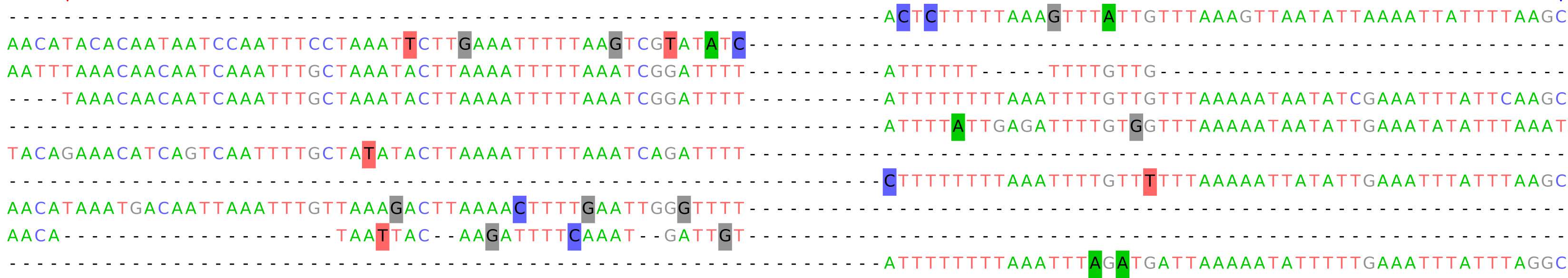


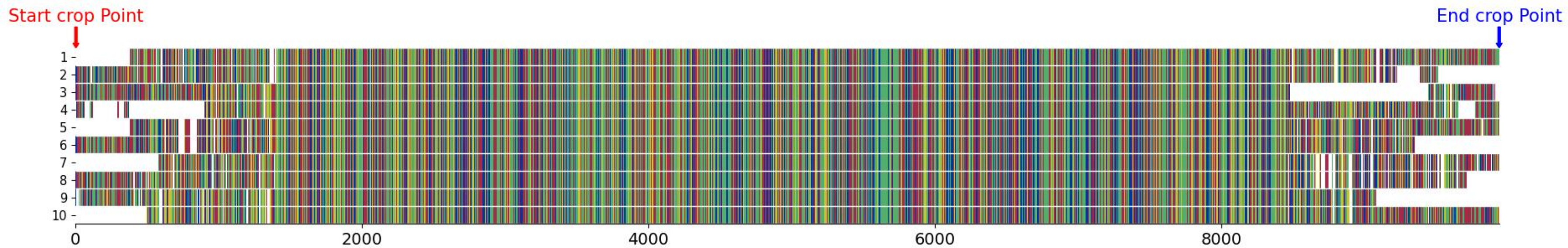
Start crop Point



