

Start crop Point

End crop Point

0

10

0

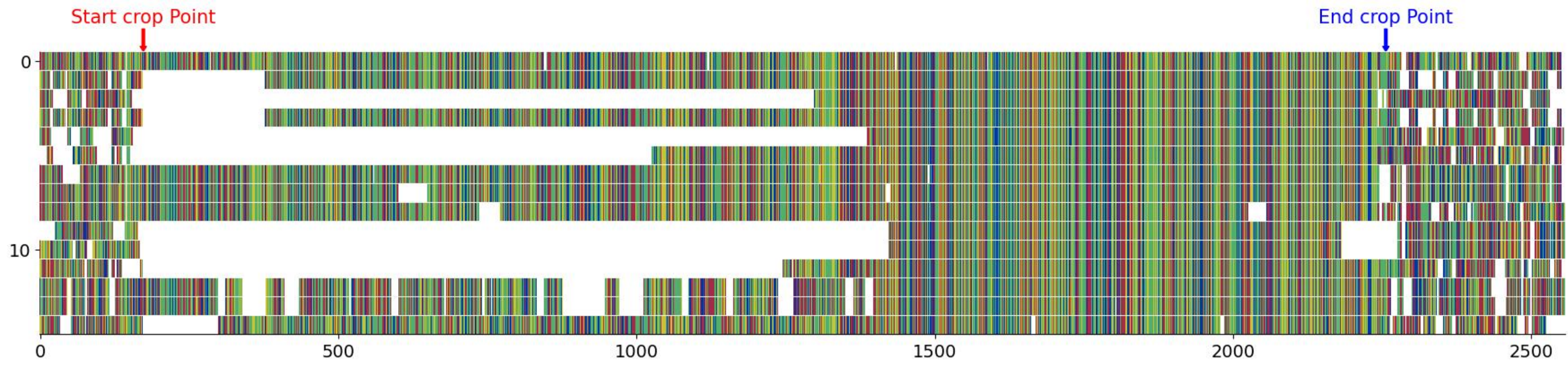
500

