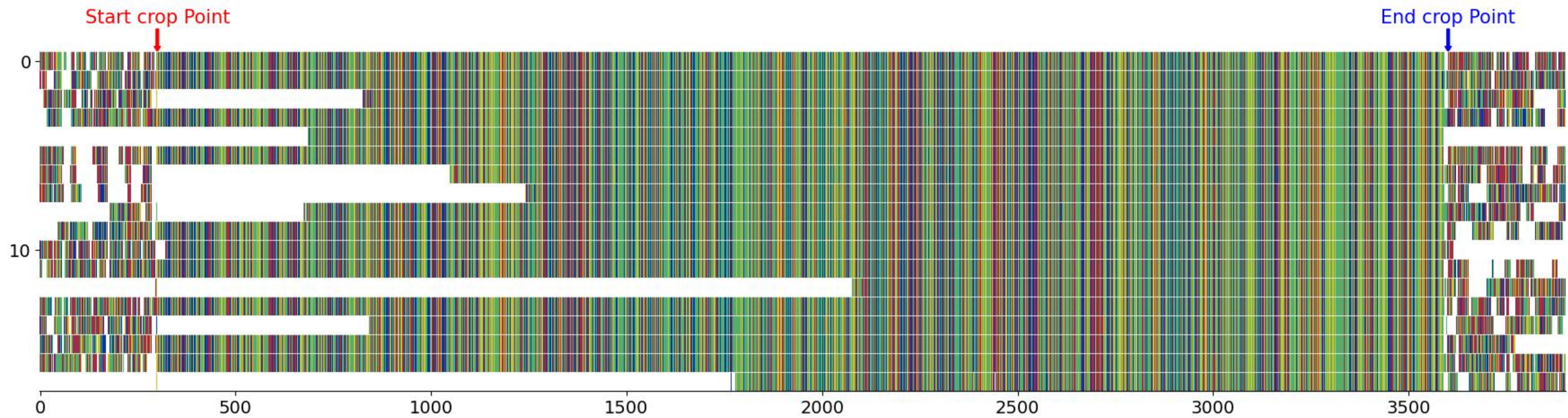


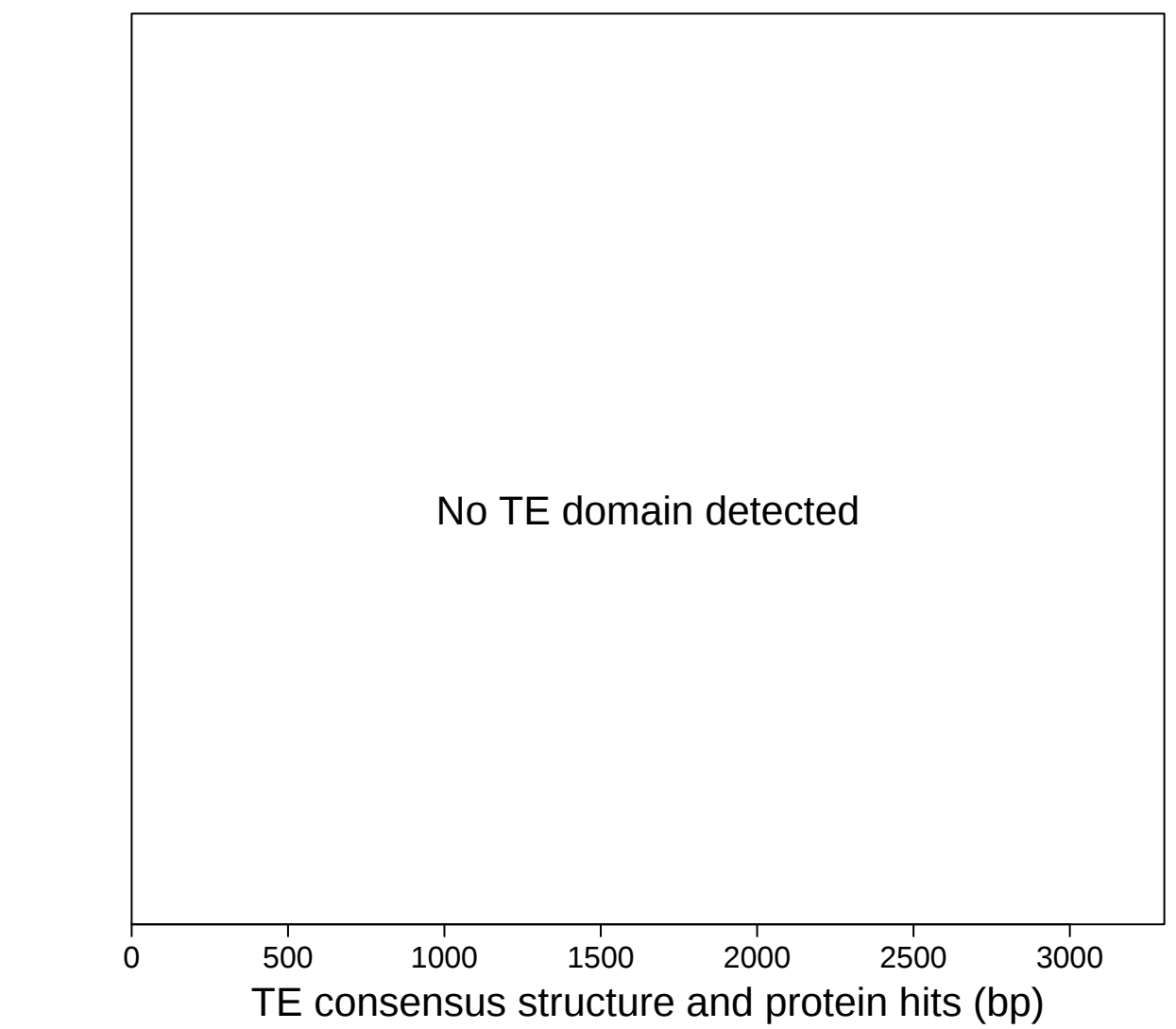
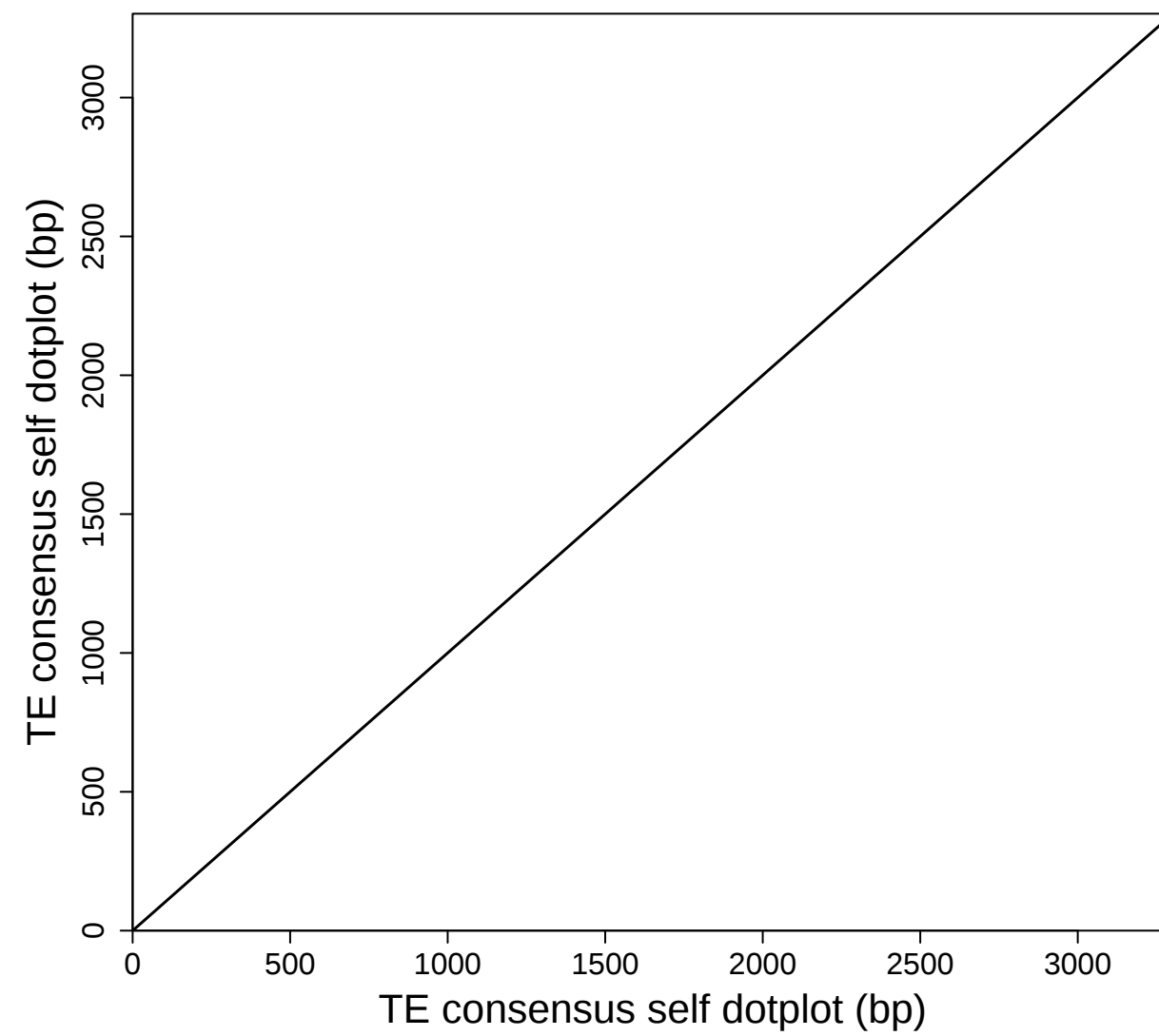
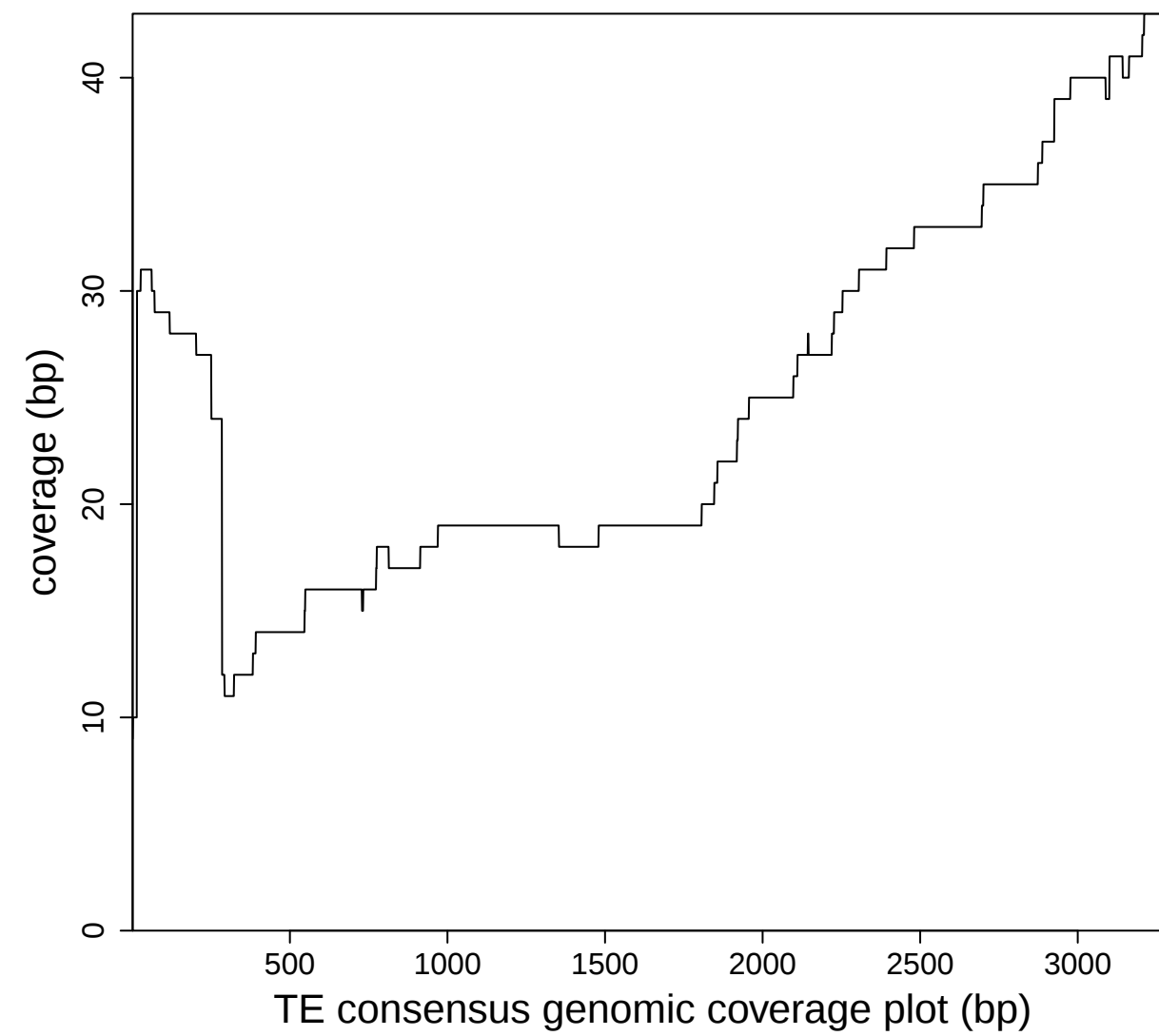
