

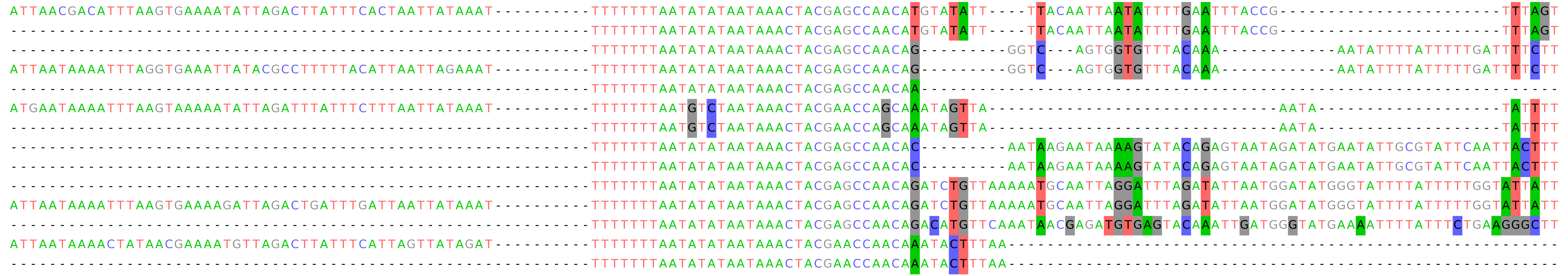
Start crop Point

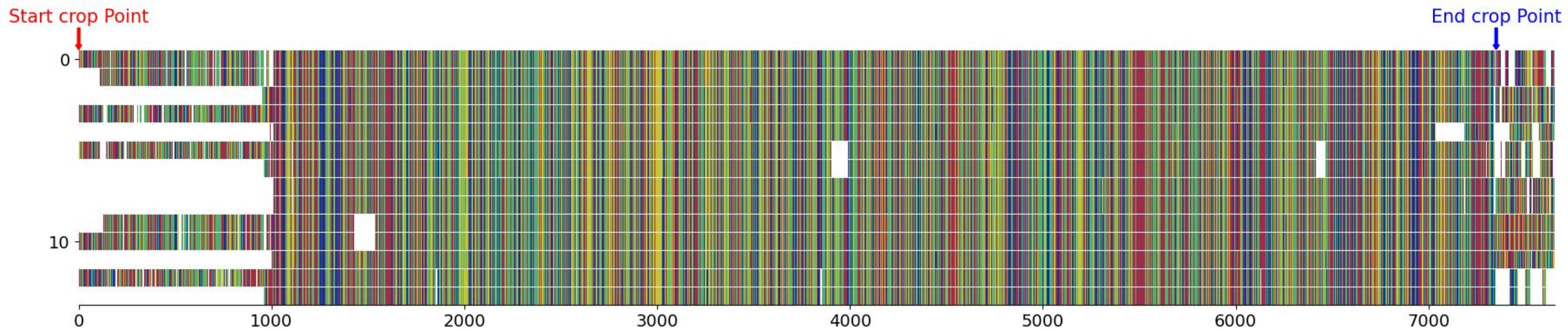


End crop Point



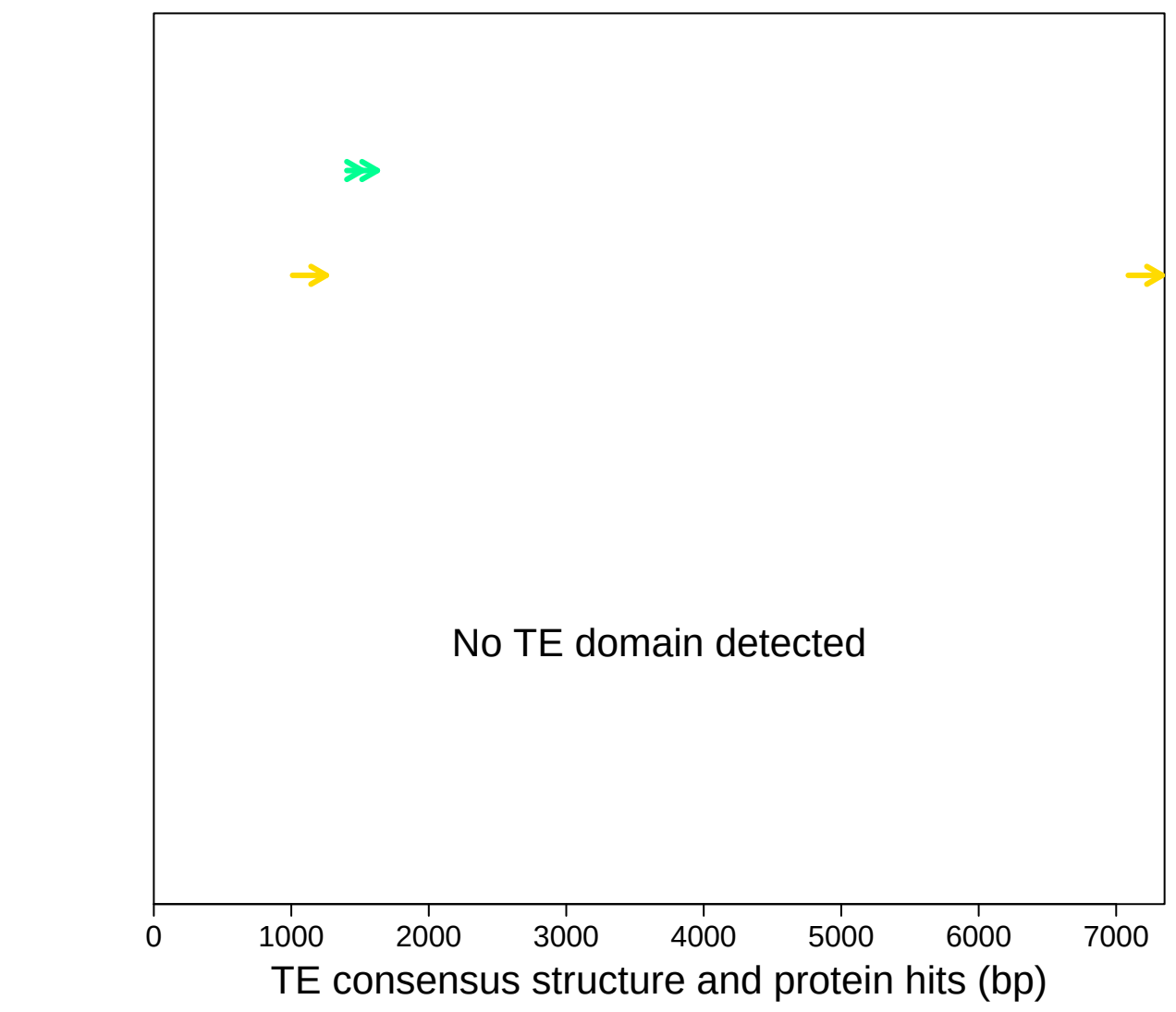
