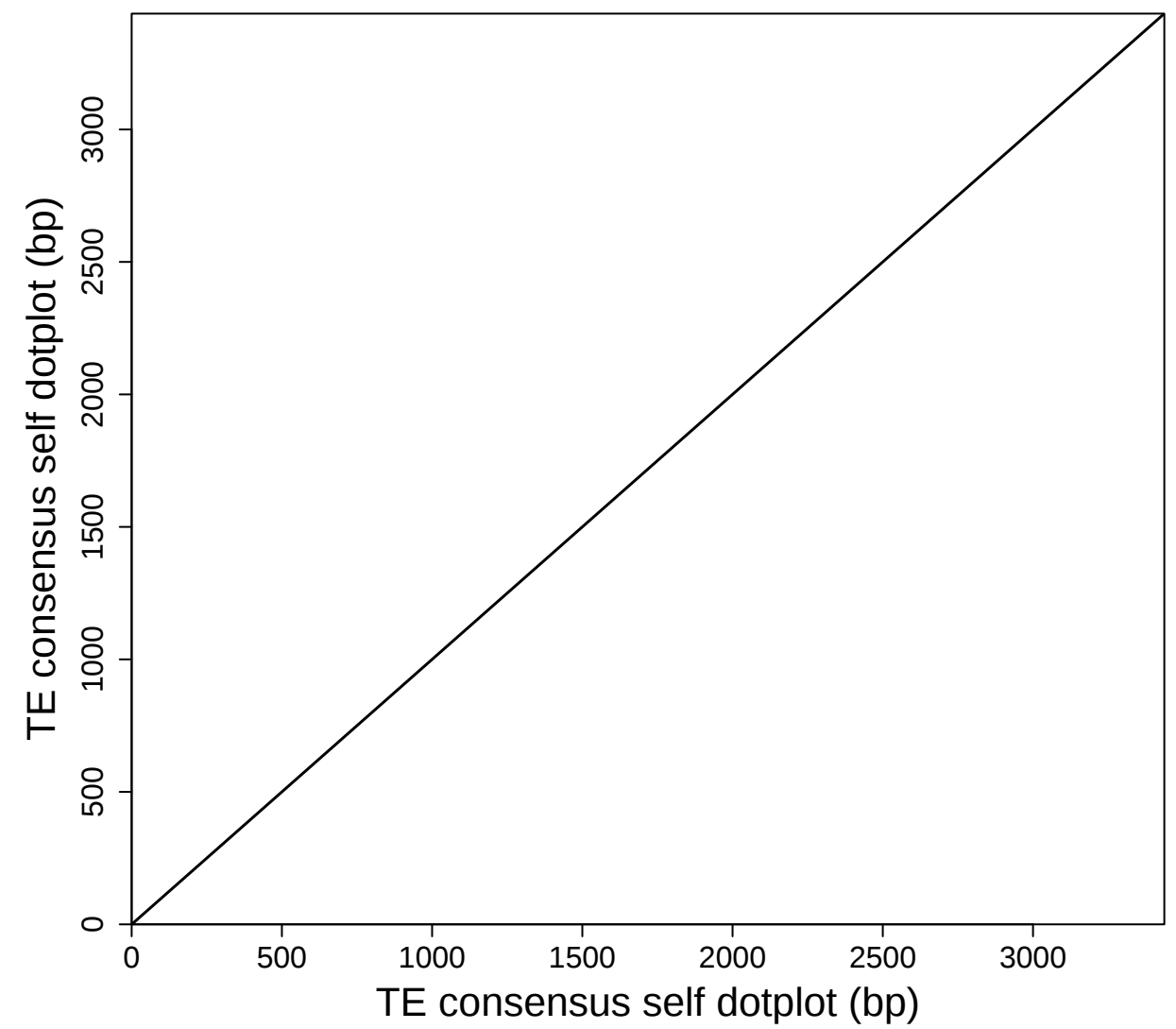
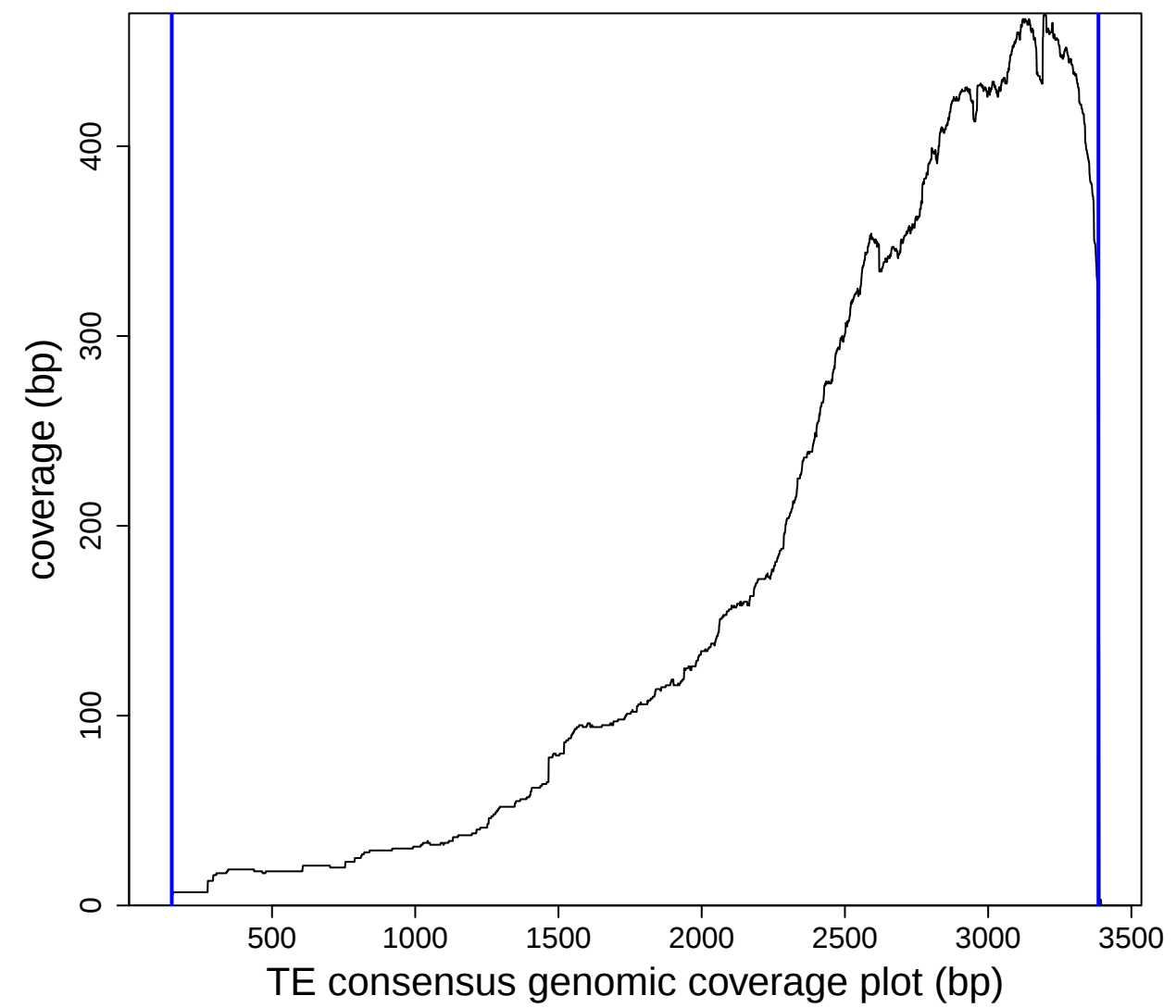
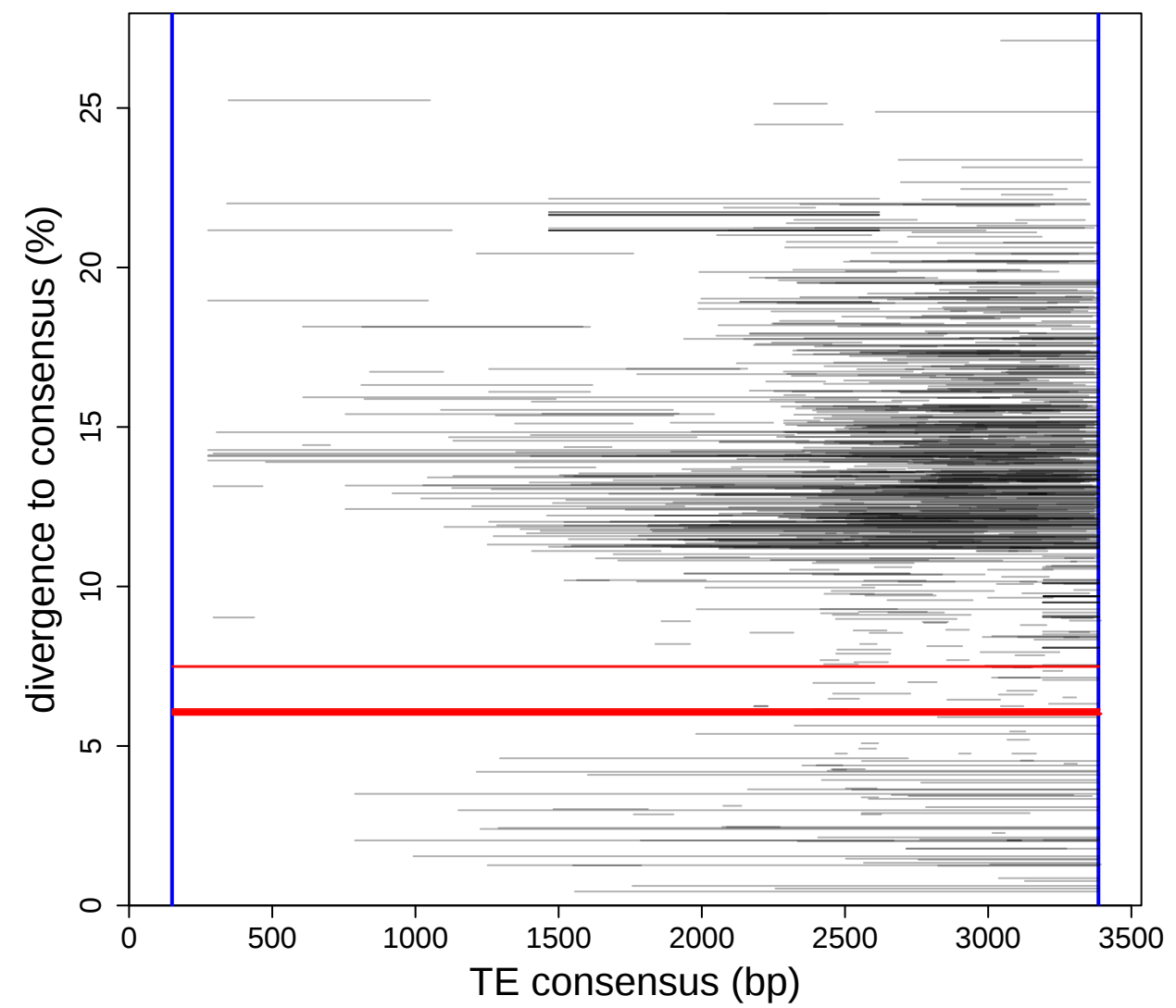
