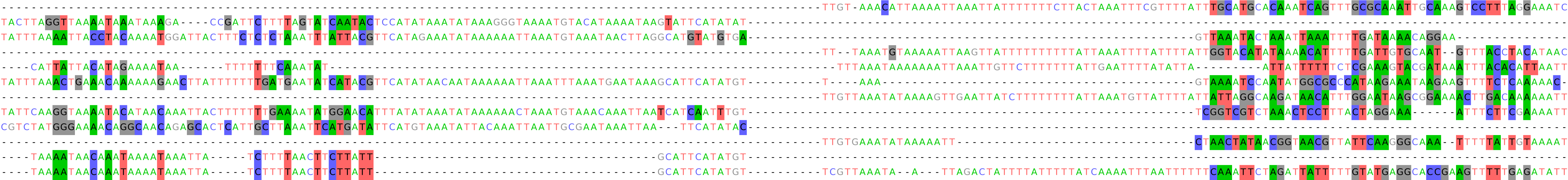
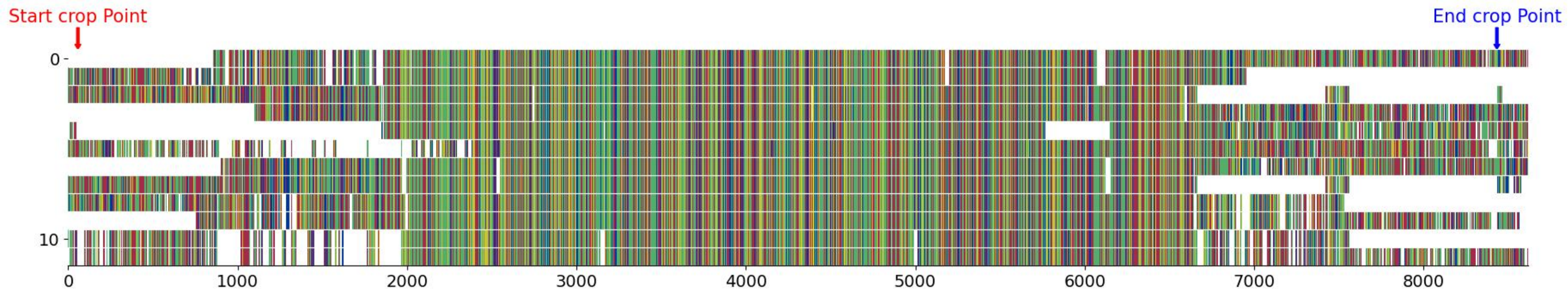


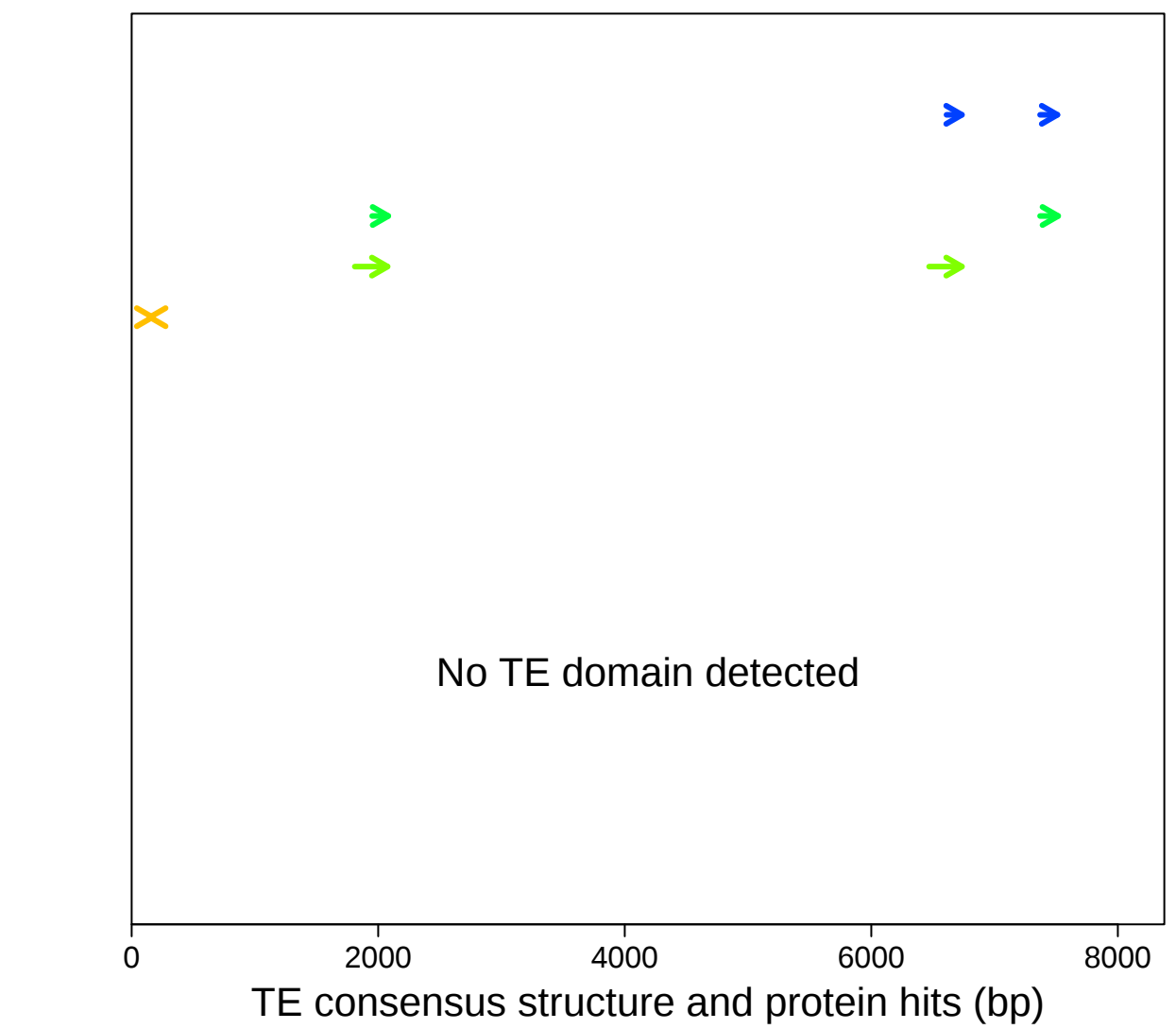