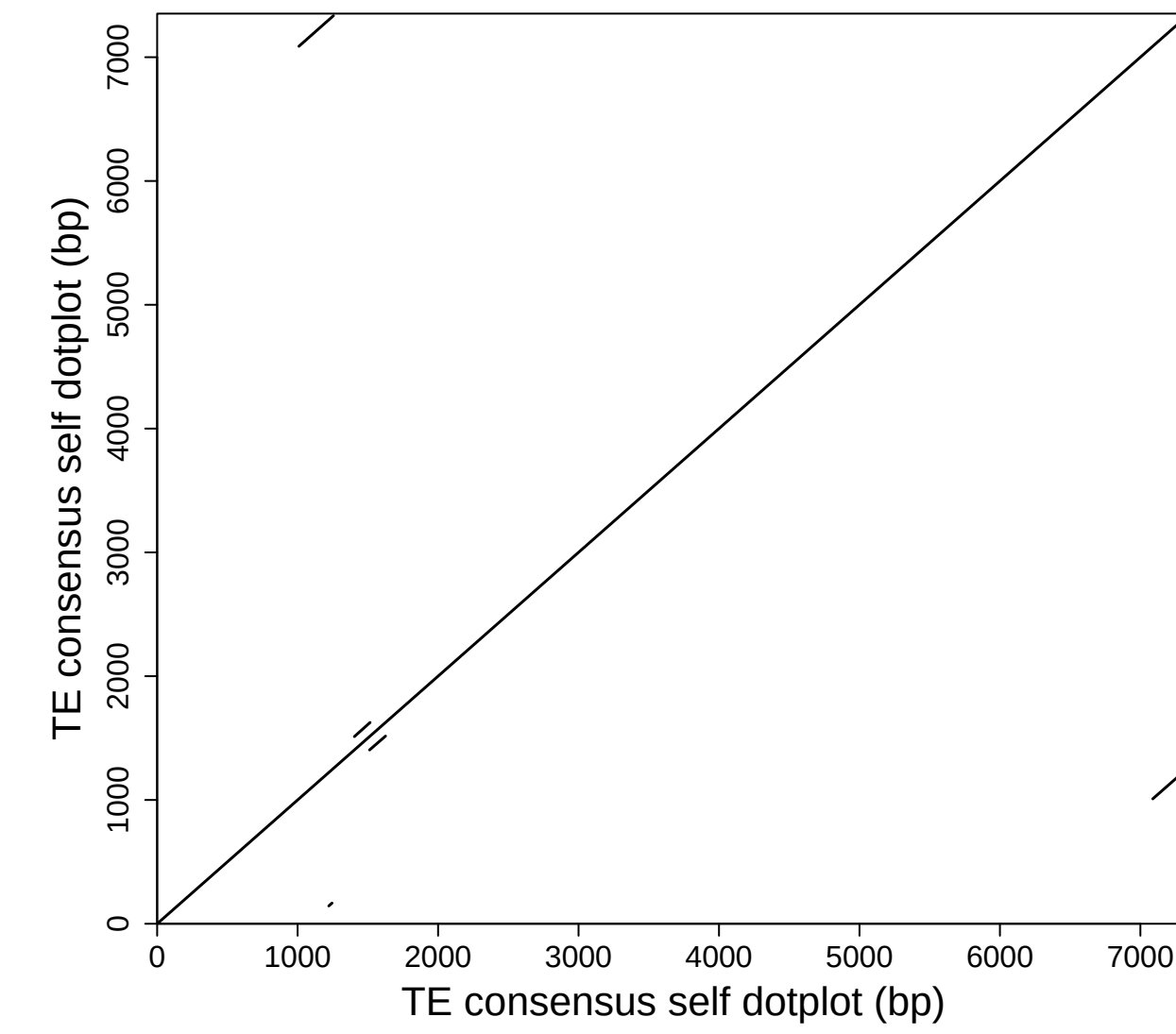
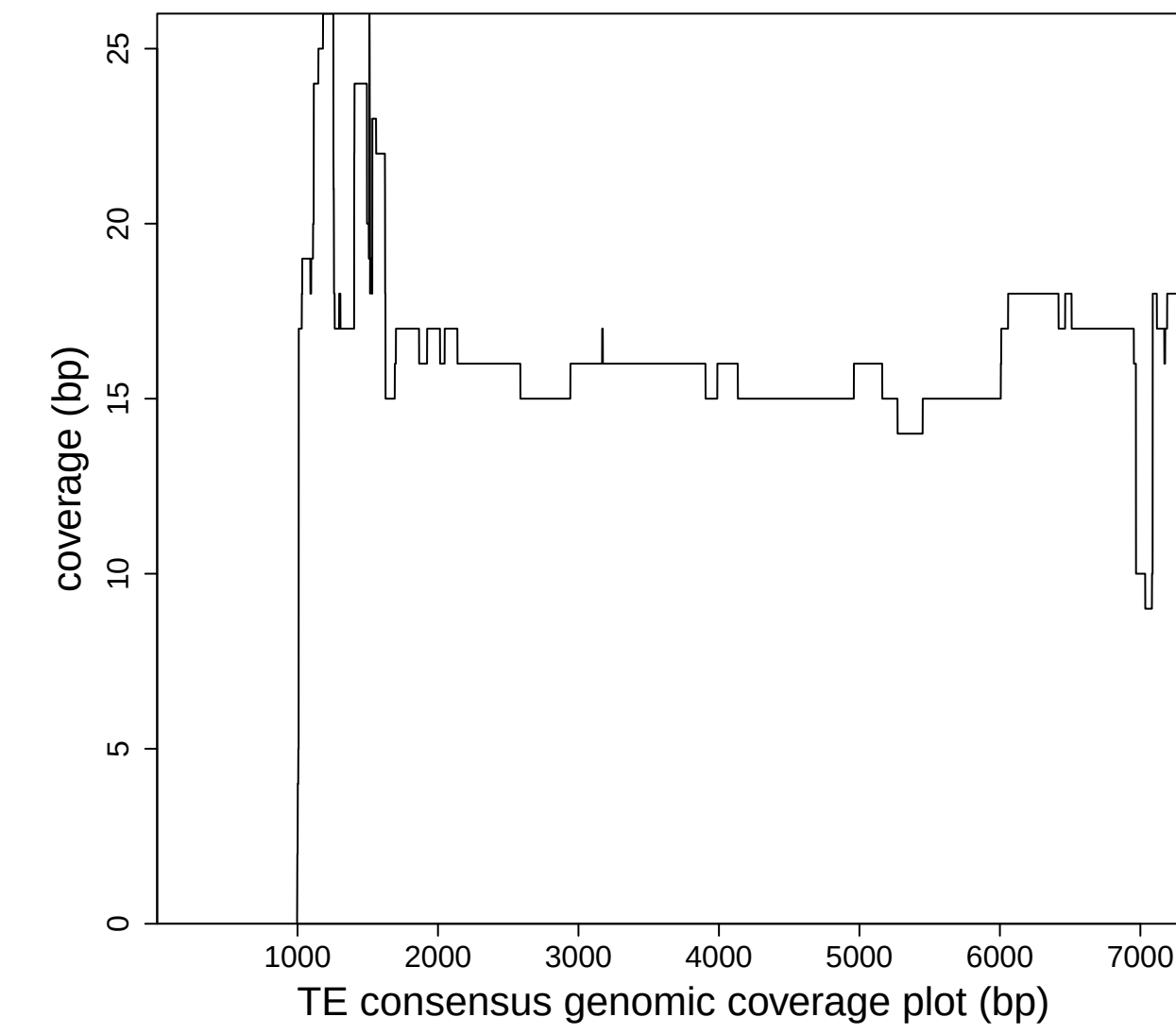
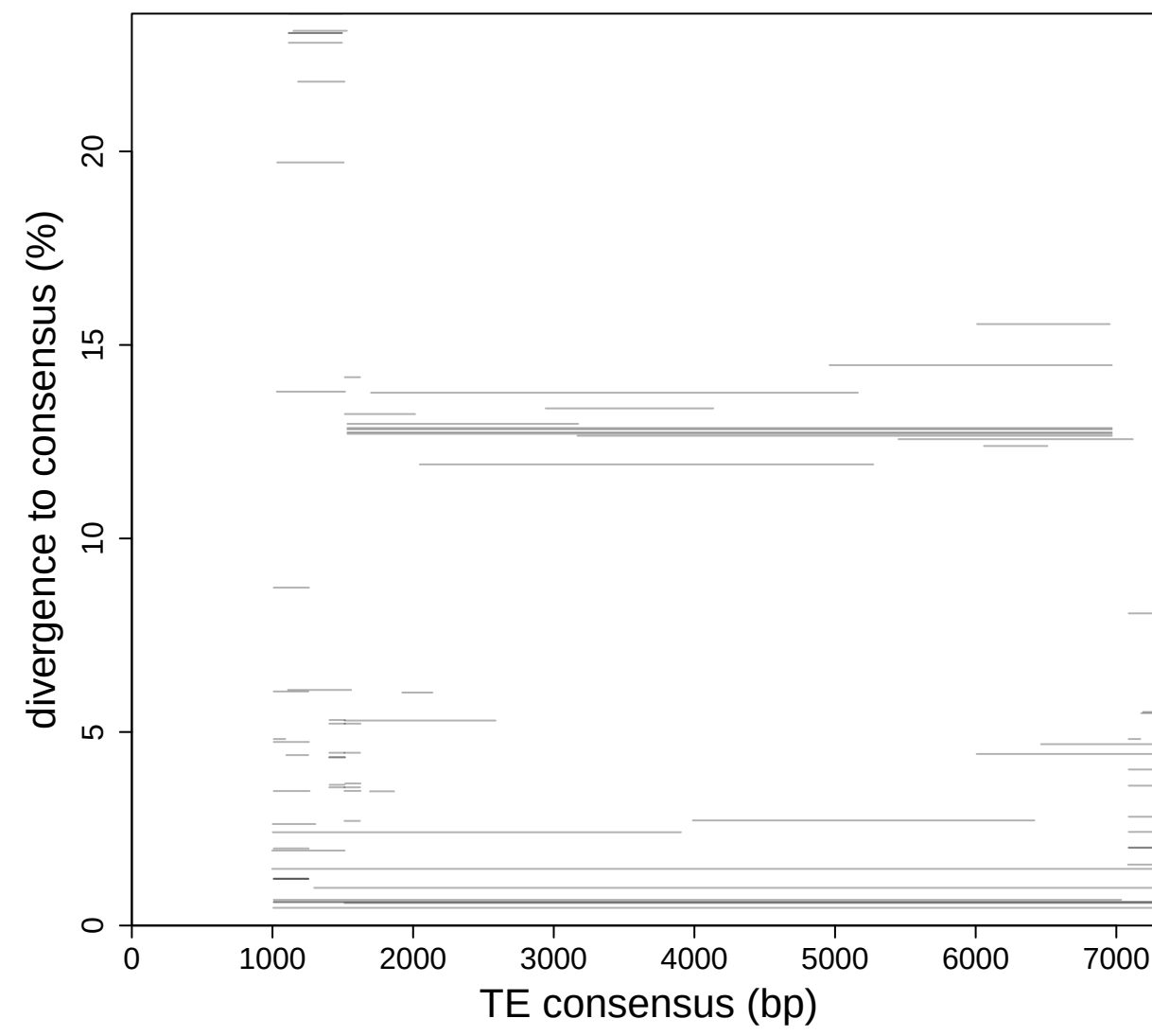
MSA length = 7352