ed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 3225bp; fragments: 860; full length: 14 (≥ 2902.5 bp)

After TEtrimmer 3225 bp

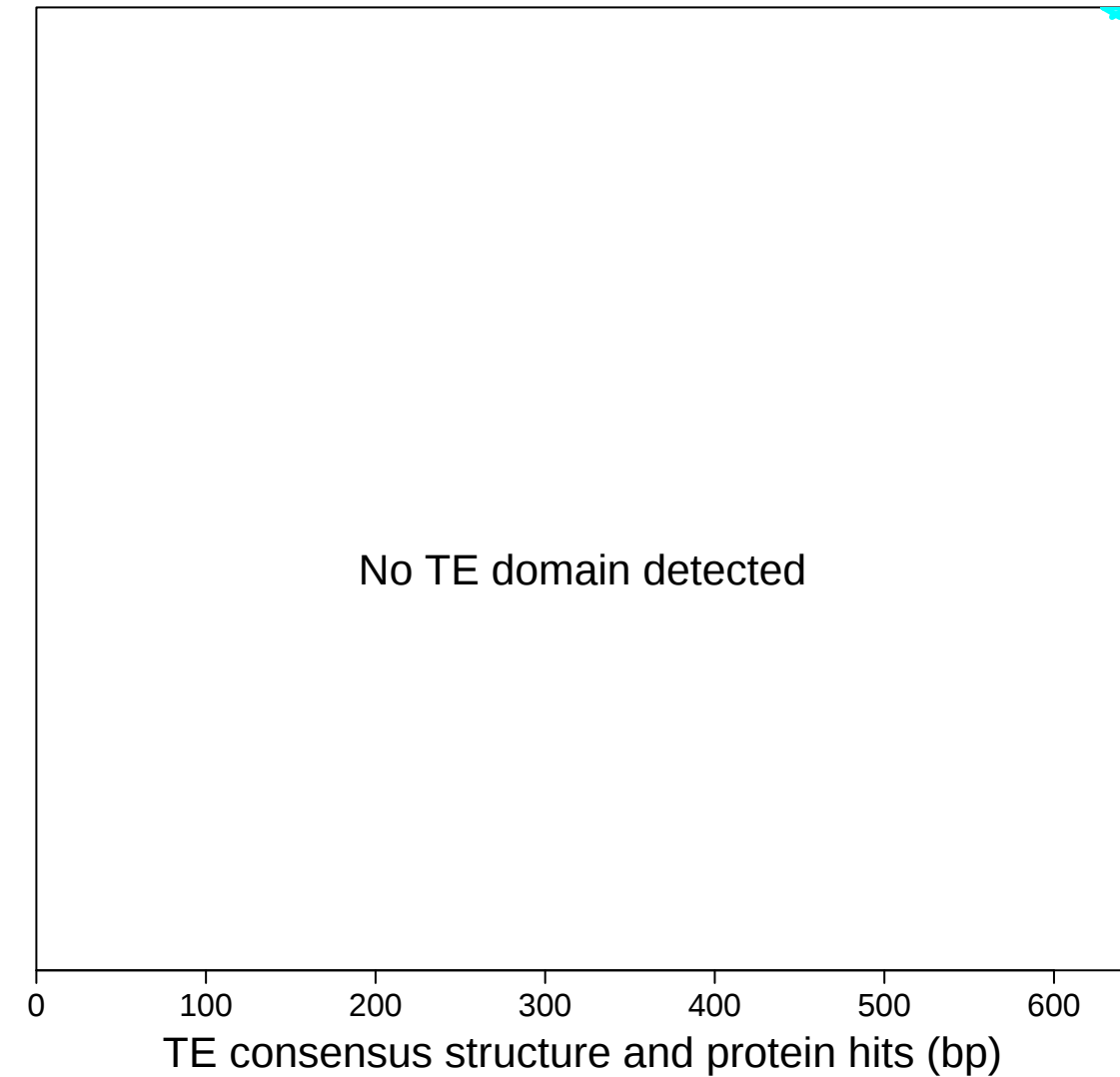
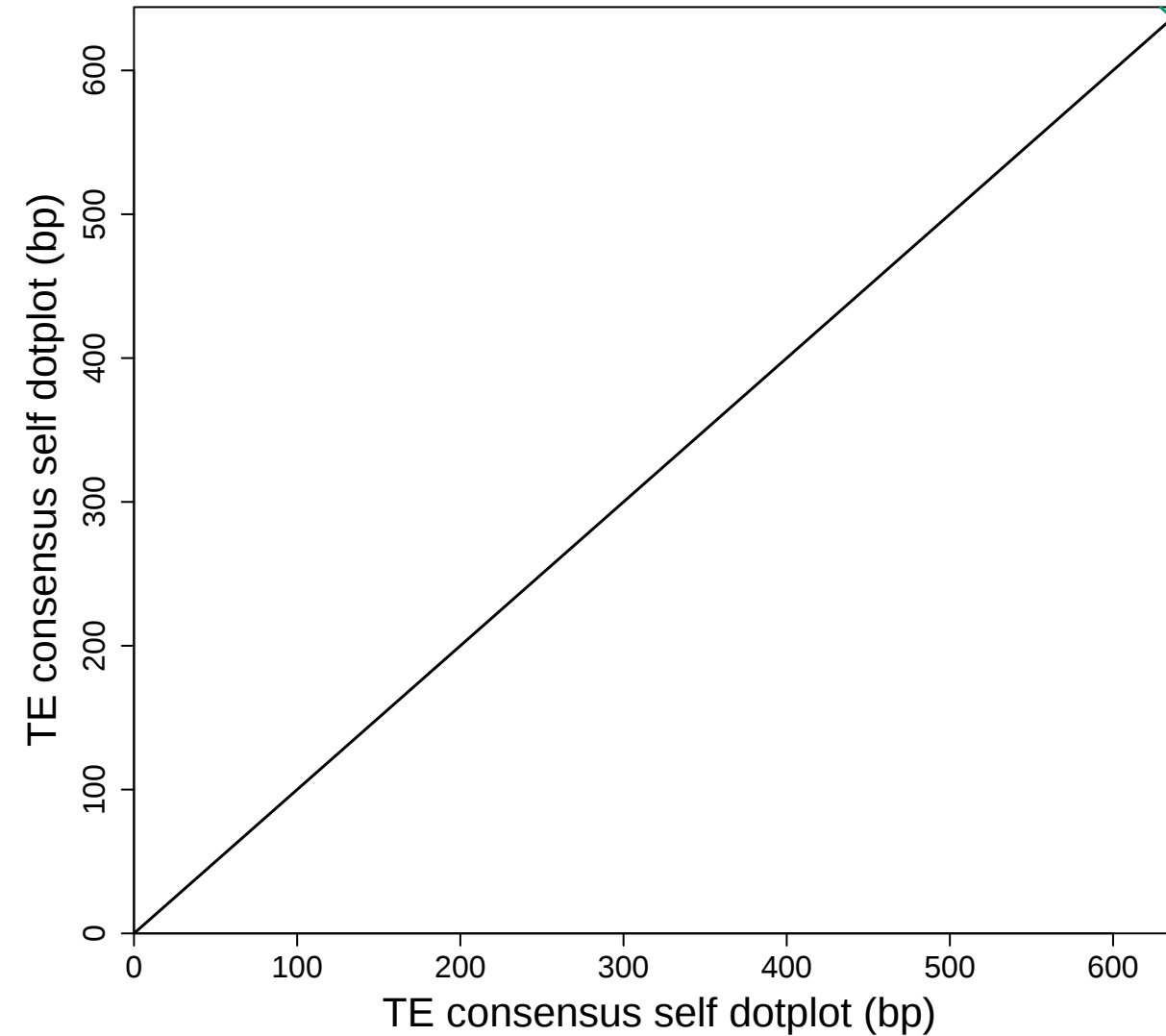