MSA length = 9937

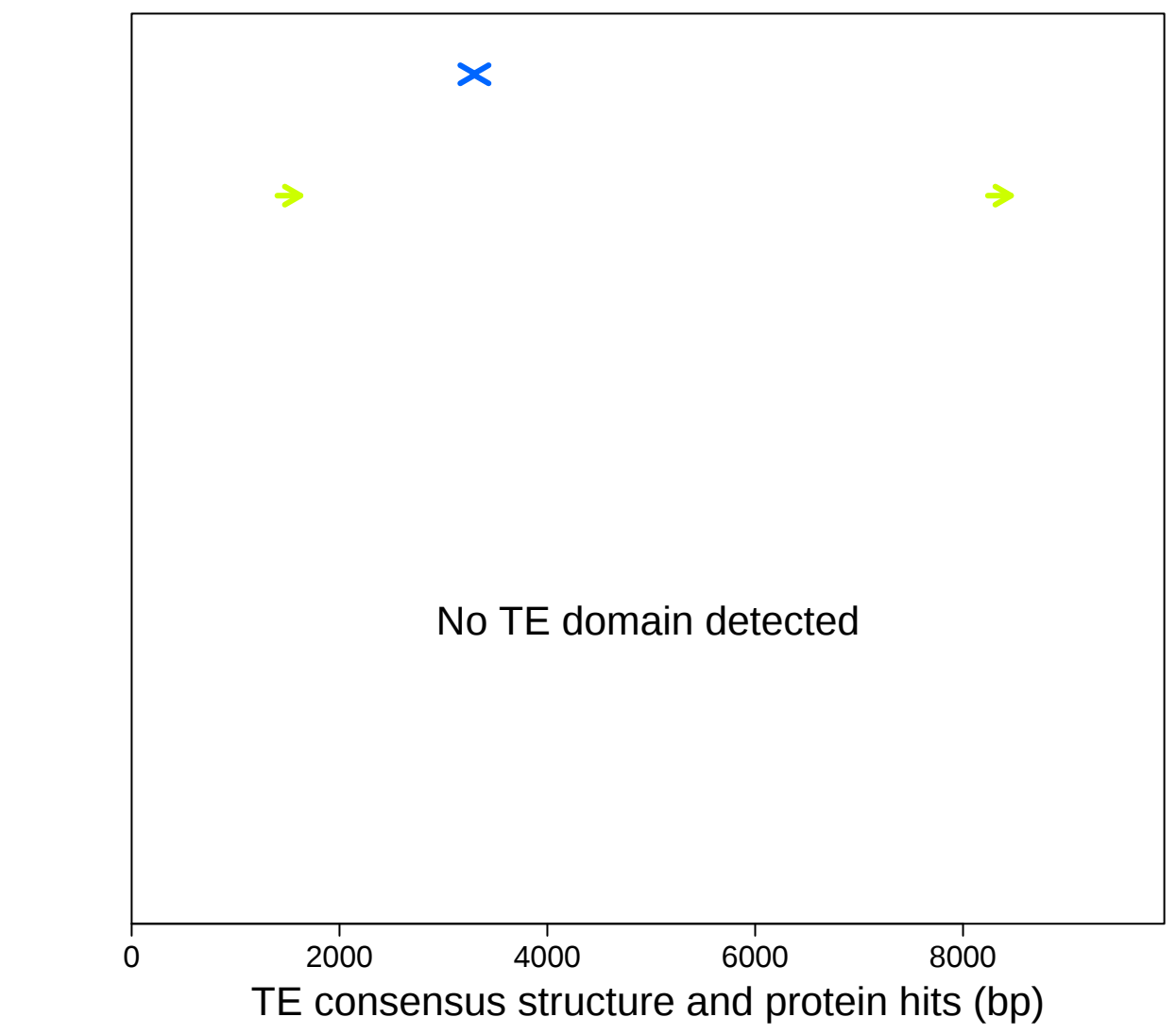
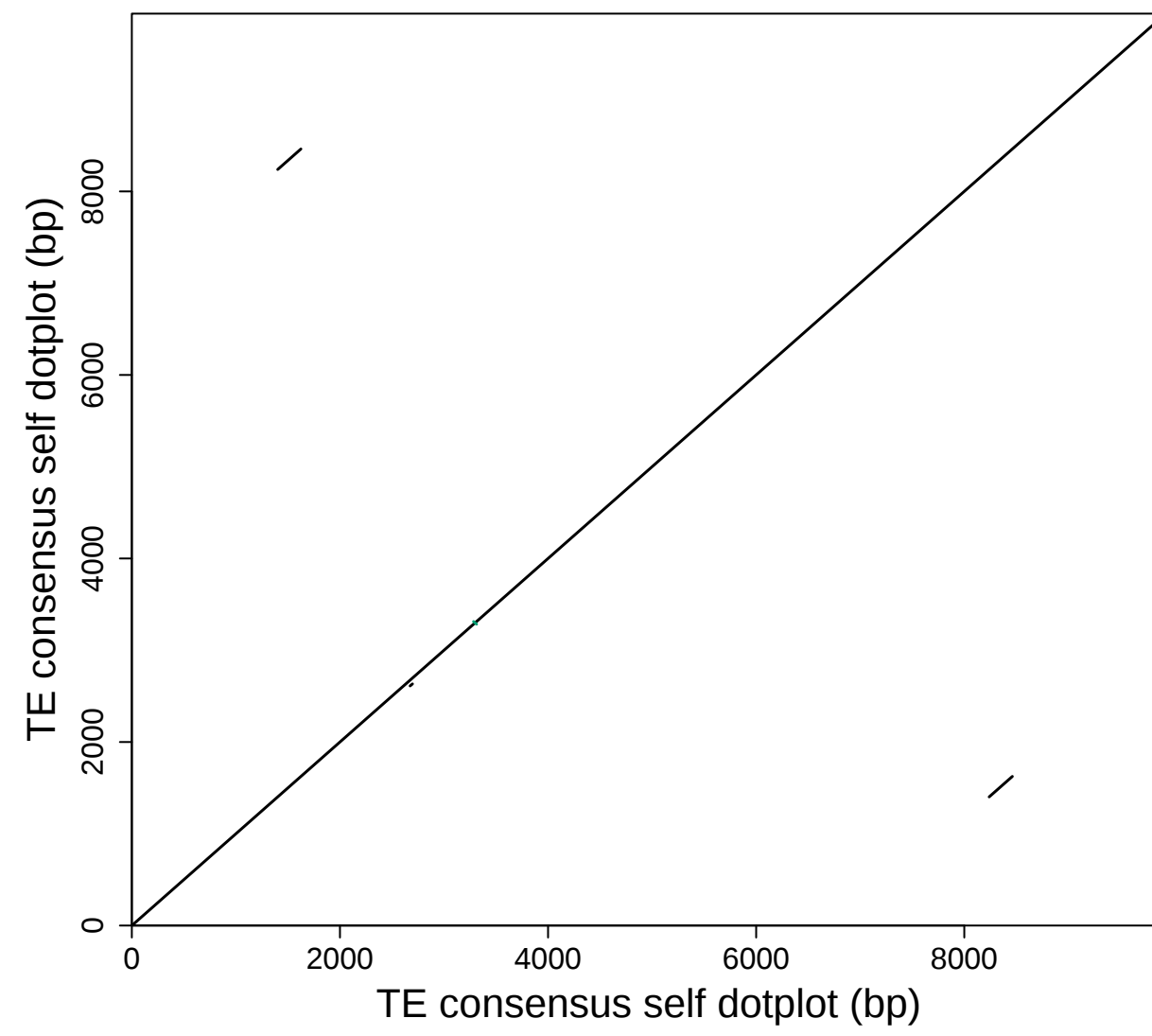