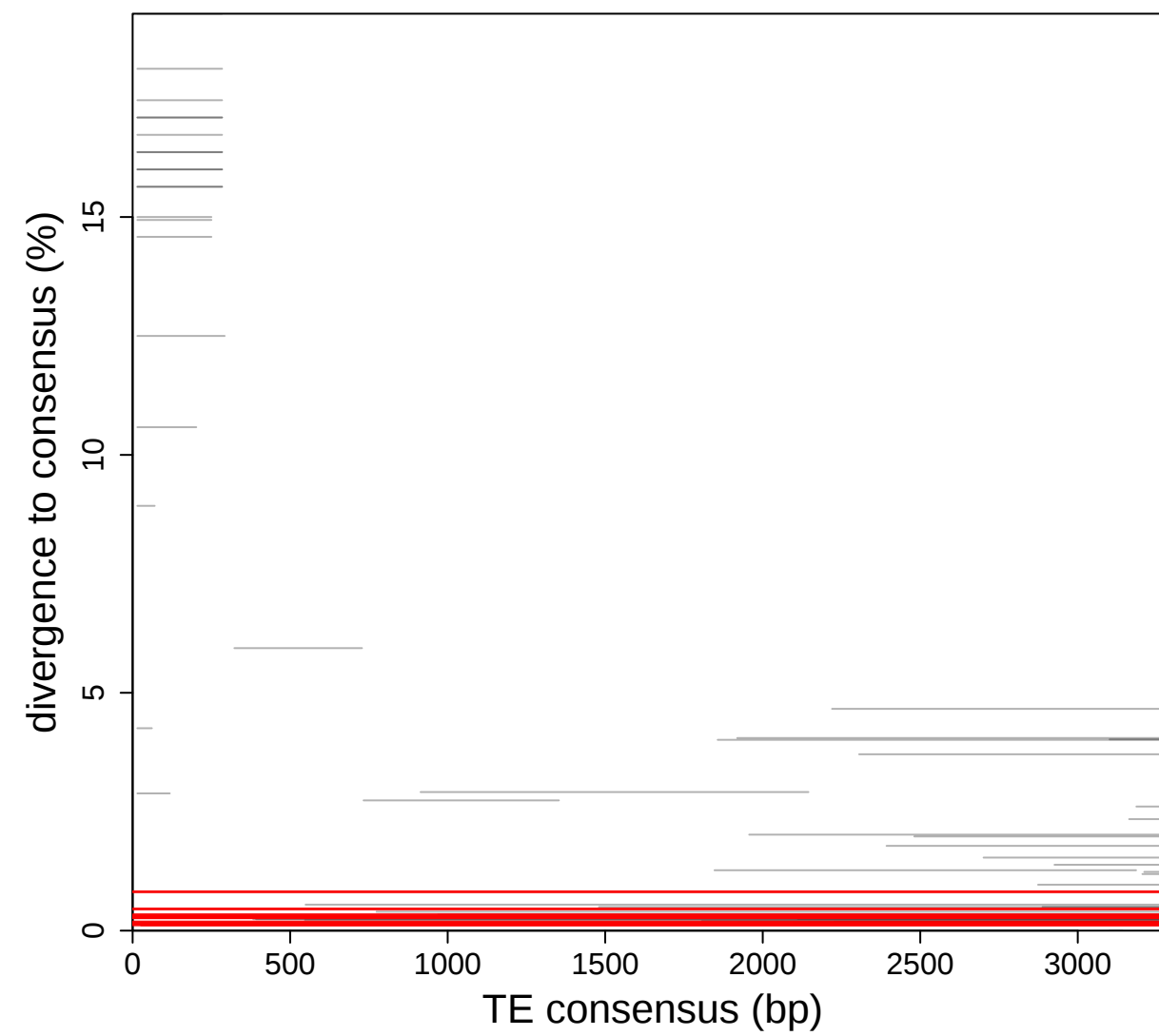
Start crop Point

End crop Point



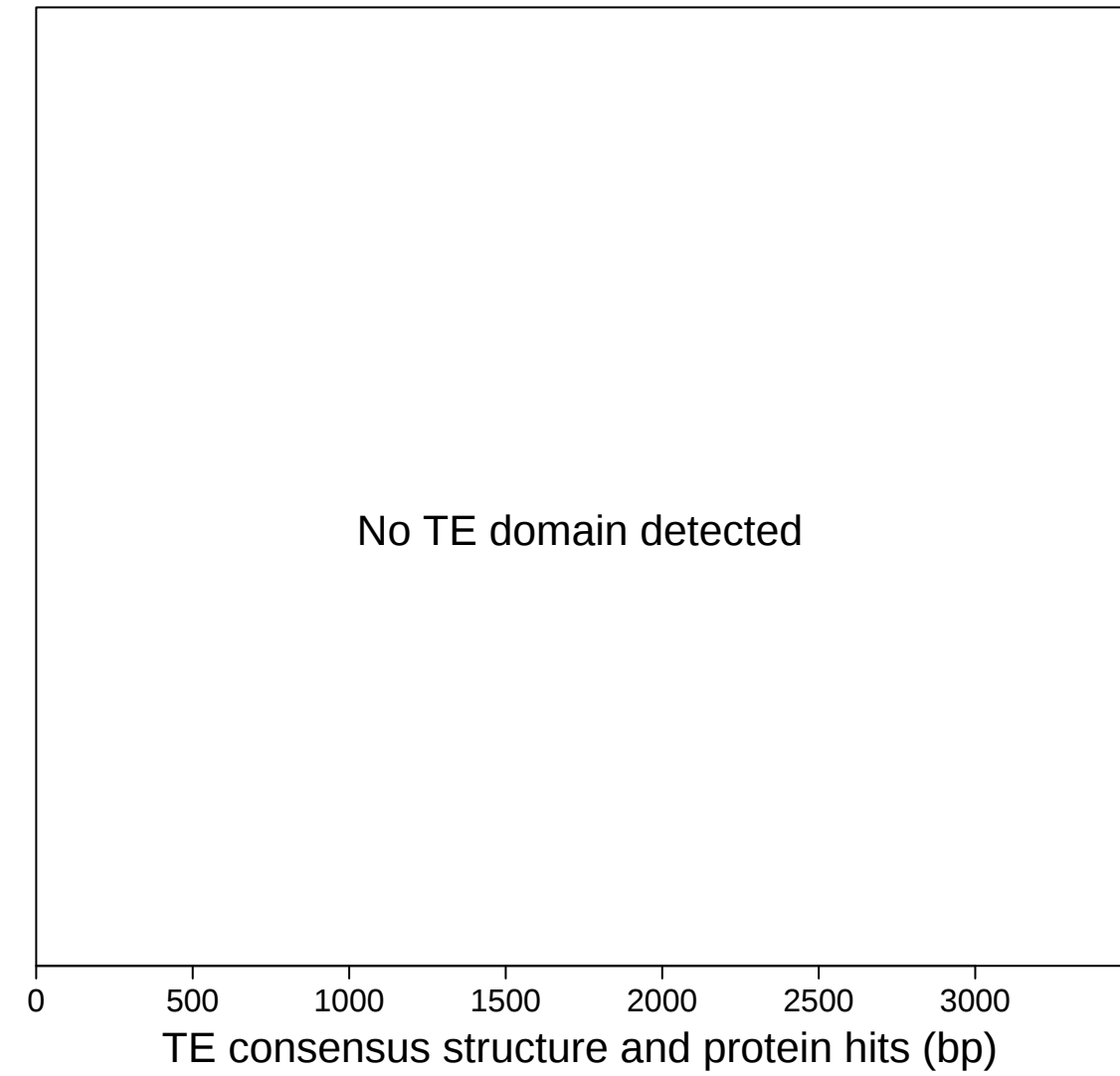