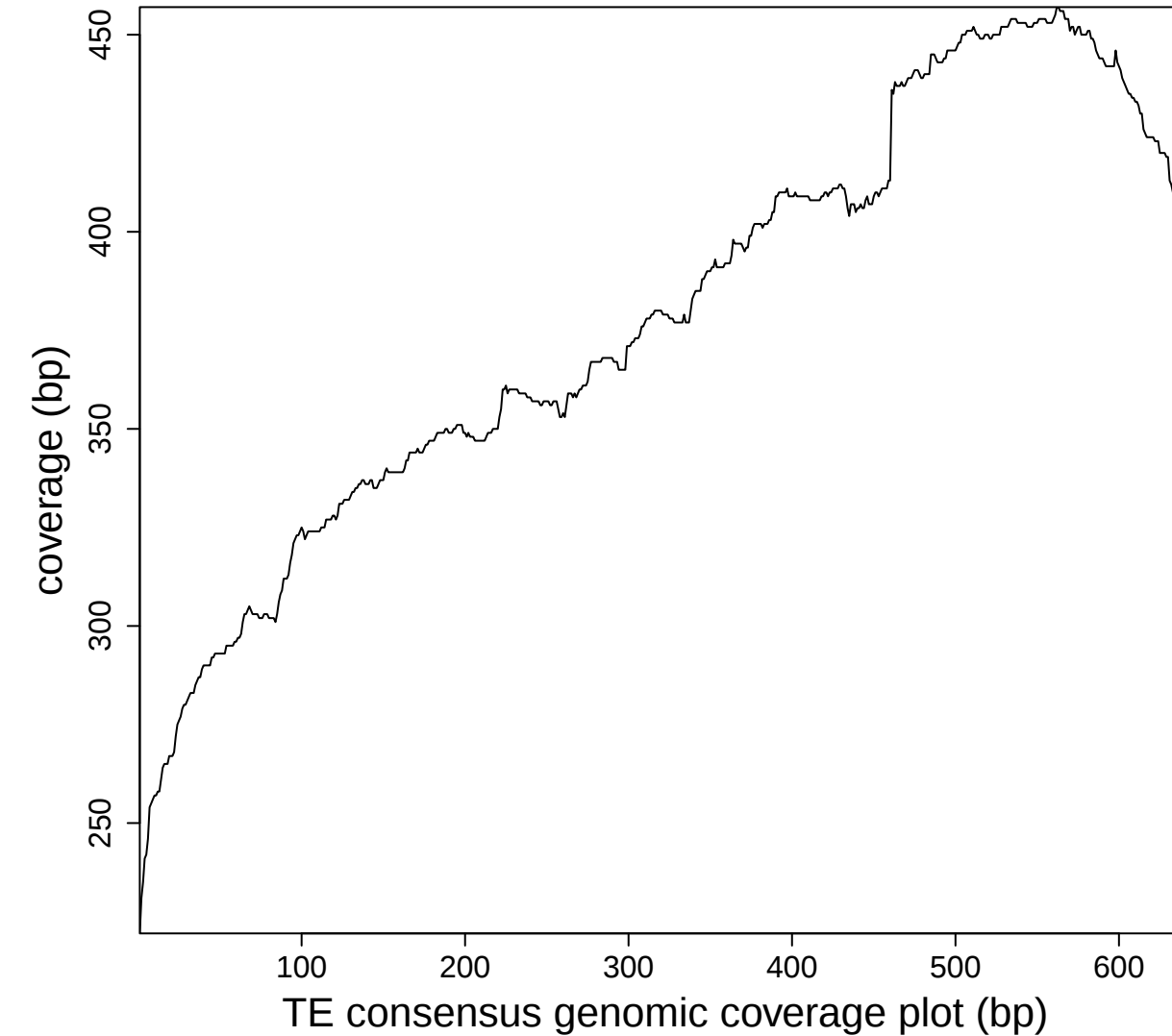
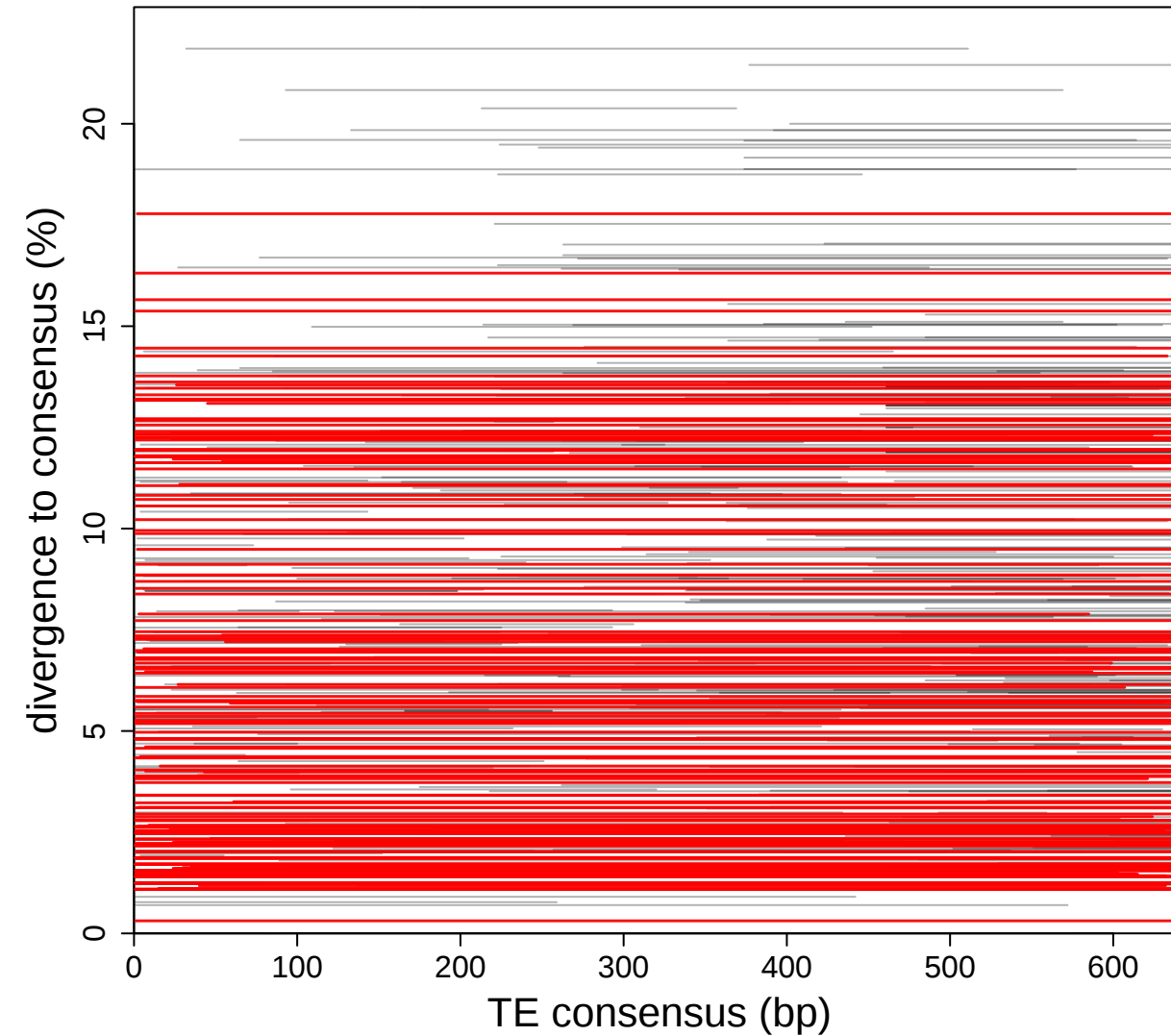


7.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