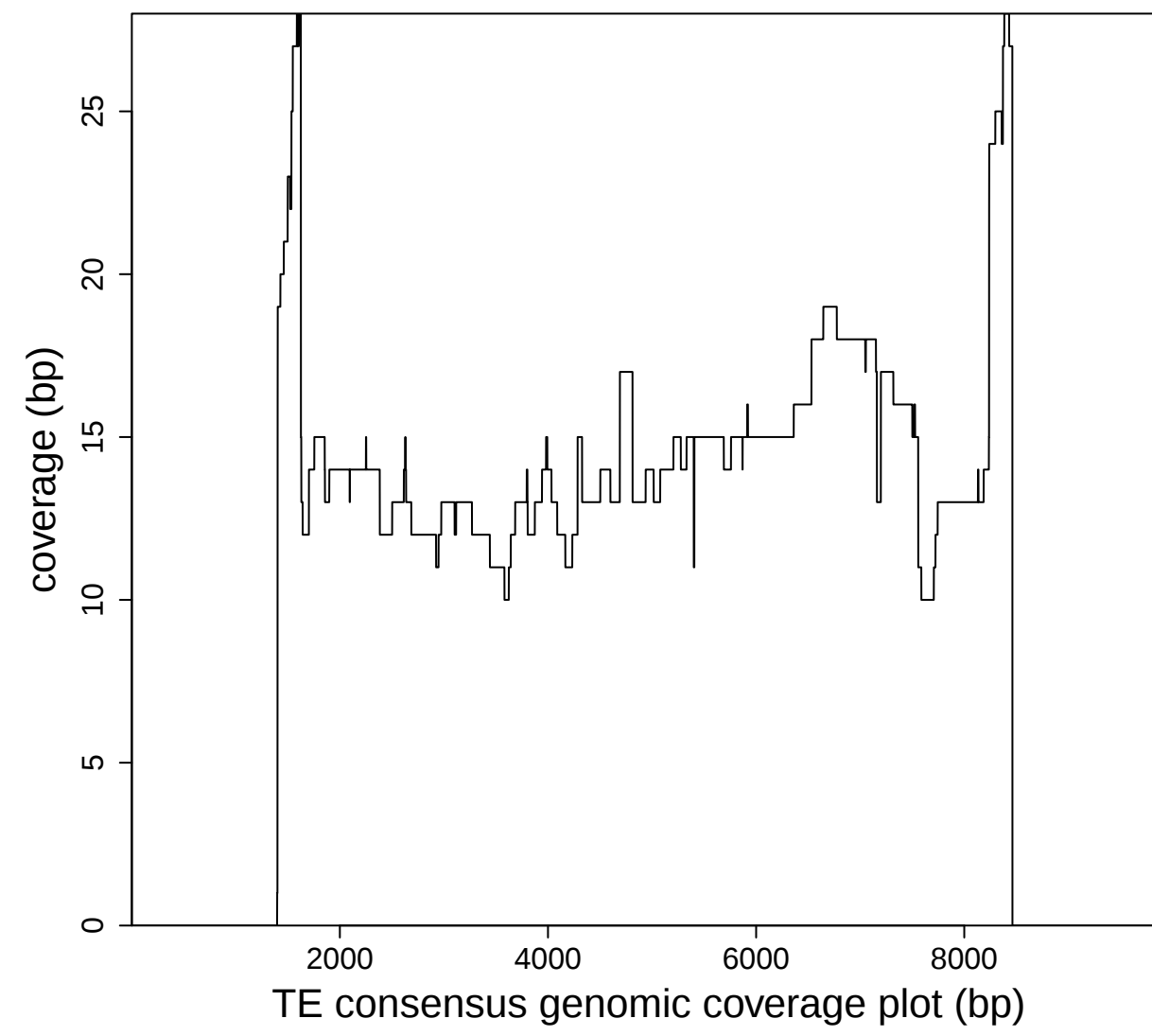
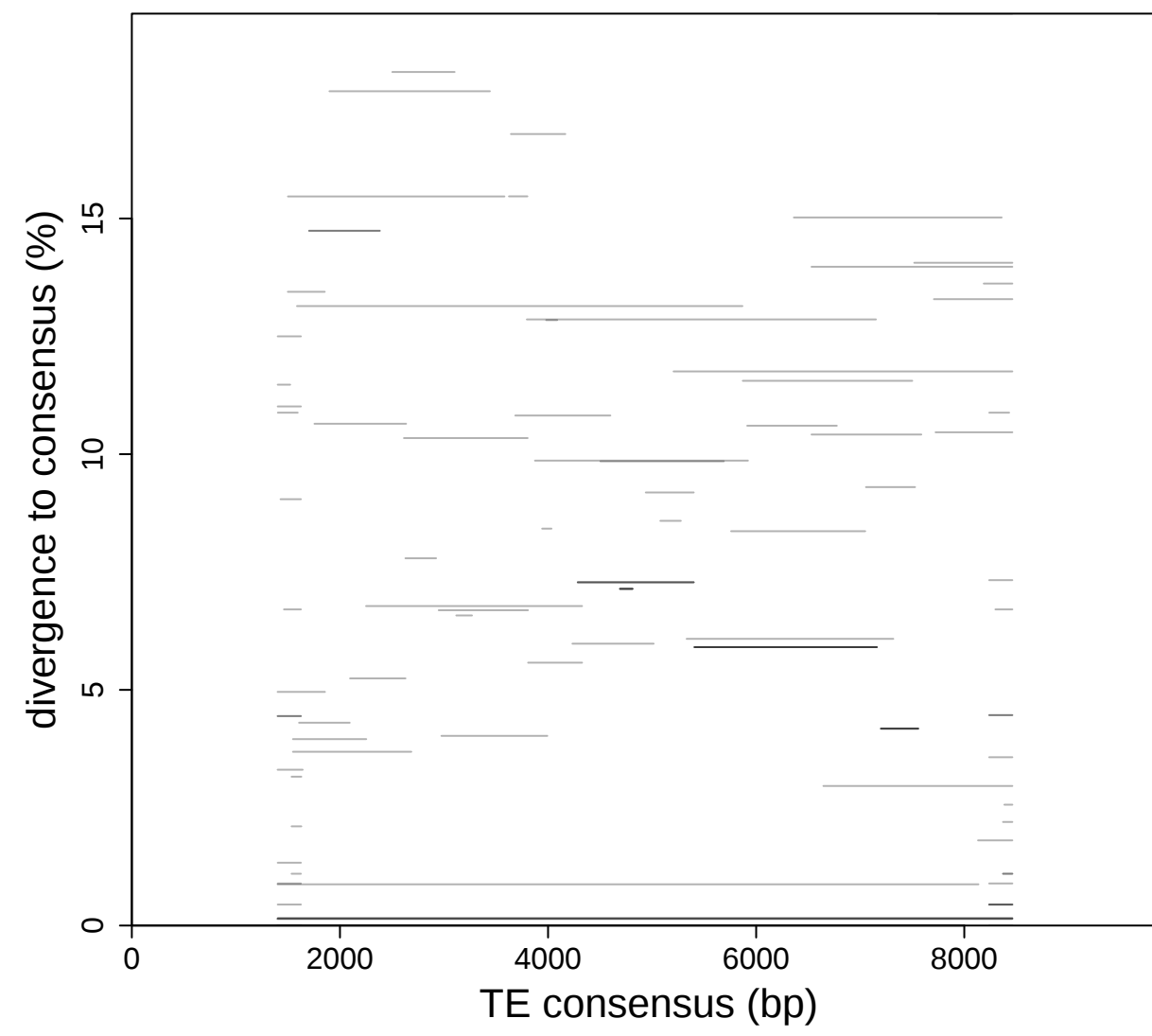
End crop Point





_