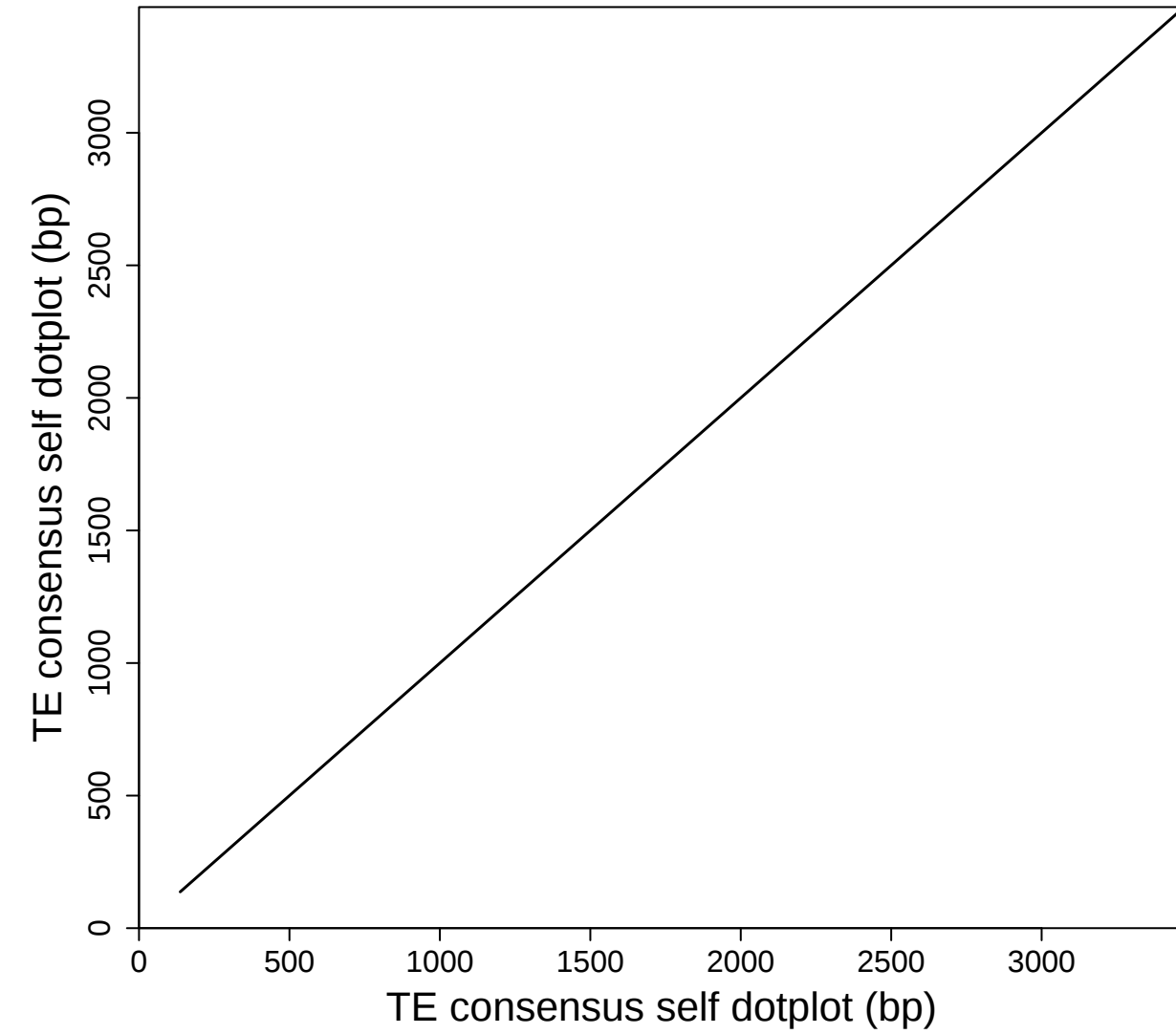
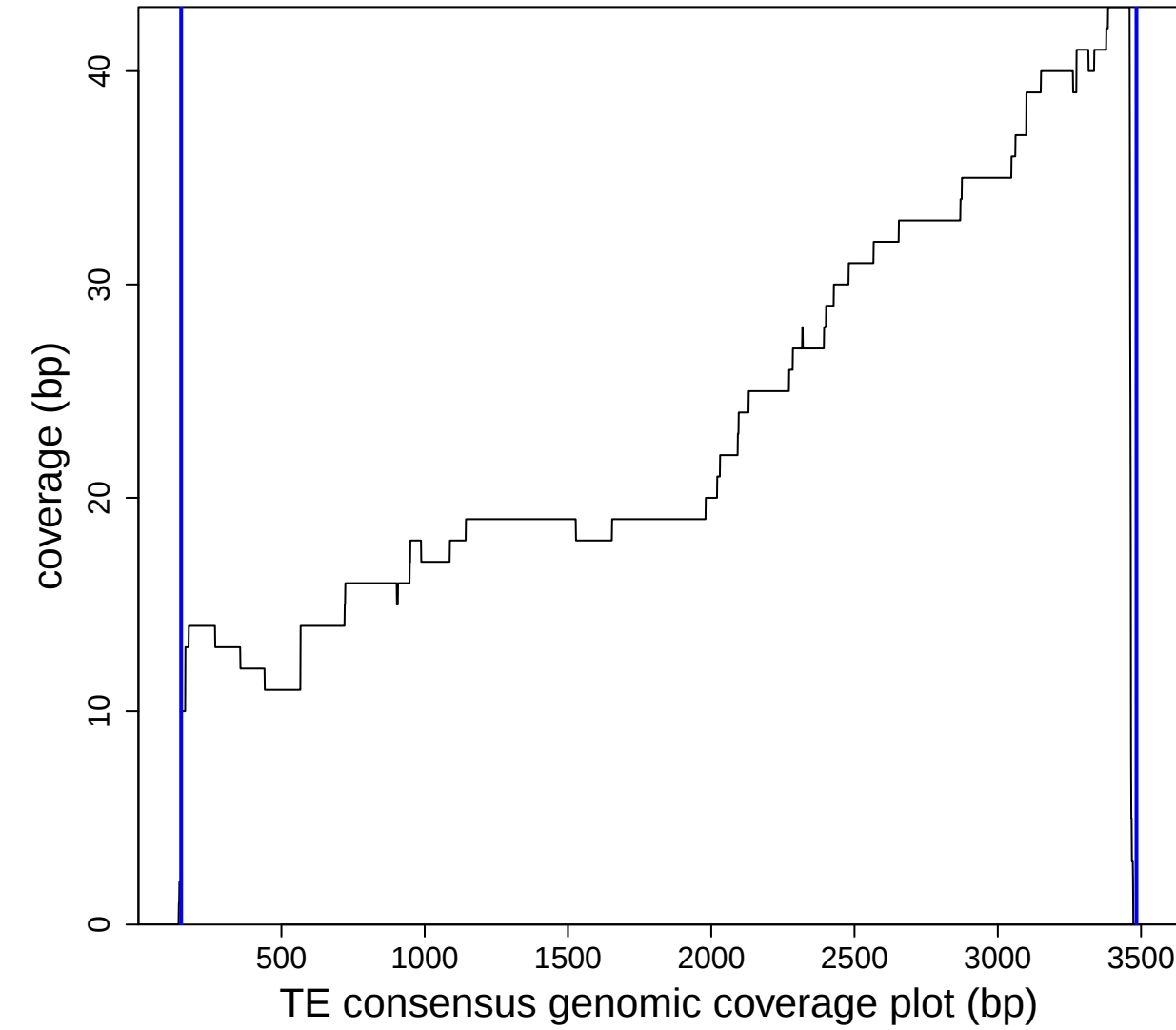
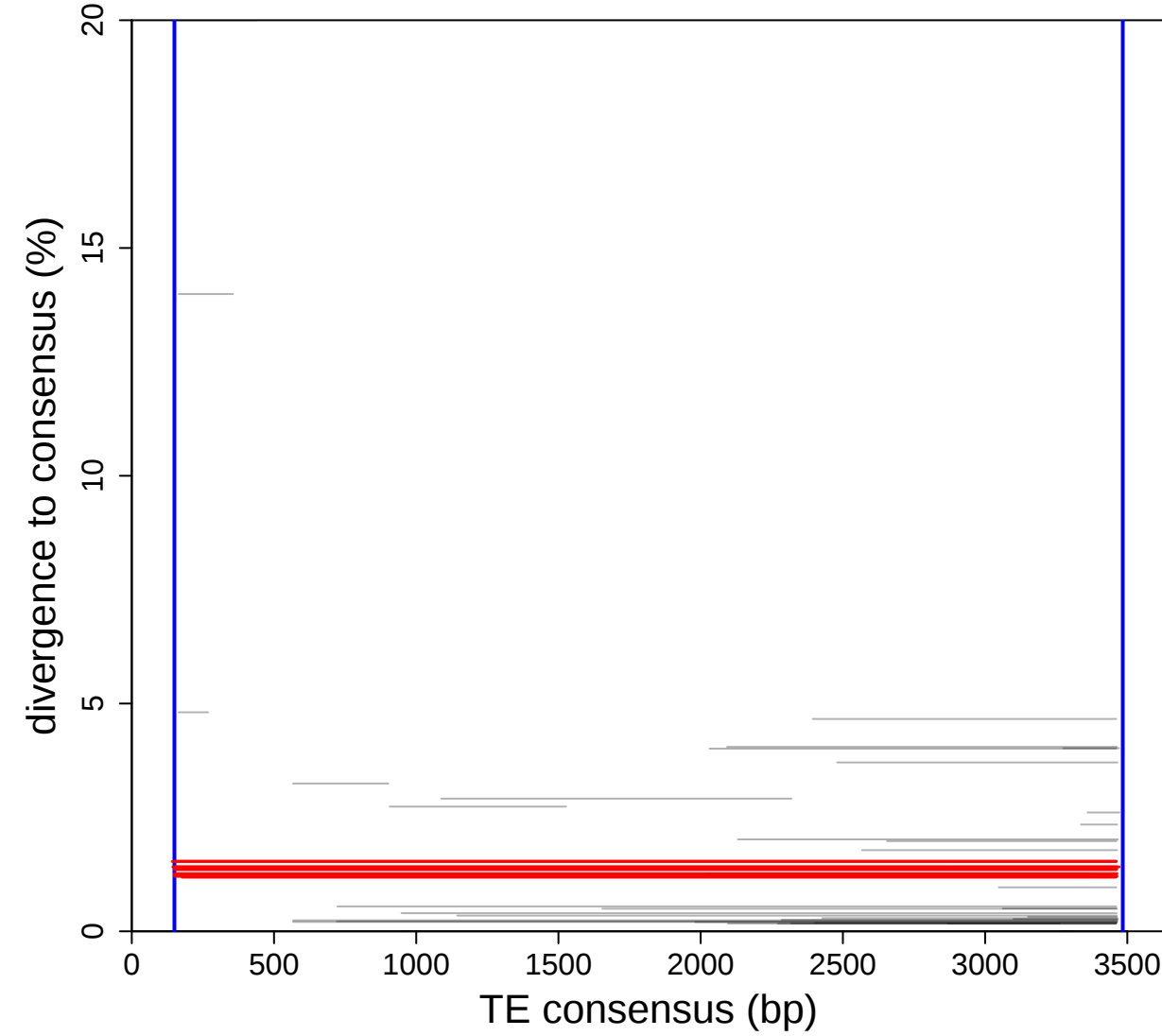
bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 