 3535bp; fragments: 842; full length: 7 (>=3181.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

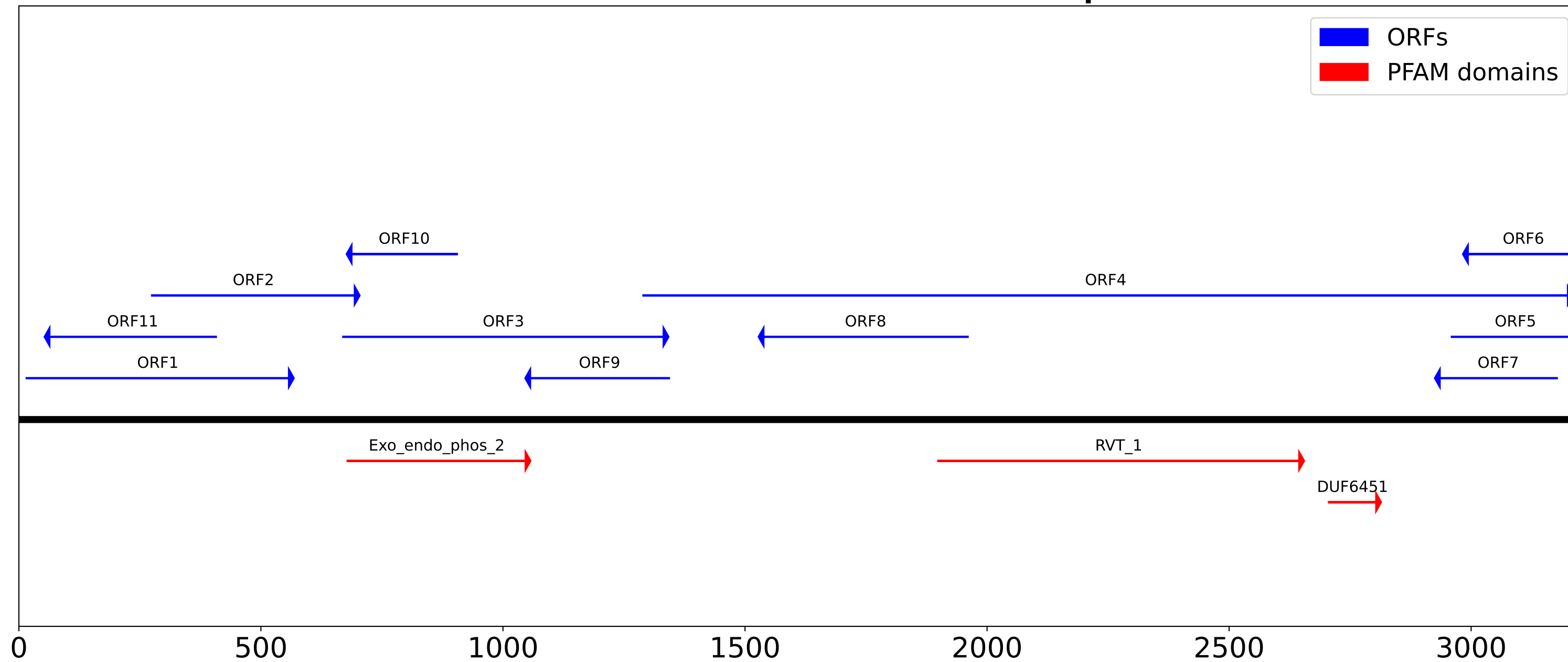


TE: rnd_1_family_977
size: 644bp; fragments: 584; full length: 