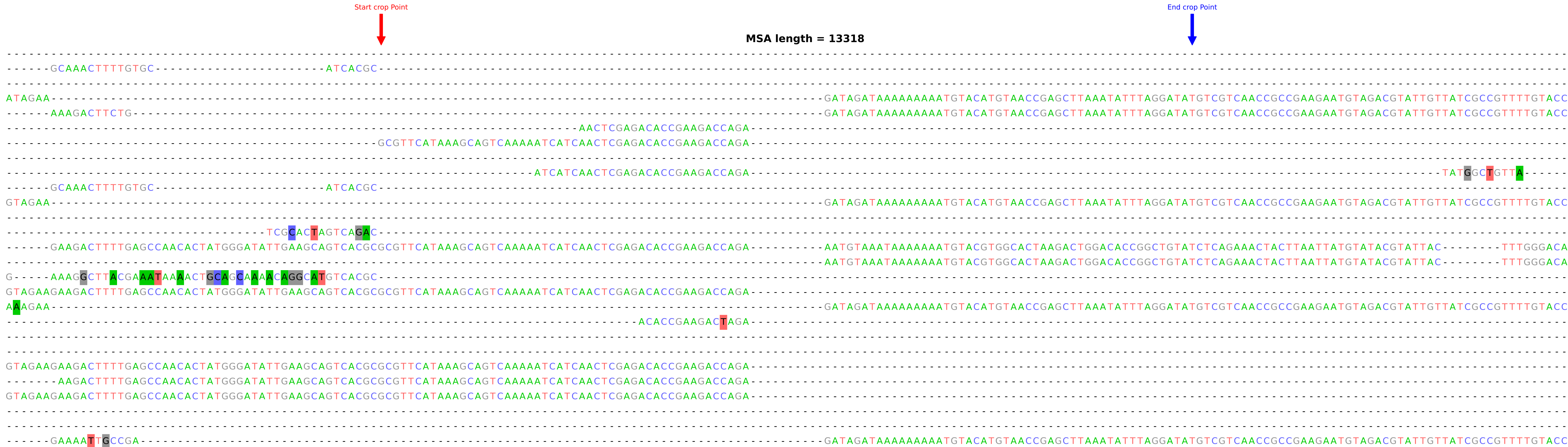
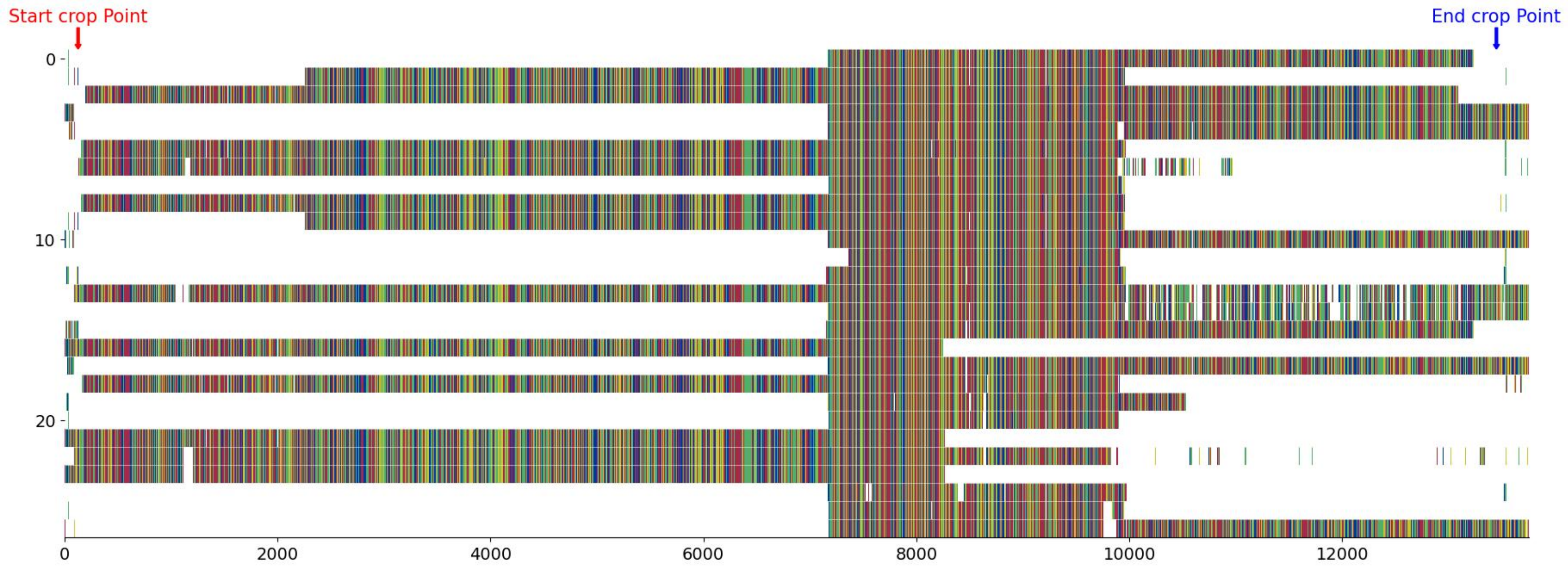
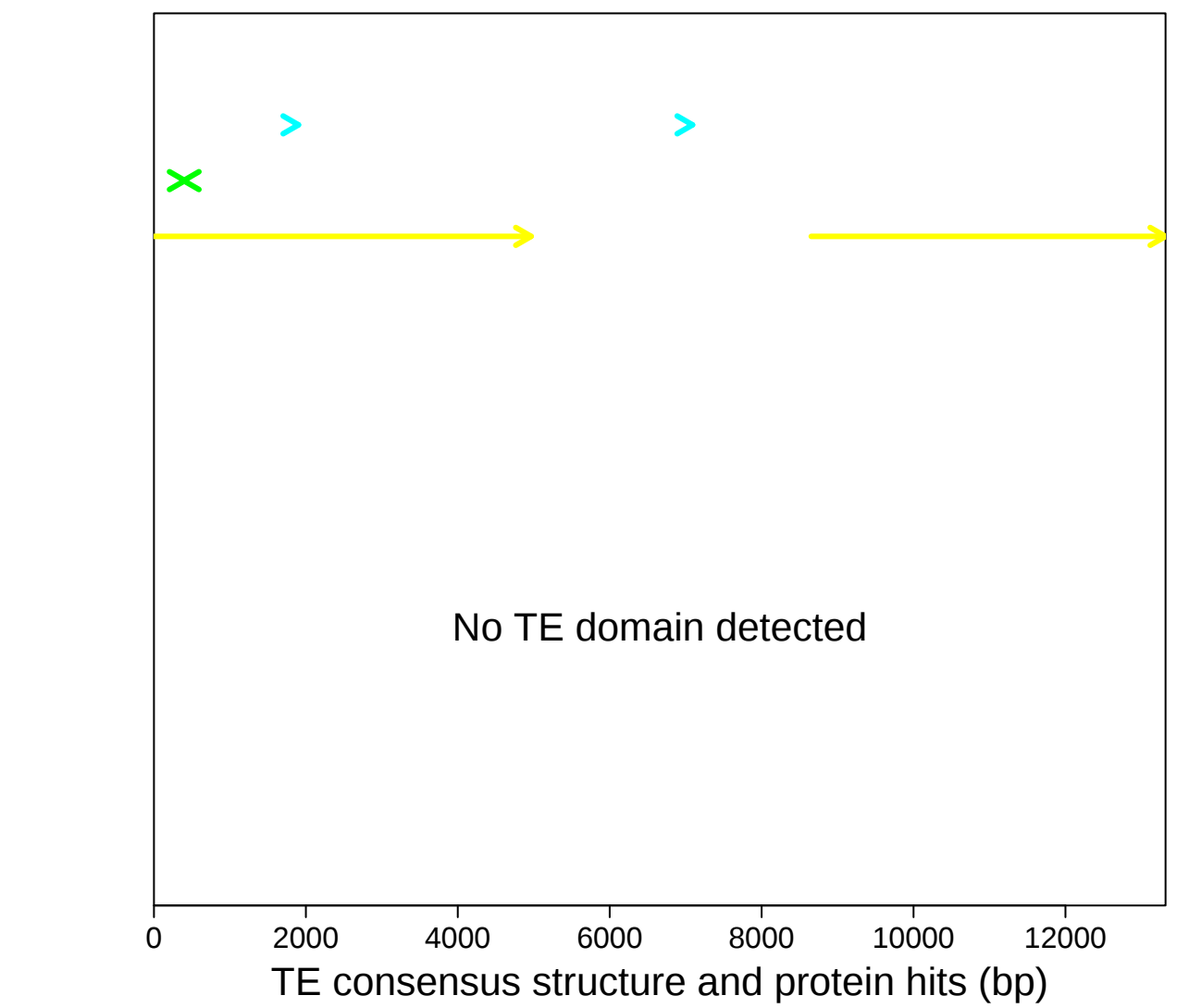