3302bp; fragments: 70; full length: 11 (>=2971.8bp)

After TETrimmer 3302 bp



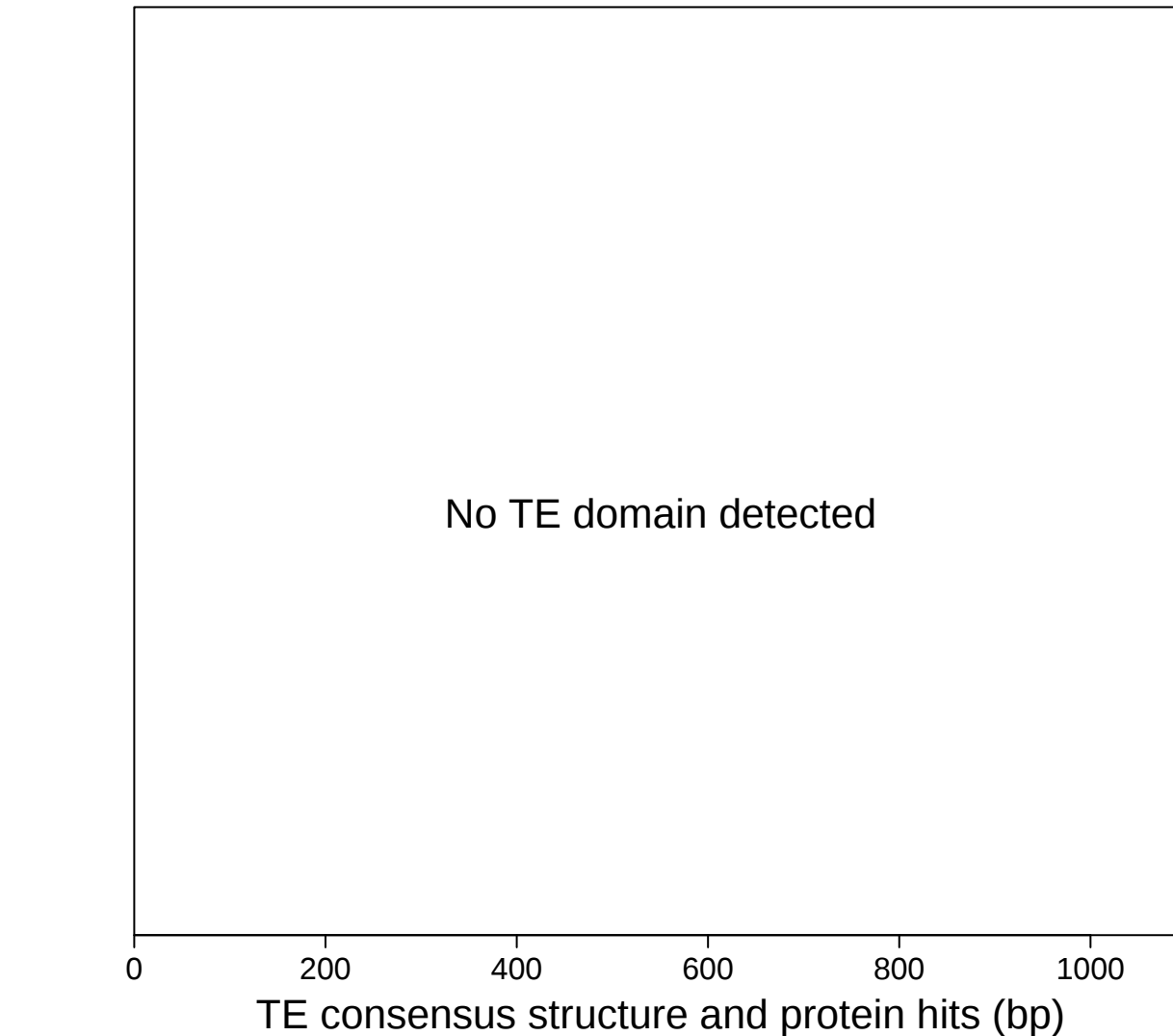