1000

1500

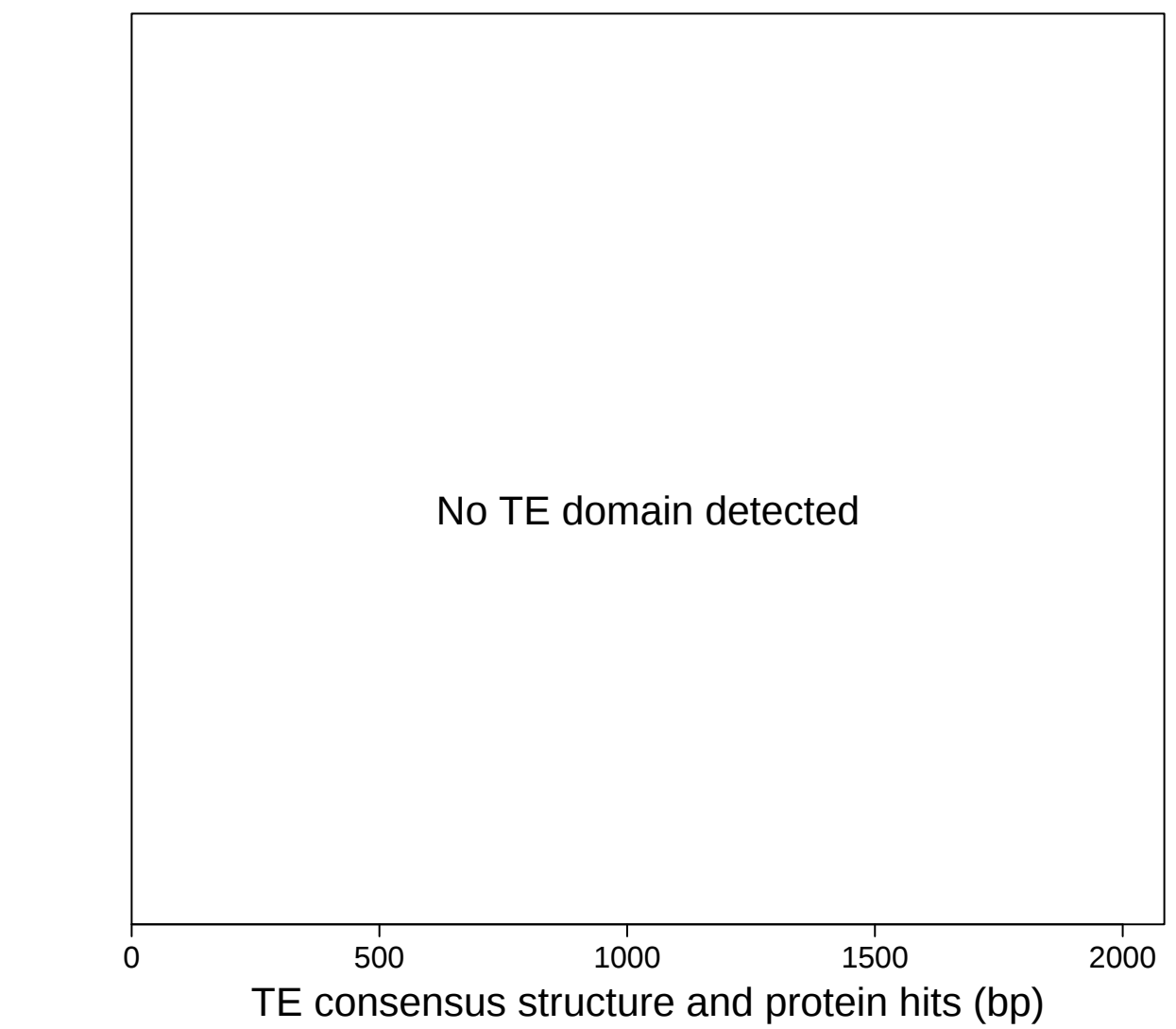
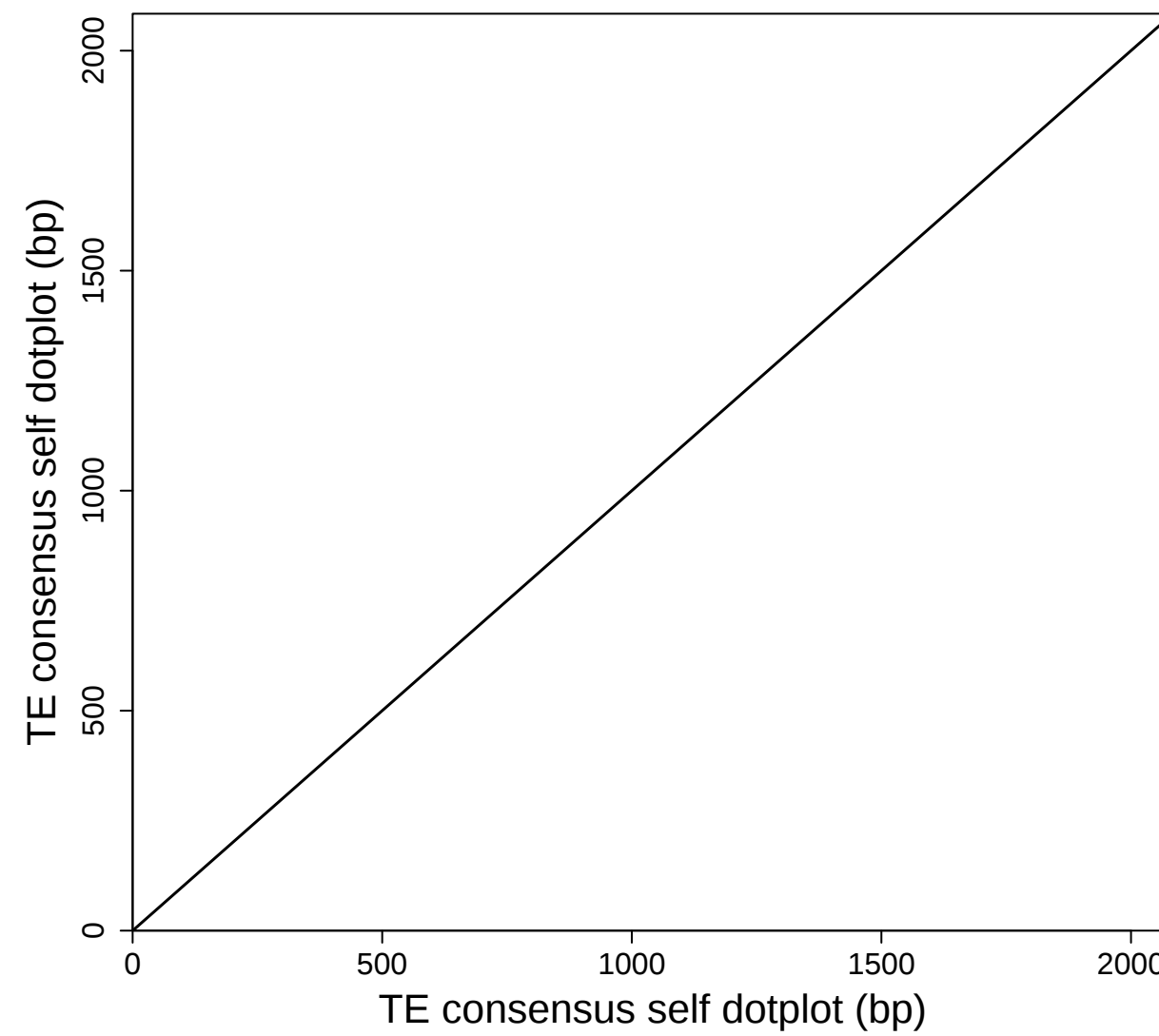