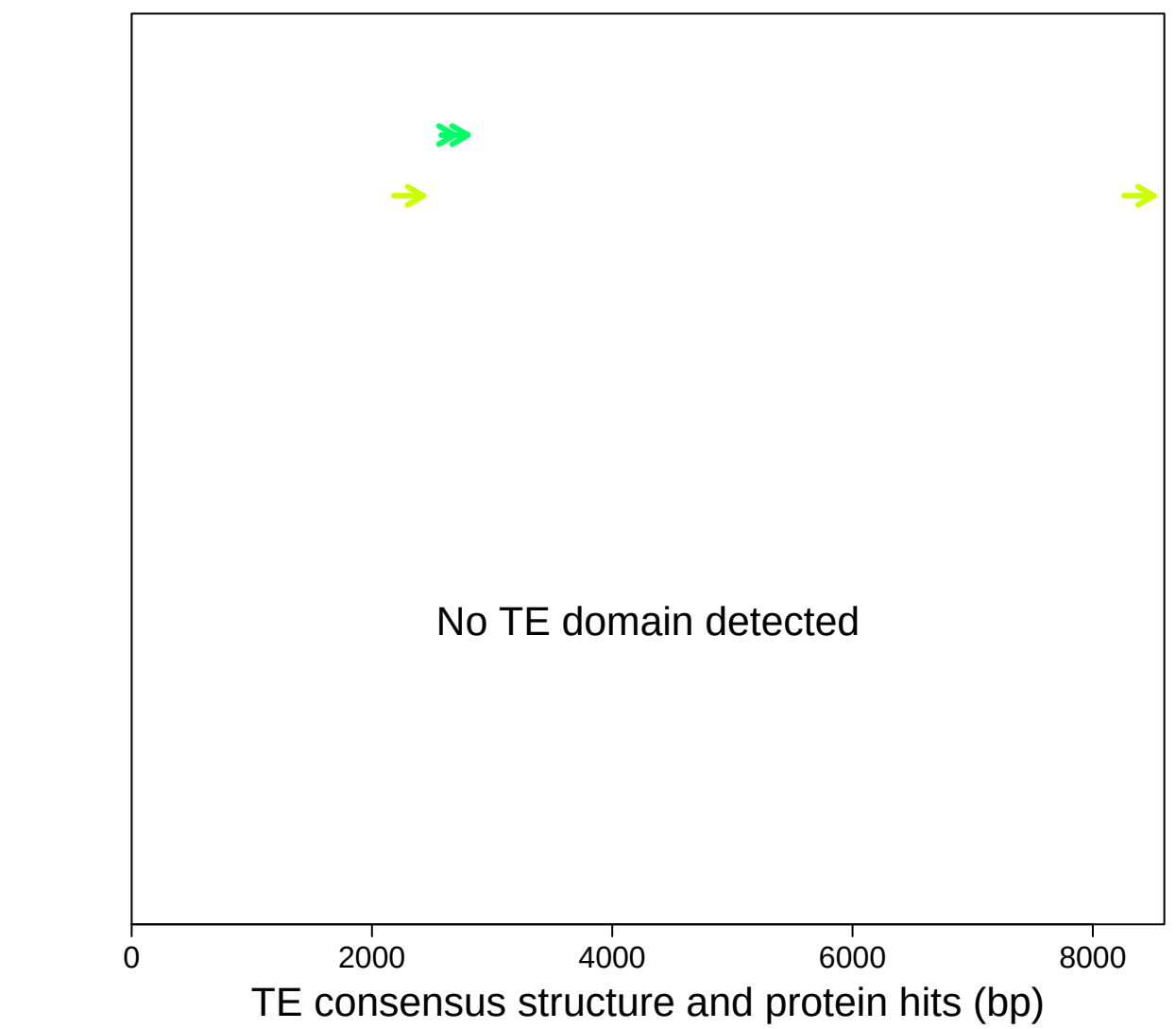
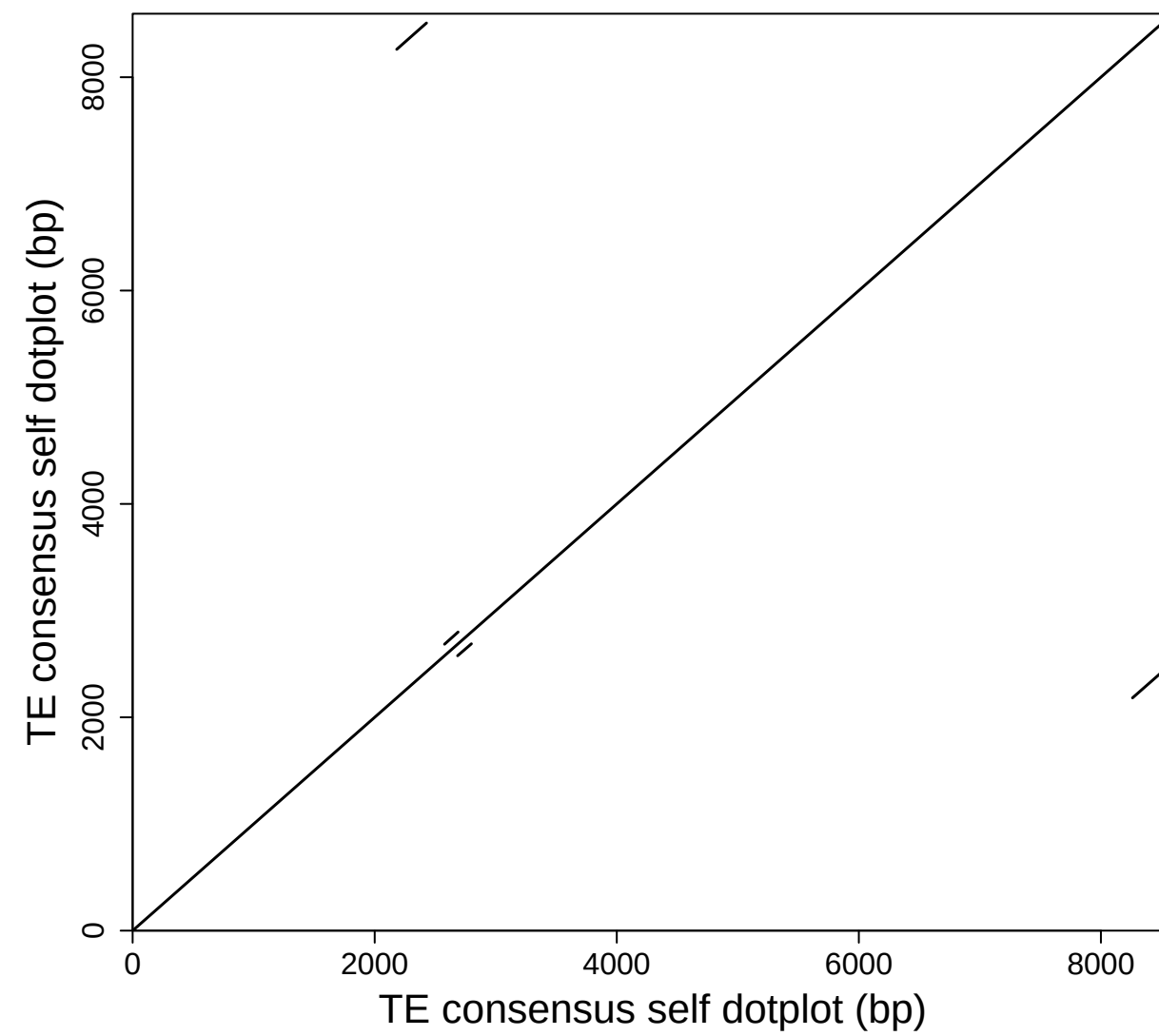
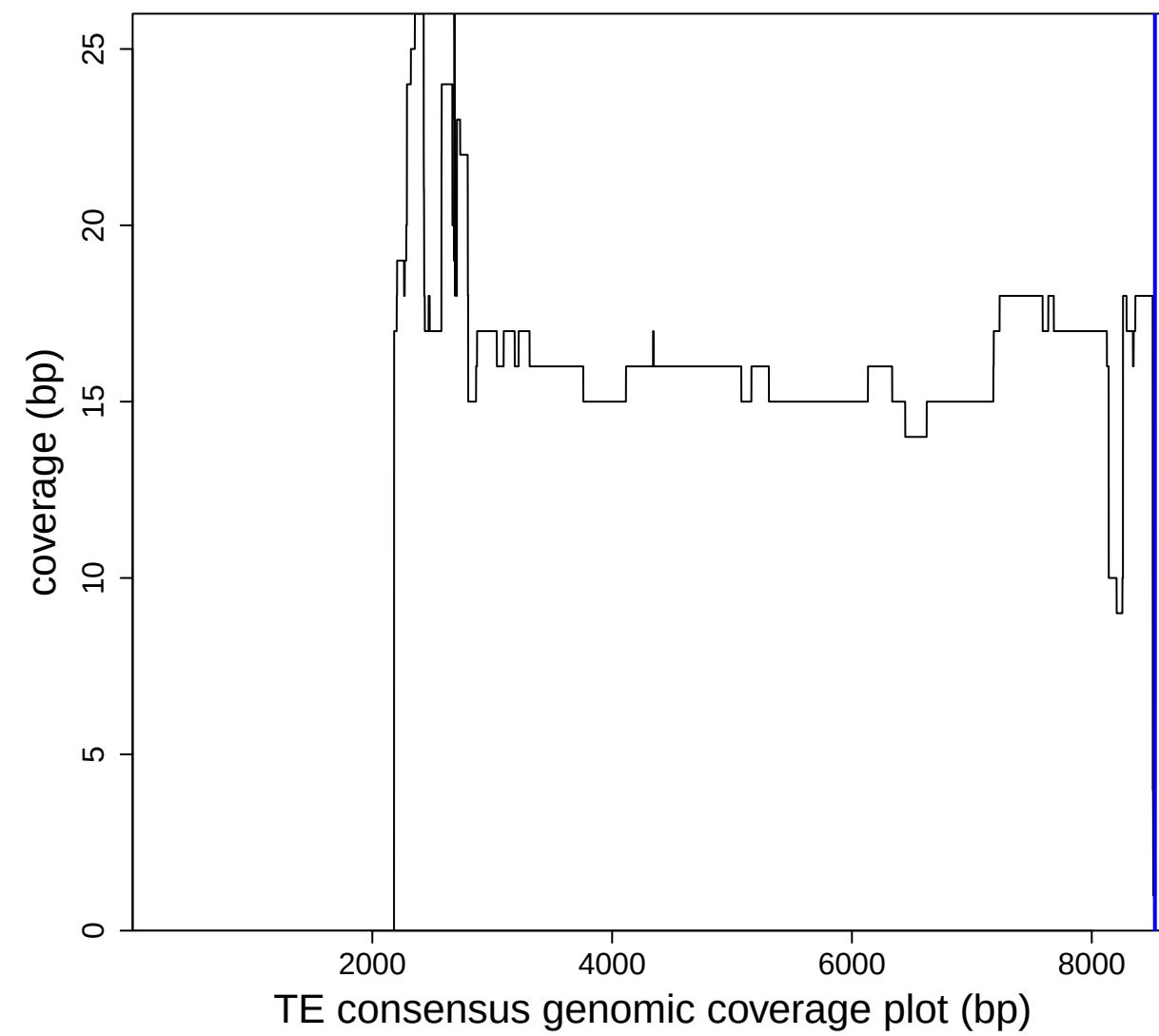