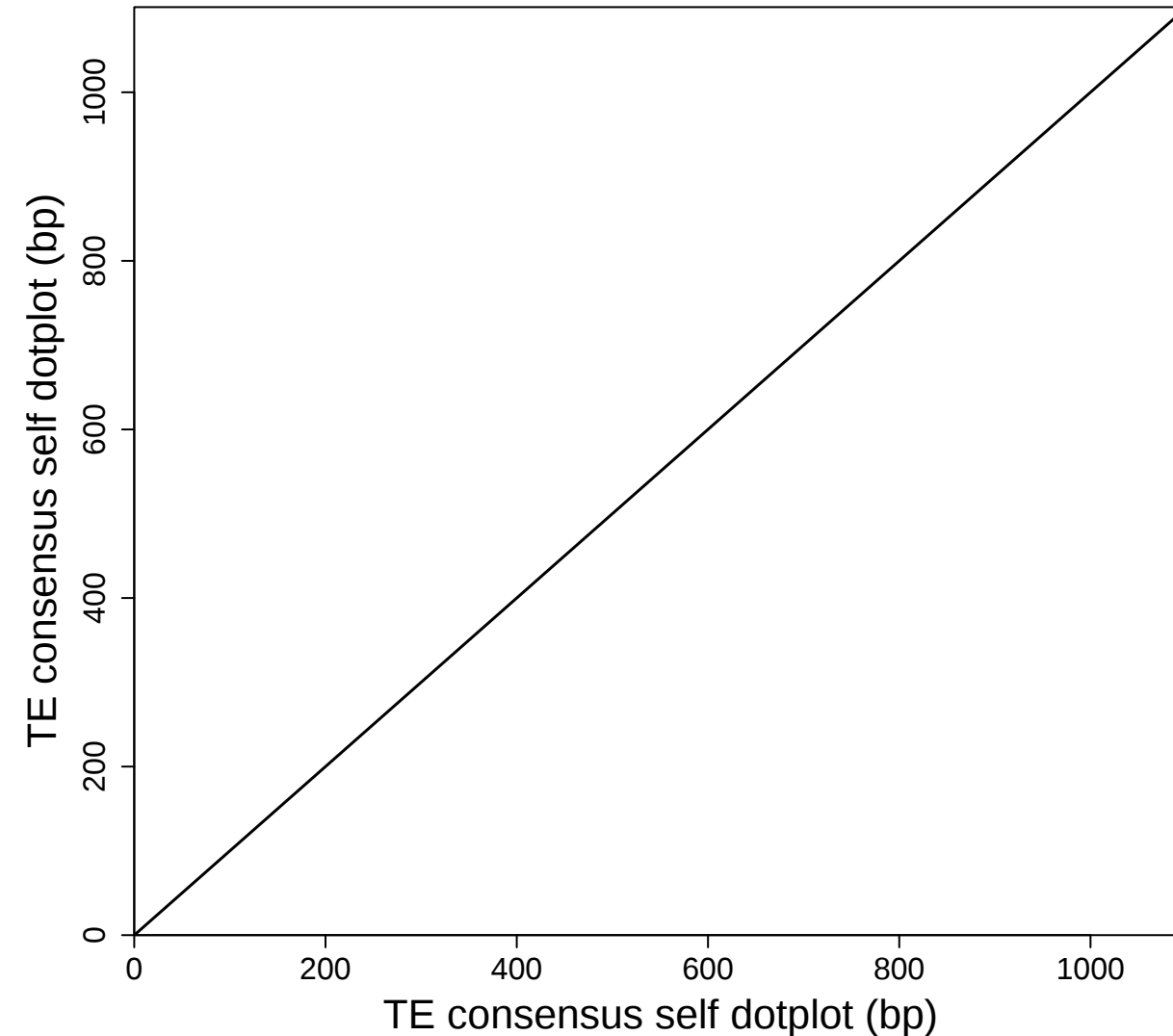
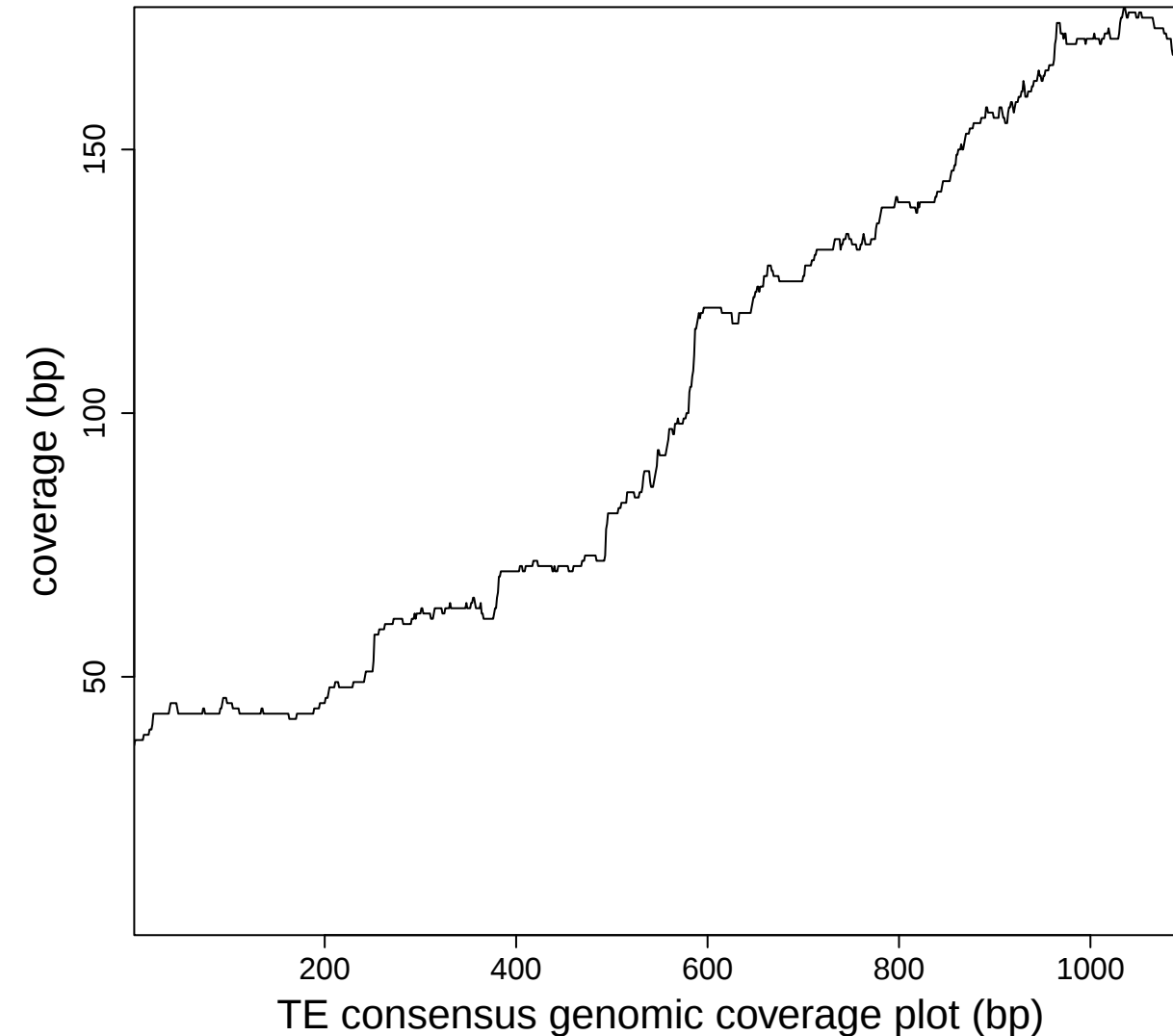
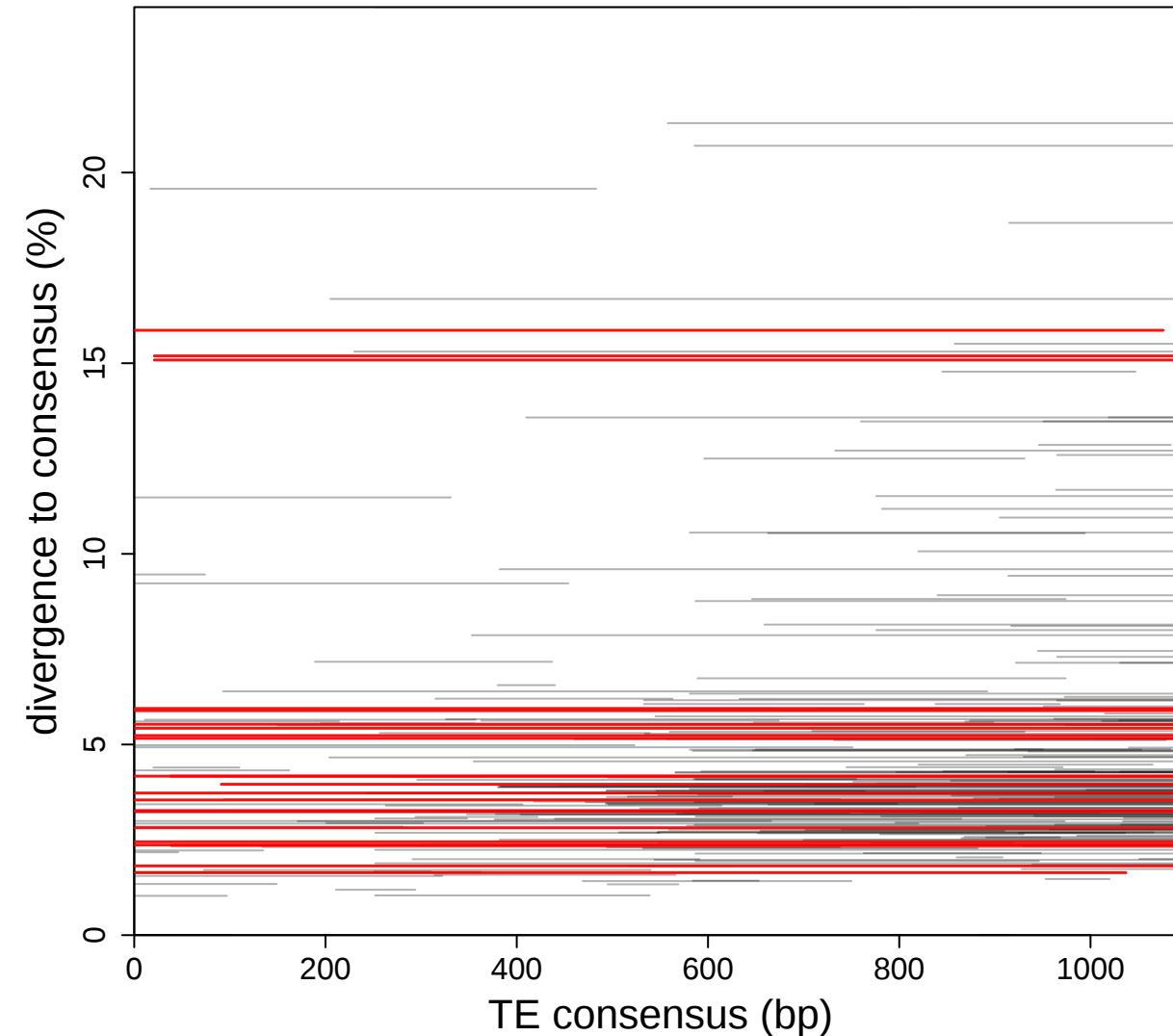
3.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.f