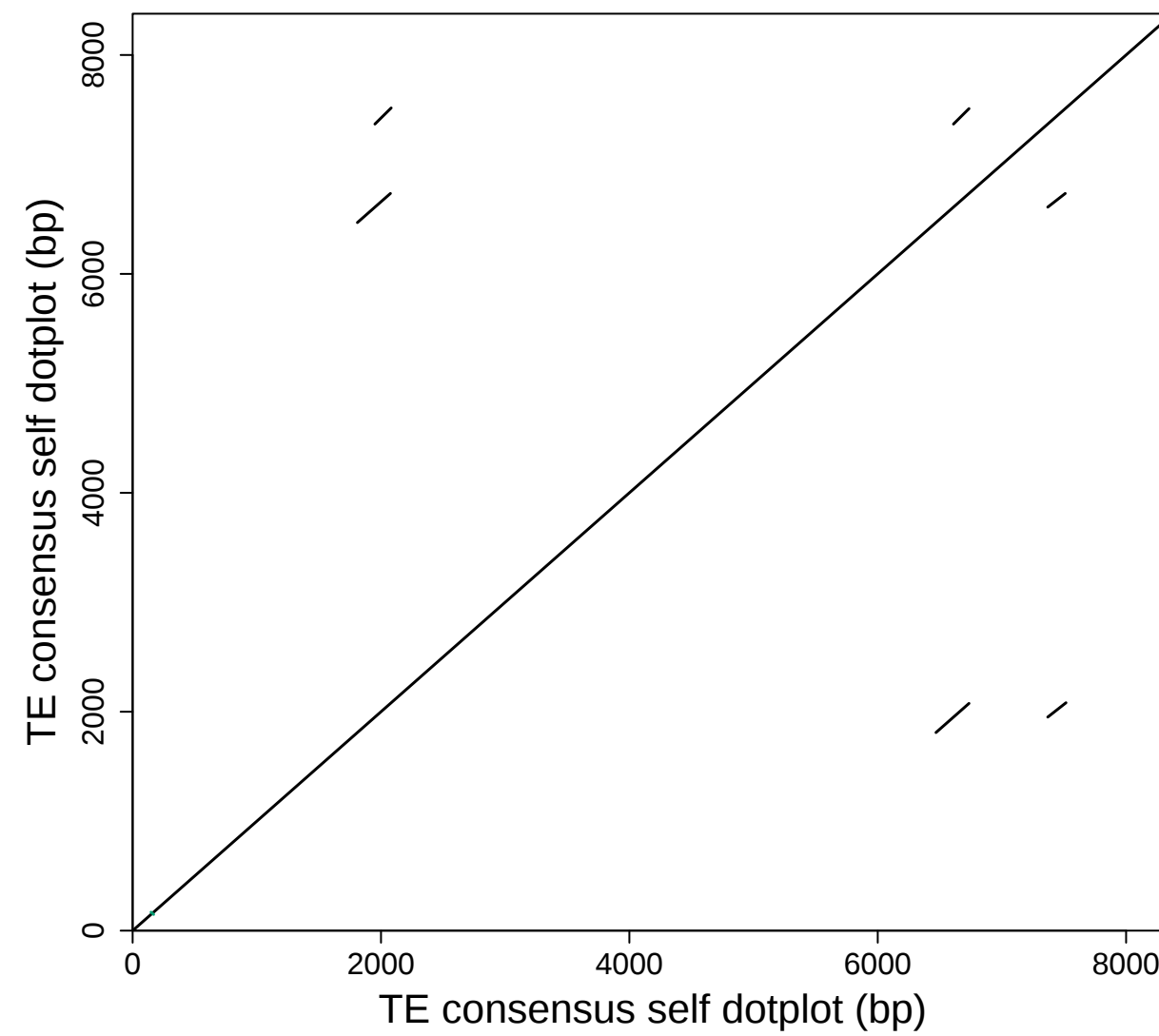
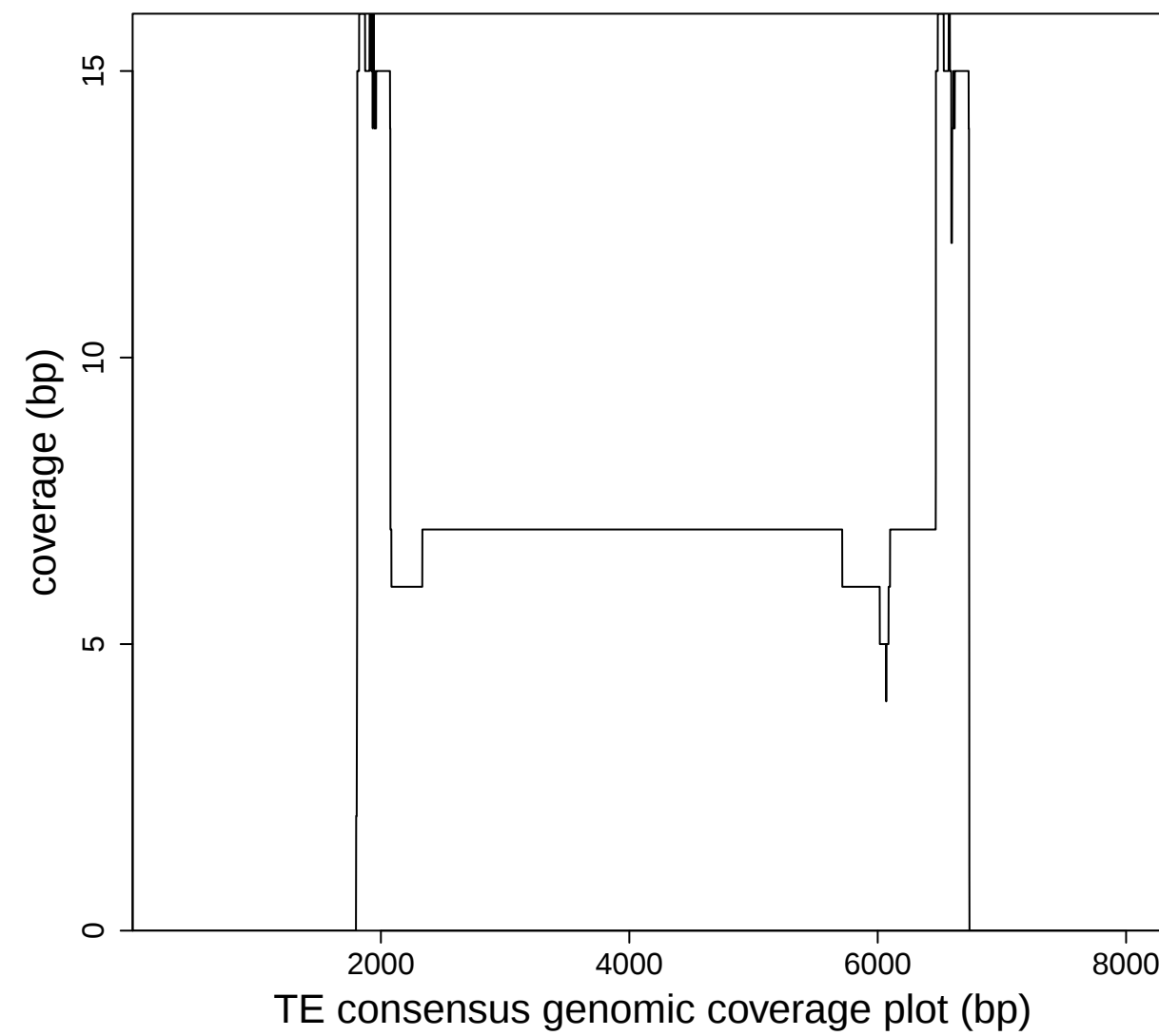