213 (≥ 579.6 bp)

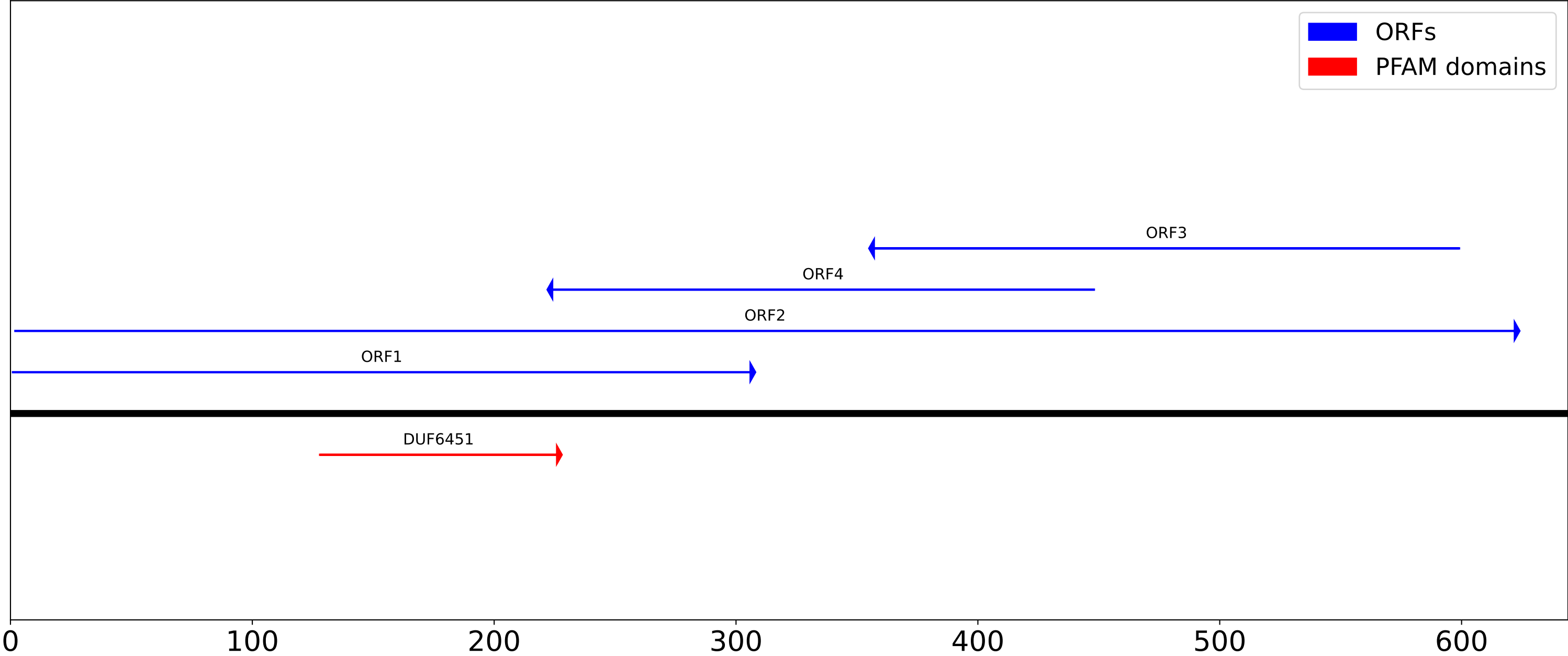


Before TEtrimmer 644 bp

After TEtrimmer ORF and