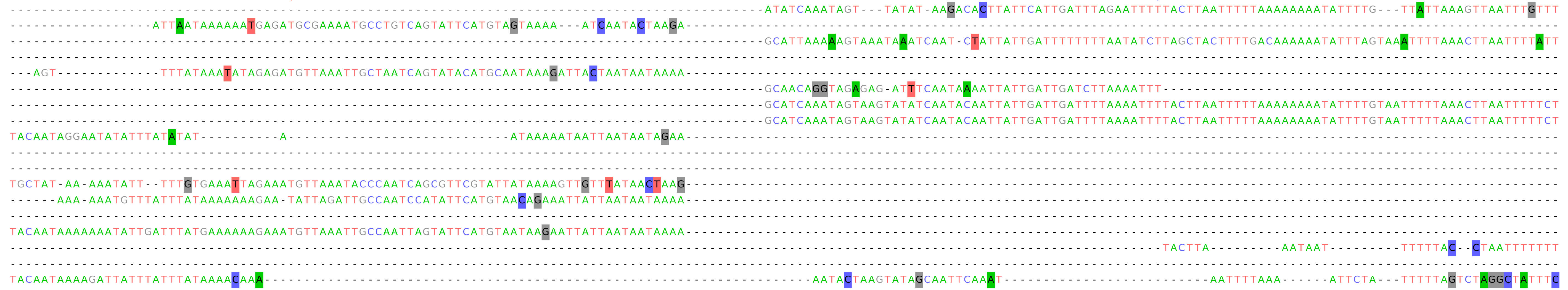


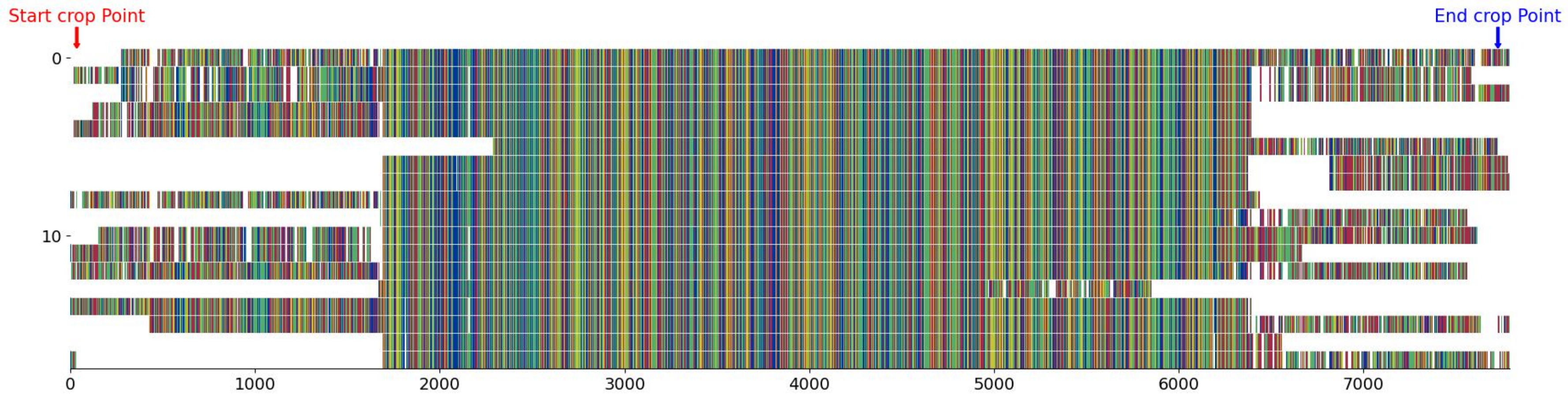
MSA length = 7694