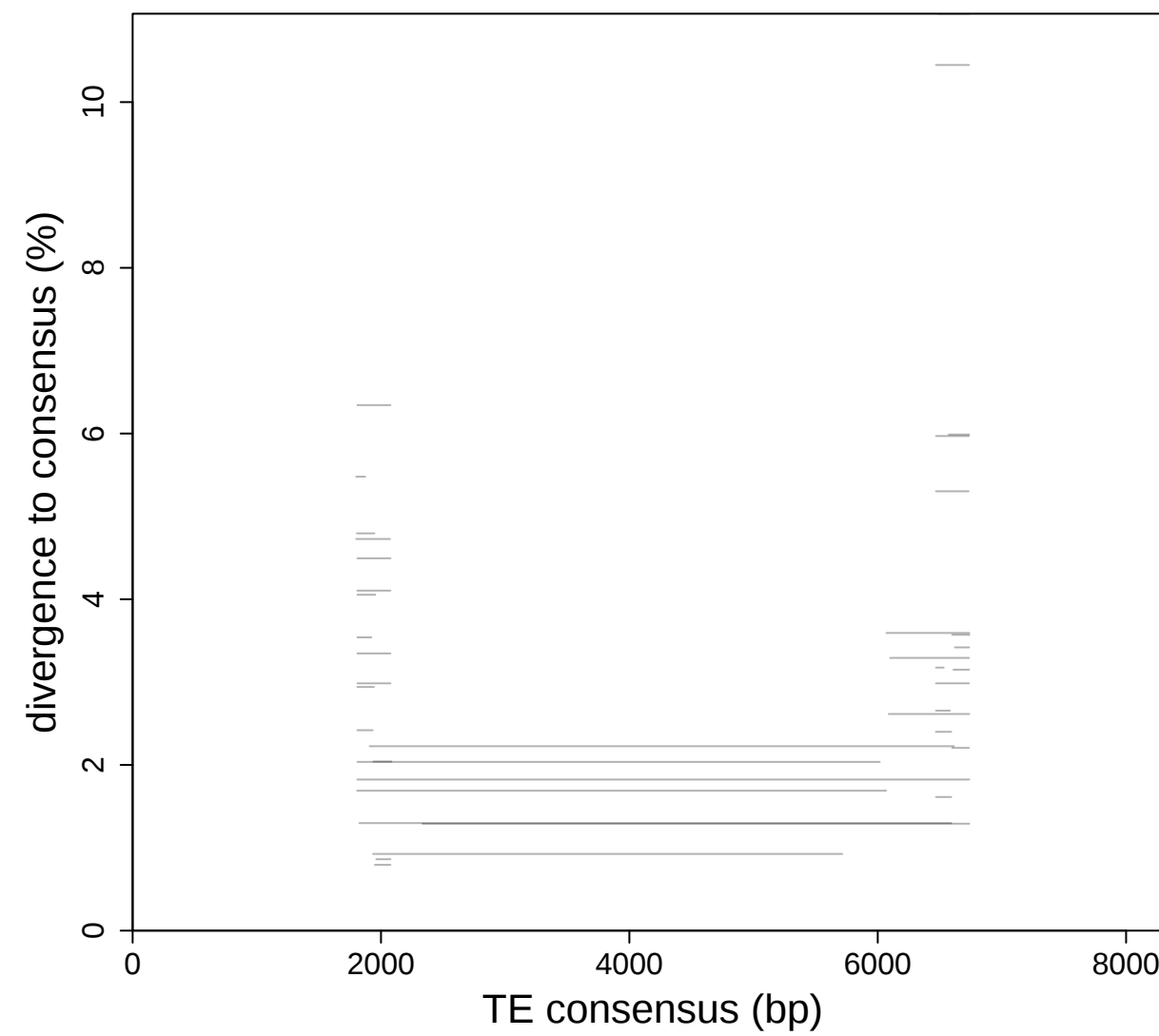
MSA length = 8377

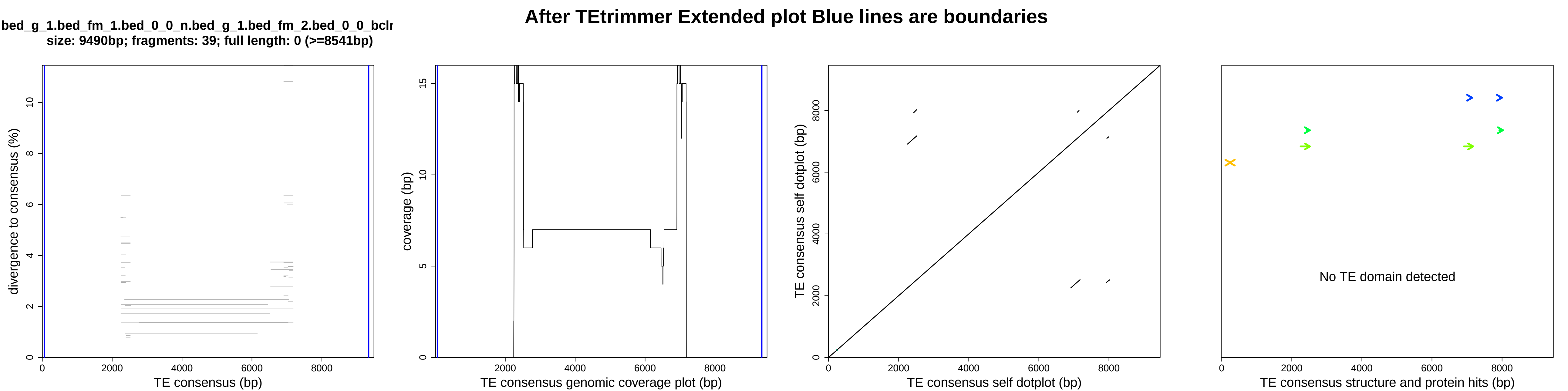