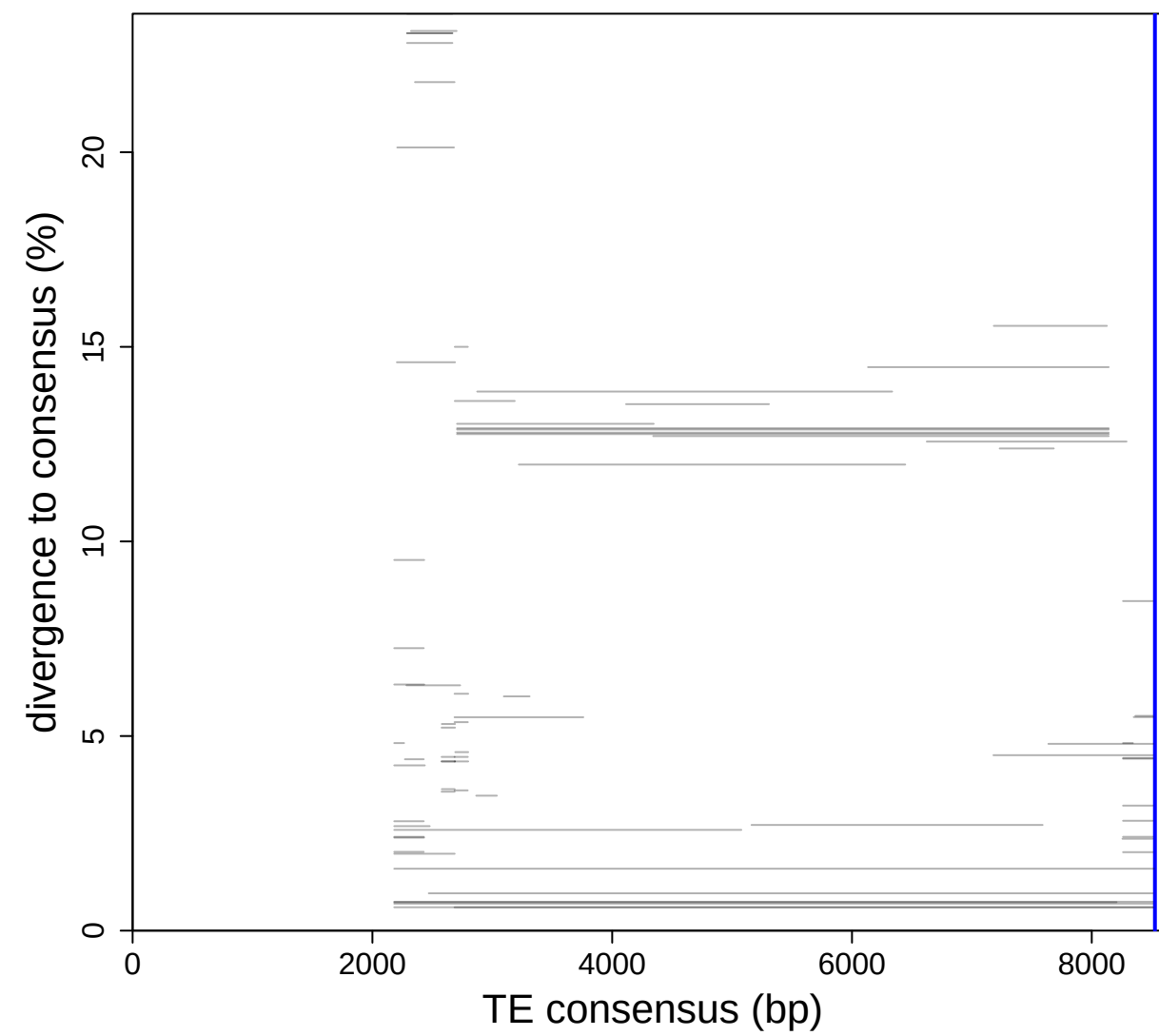
1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcIn.fa_alr
size: 7352bp; fragments: 74; full length: 0 (>=6616.8bp)