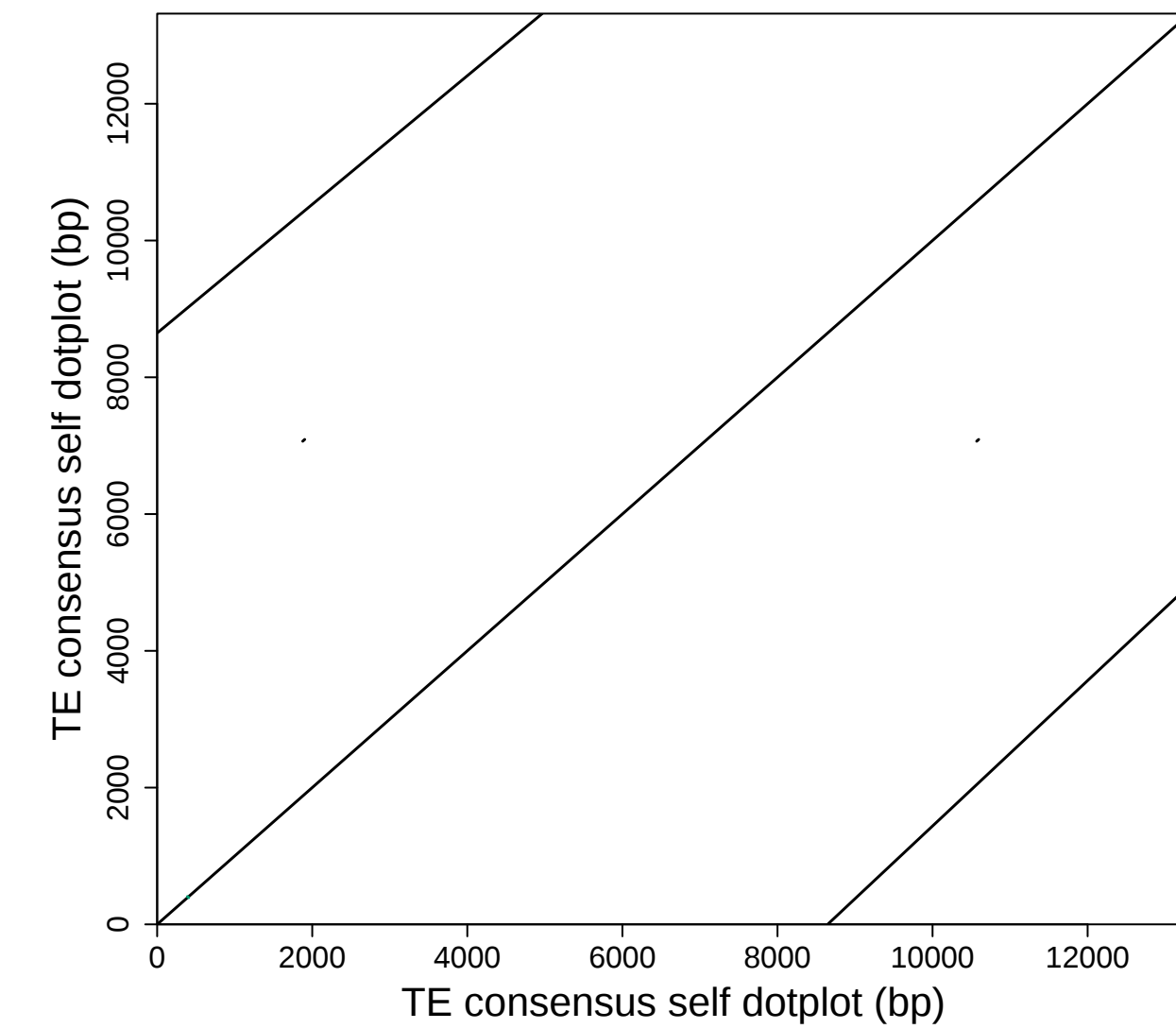
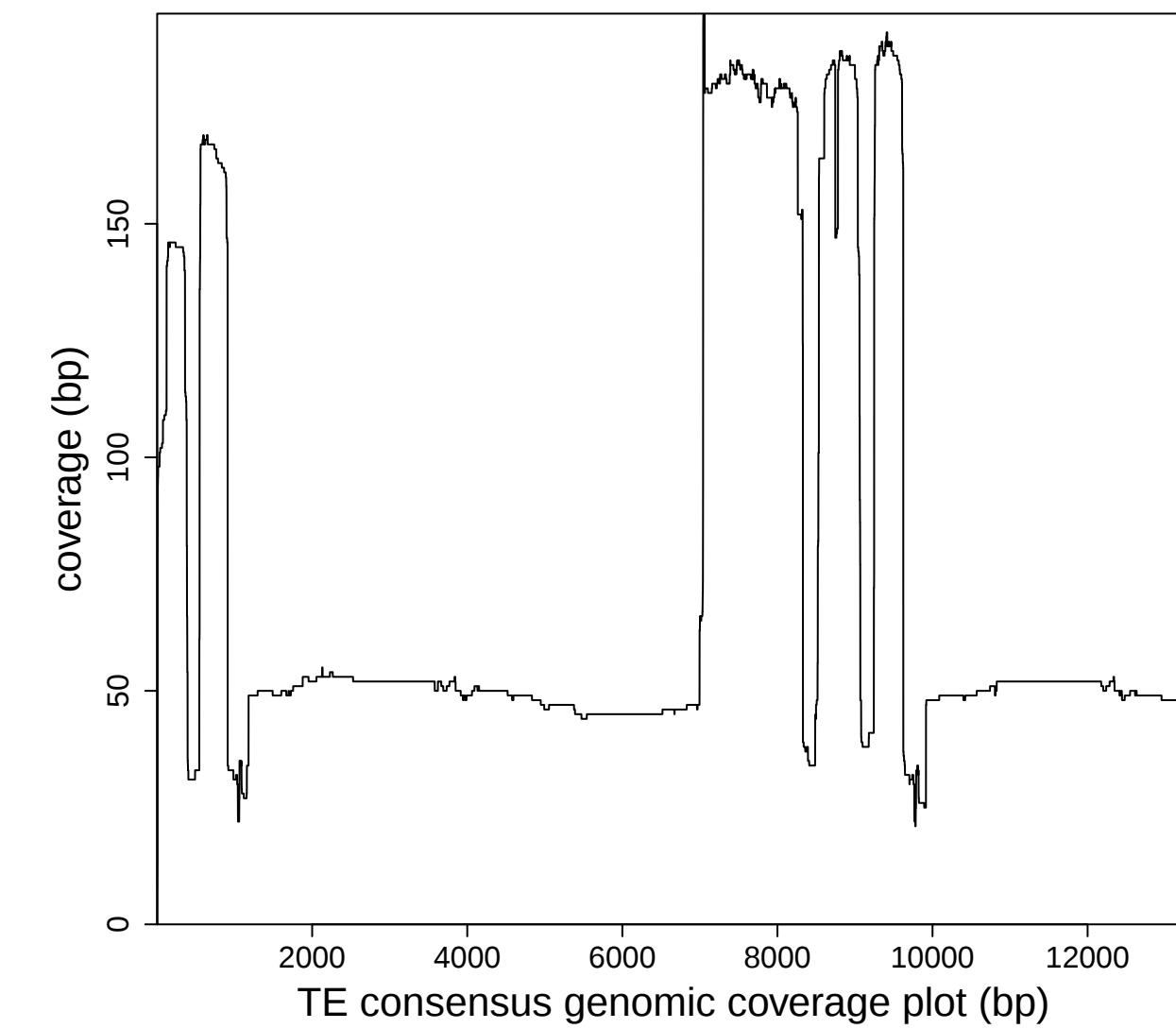
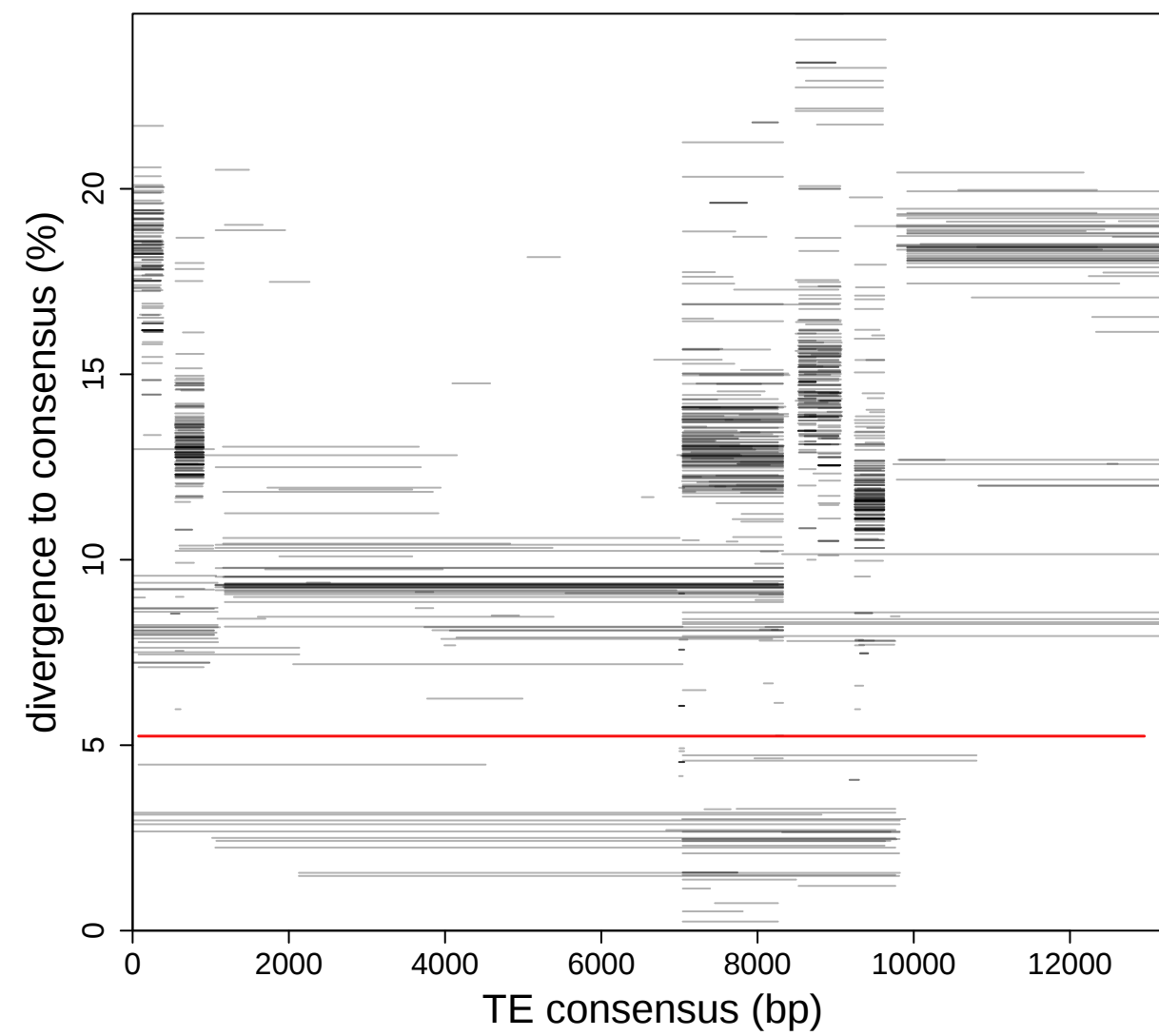






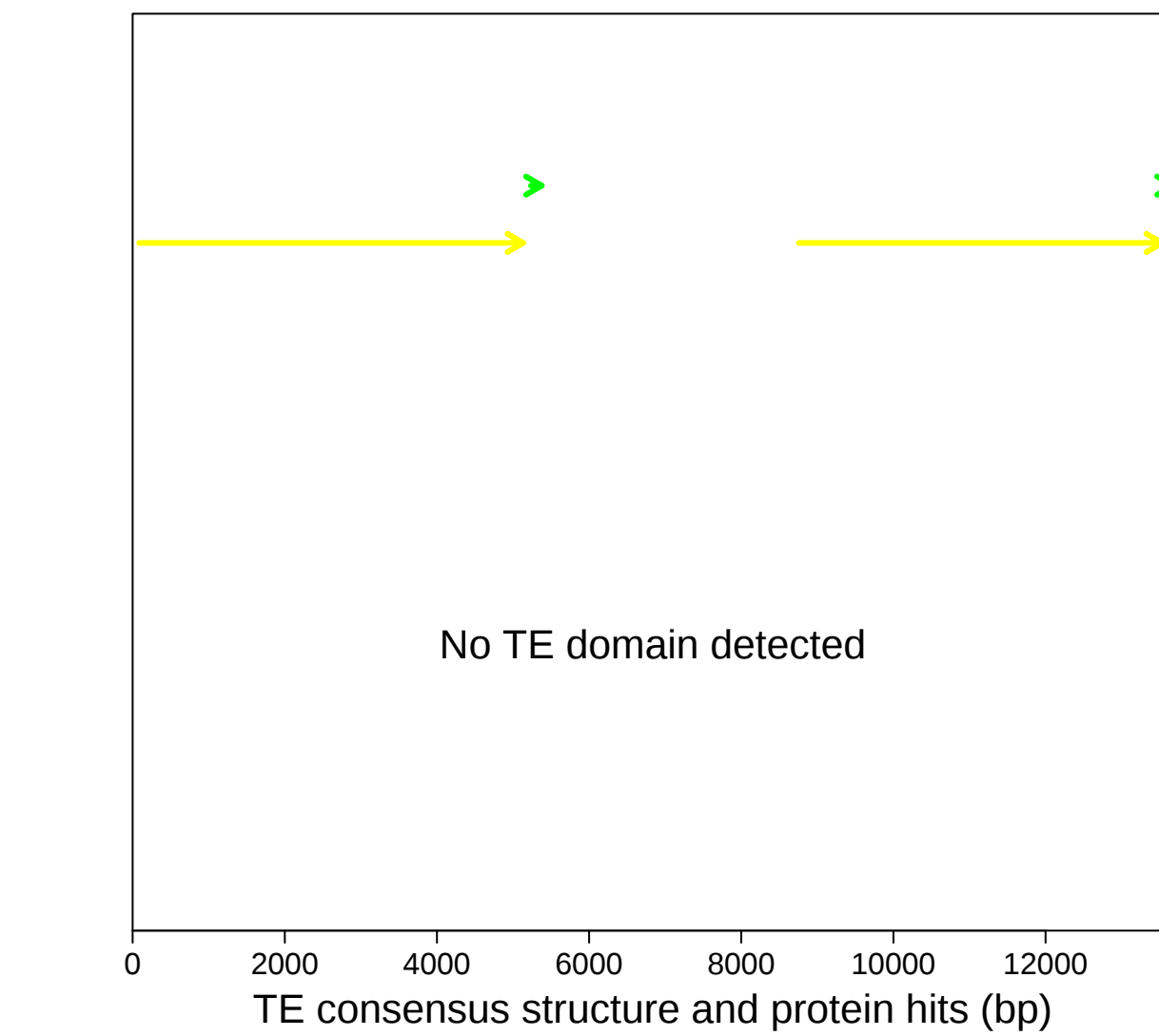