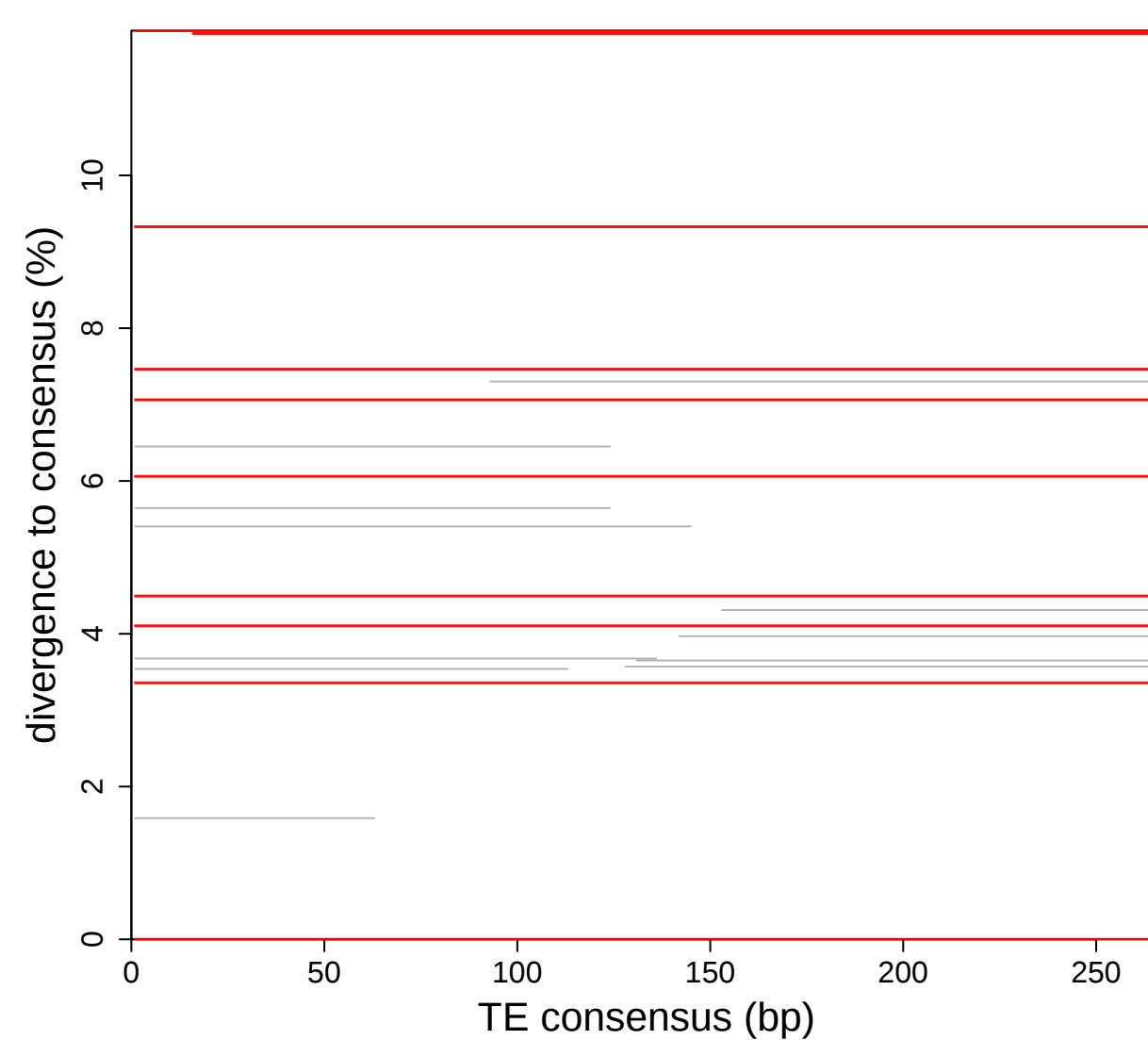
_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa_al
size: 8377bp; fragments: 