After TETrimmer 7352 bp



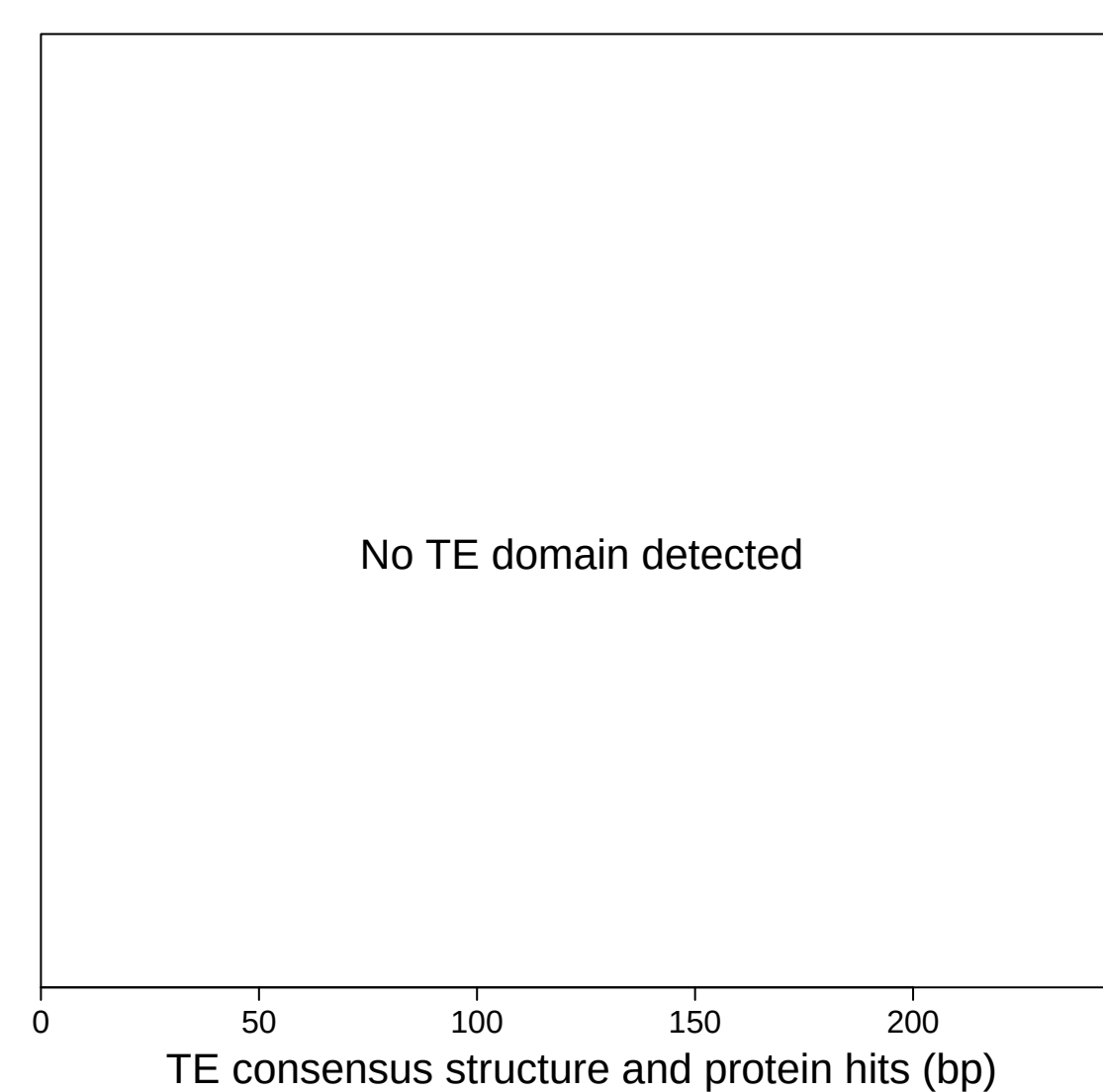
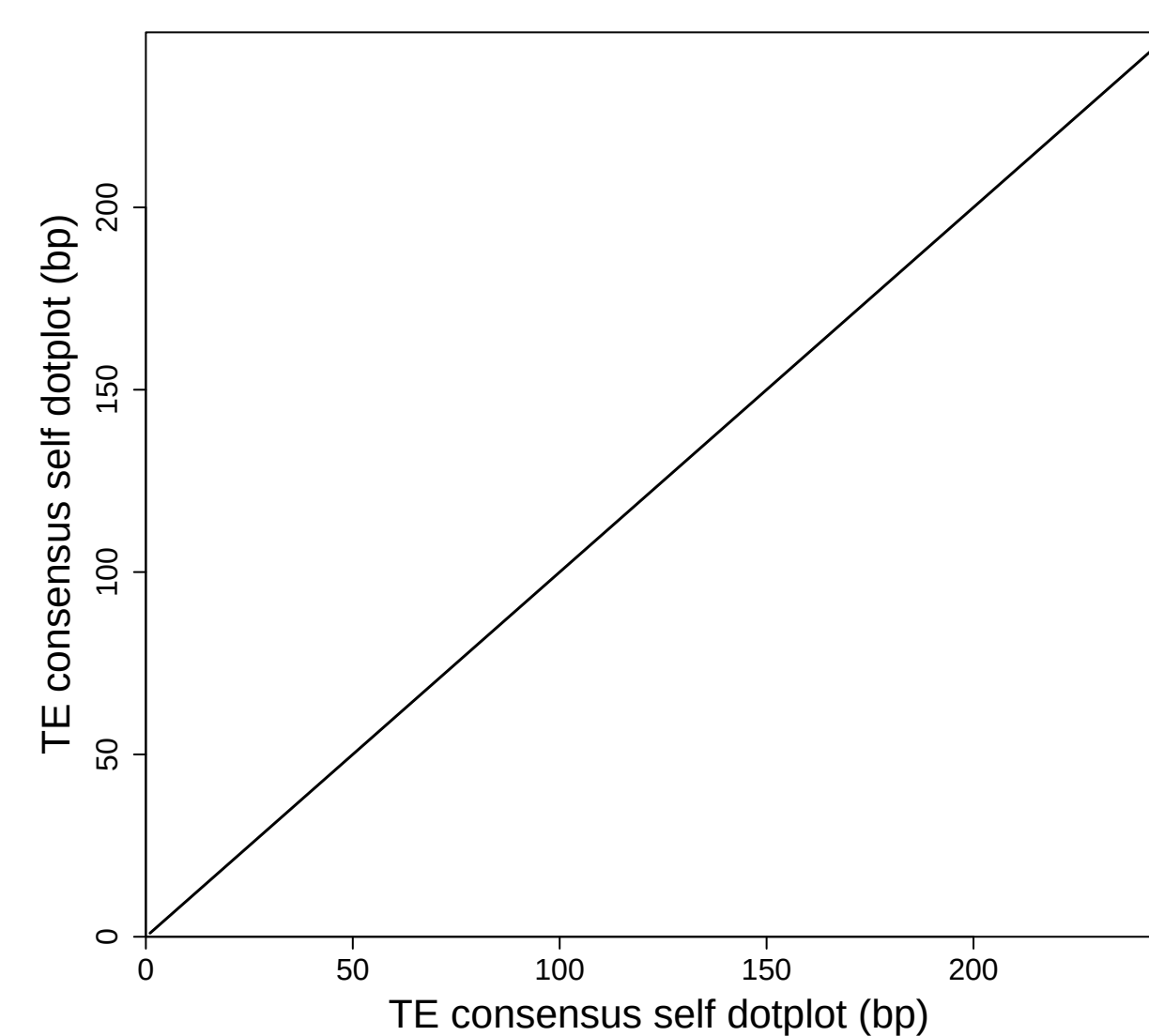
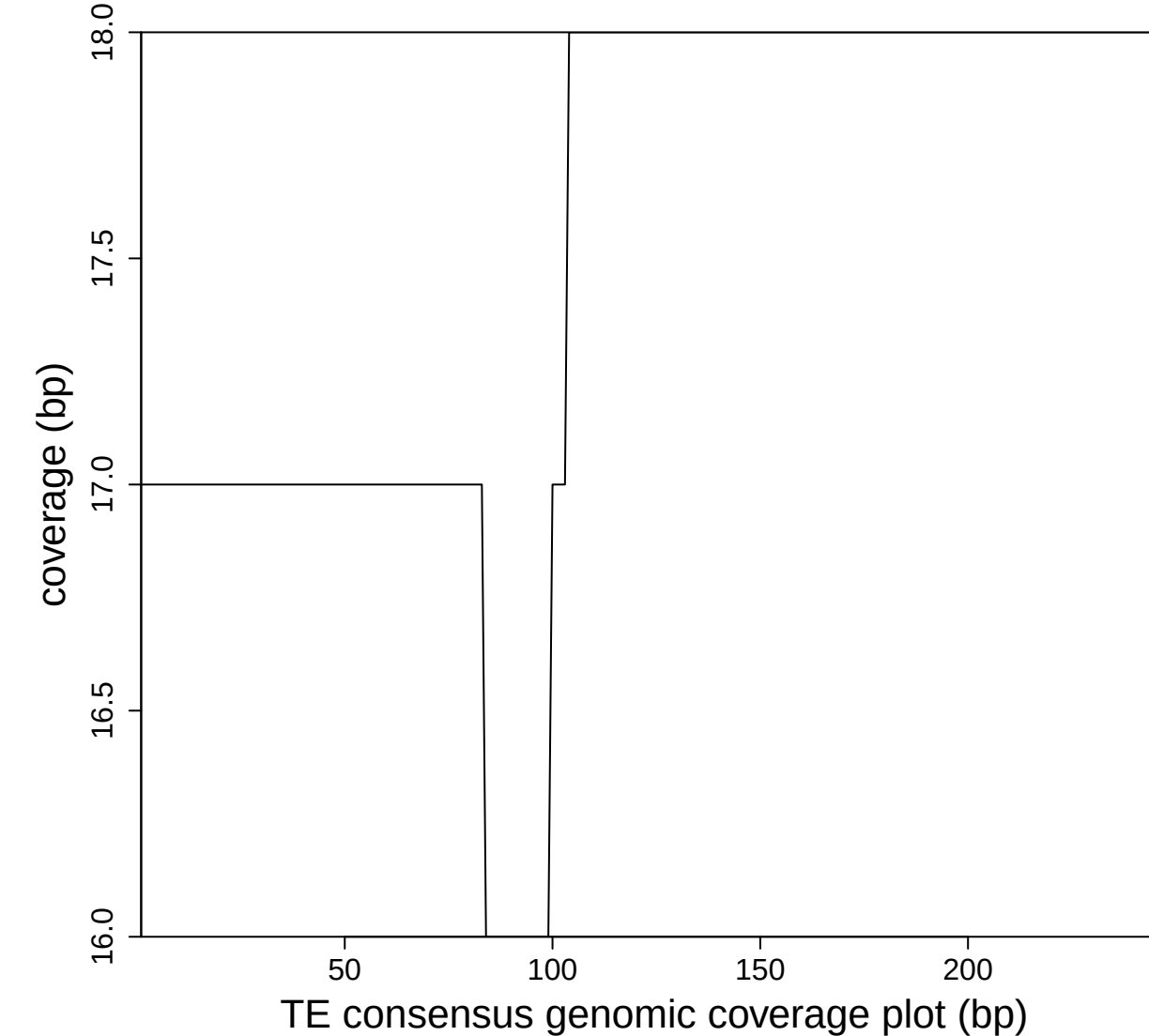
