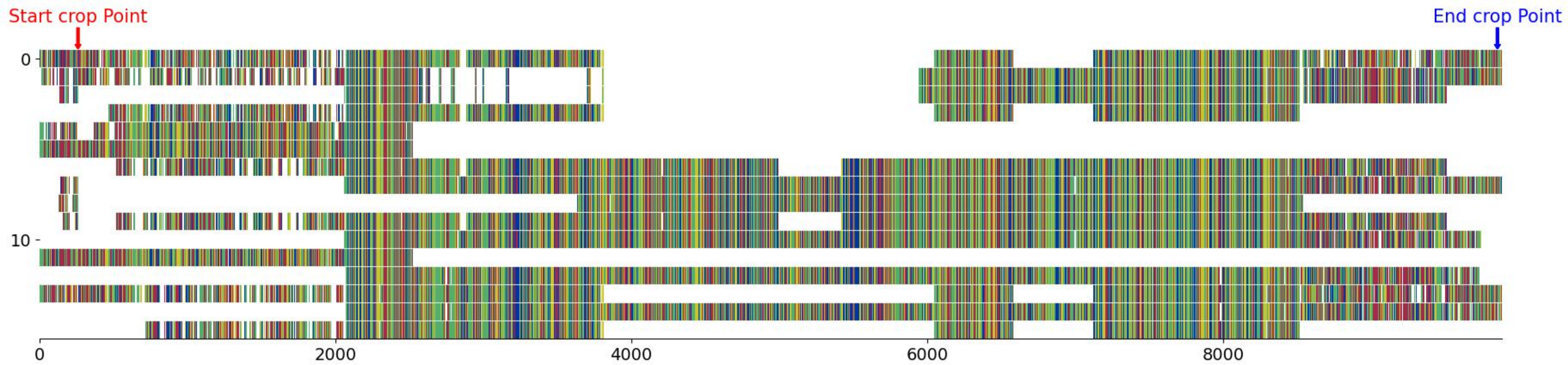


Start crop Point

End crop Point

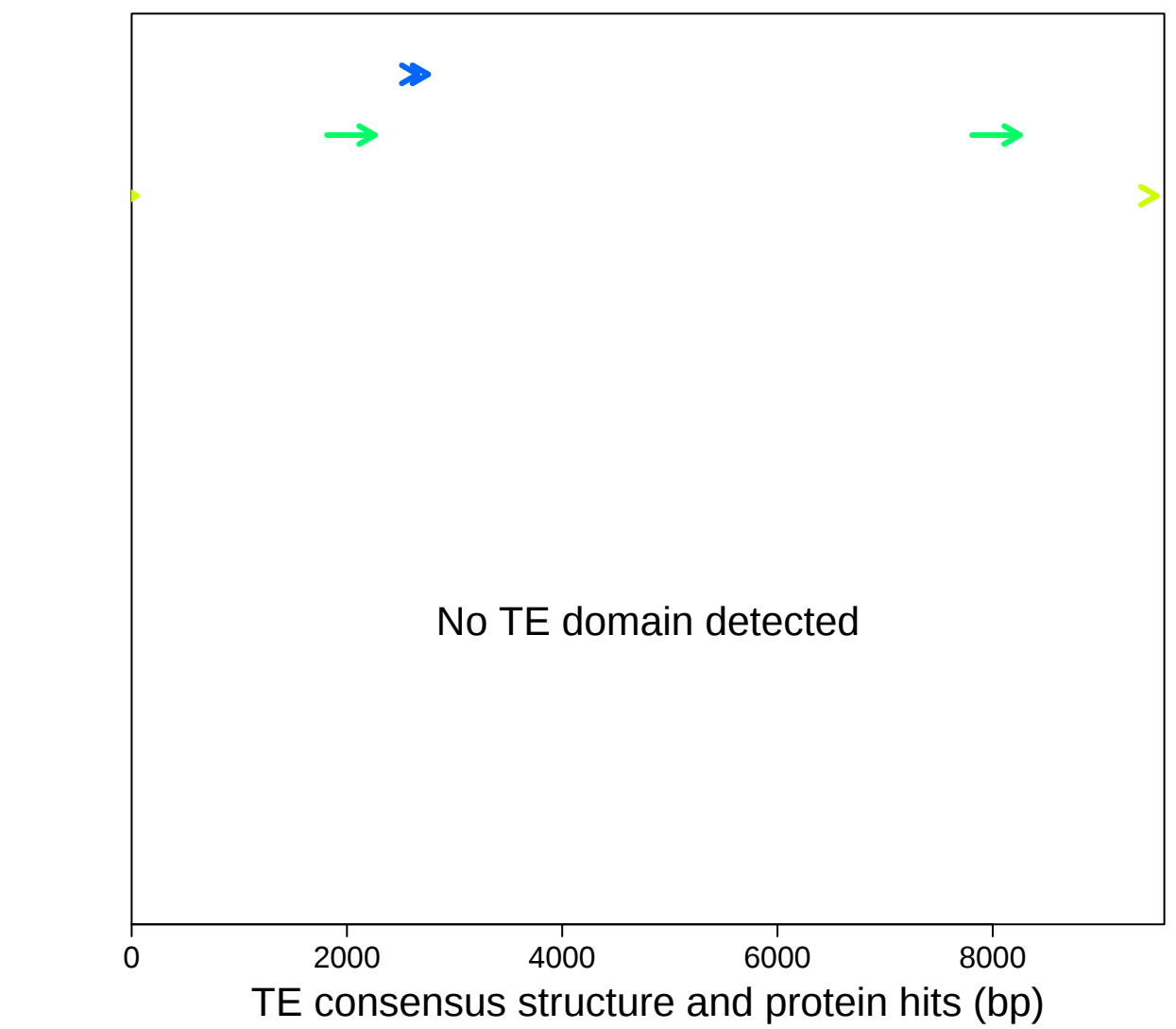
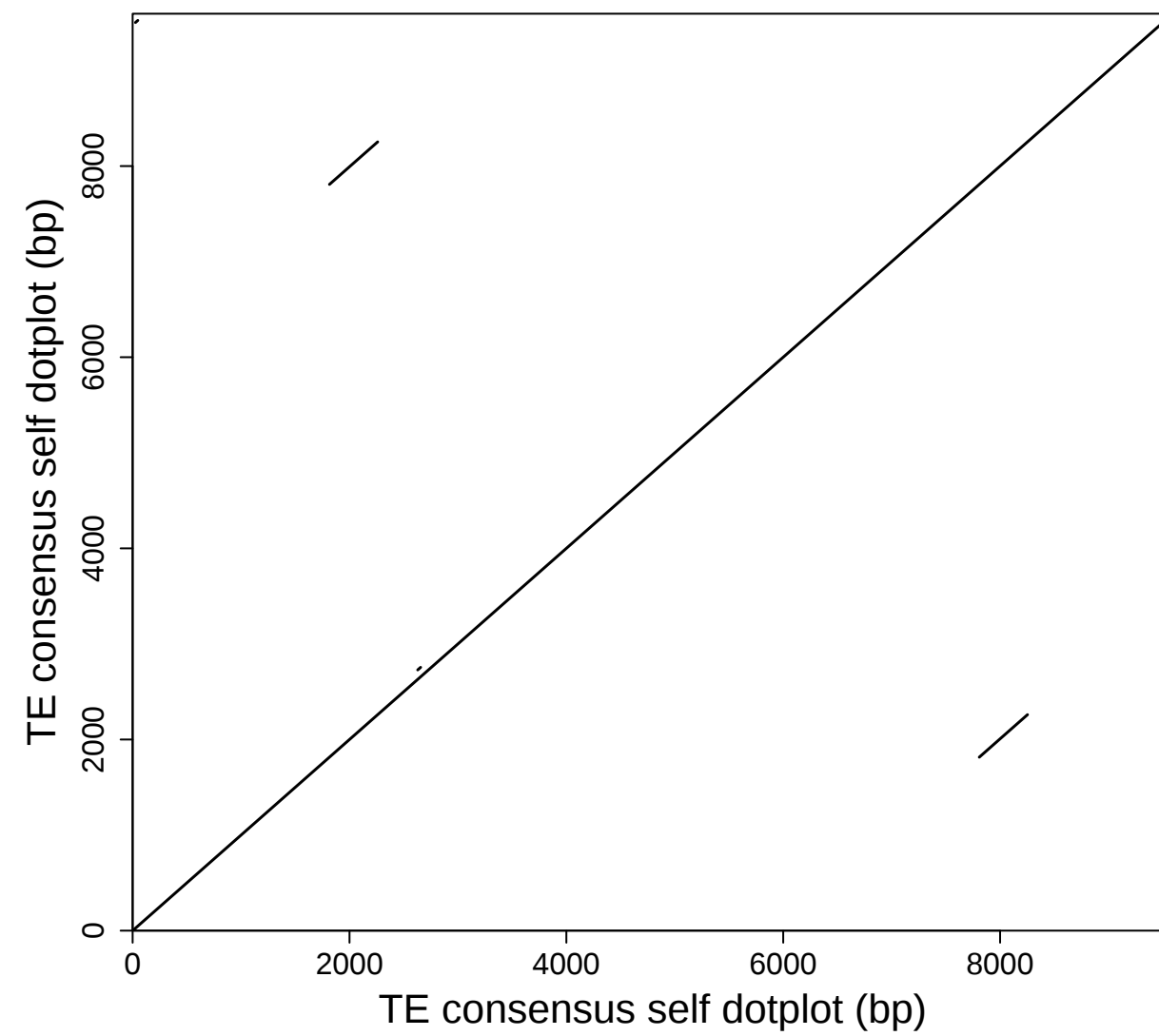