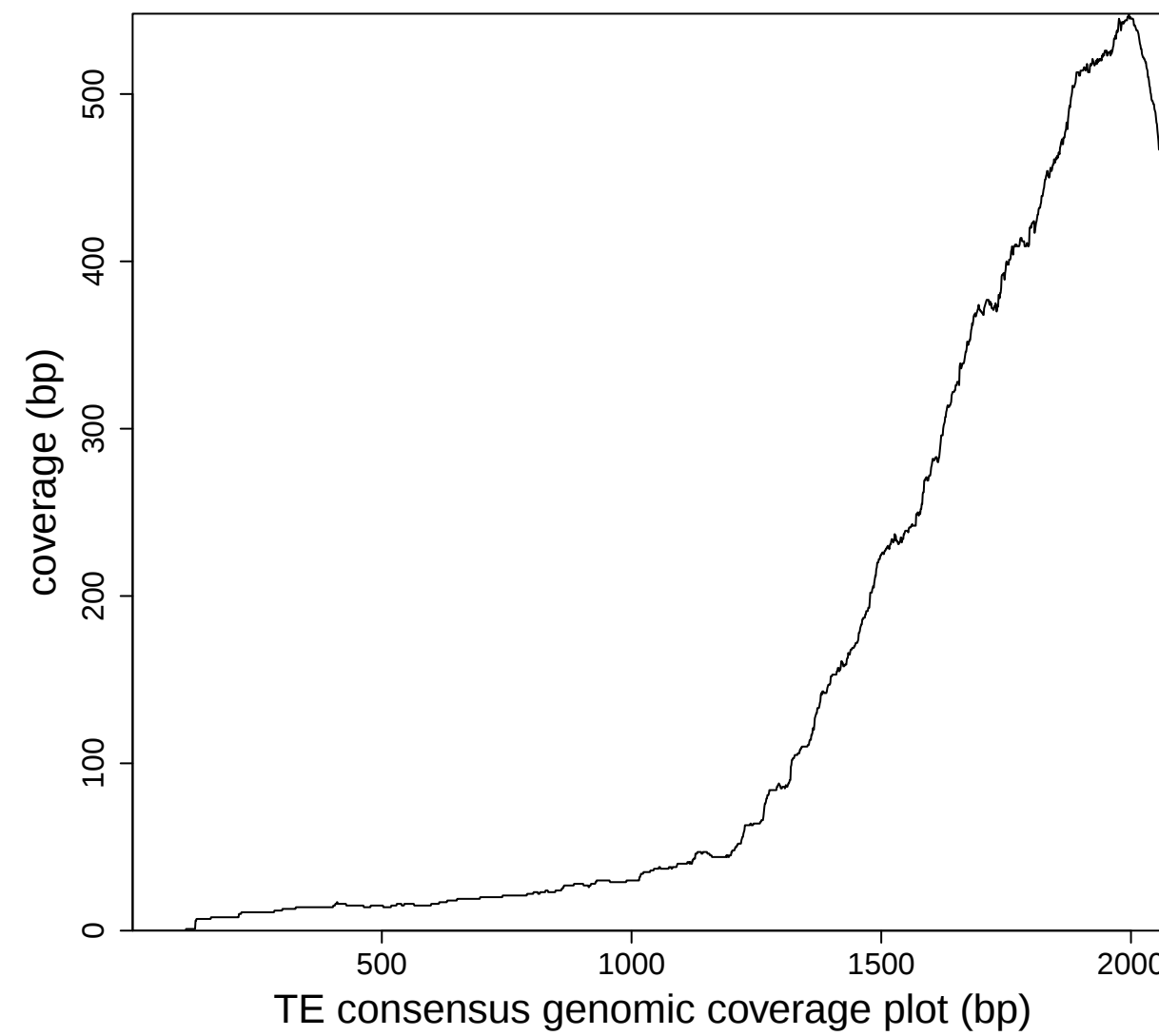
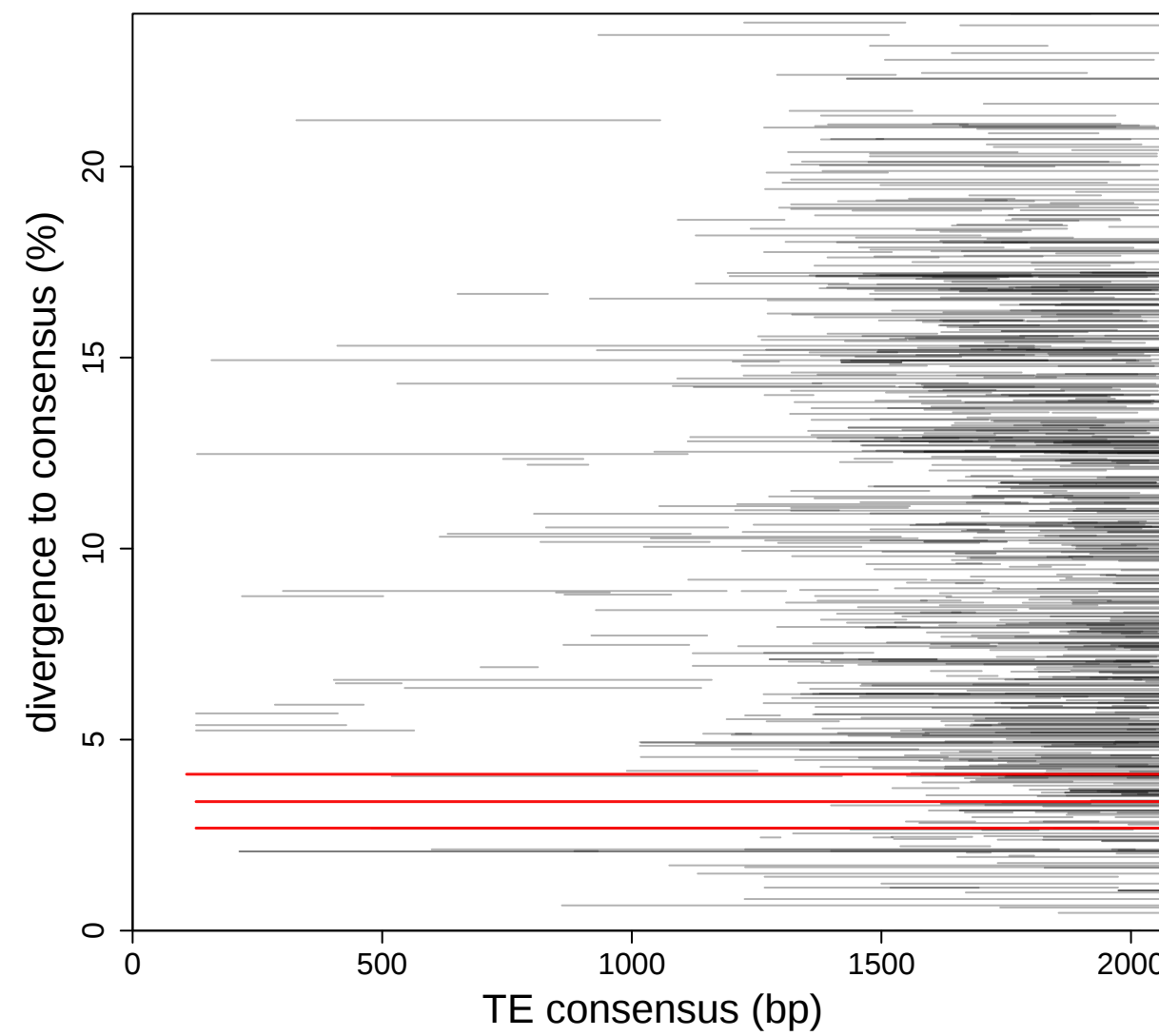
2000