size: 3634bp; fragments: 53; full length: 11 (>=3270.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

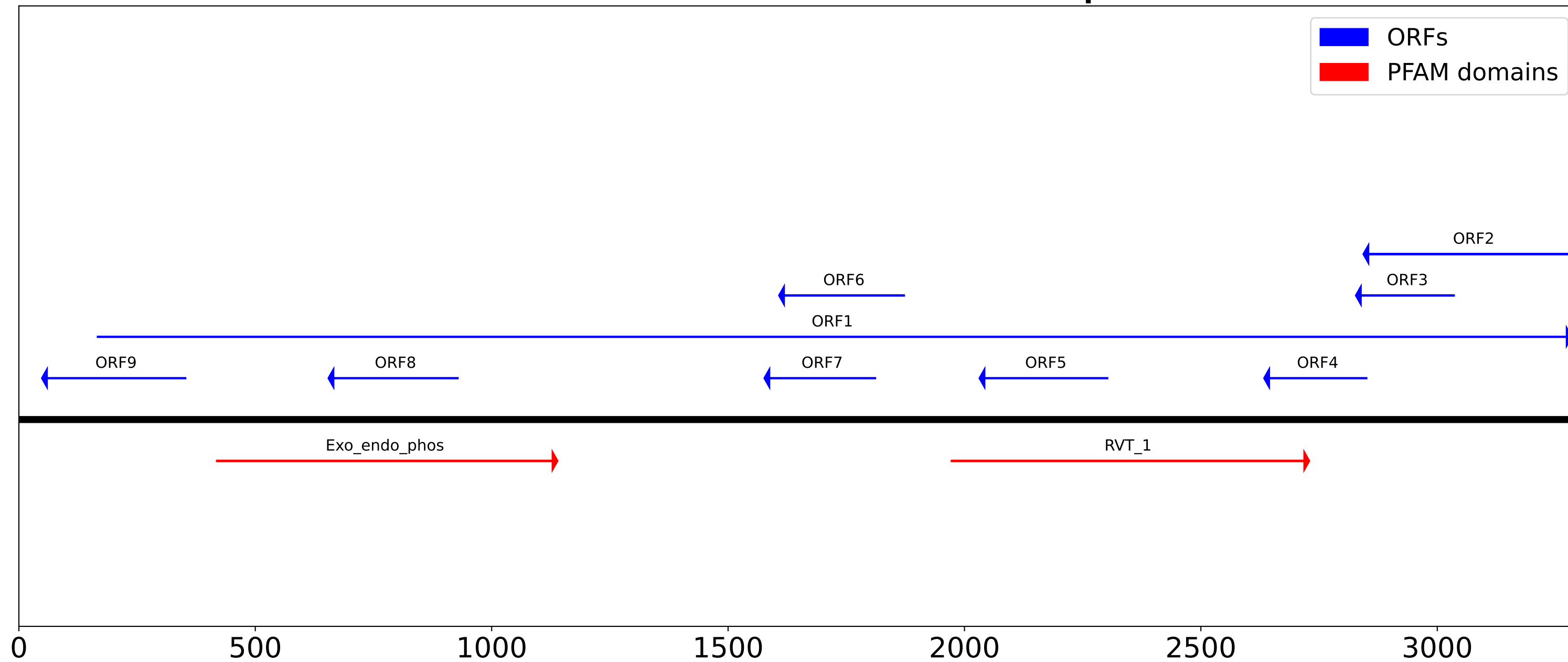