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_al
size: 9937bp; fragments: 100; full length: 0 (>=8943.3bp)

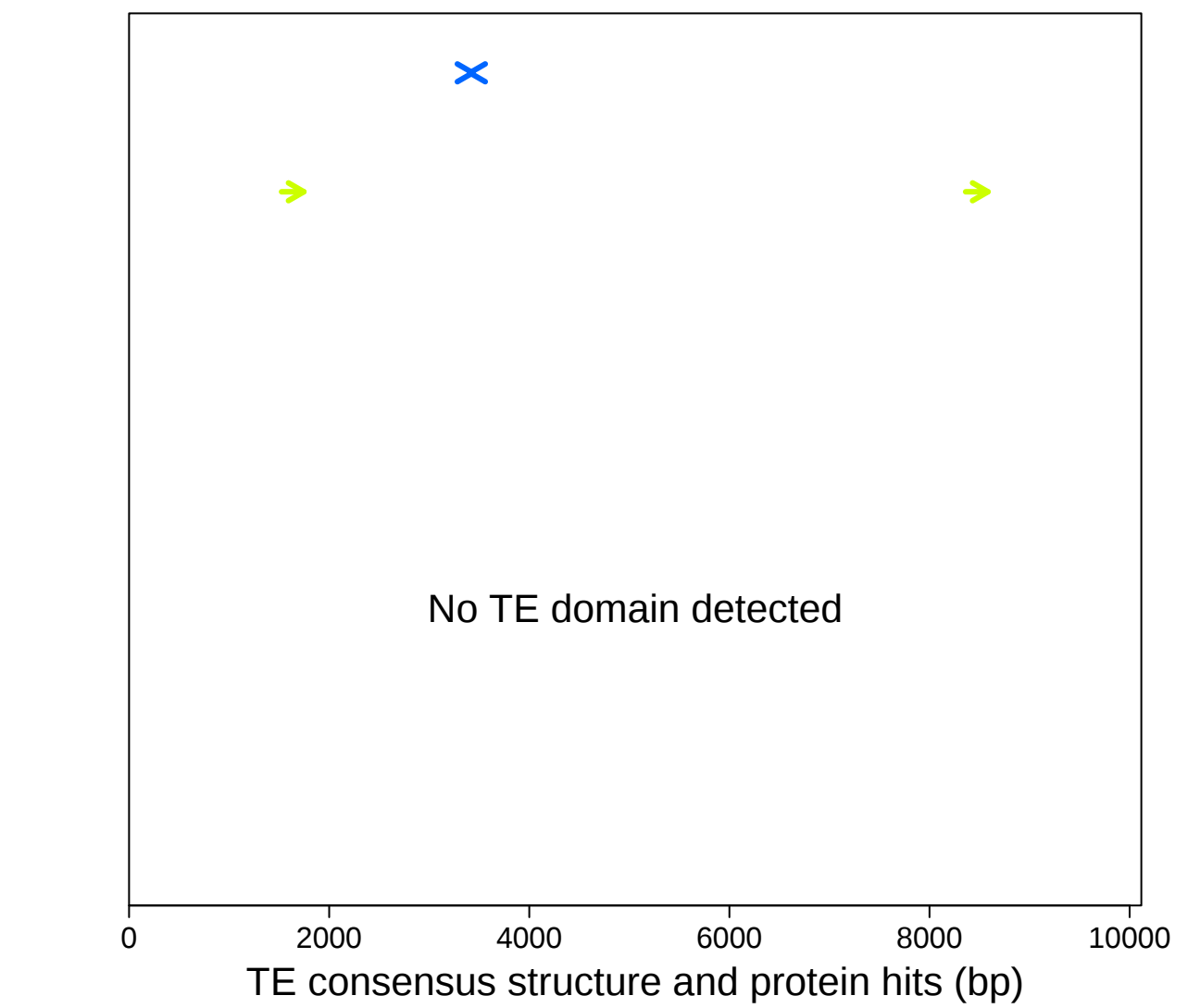
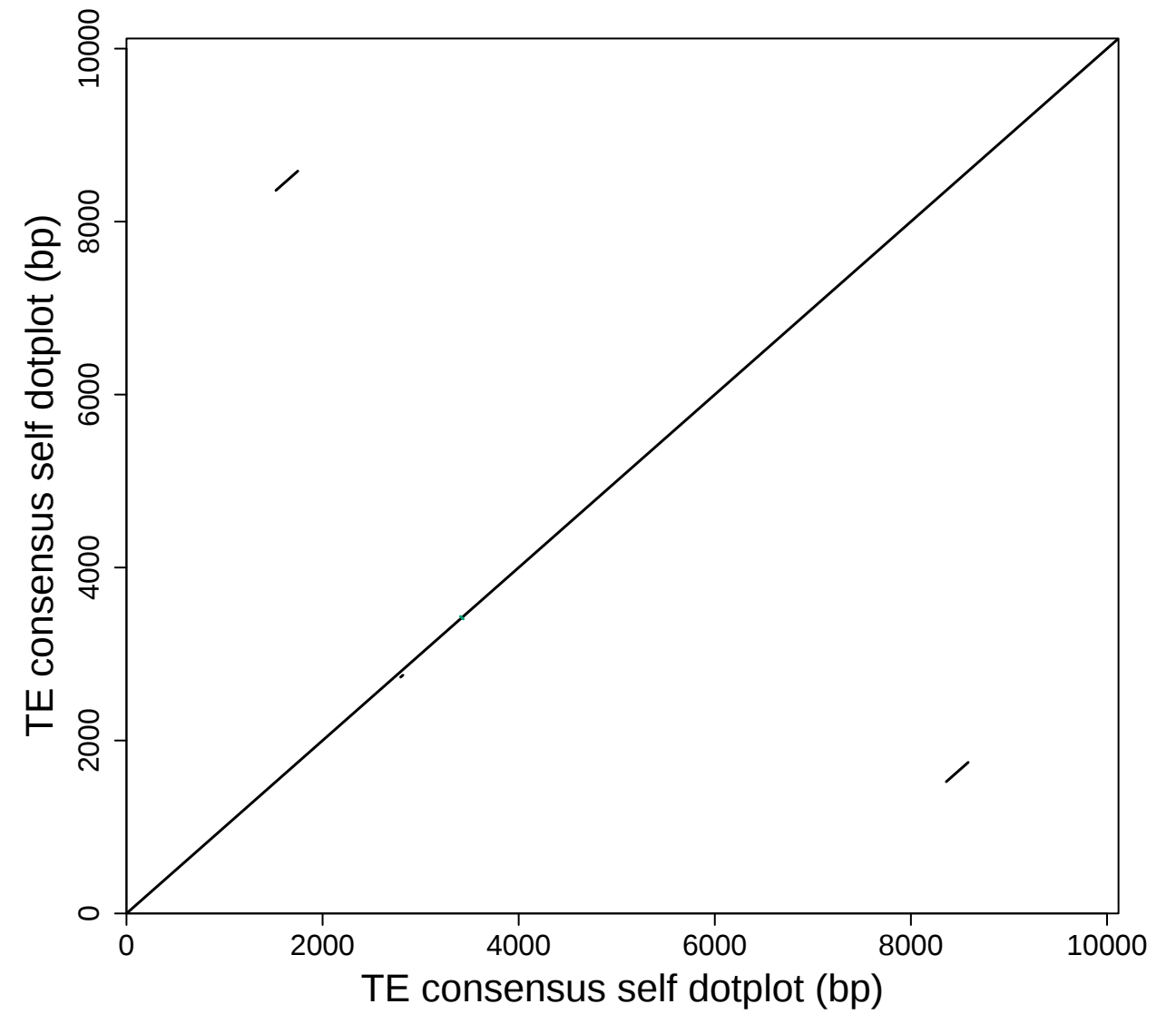
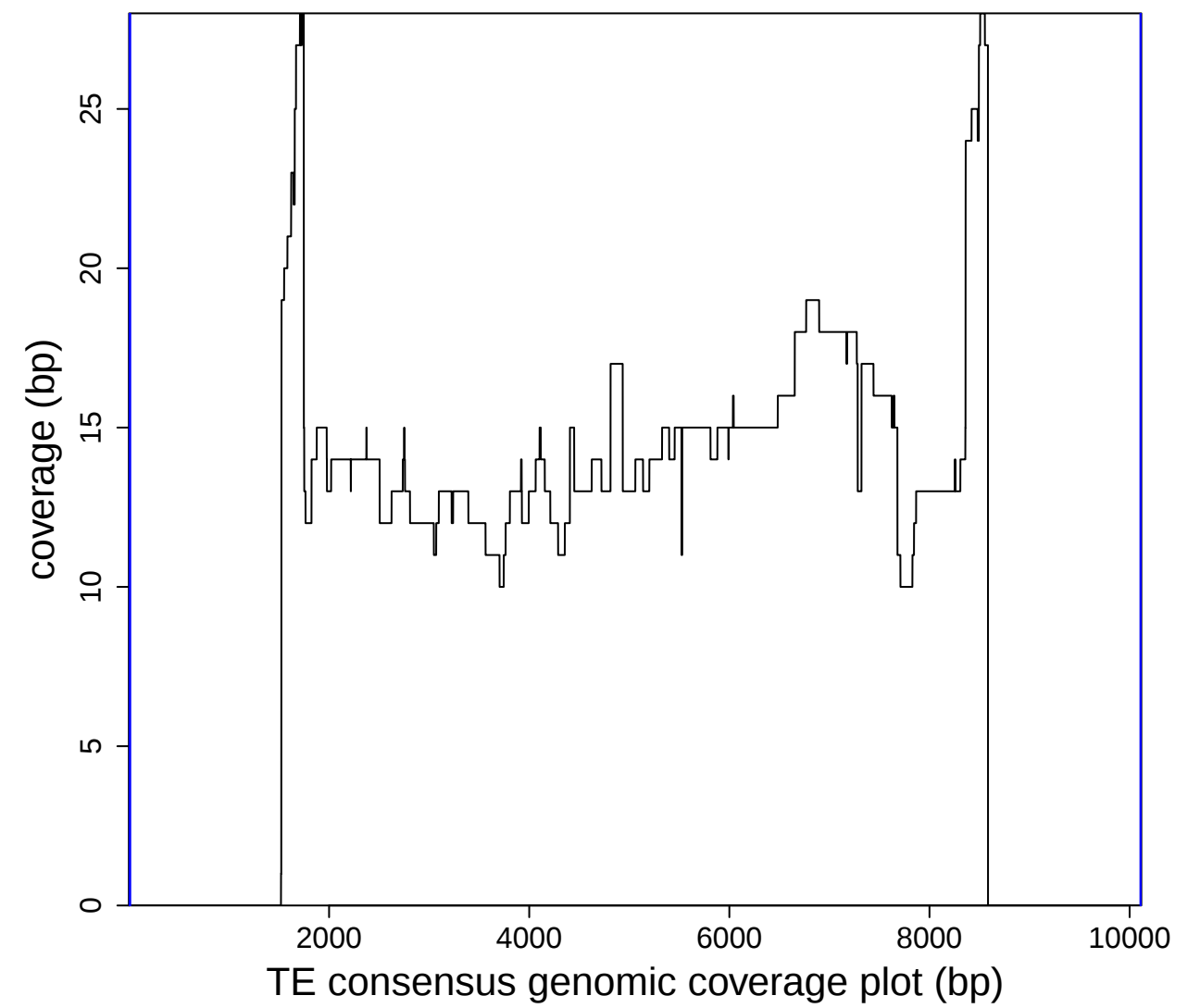