39; full length: 0 (>=7539.3bp)

After TEtrimmer 8377 bp

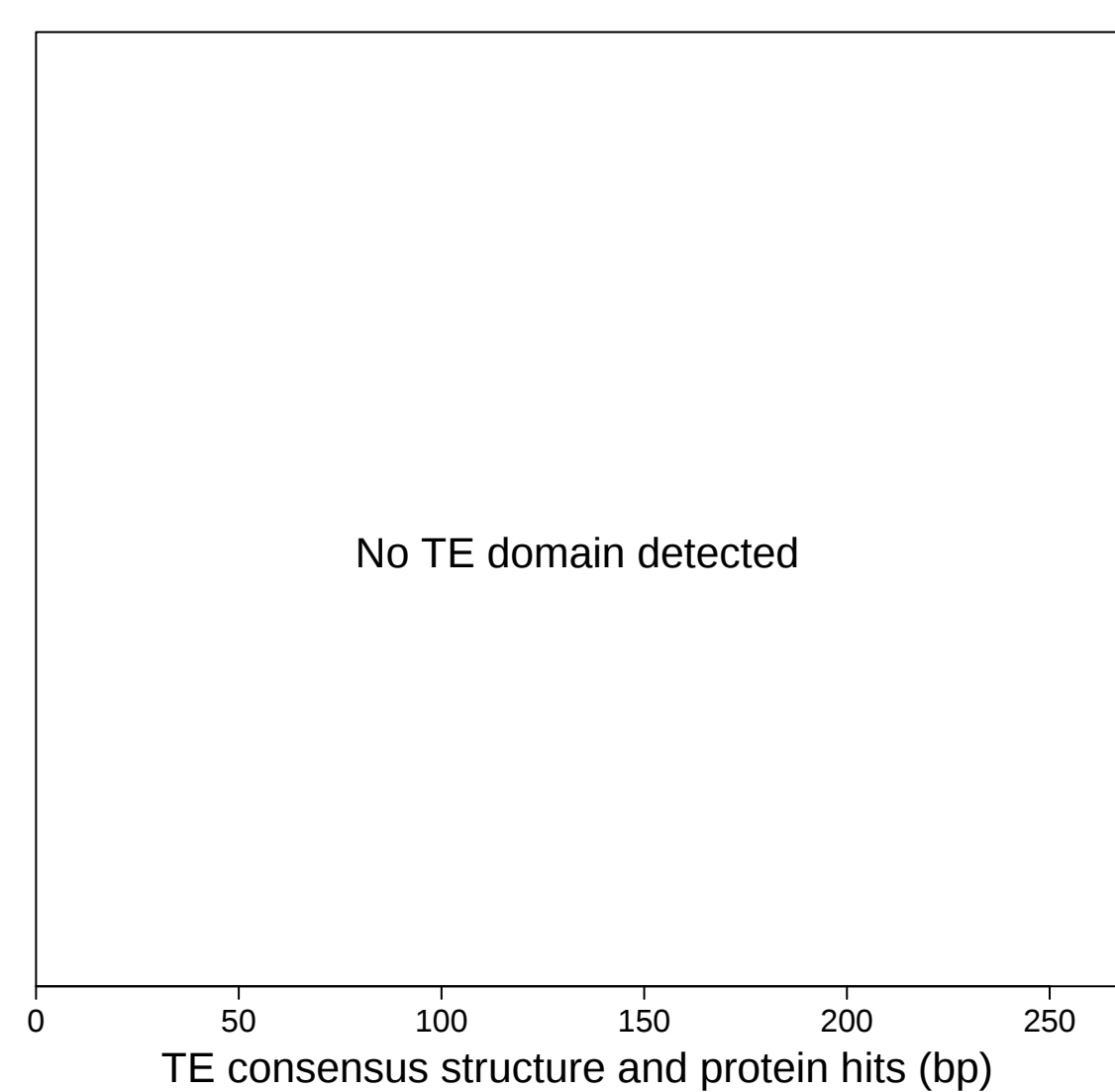