Start crop Point



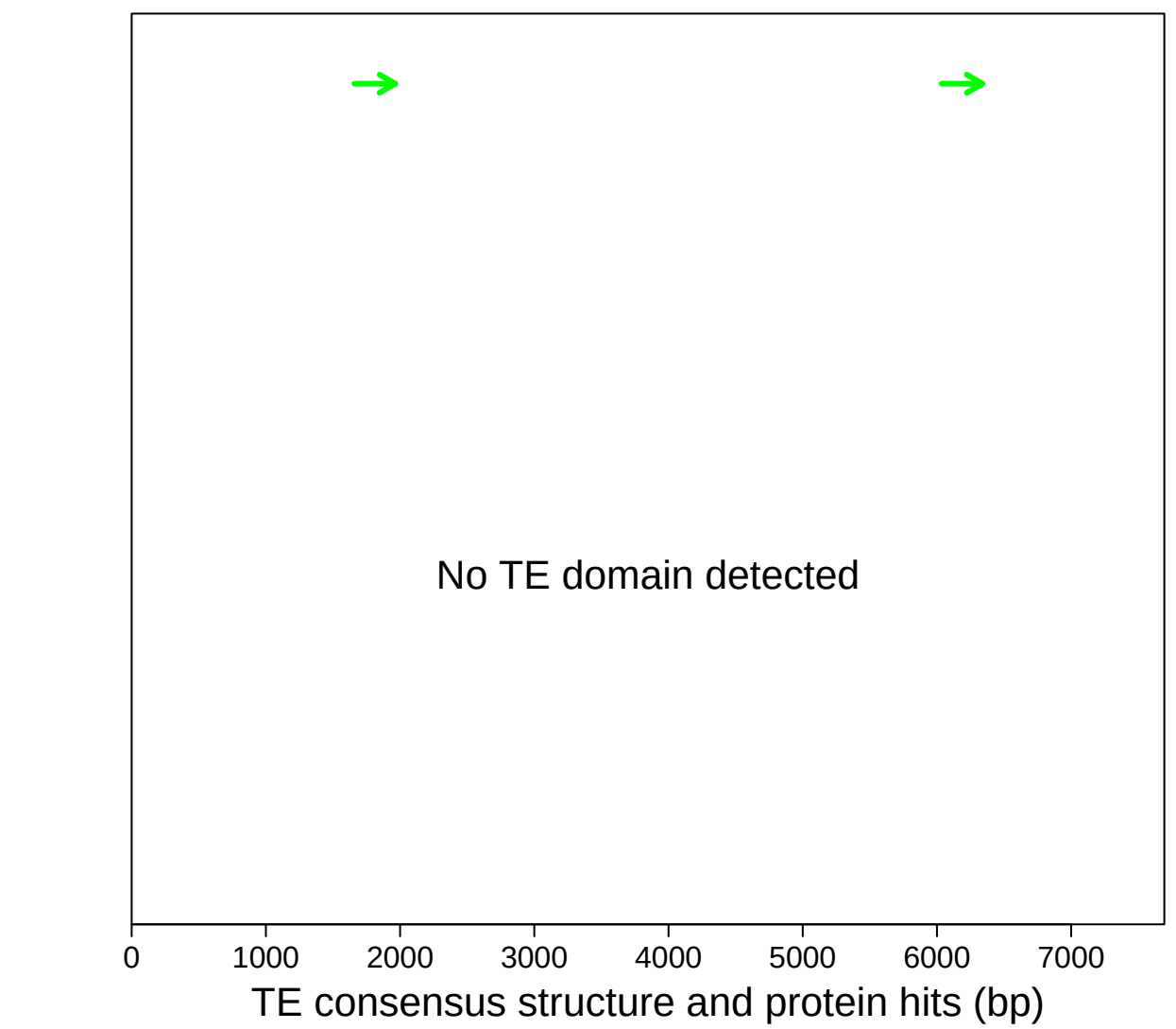
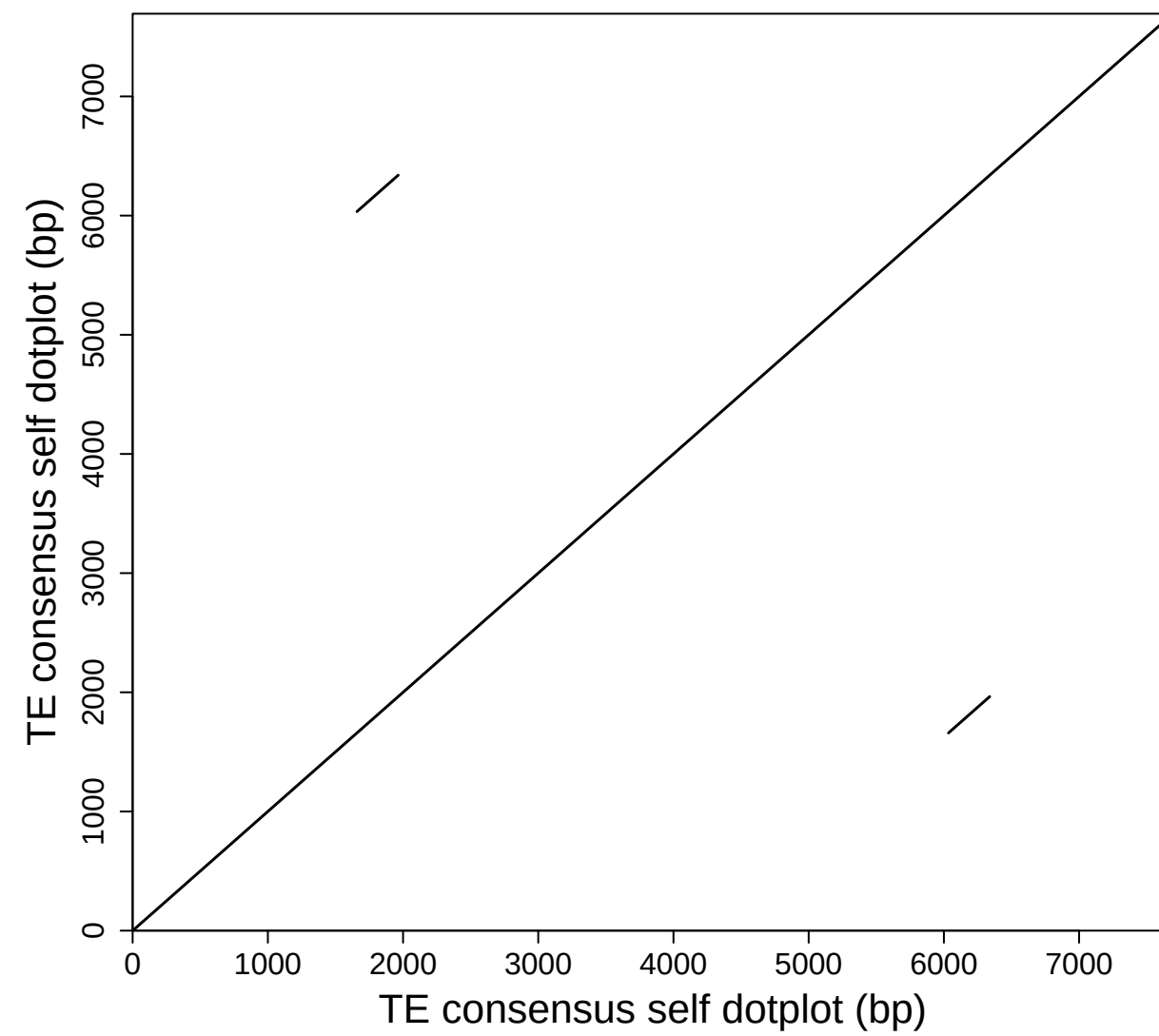
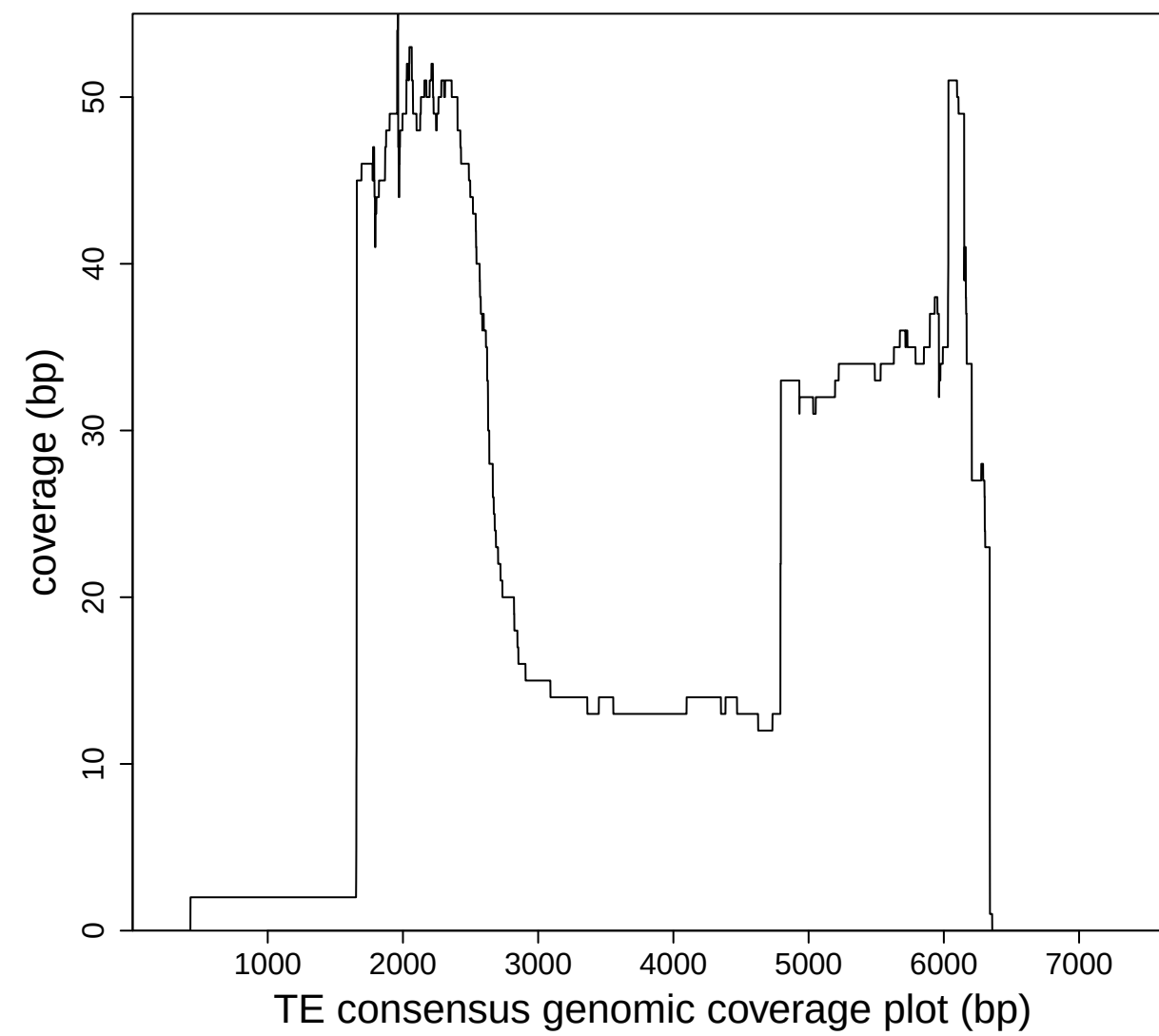