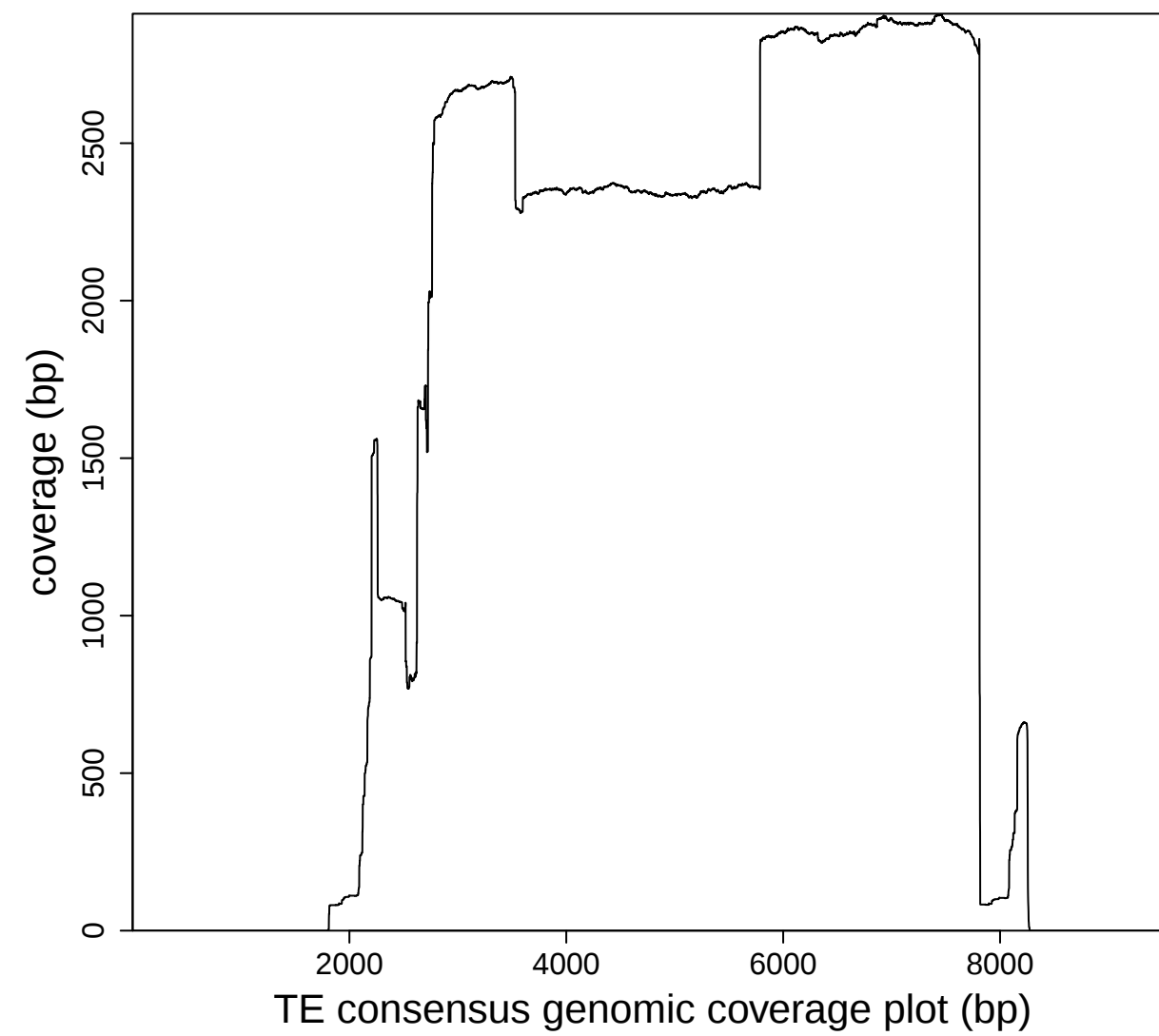
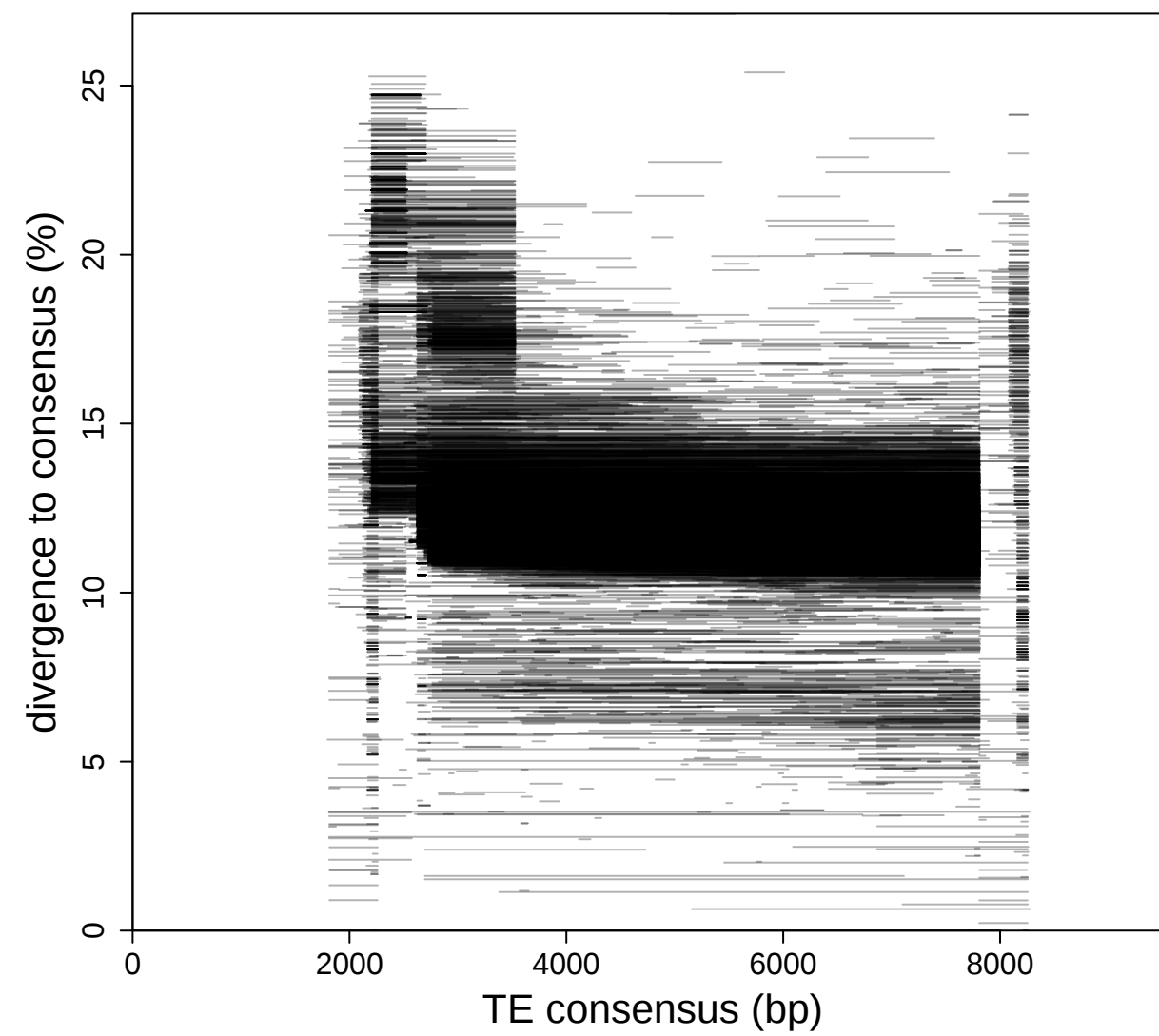
MSA length = 9594





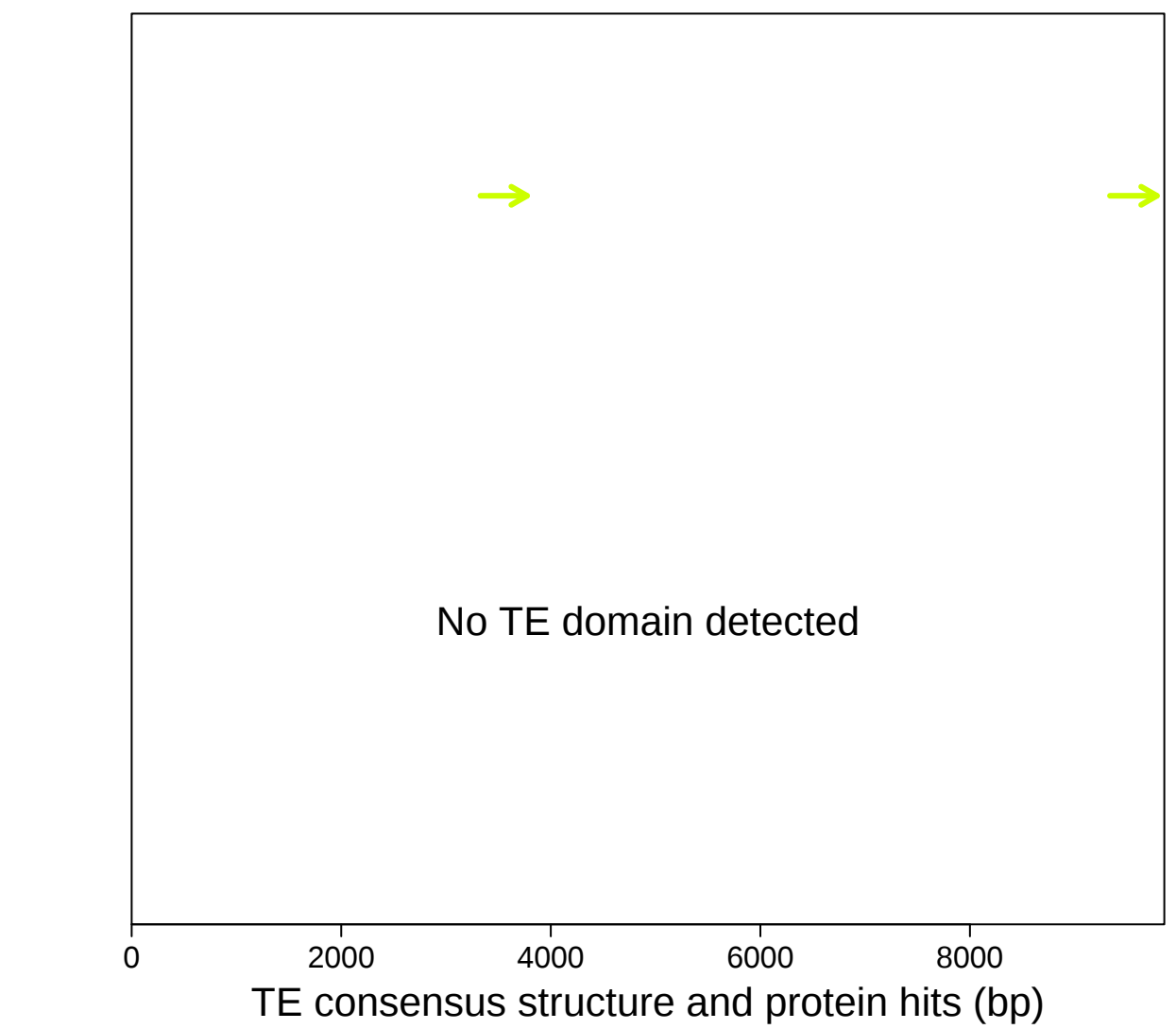