TE: rnd_5_family_1163
size: 1101bp; fragments: 261; full length: 22 (≥ 990.9 bp)

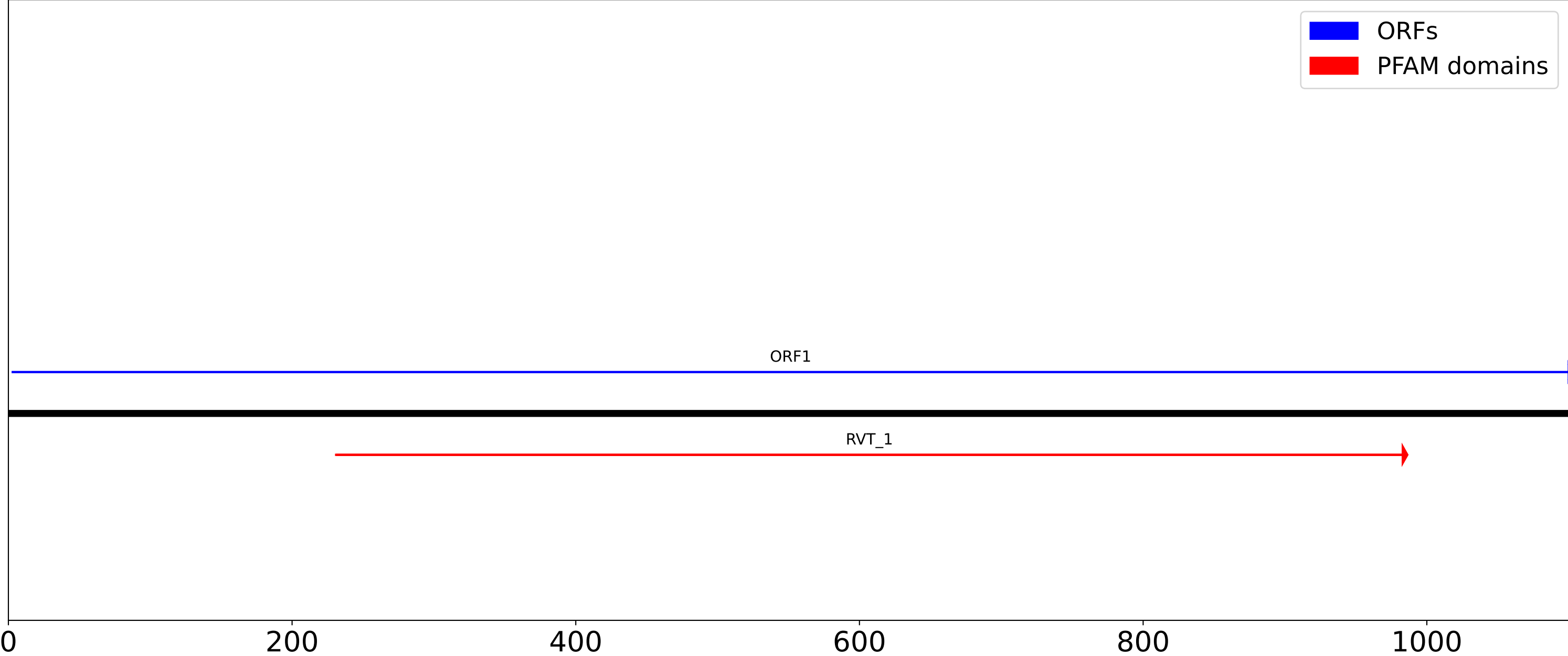
Before TEtrimmer 1101 bp