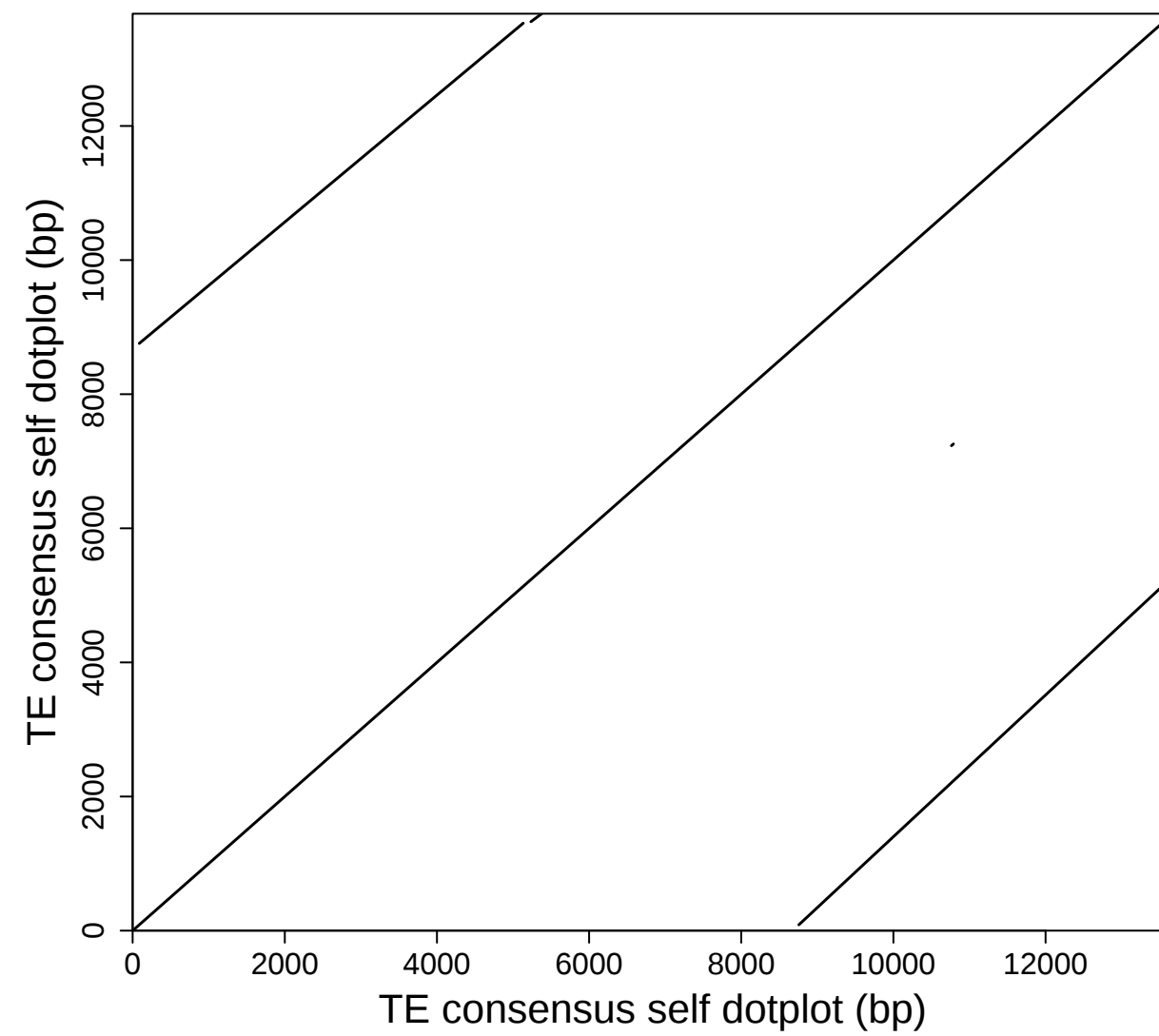
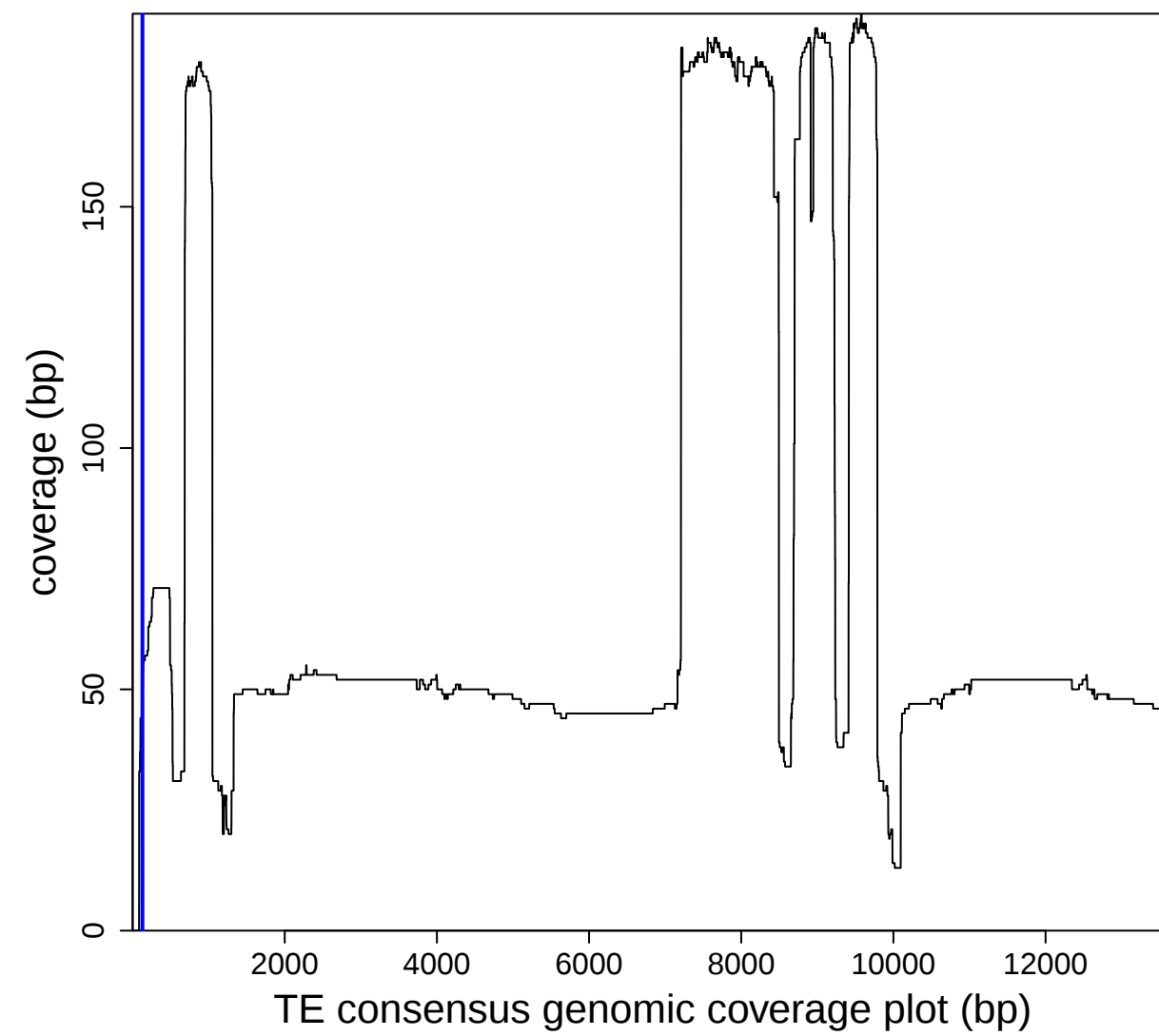
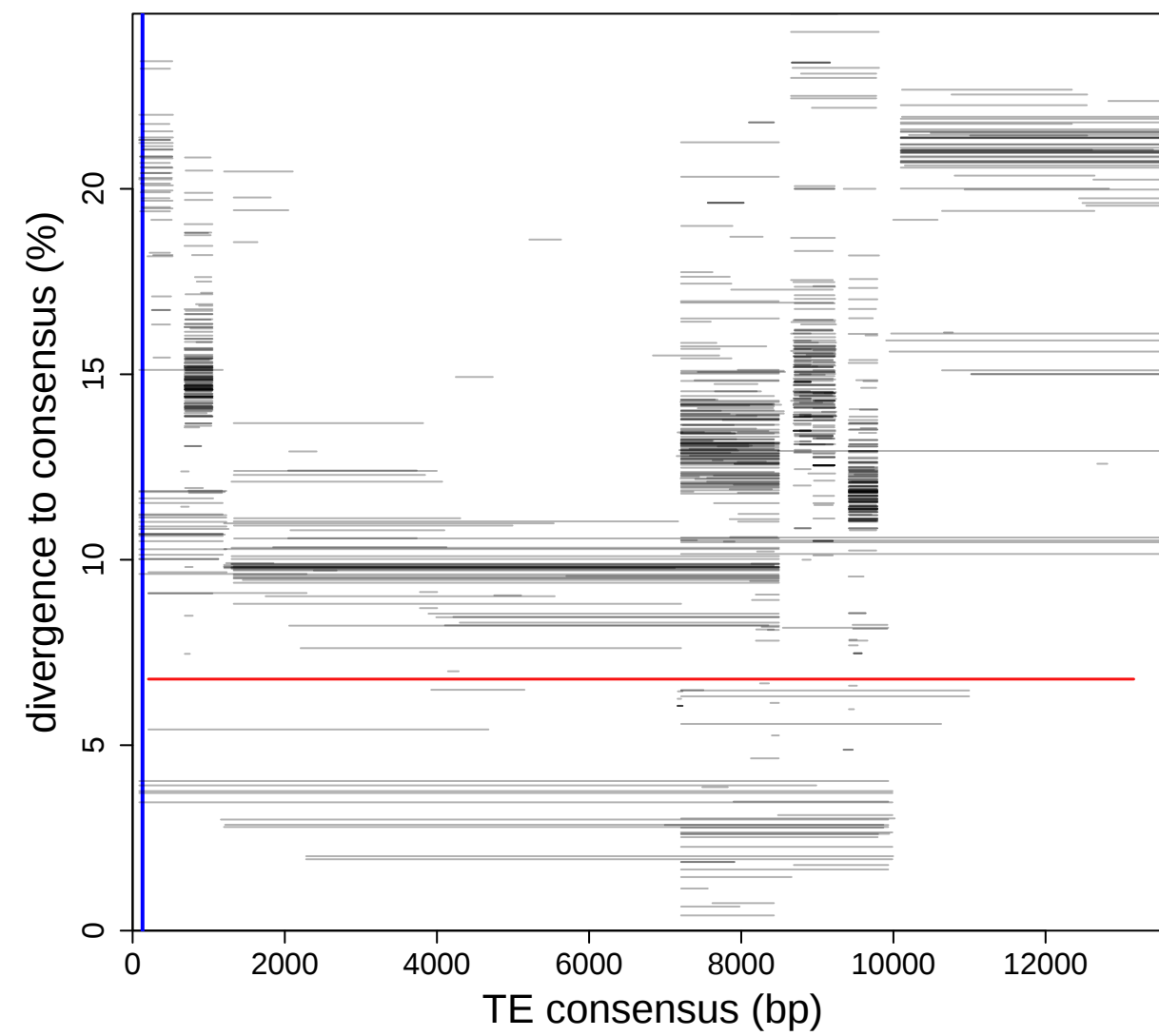