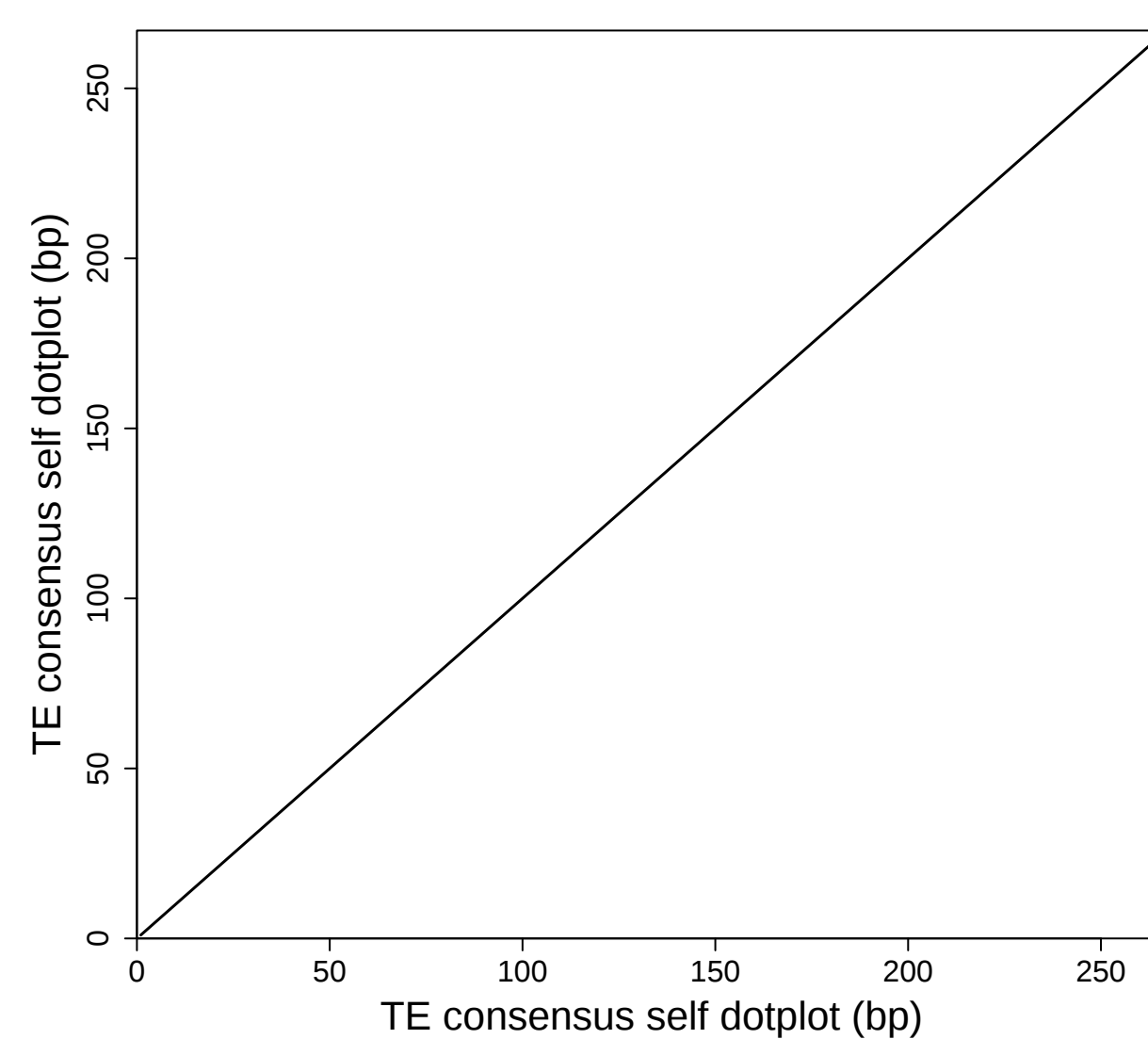
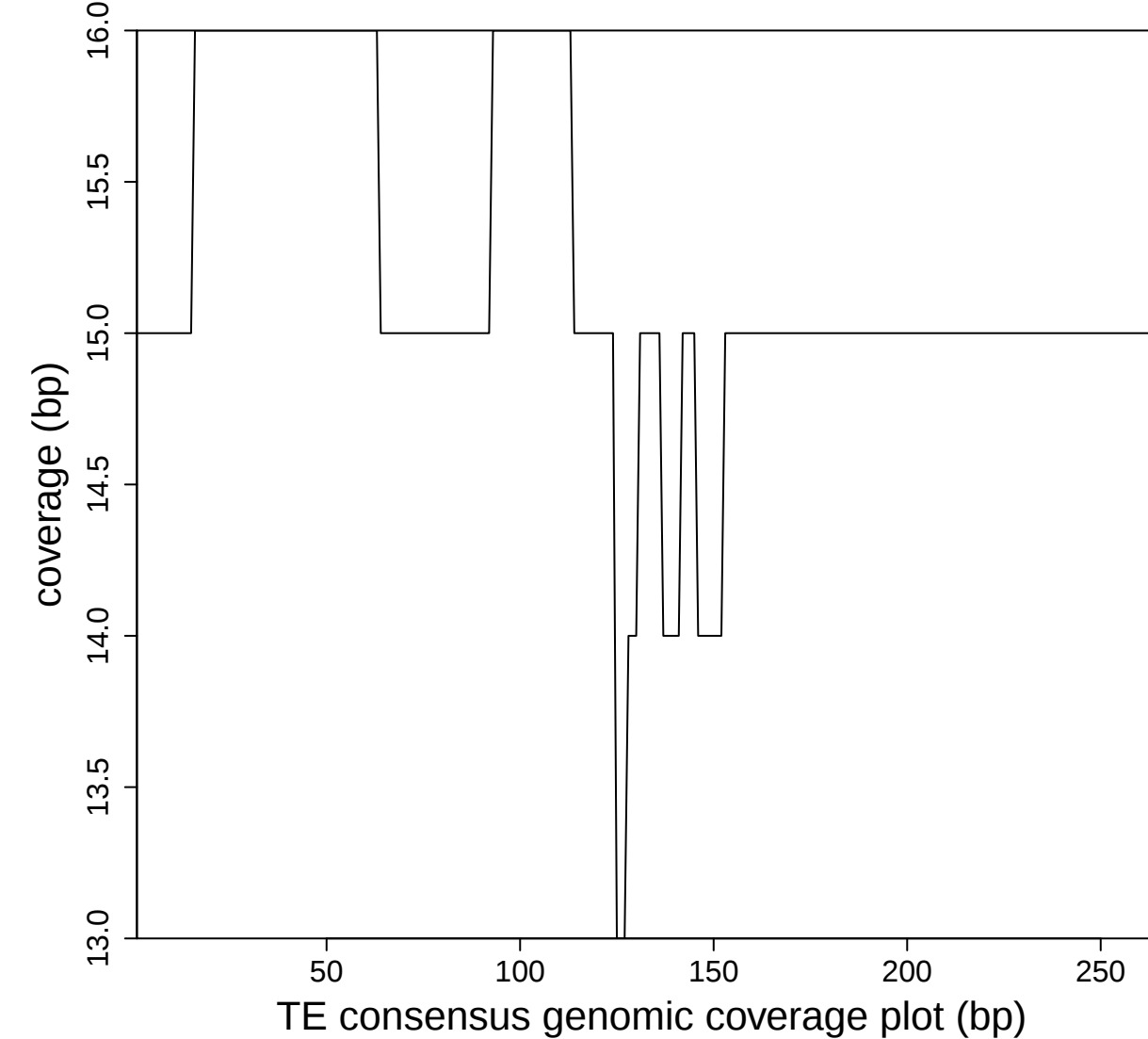




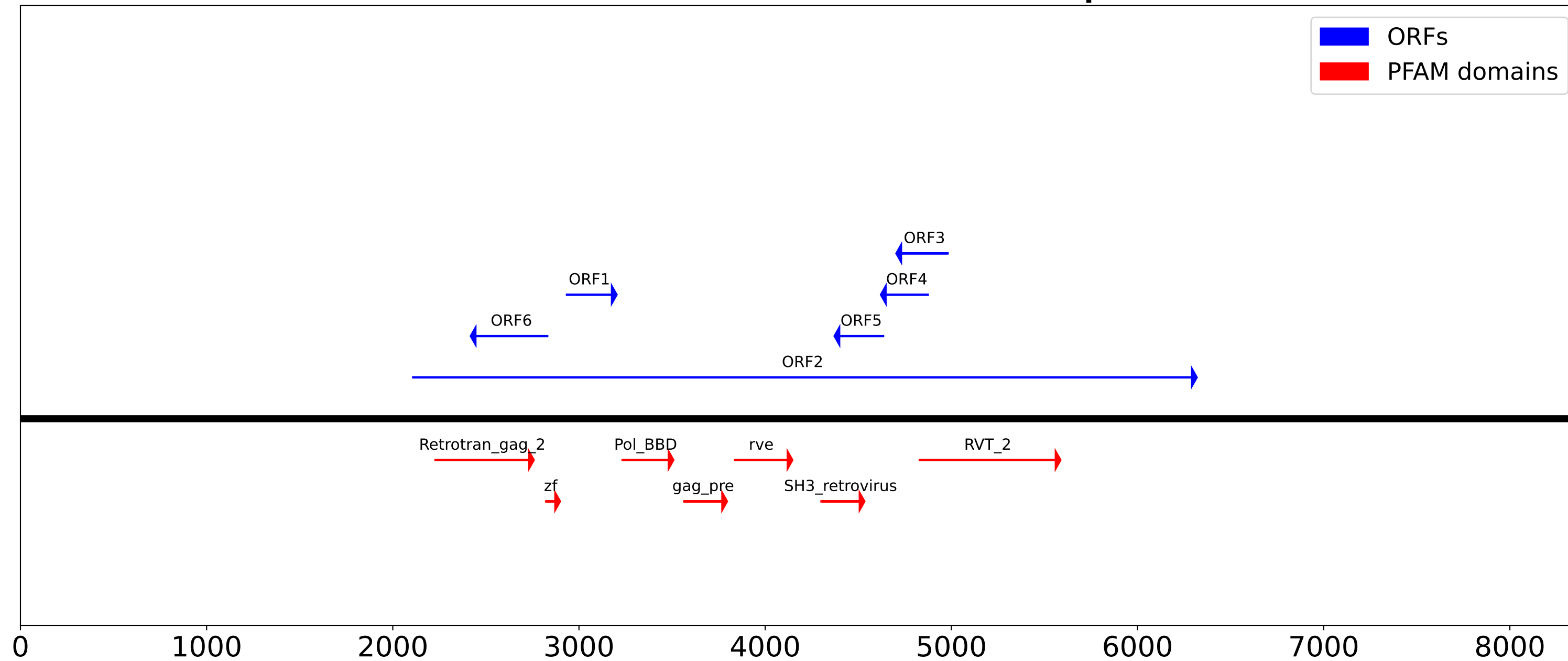
TE: ltr_1_family_115
size: 267bp; fragments: 21; full length: 10 (≥ 240.3 bp)



Before TEtrimmer 267 bp



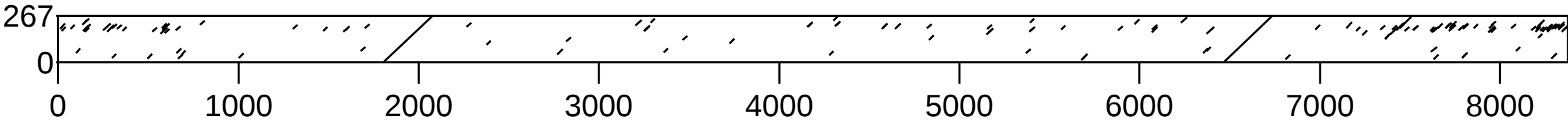
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)