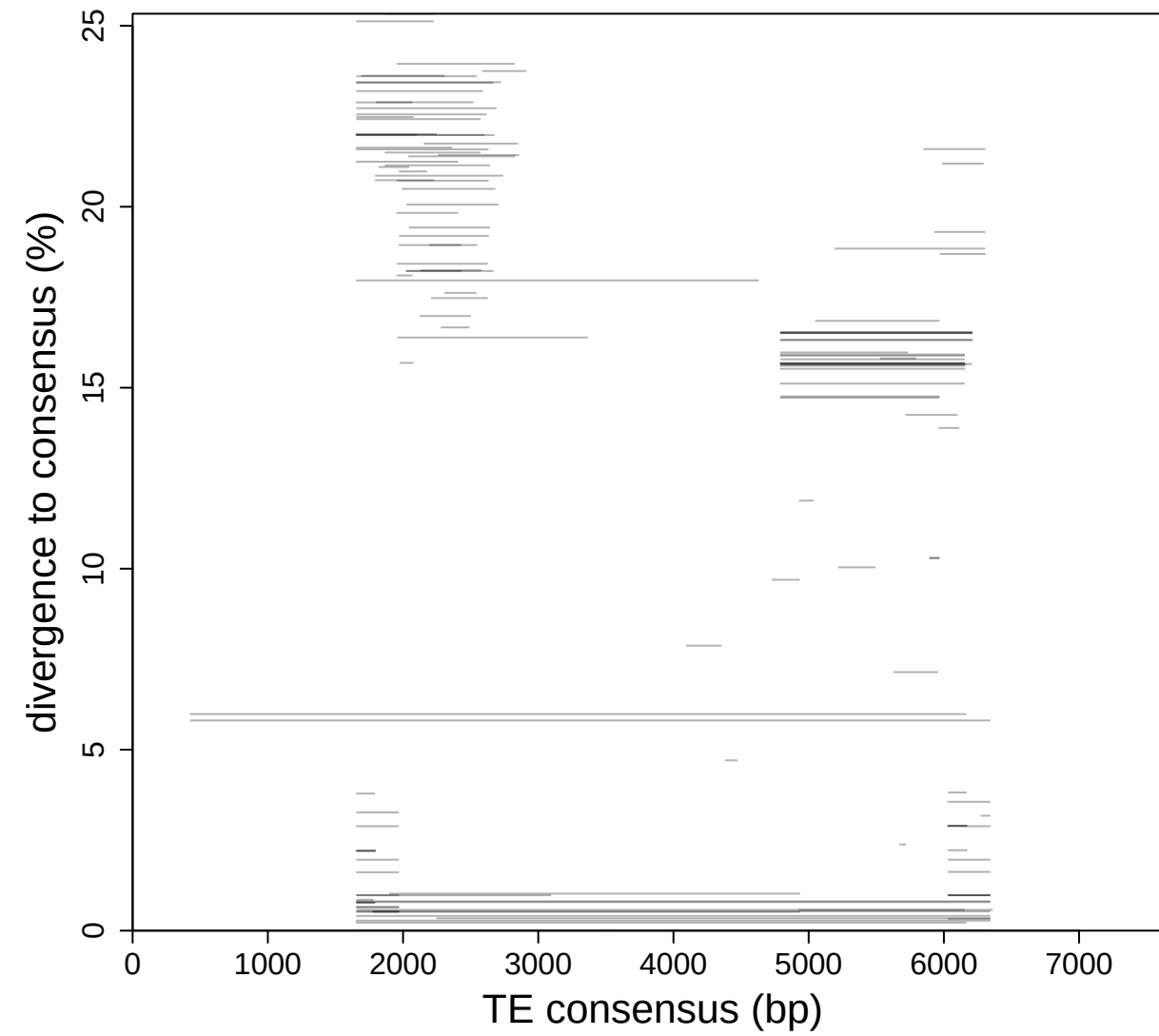
End crop Point

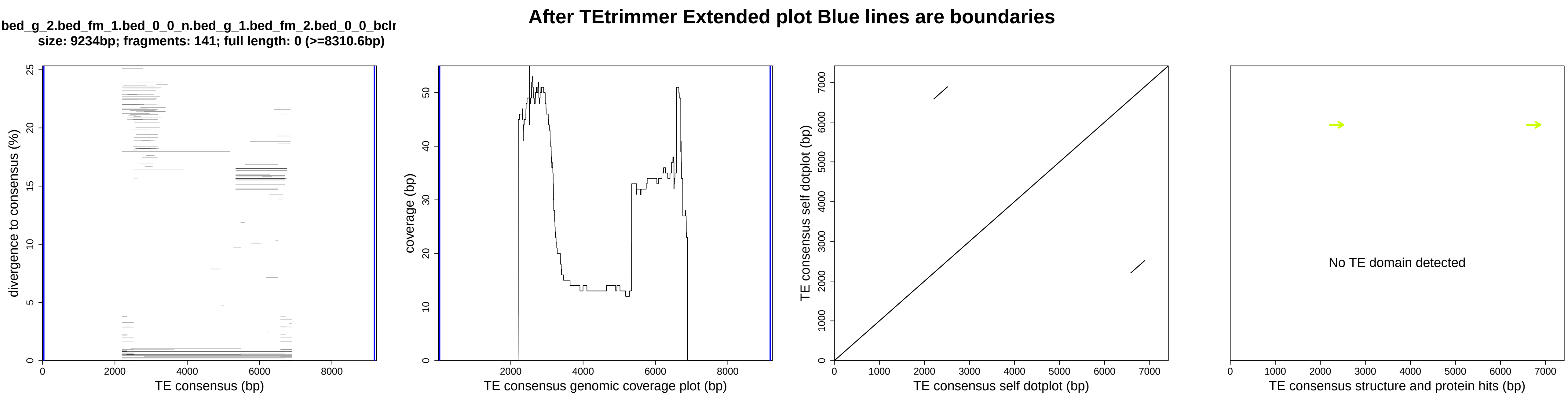