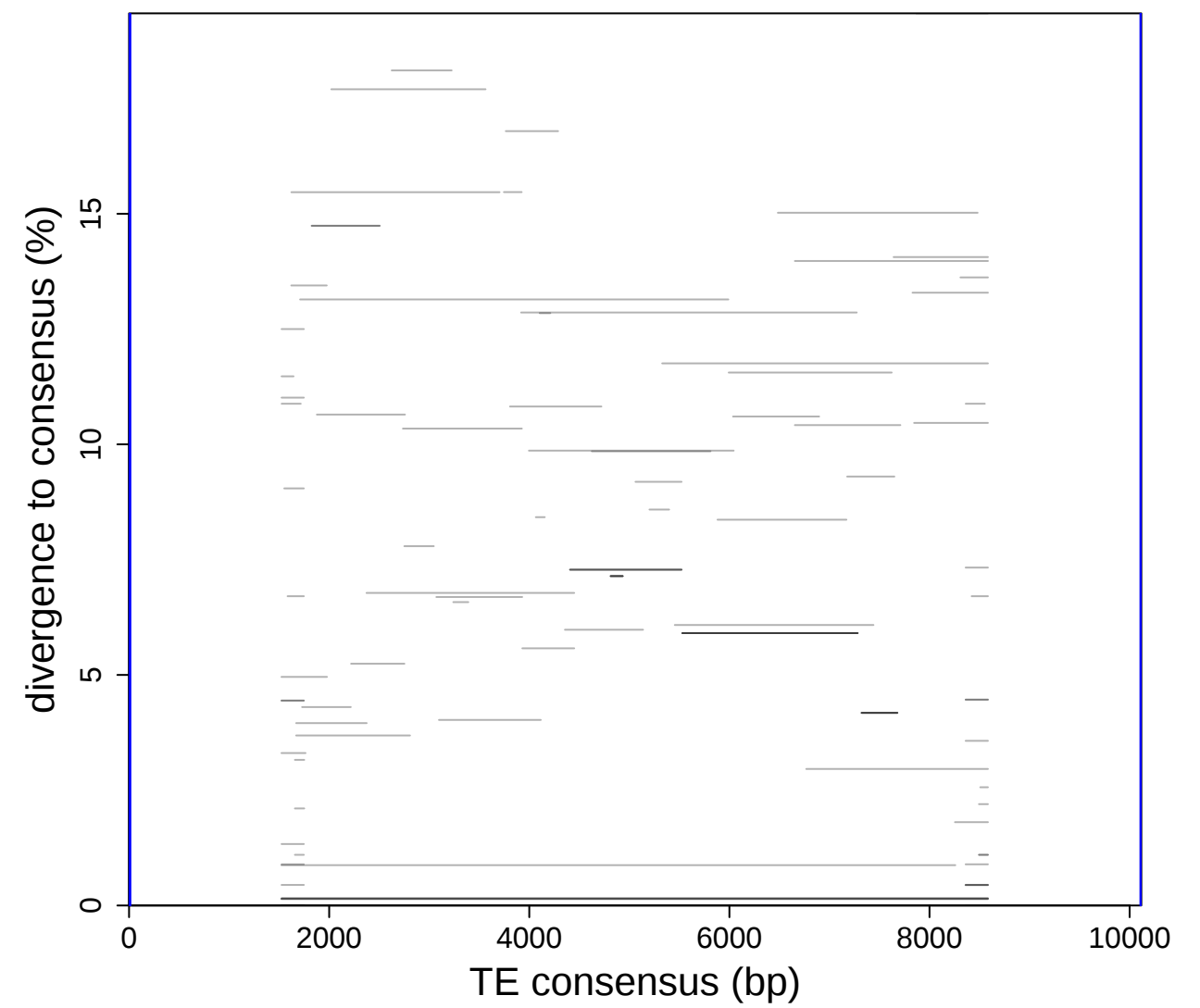
After TEtrimmer 9937 bp



bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bclr