PFAM domain plot



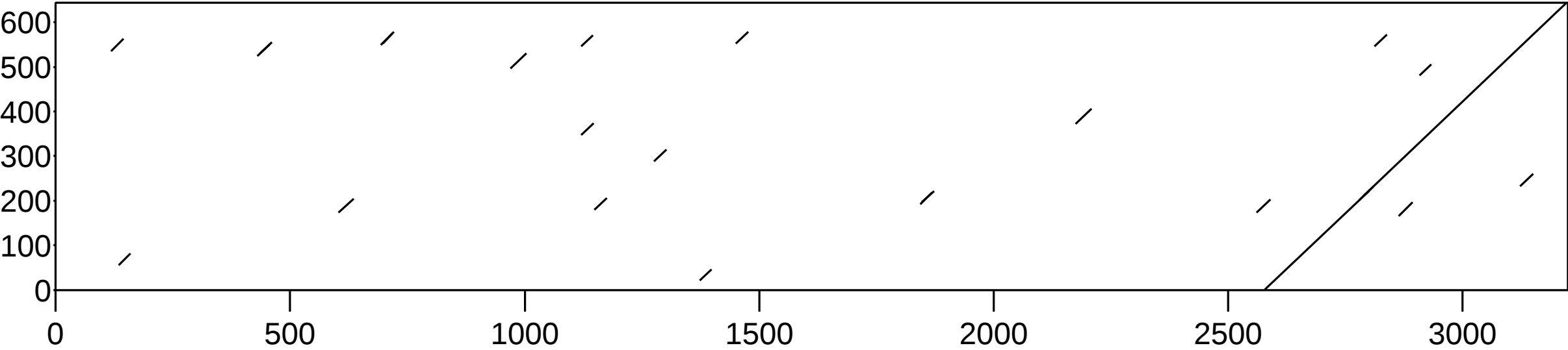
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)