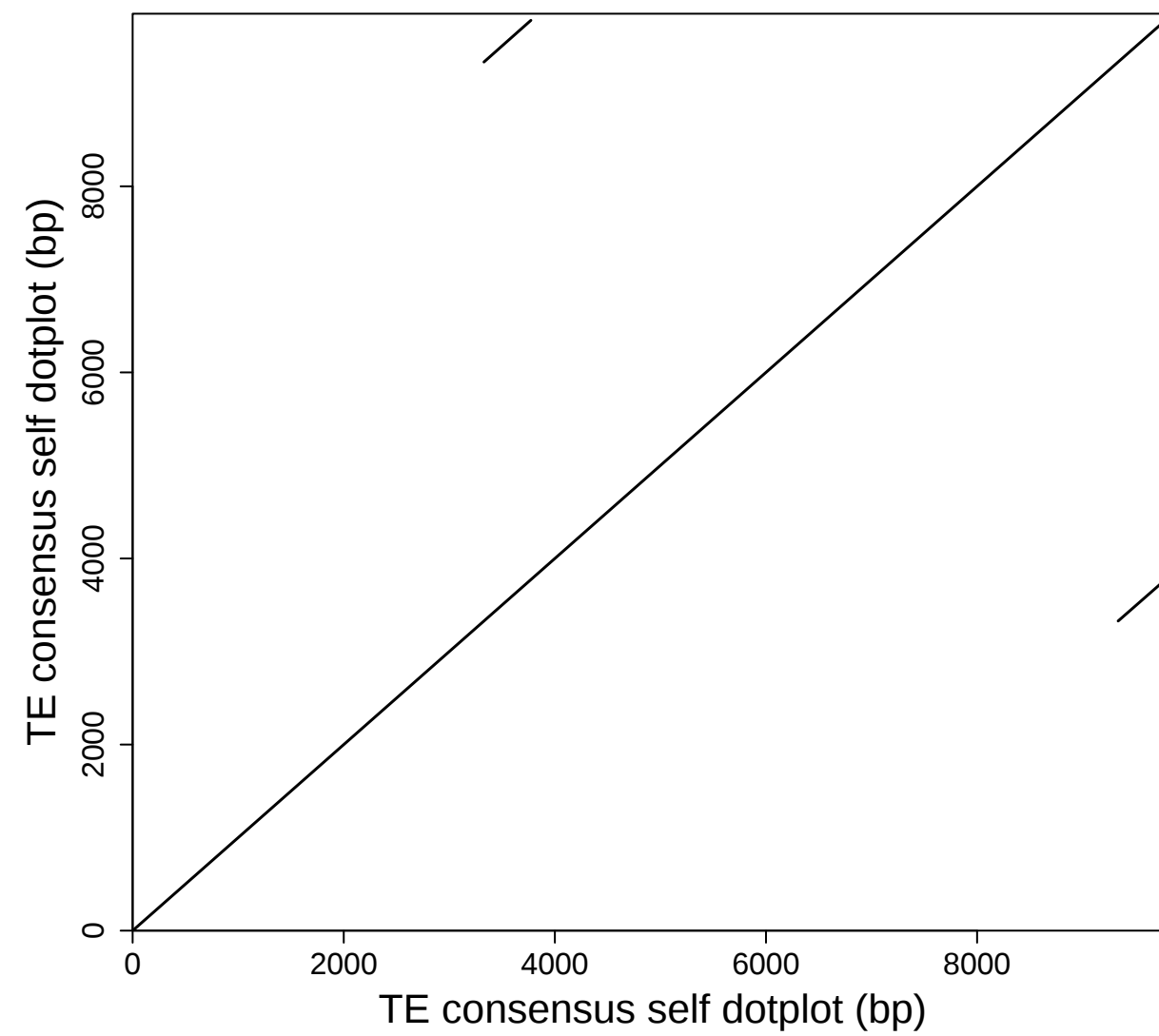
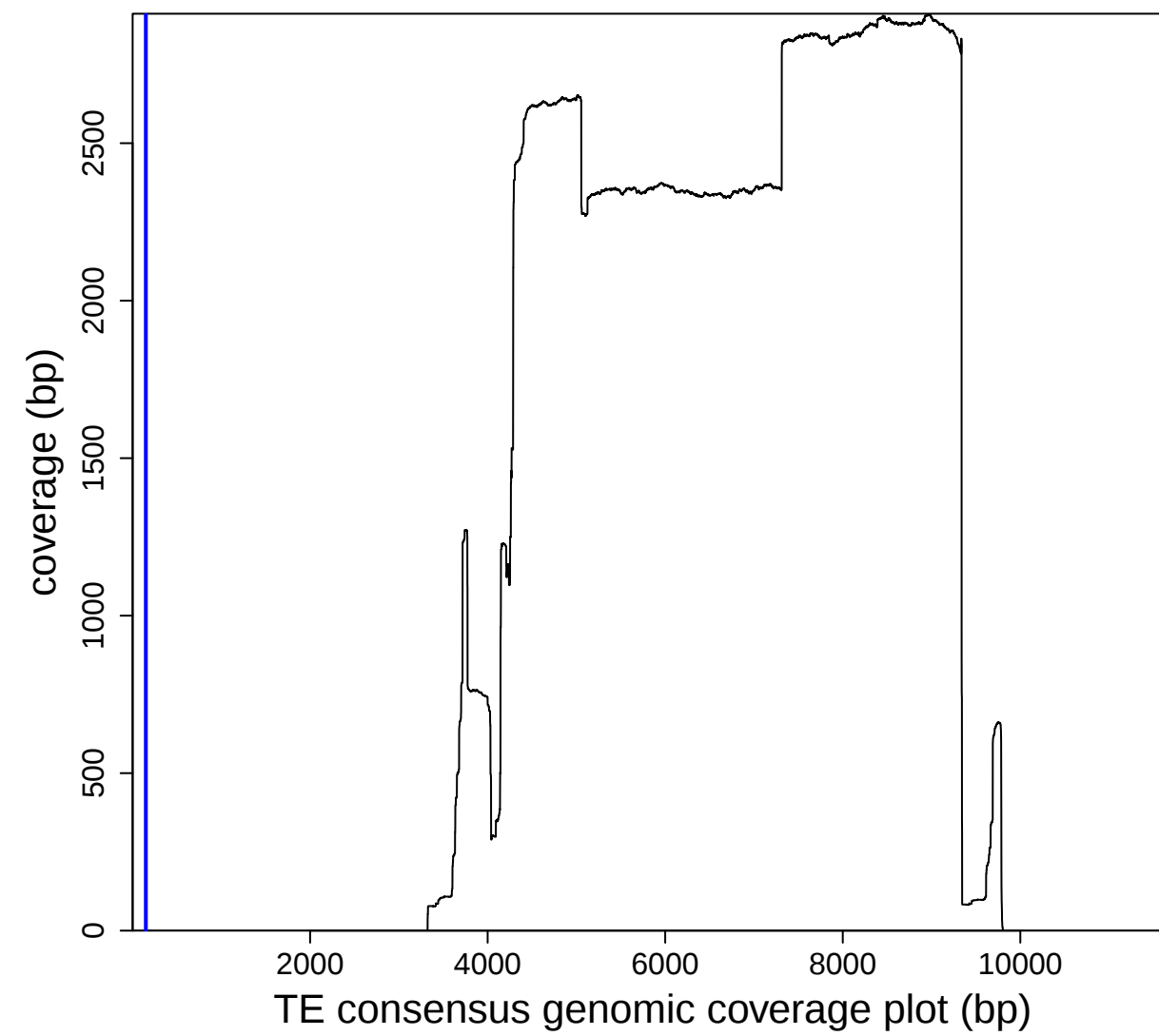
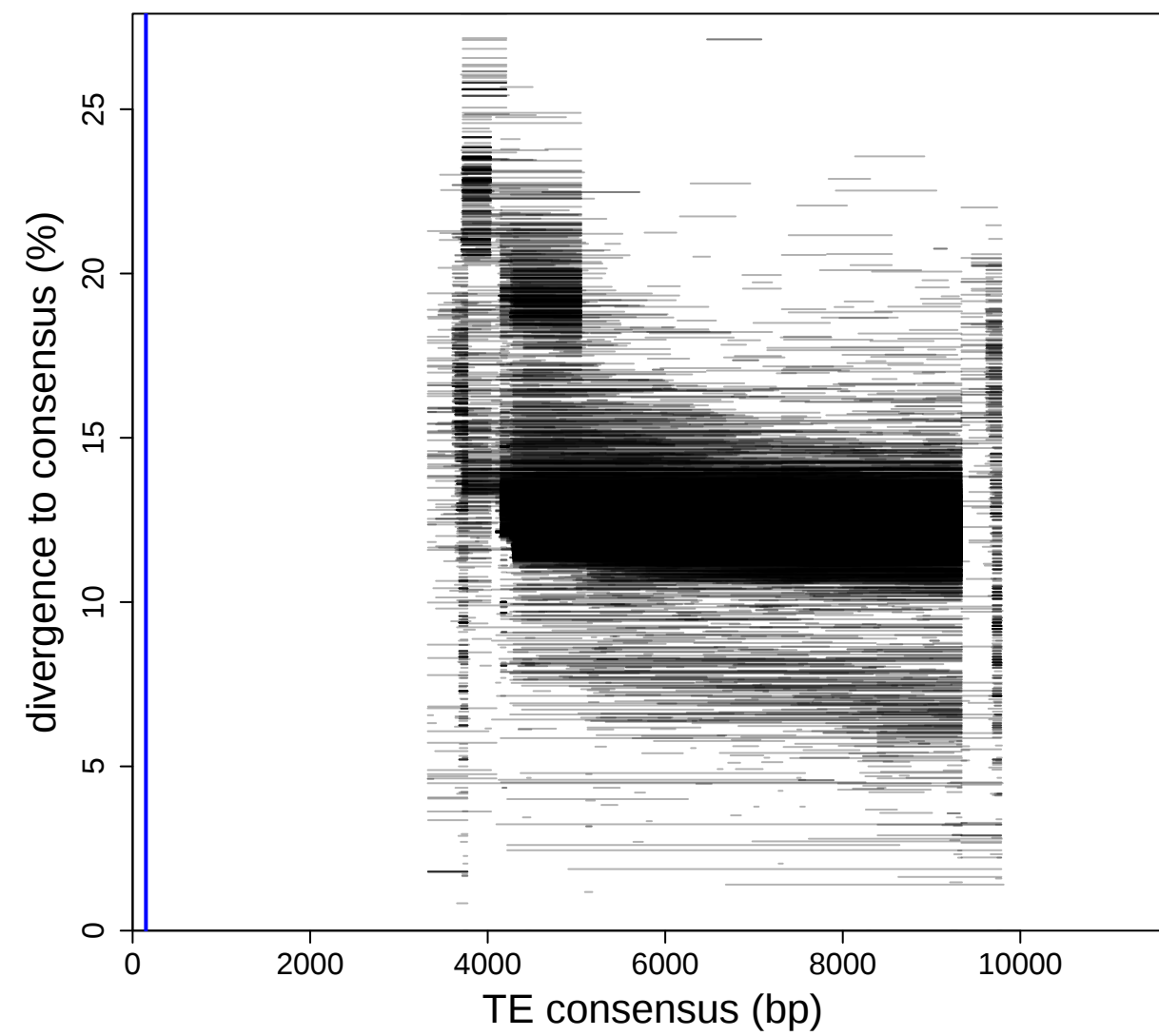
g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_a