2500



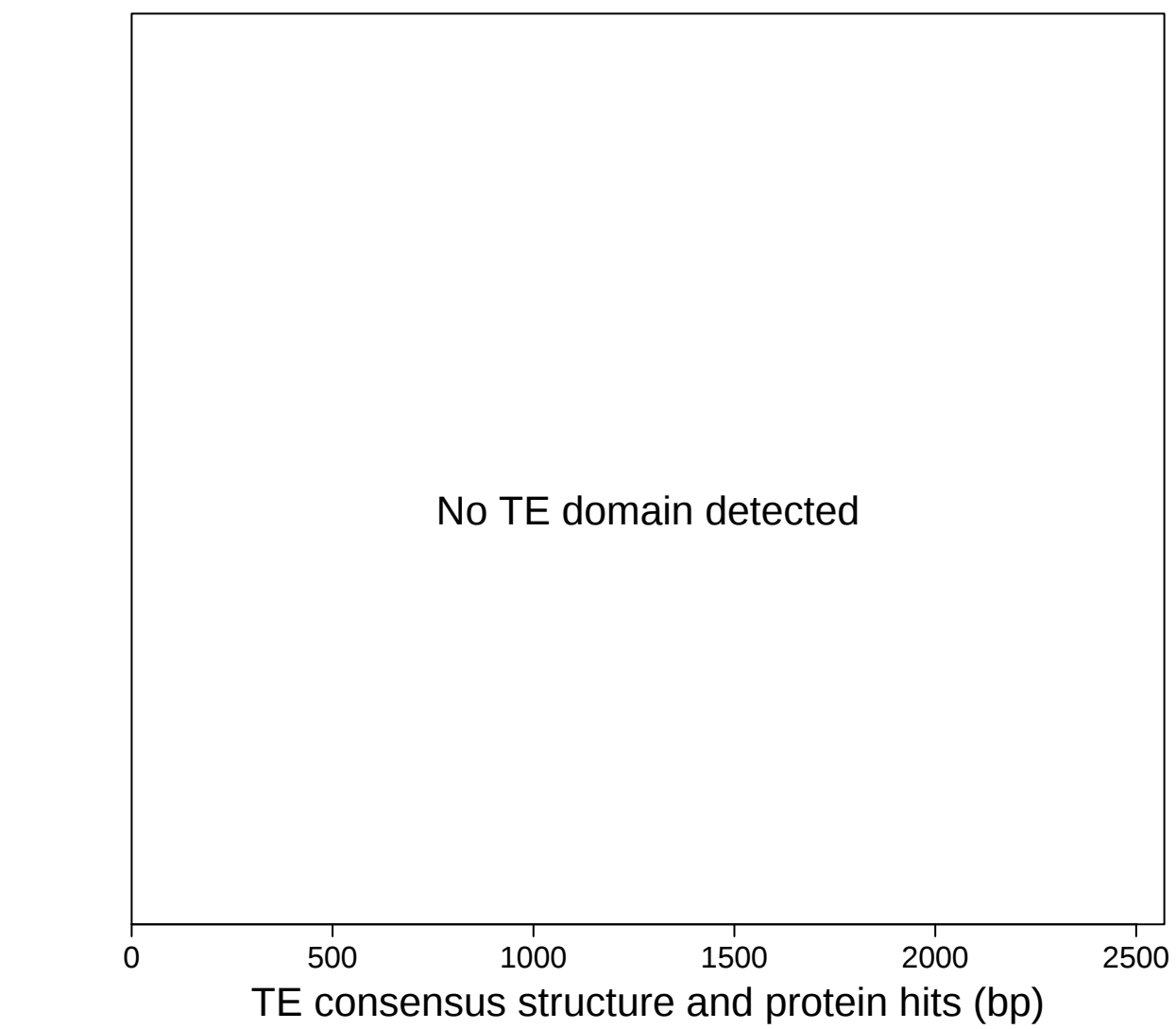
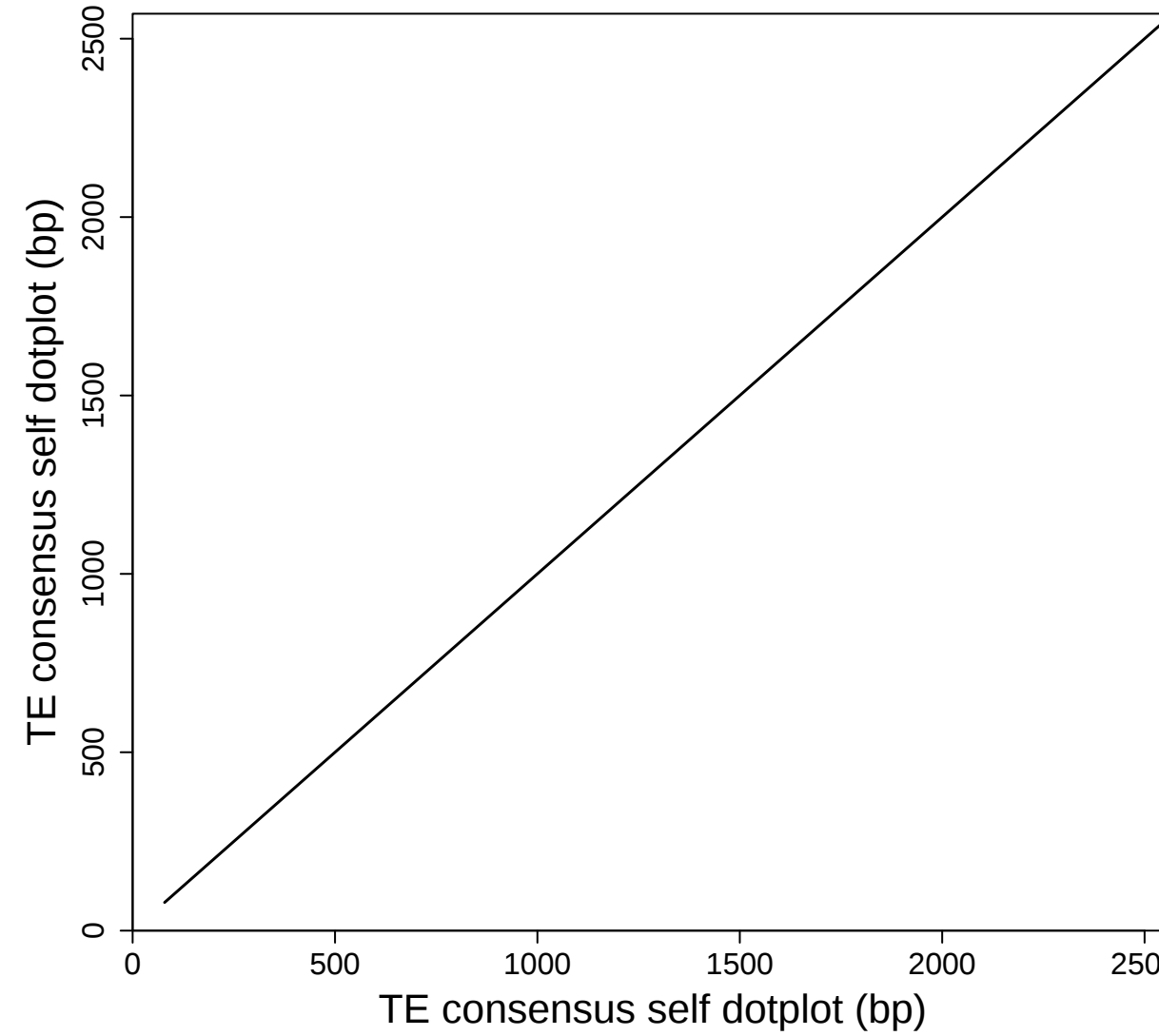
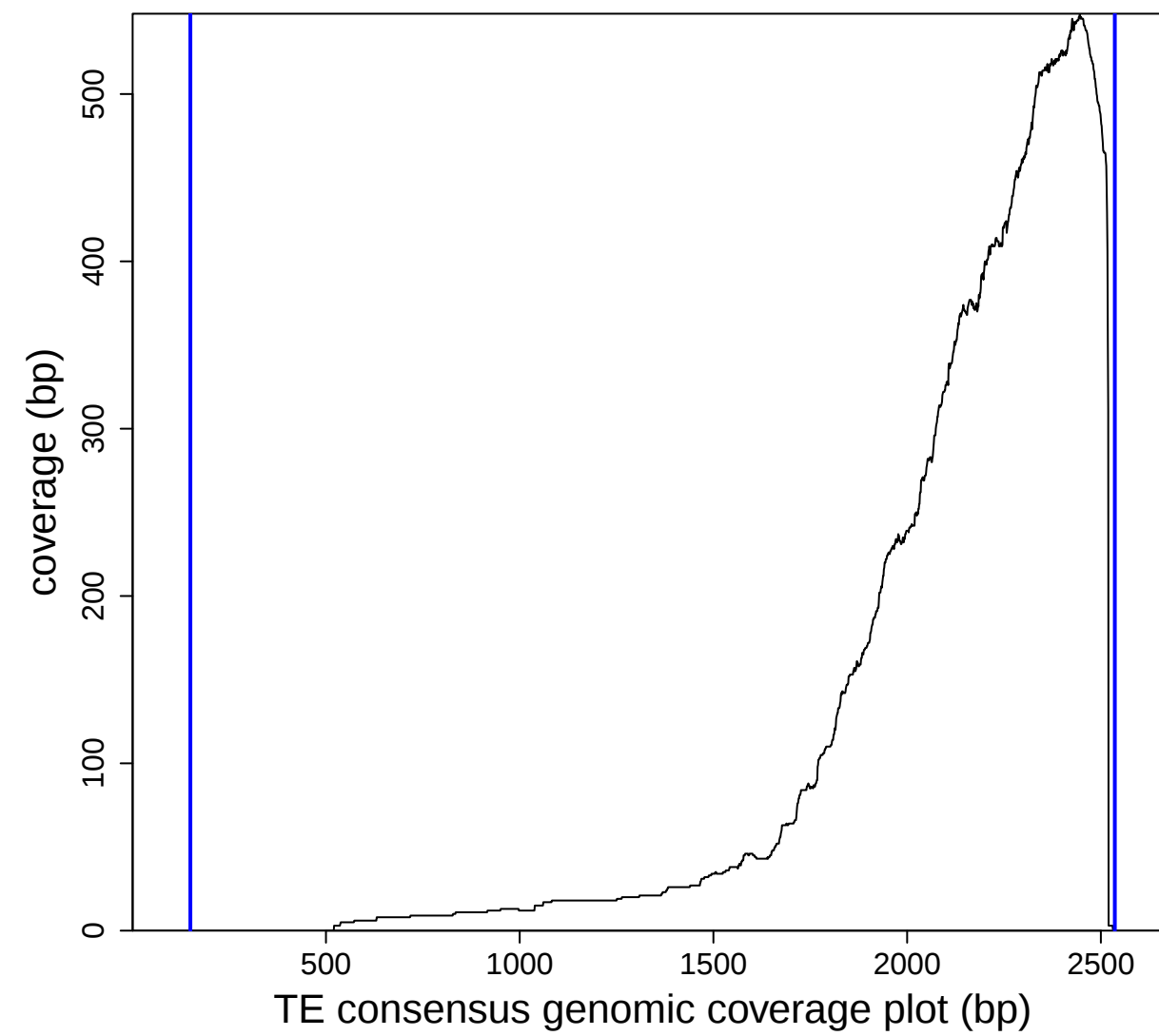