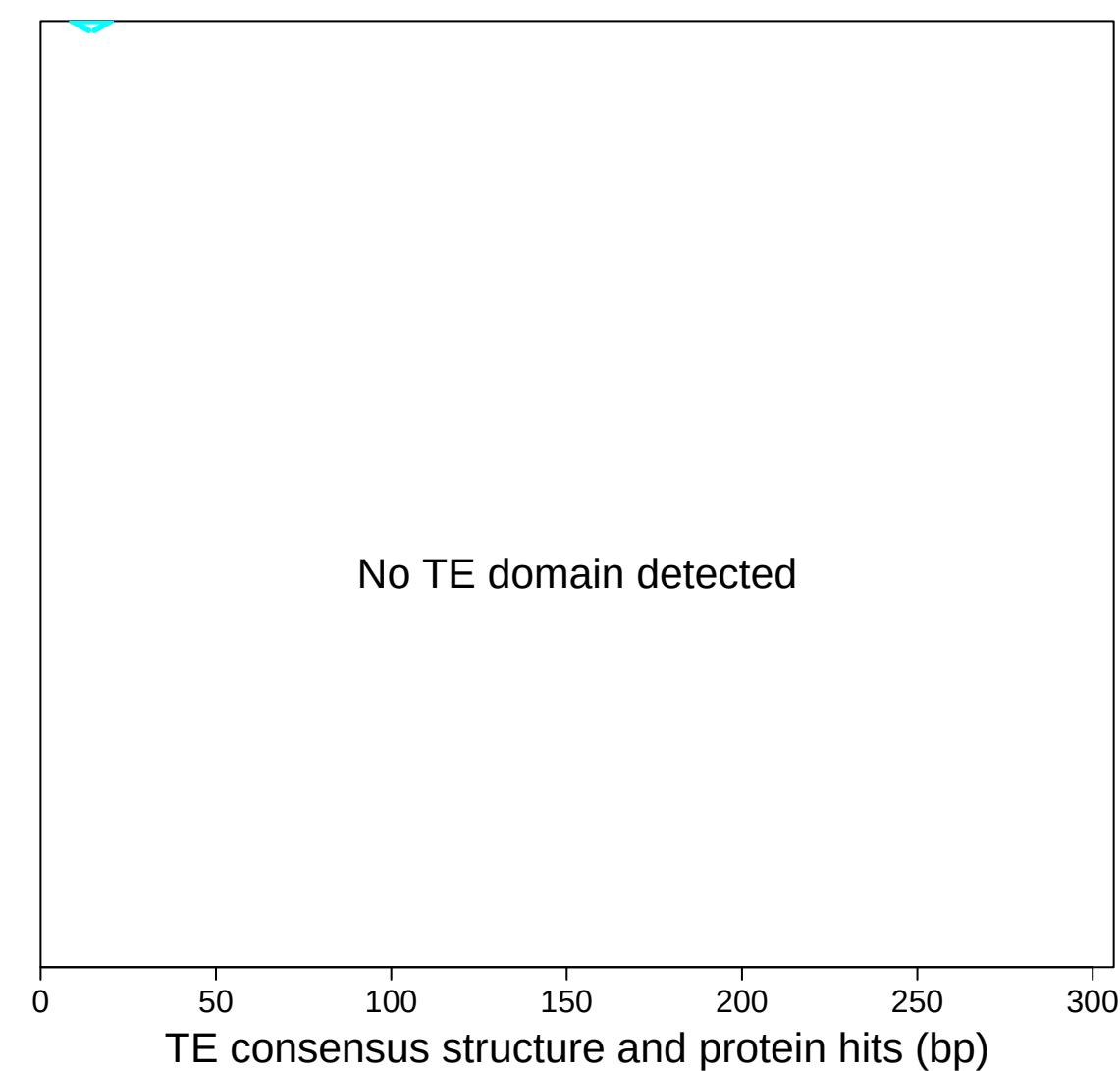
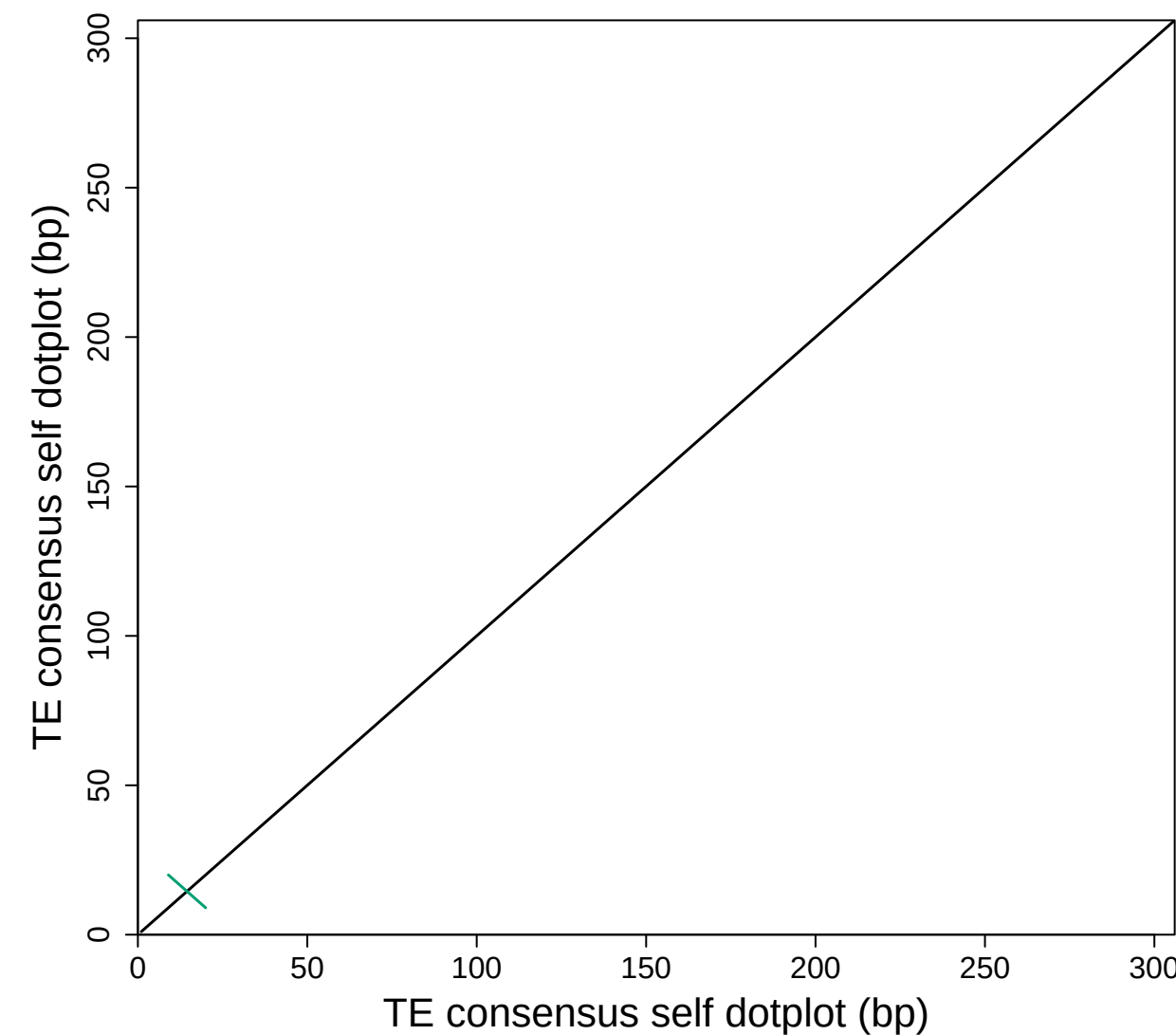
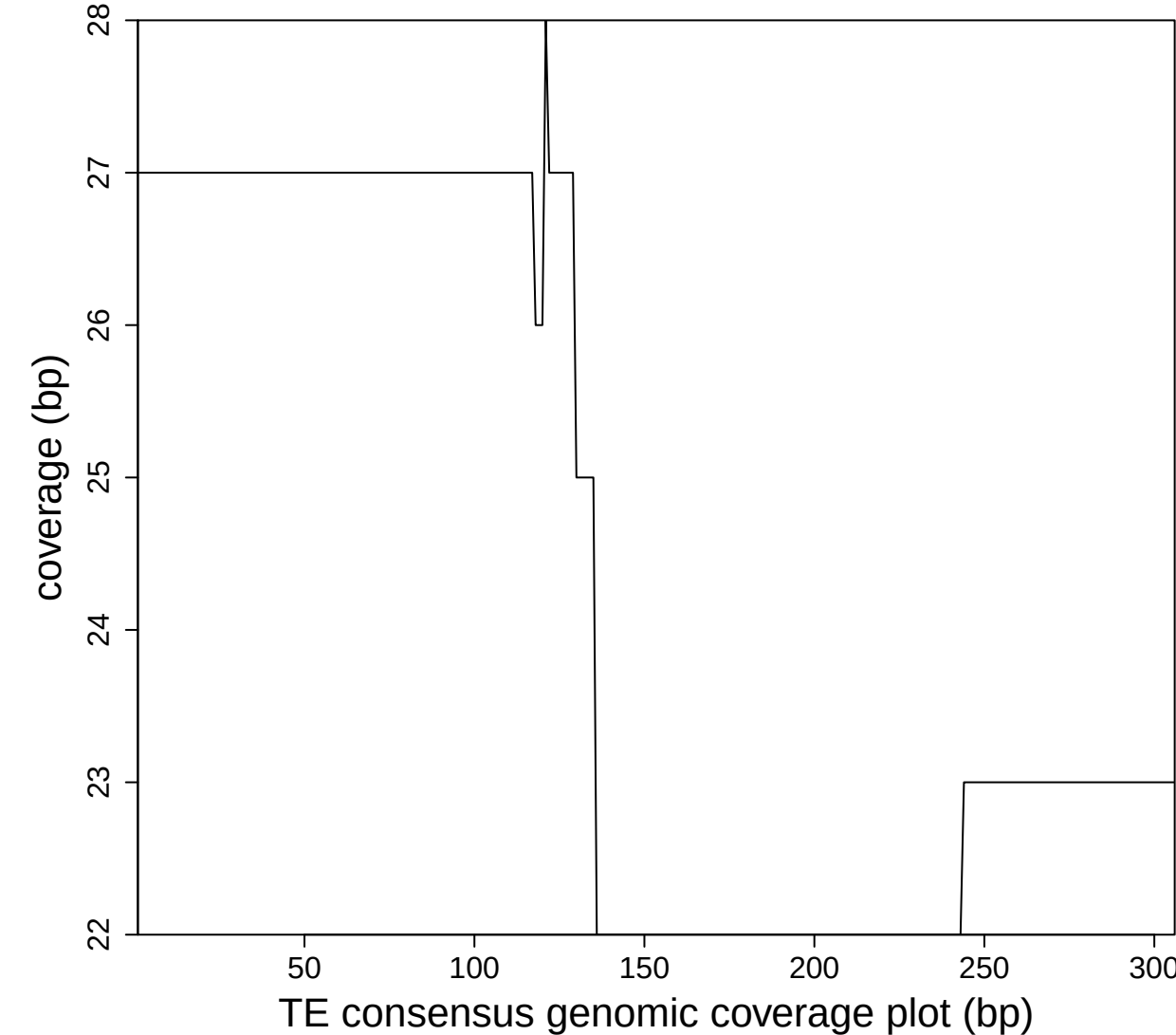