size: 9594bp; fragments: 7045; full length: 0 (>=8634.6bp)

After TEtrimmer 9594 bp

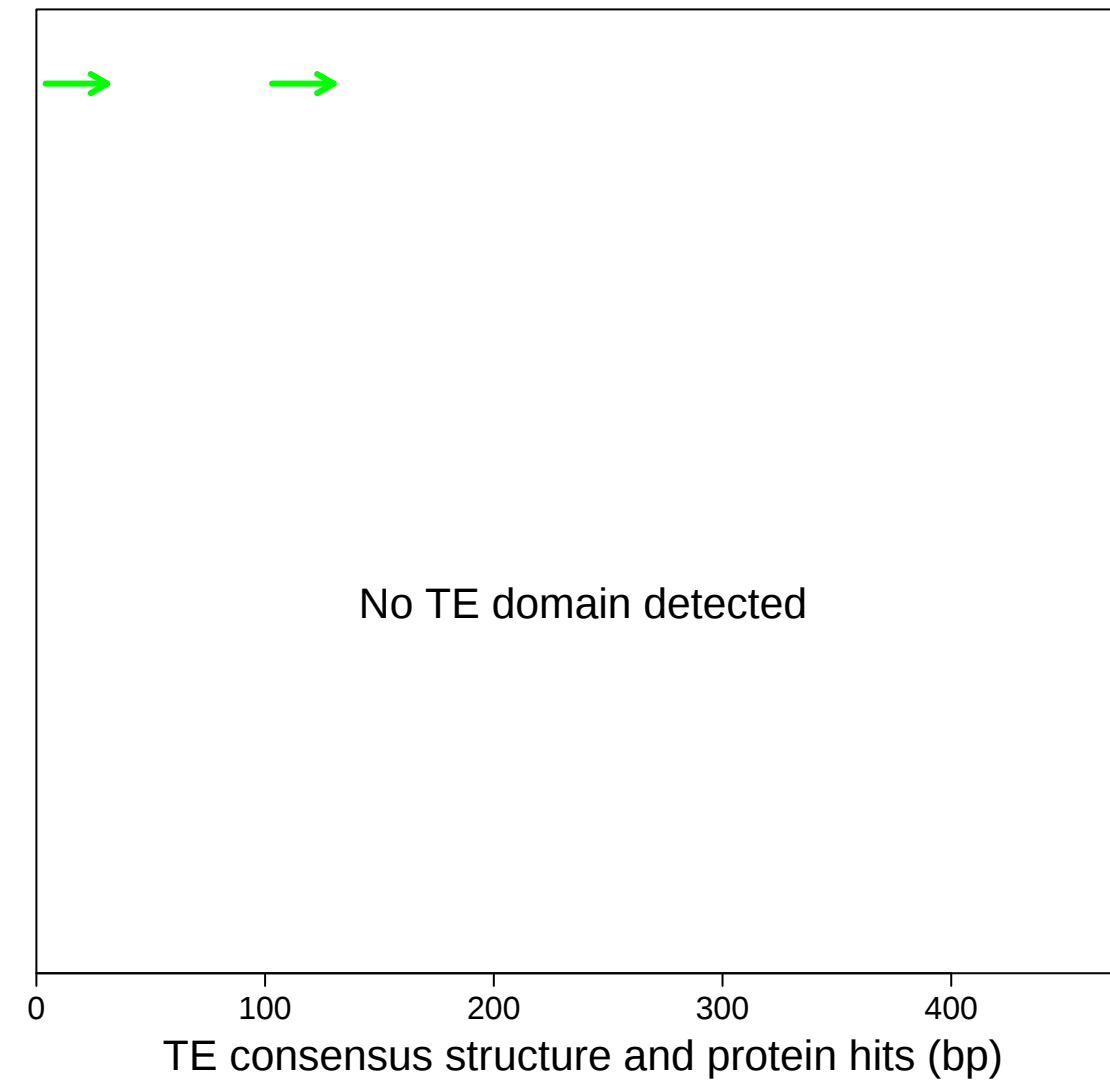