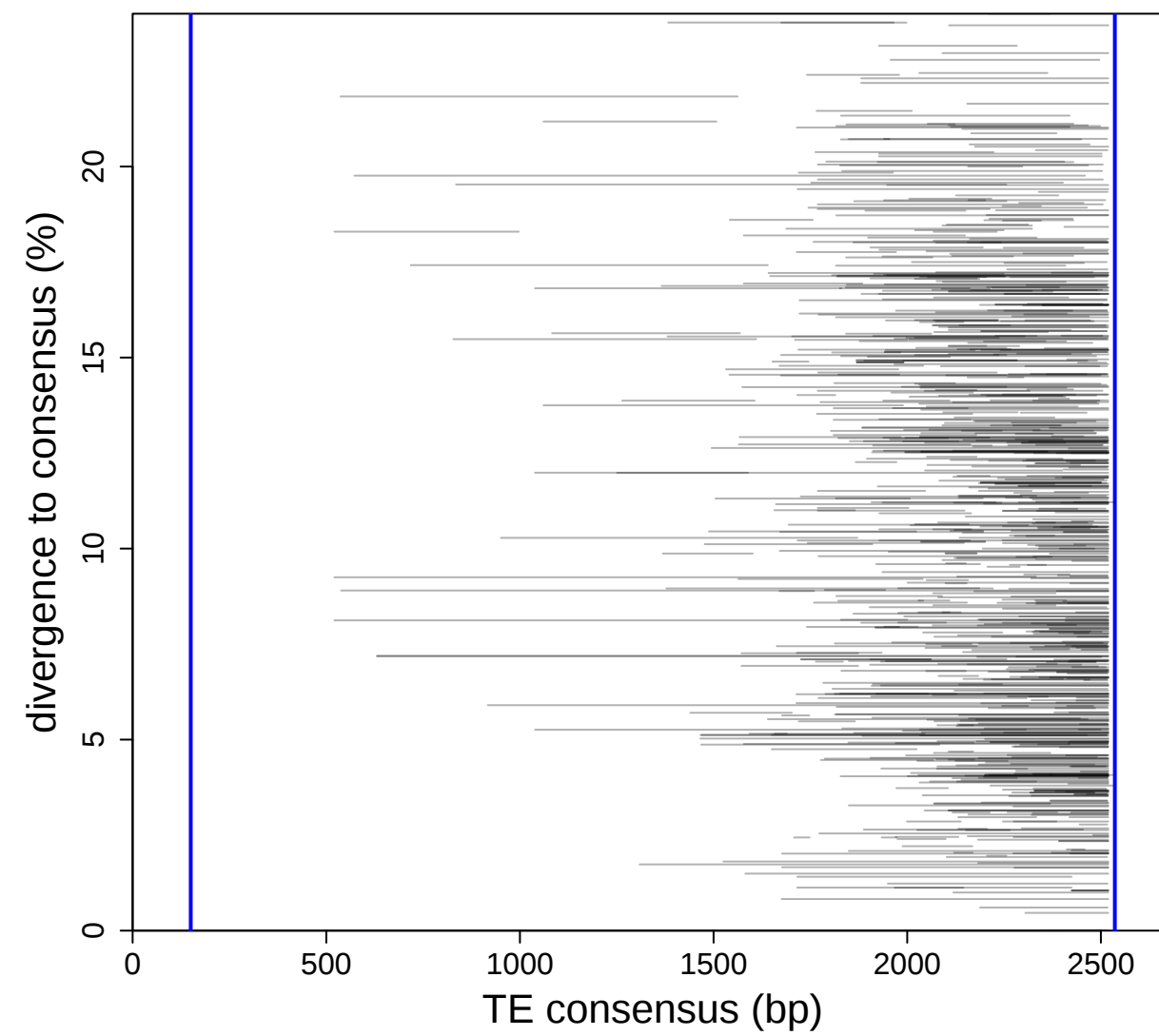
bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 2084bp; fragments: 838; full length: 3 (>=1875.6bp)