After TEtrimmer ORF and PFAM domain plot



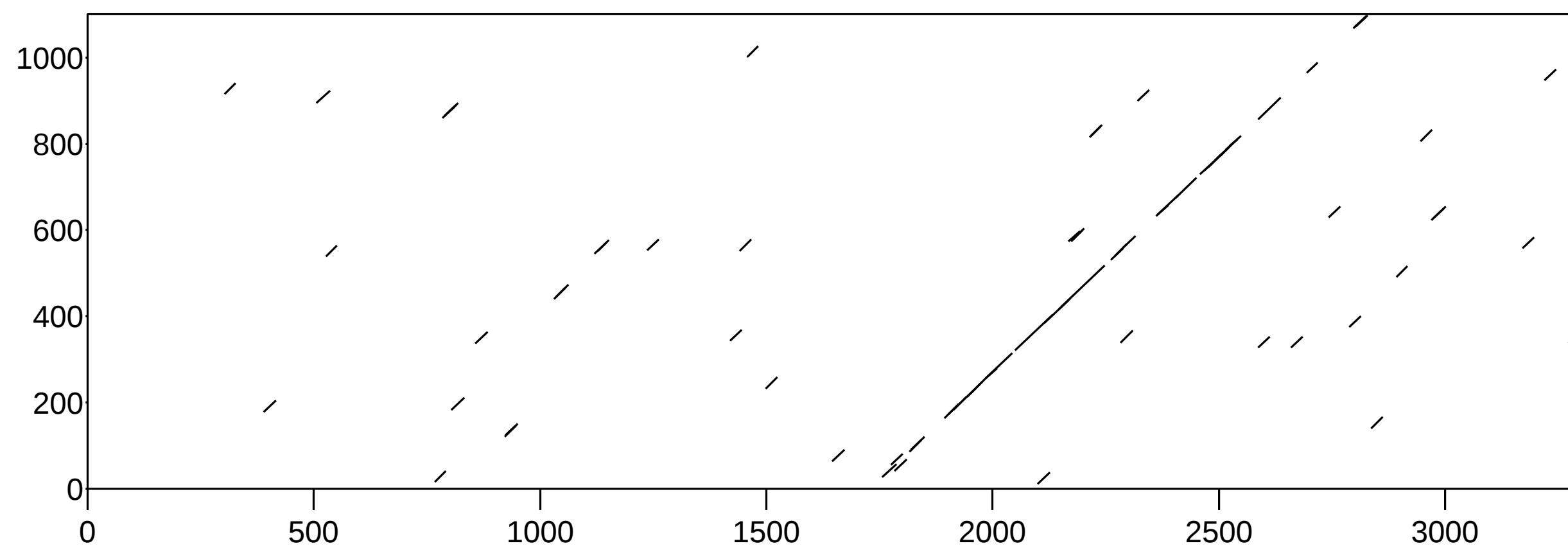
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)