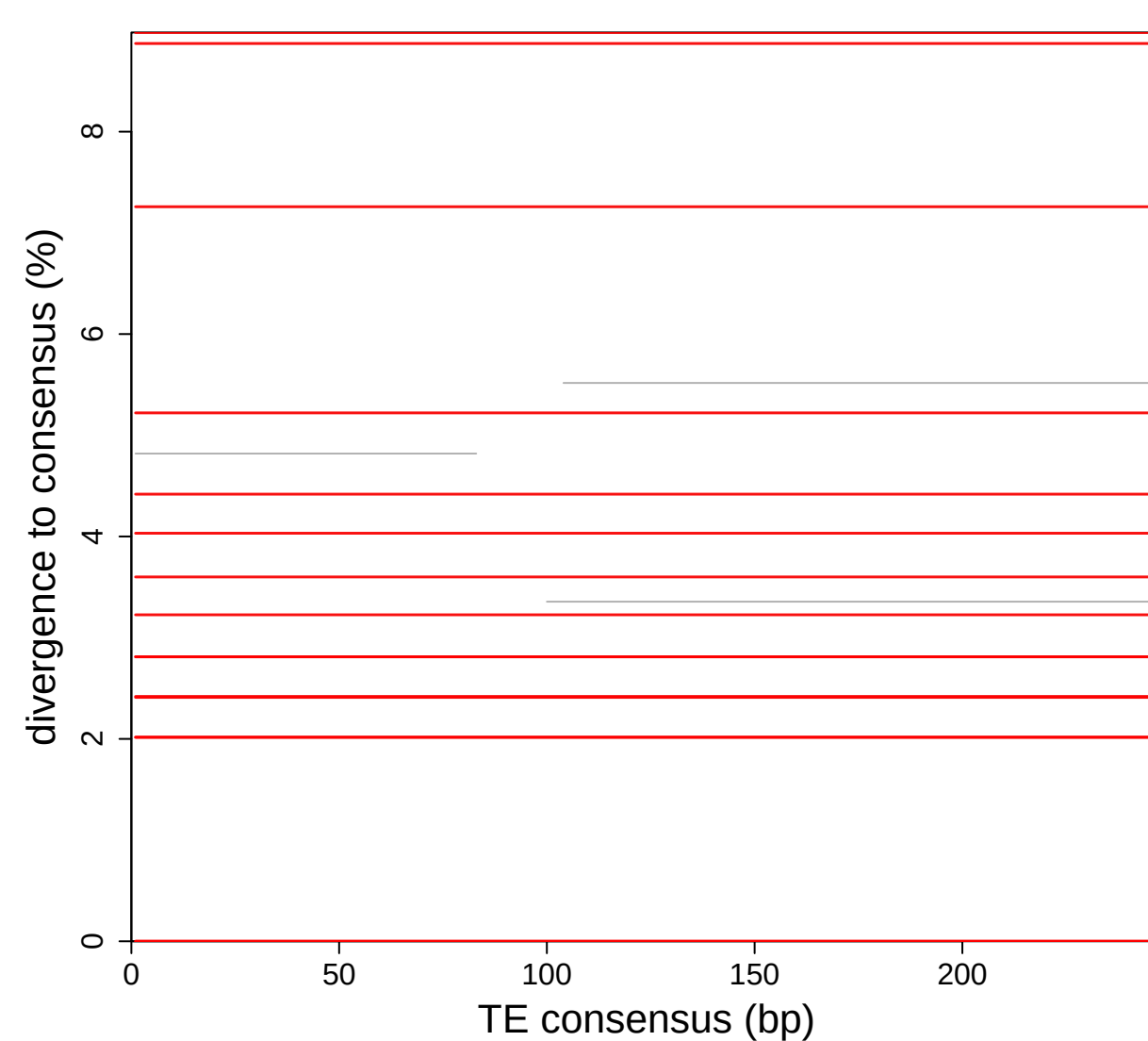
bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln
size: 8677bp; fragments: 74; full length: 0 (>=7809.