After TEtrimmer 2084 bp

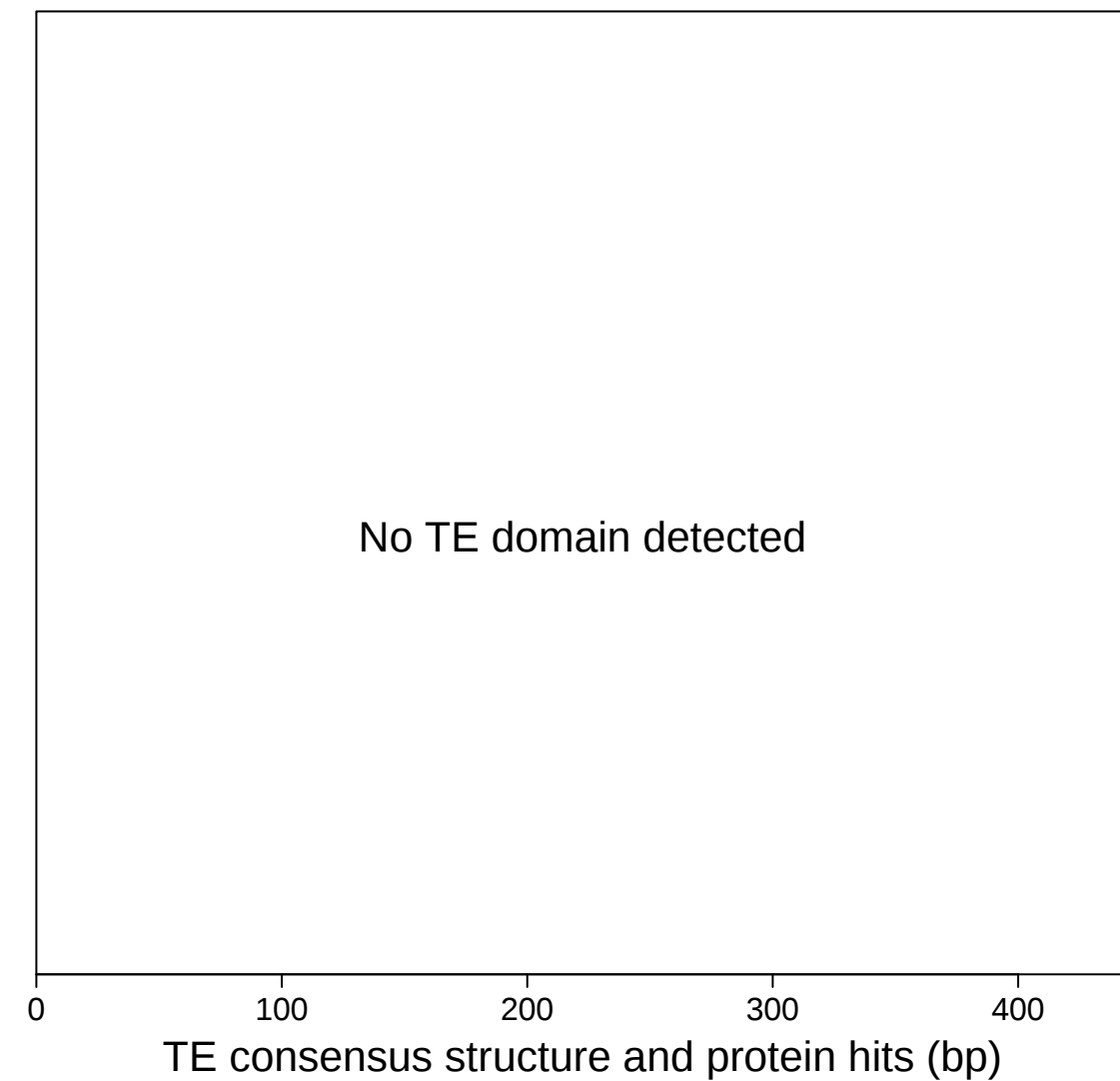
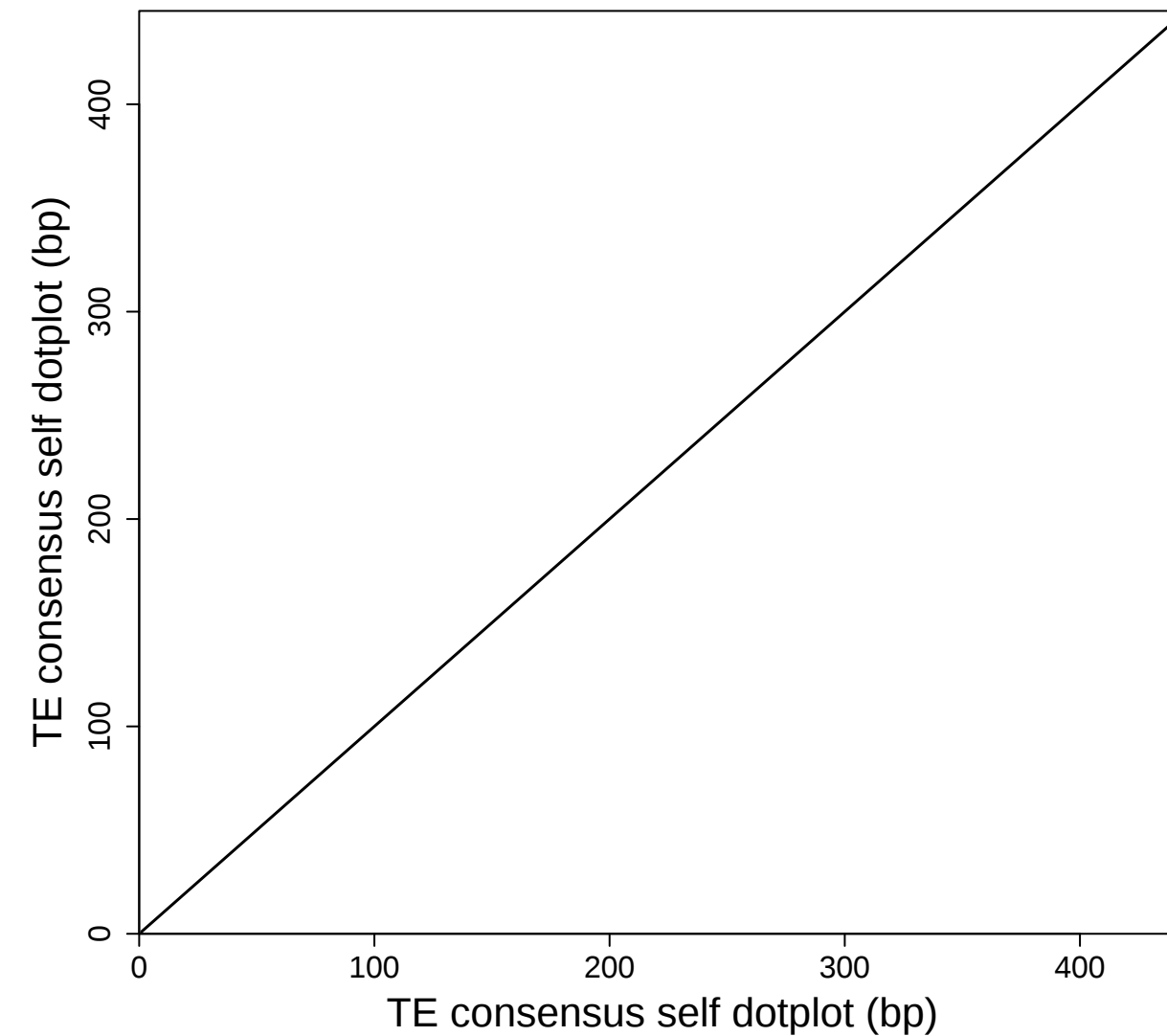