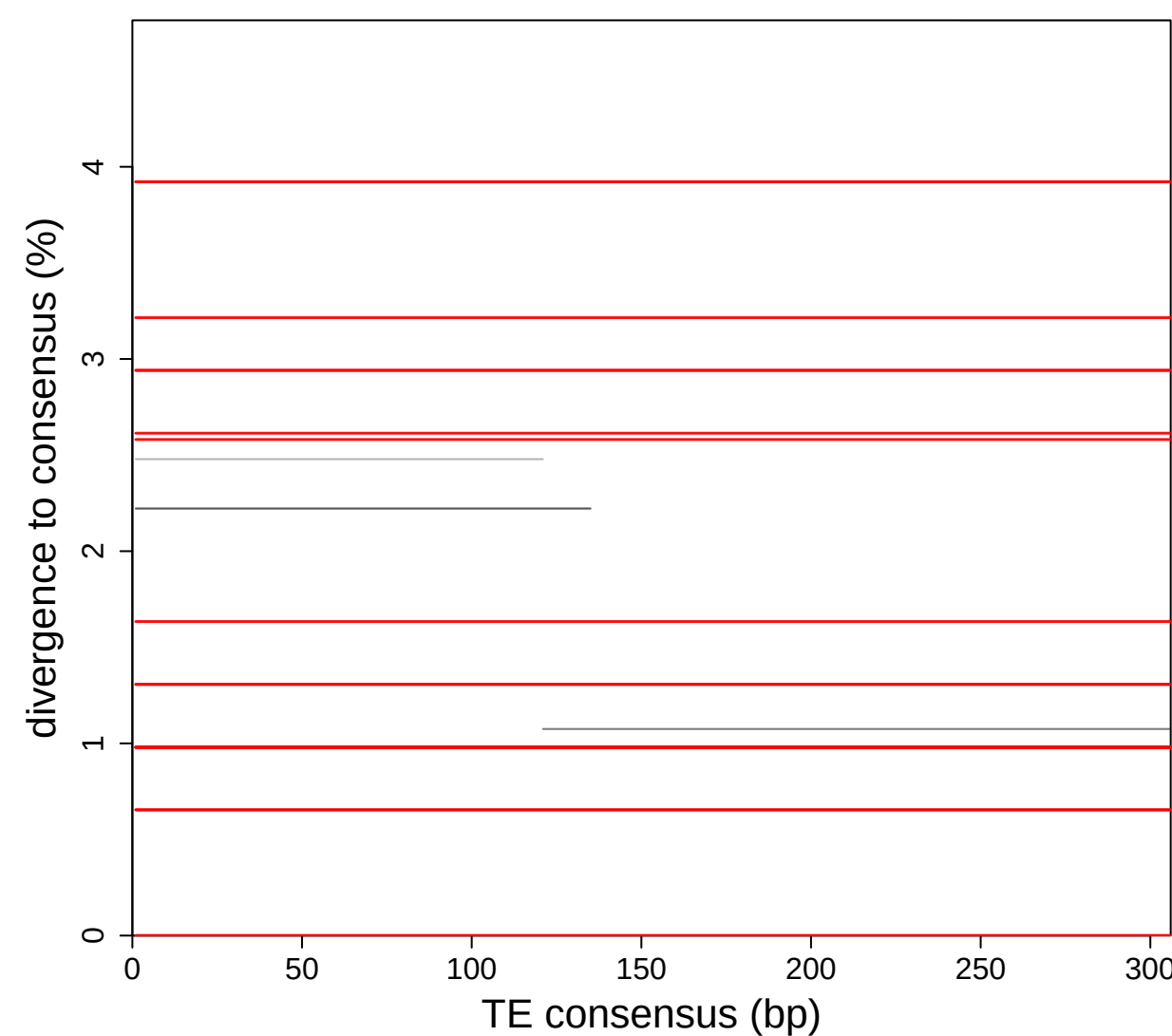
_2.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_al