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 13318bp; fragments: 971; full length: 1 (>=11986.2bp)

After TEtrimmer 13318 bp



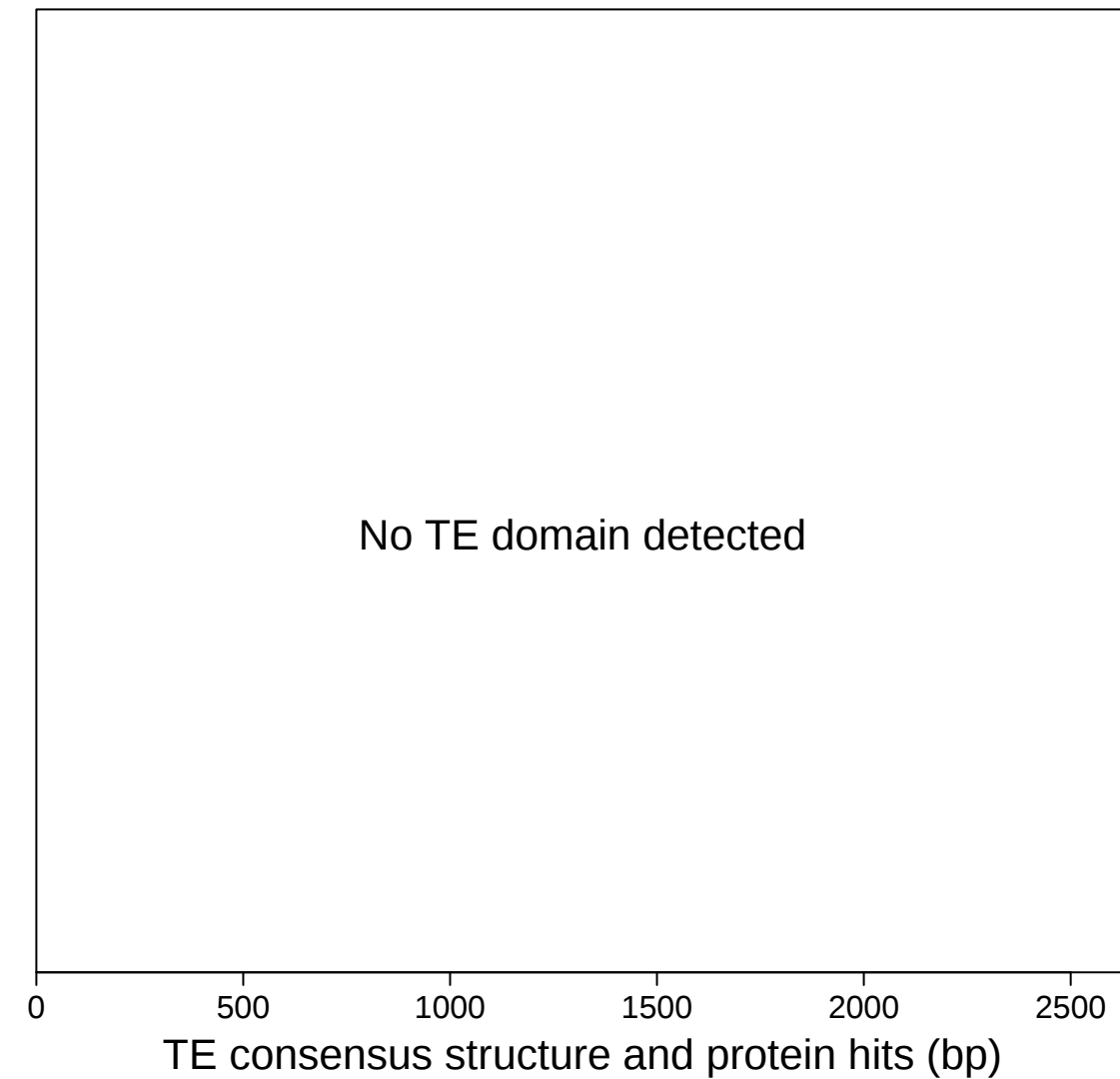
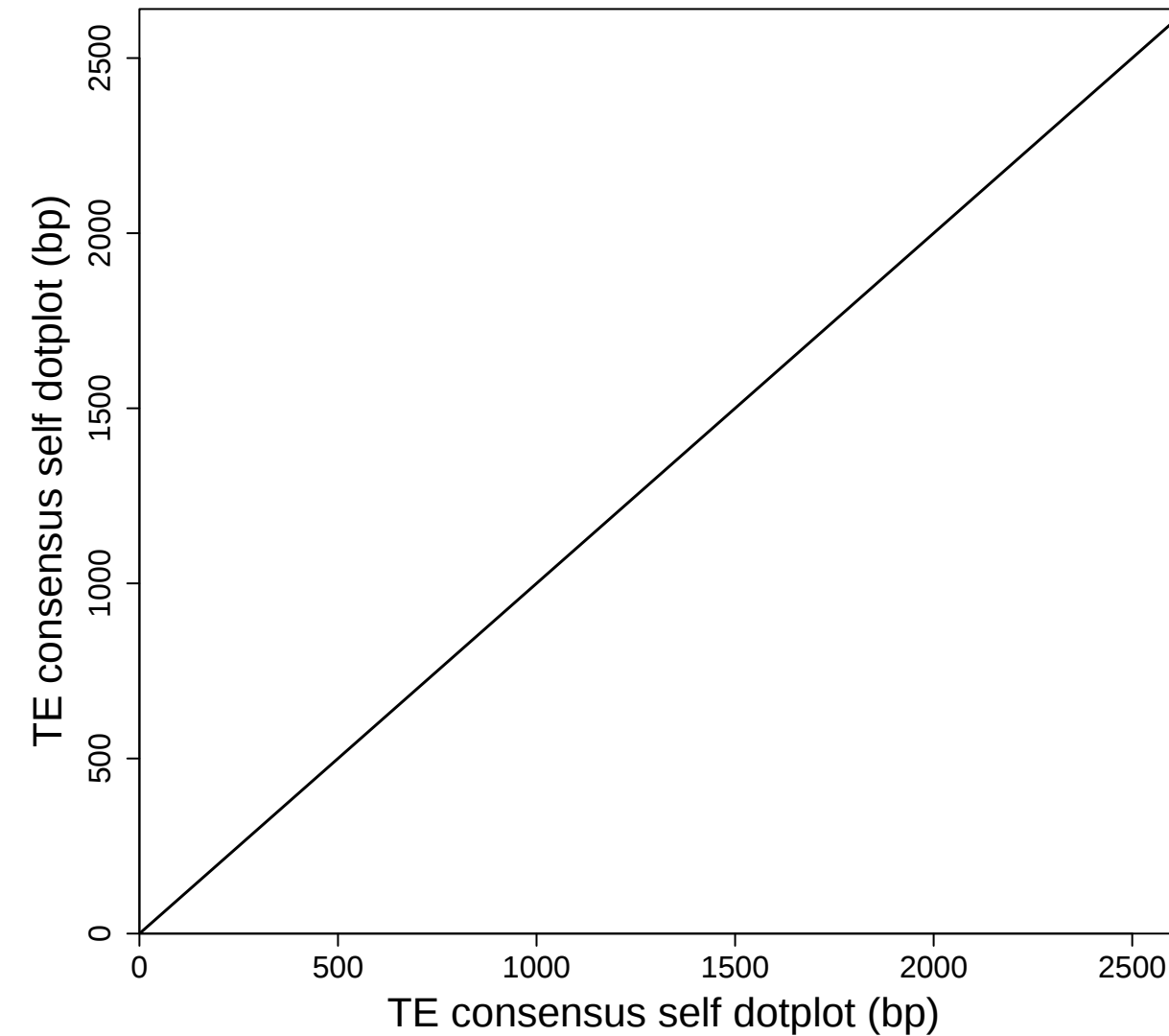
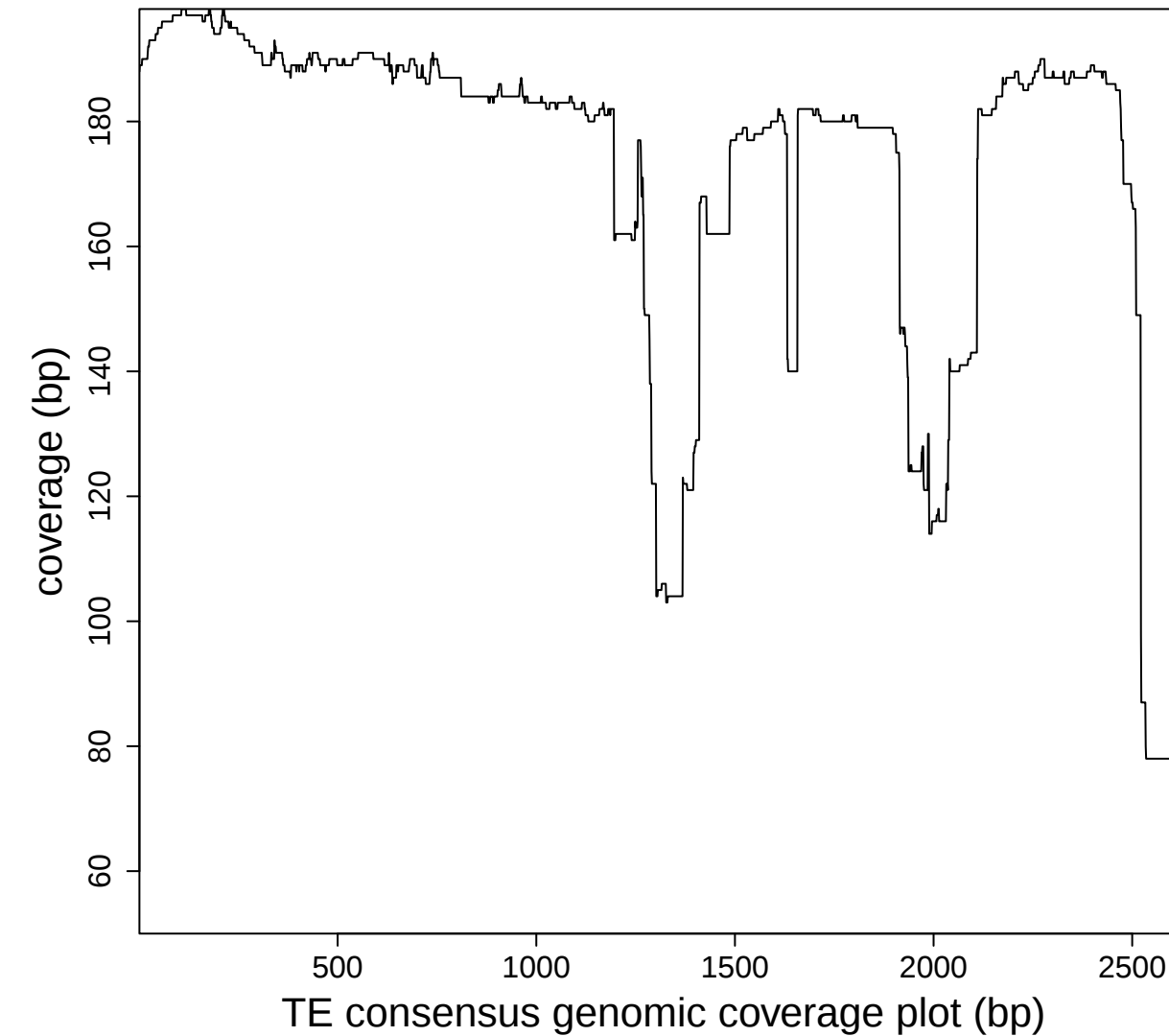
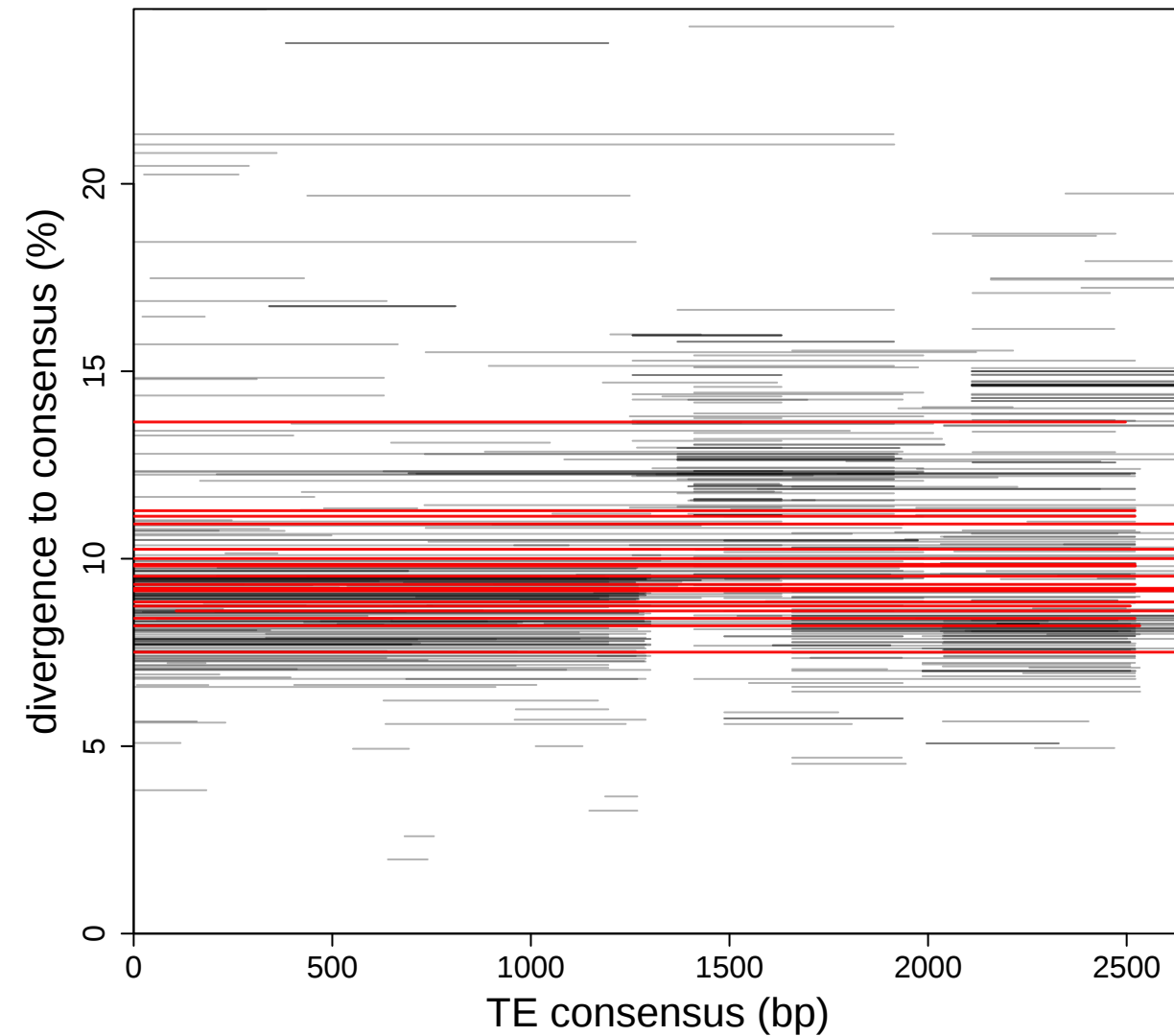