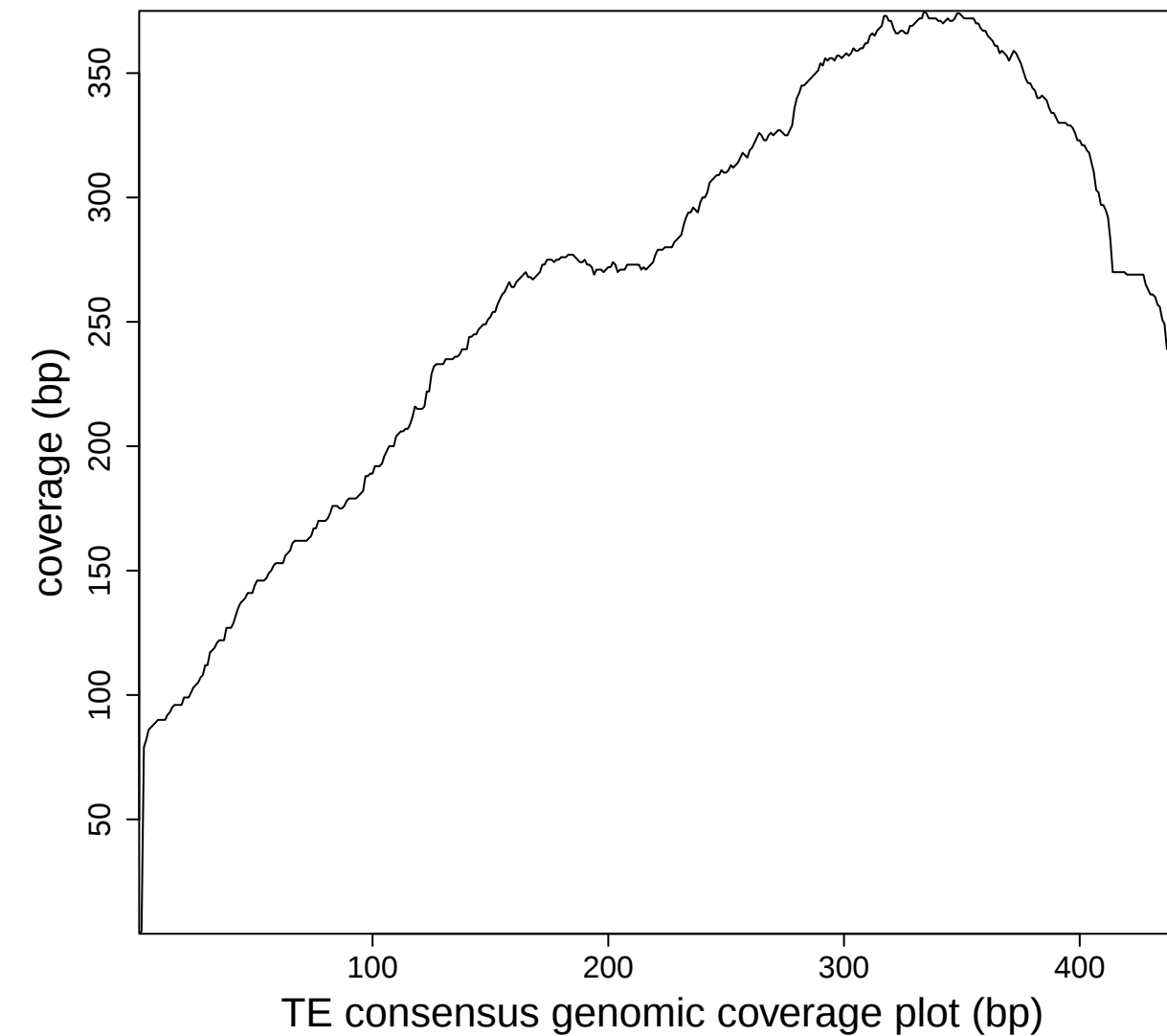
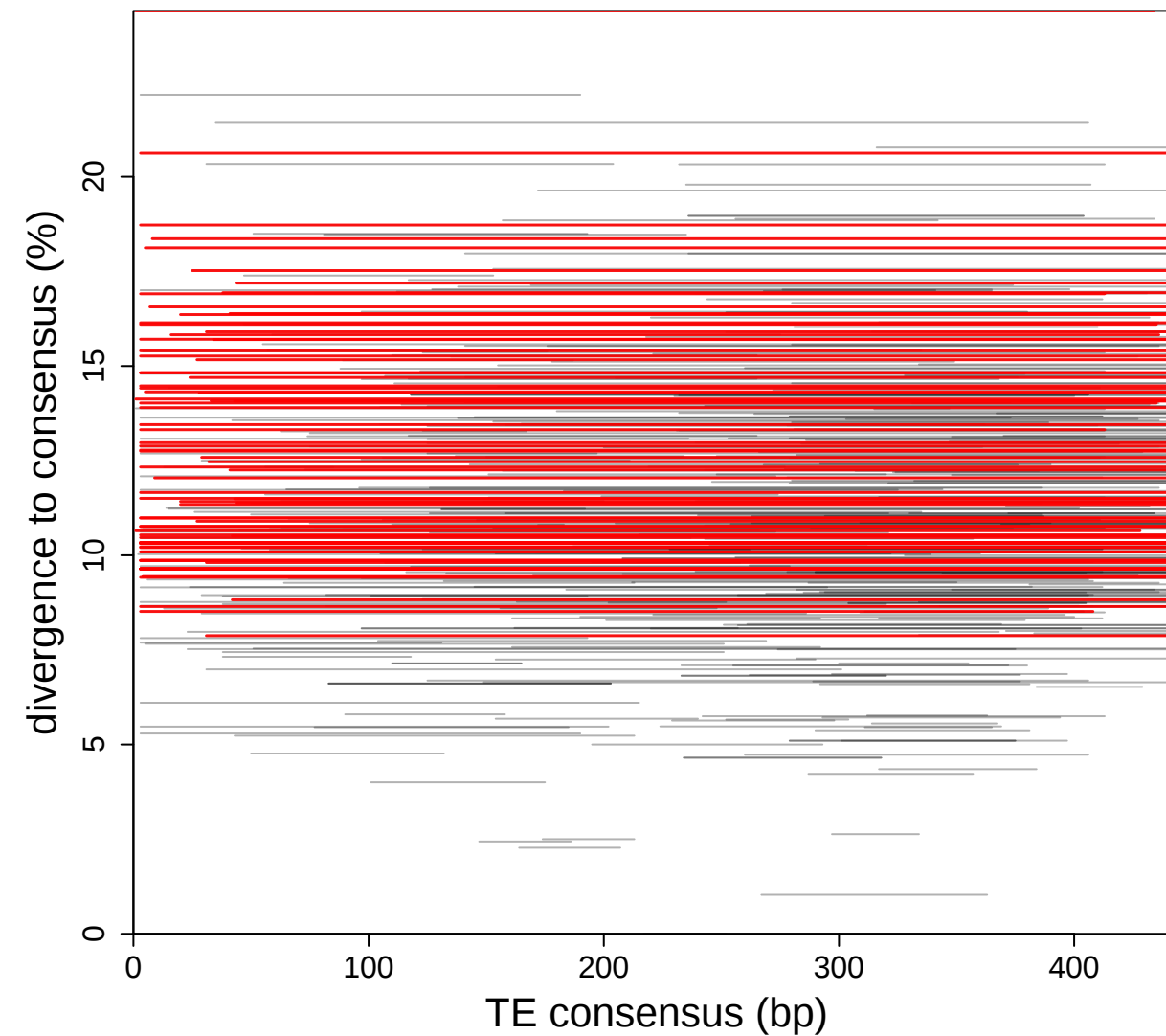