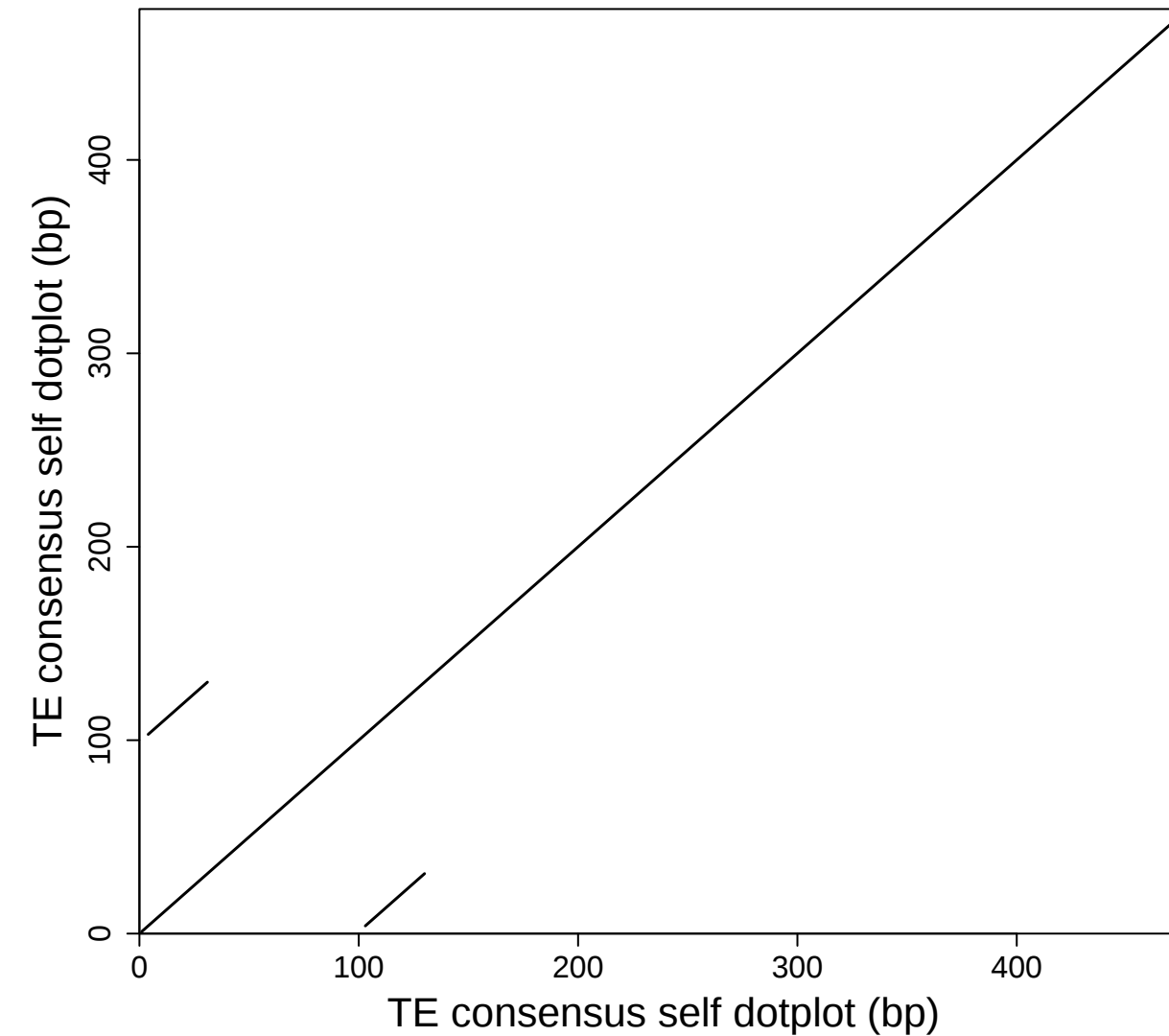
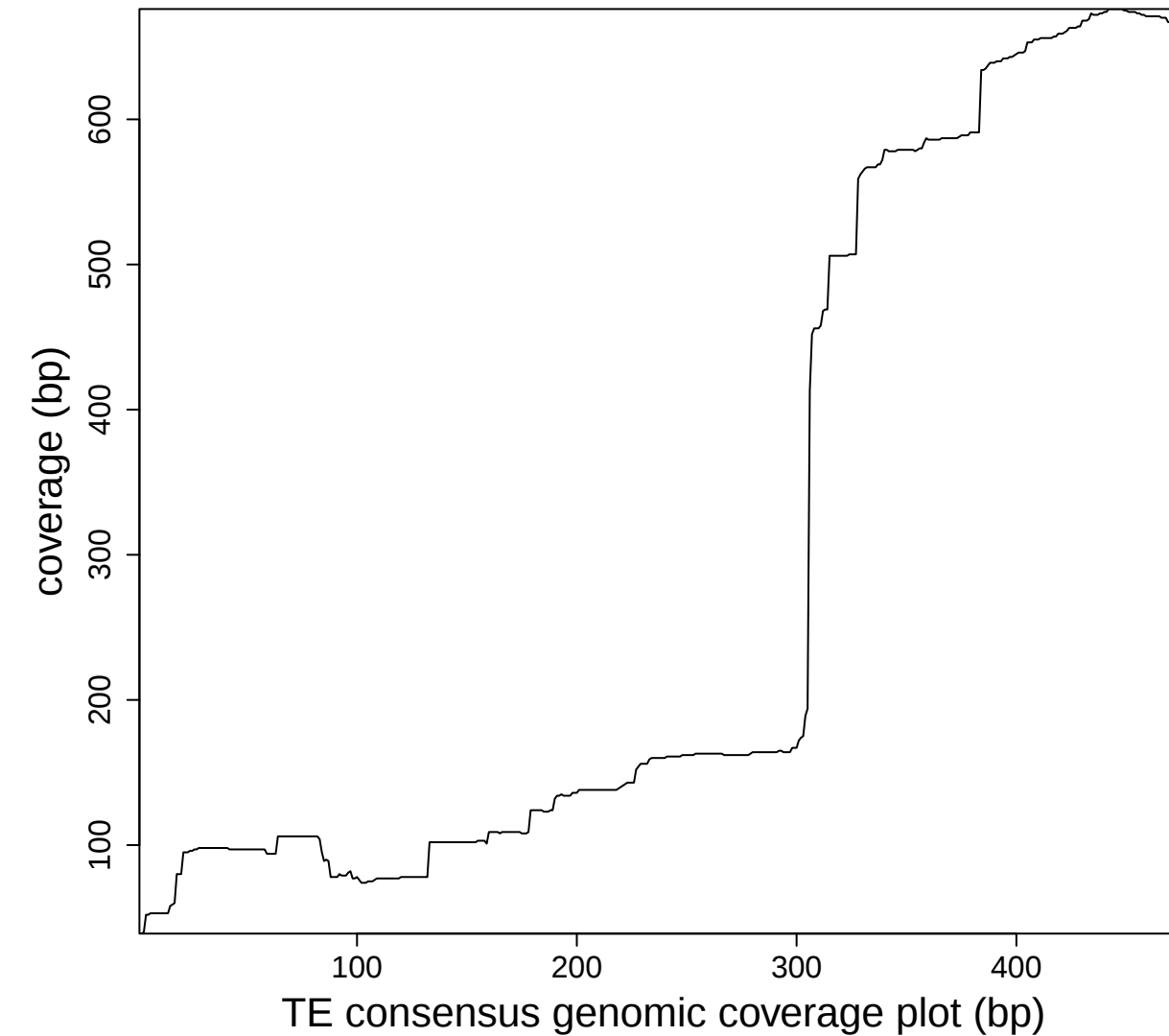
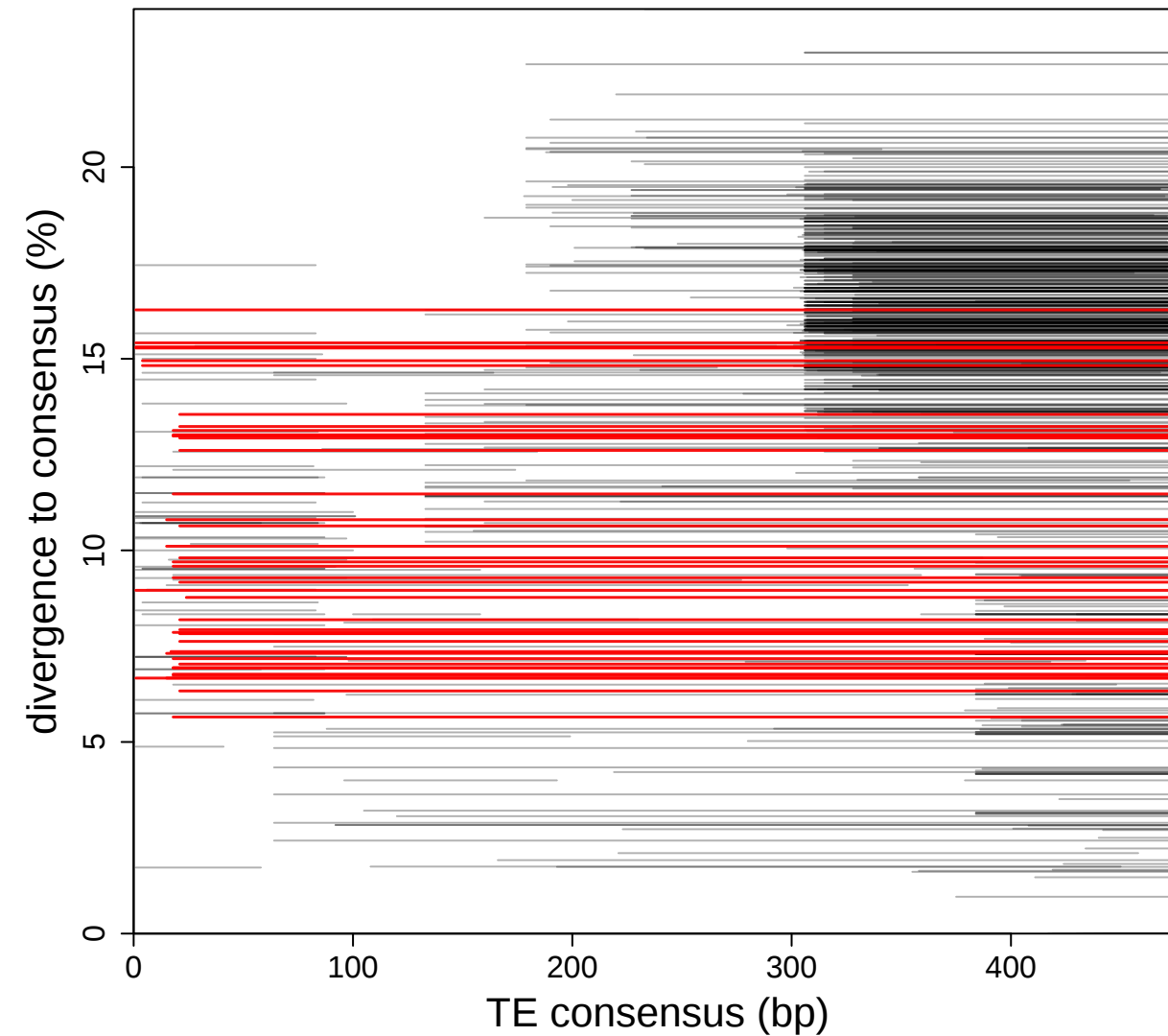


f.bed_g_