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

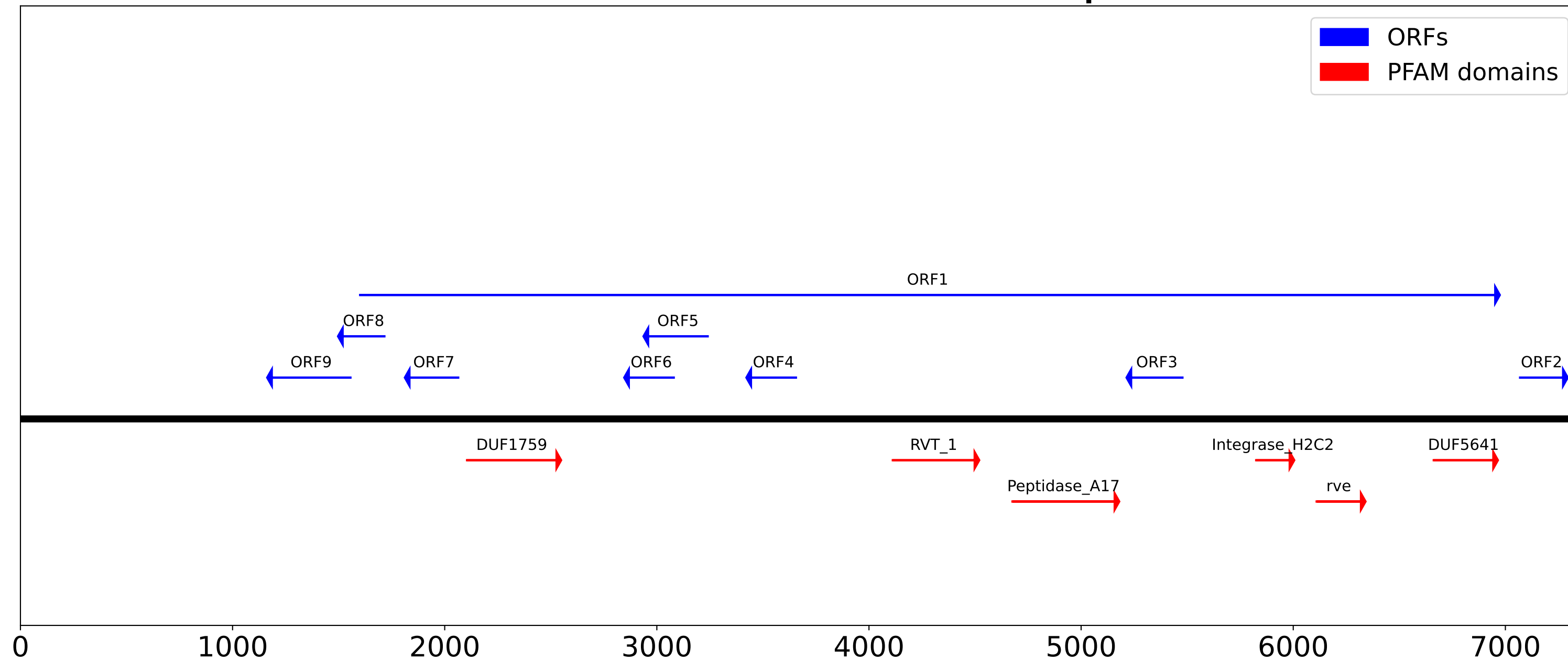


Before TEtrimmer 248 bp

TE: ltr_1_family_92
size: 248bp; fragments: 19; full length: 16 (≥ 223.2 bp)



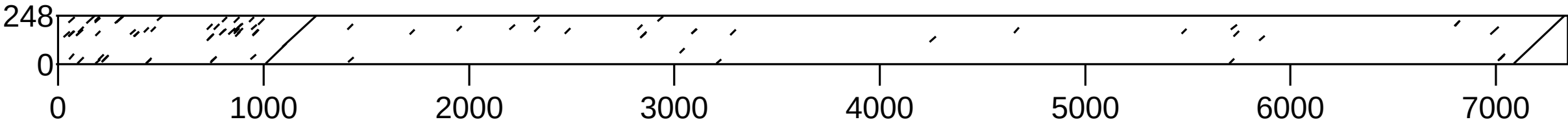
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)