ly_9094.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.f
size: 13674bp; fragments: 910; full length: 1 (>=12306.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

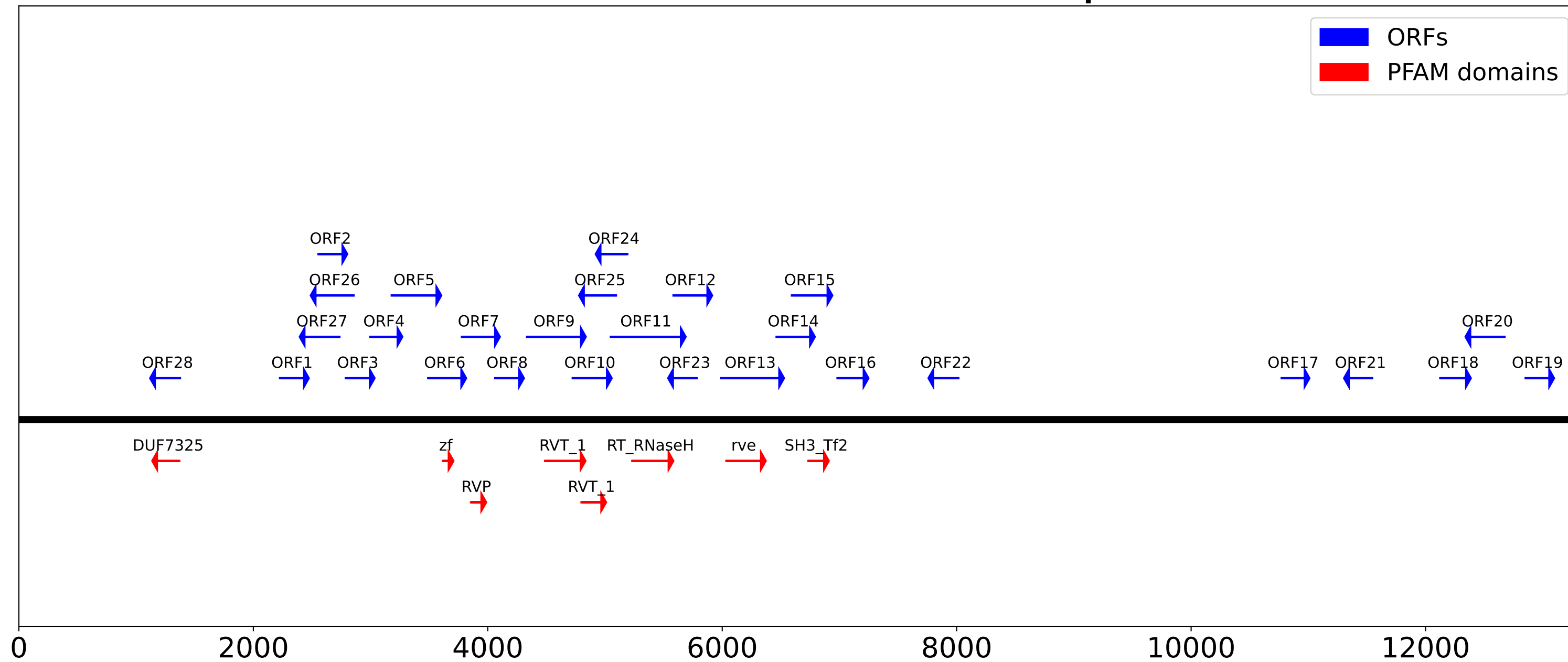


TE: rnd_5_family_9094
size: 2640bp; fragments: 529; full length: 20 (≥ 2376 bp)