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 11719bp; fragments: 7018; full length: 0 (>=10547.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

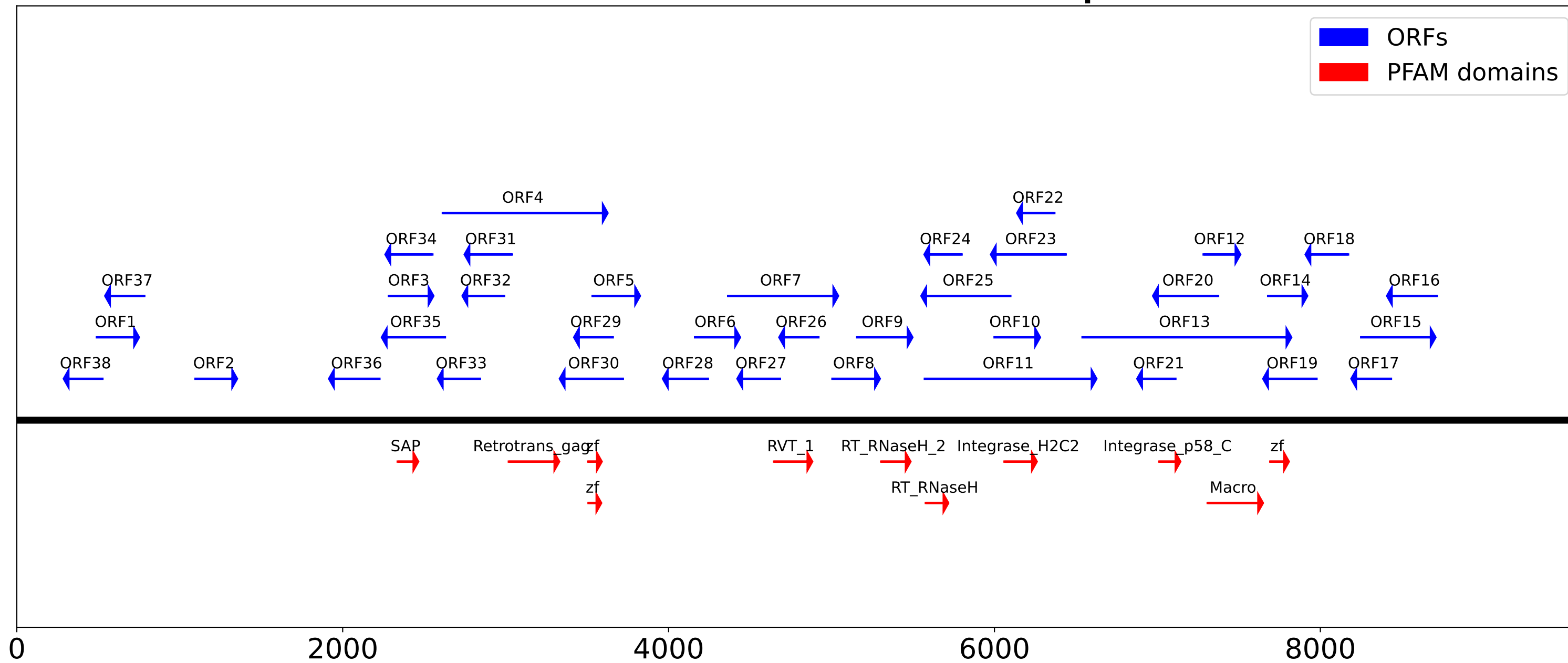


TE: rnd_5_family_747
size: 478bp; fragments: 738; full length: 43 (>=430.2bp)

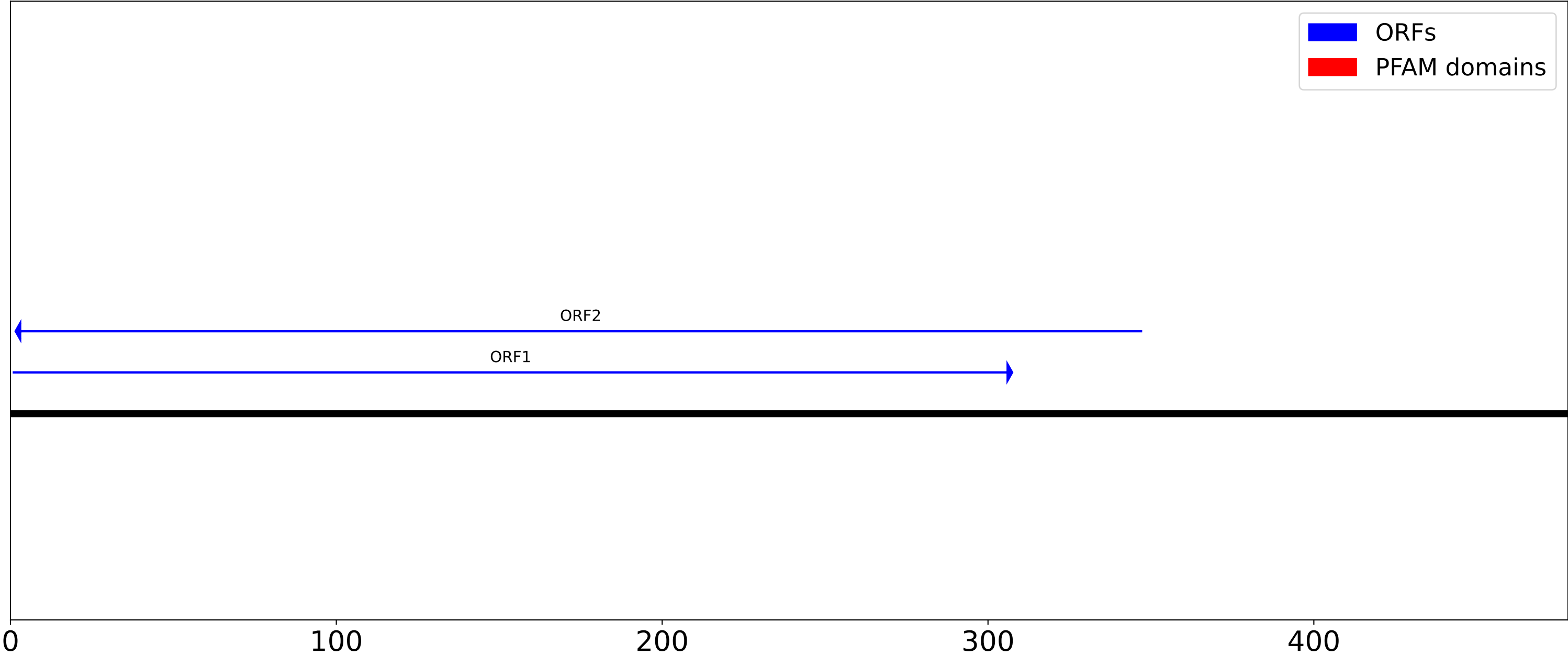


Before TEtrimmer 478 bp

After TETrimmer ORF and PFAM domain plot



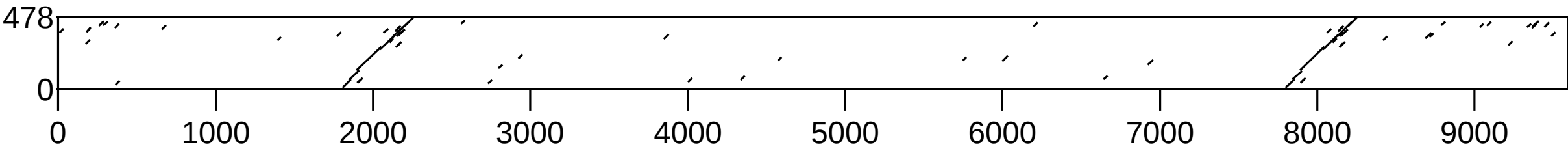
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)