4.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 2686bp; fragments: 824; full length: 0 (>=2417.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

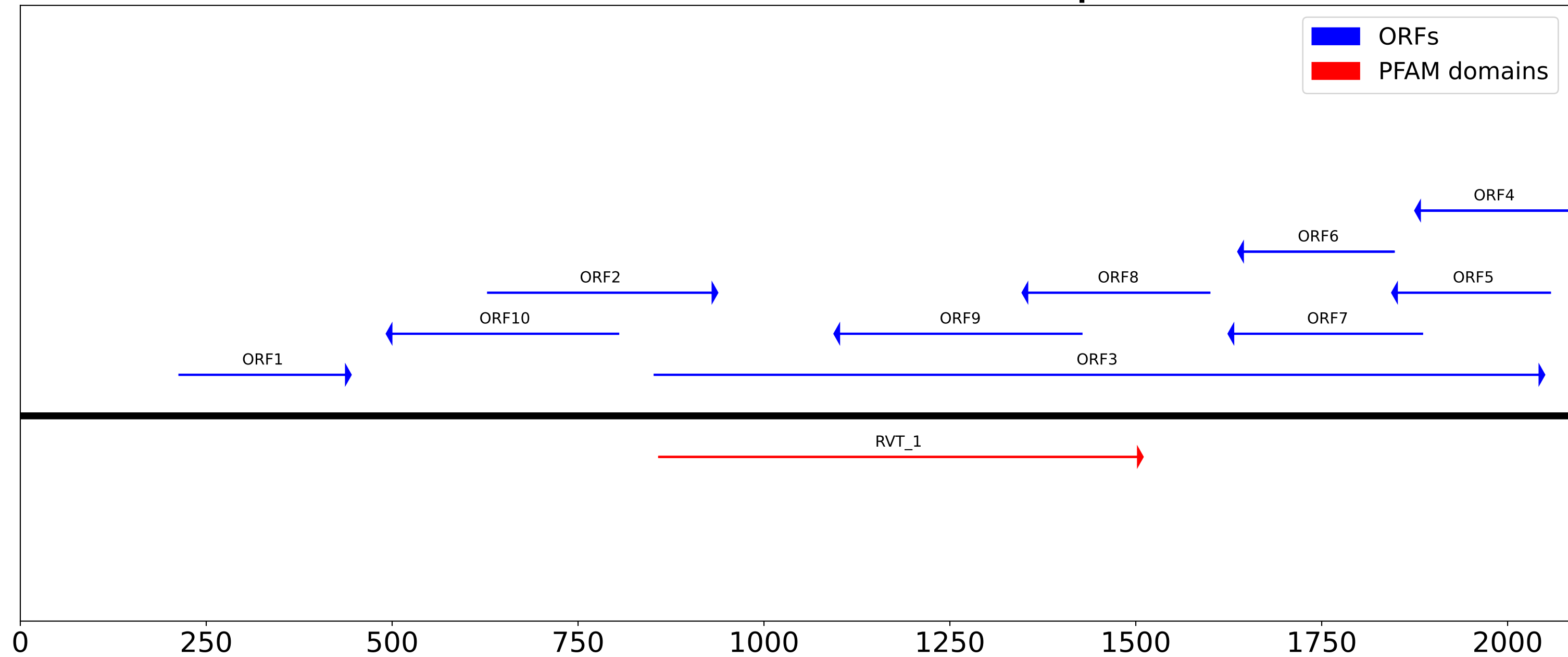


TE: rnd_5_family_8974
size: 445bp; fragments: 473; full length: 86 (≥ 400.5 bp)

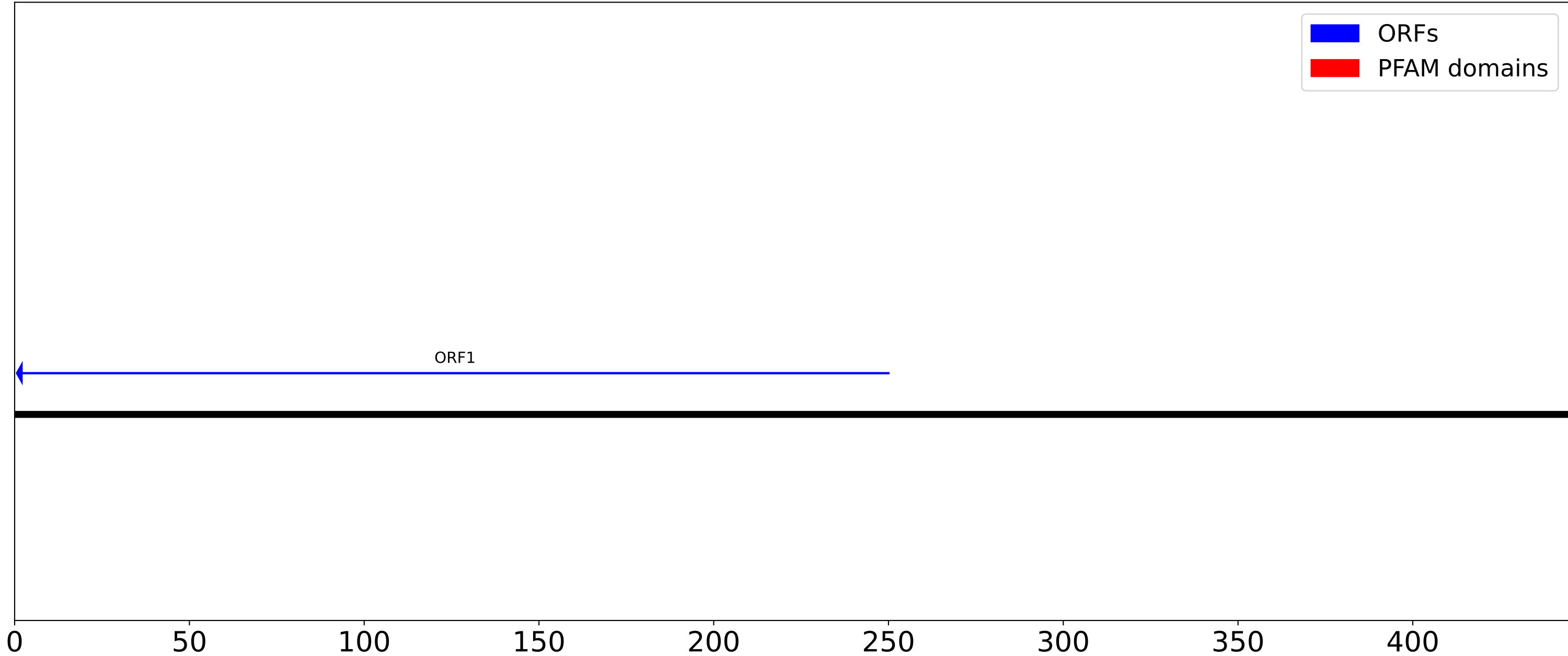


Before TEtrimmer 445 bp

After TEtrimmer ORF and PFAM domain plot



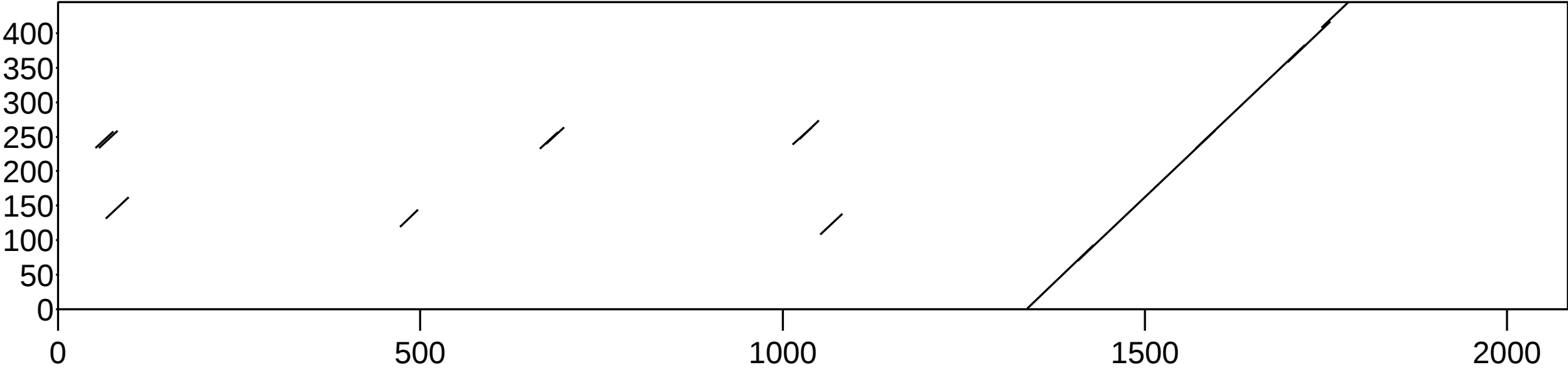
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

onsensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)