Before TEtrimmer 2640 bp



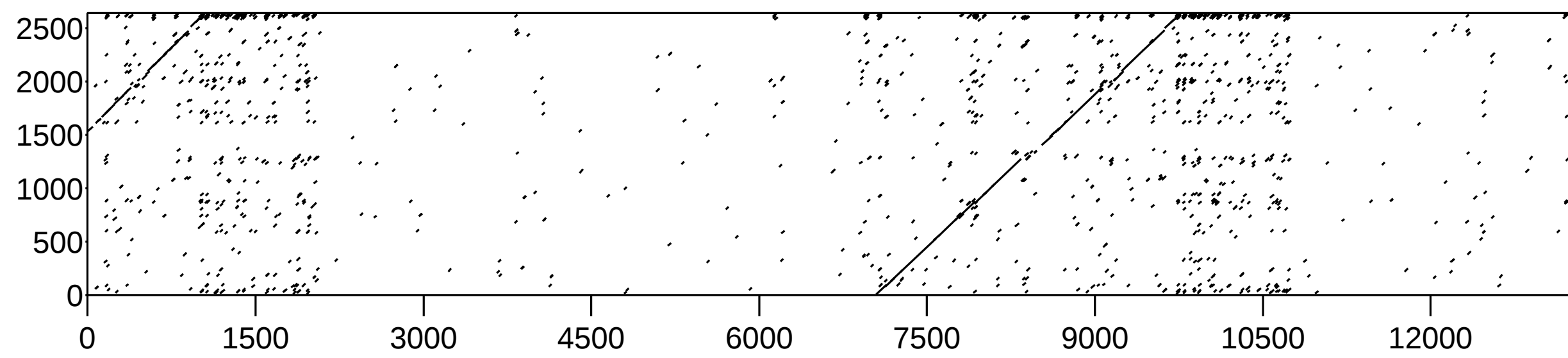
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)