size: 7694bp; fragments: 141; full length: 0 (>=6924.6bp)

After TEtrimmer 7694 bp



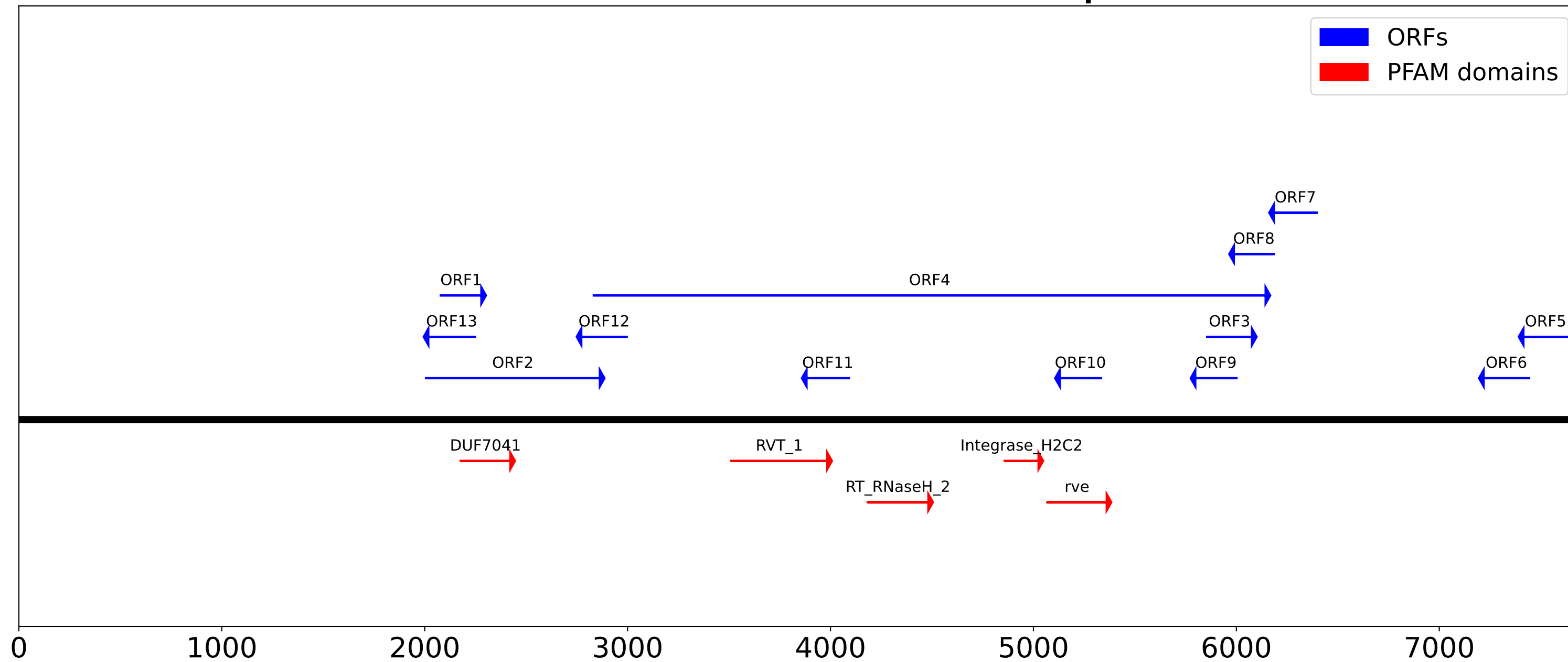


TE: ltr_1_family_137
size: 306bp; fragments: 30; full length: 20 (≥ 275.4 bp)



Before TEtrimmer 306 bp

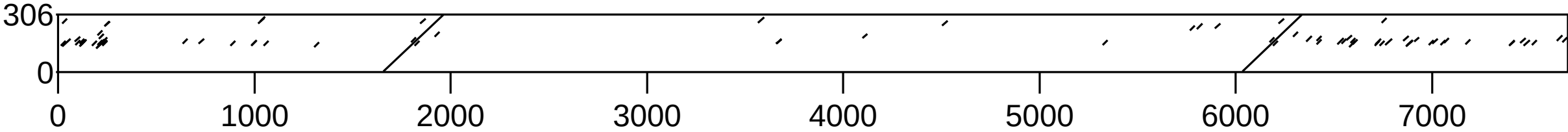
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)