size: 10117bp; fragments: 100; full length: 0 (>=9105.3bp)

After TETrimmer Extended plot

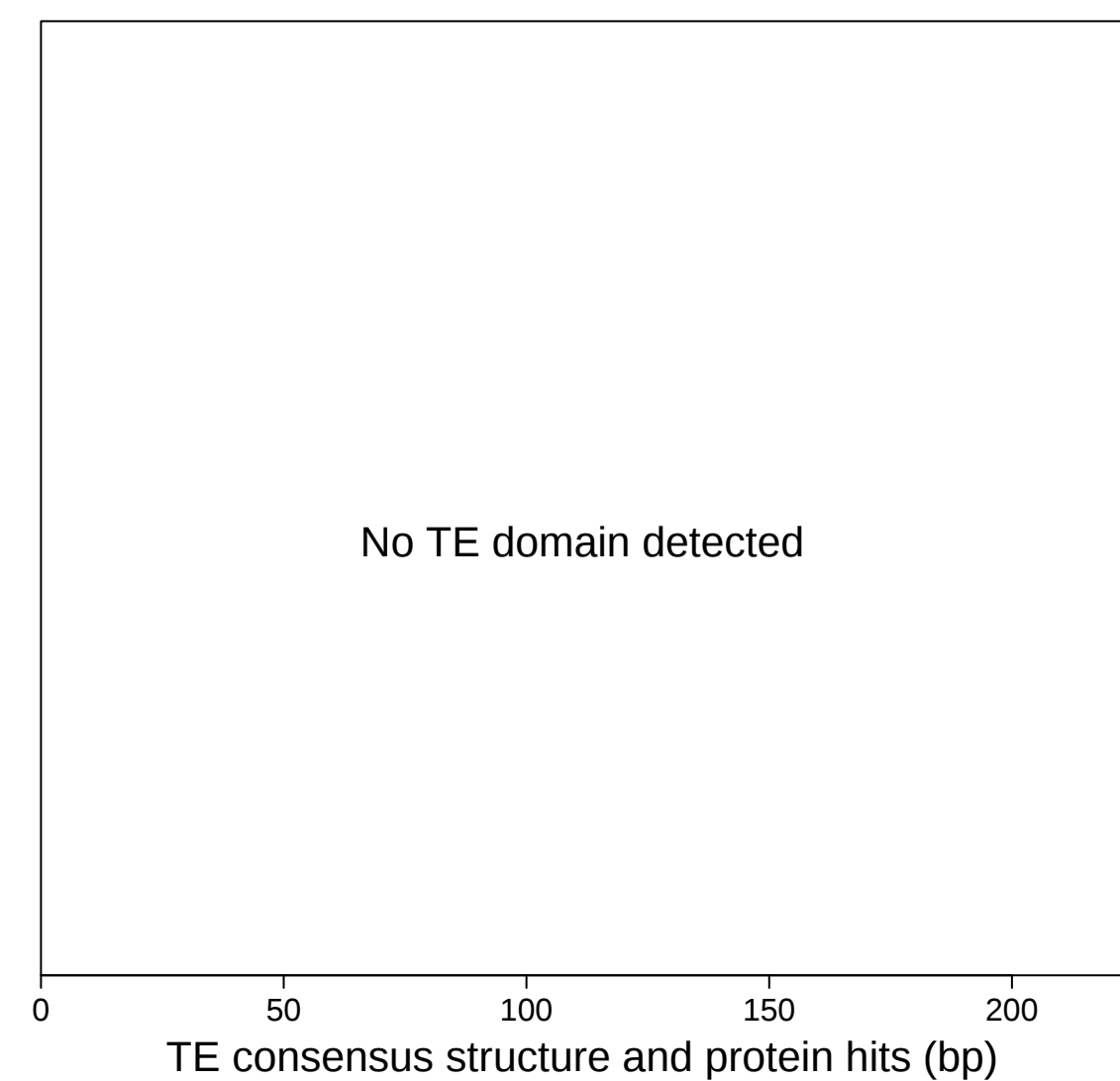
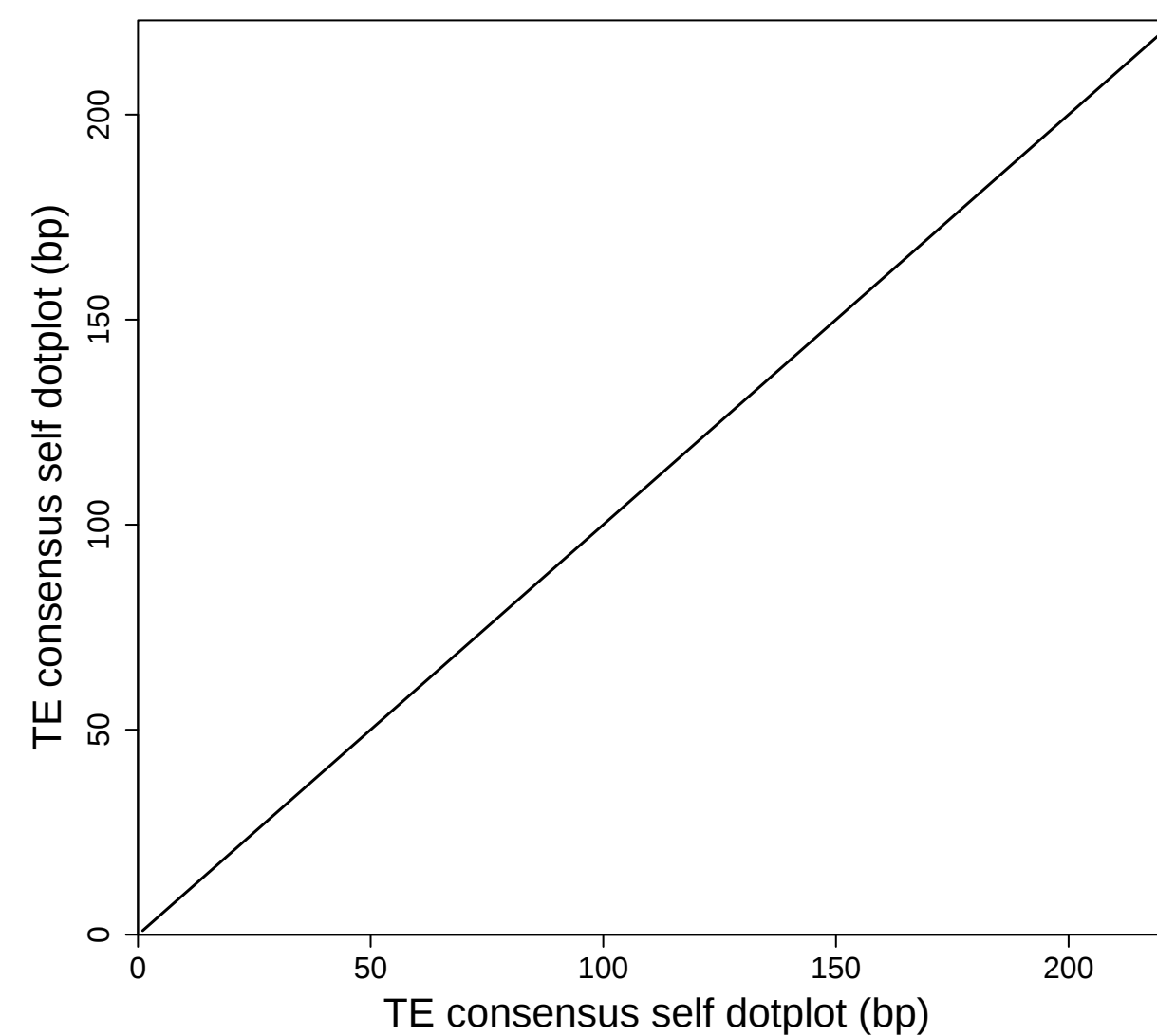
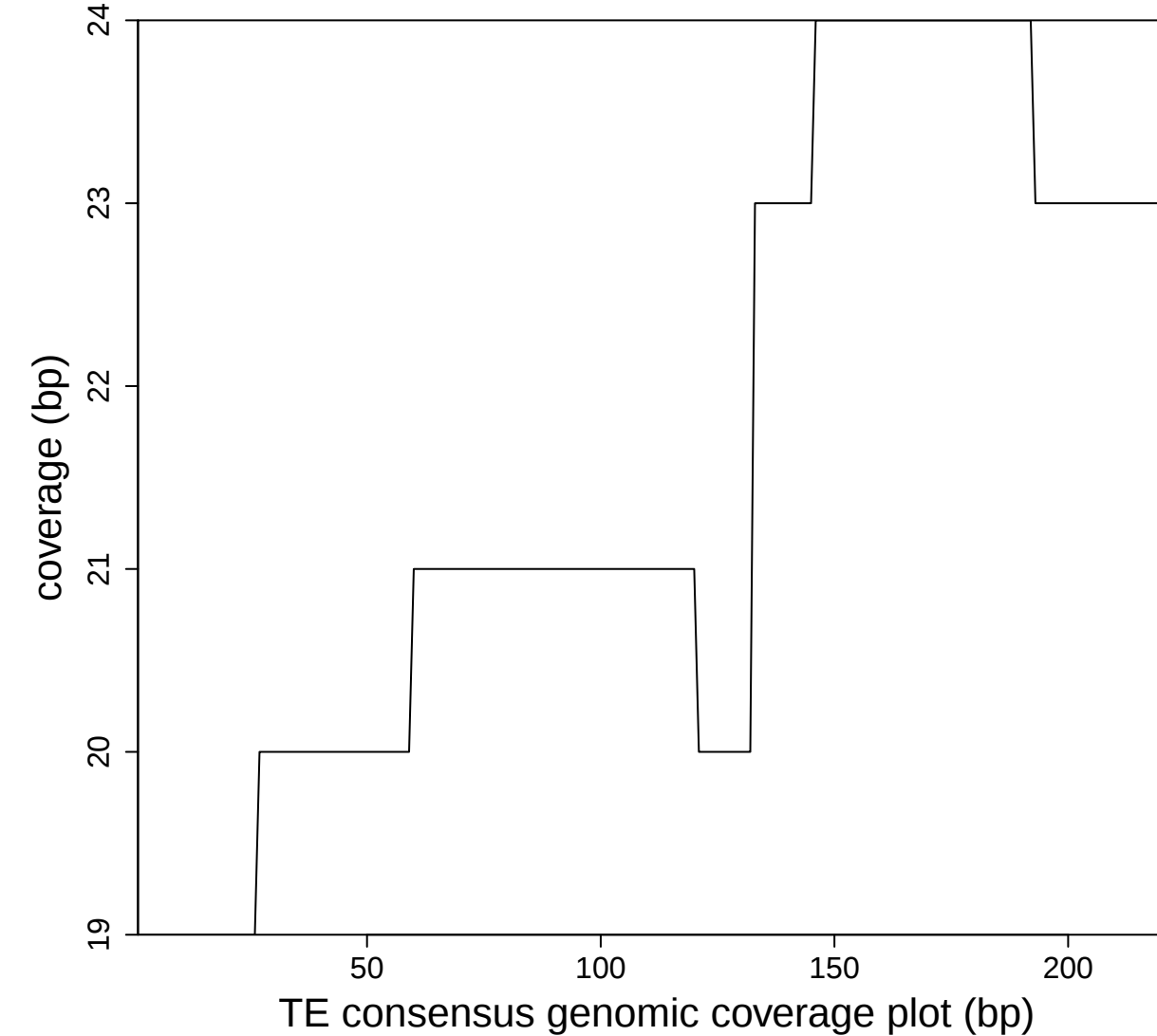
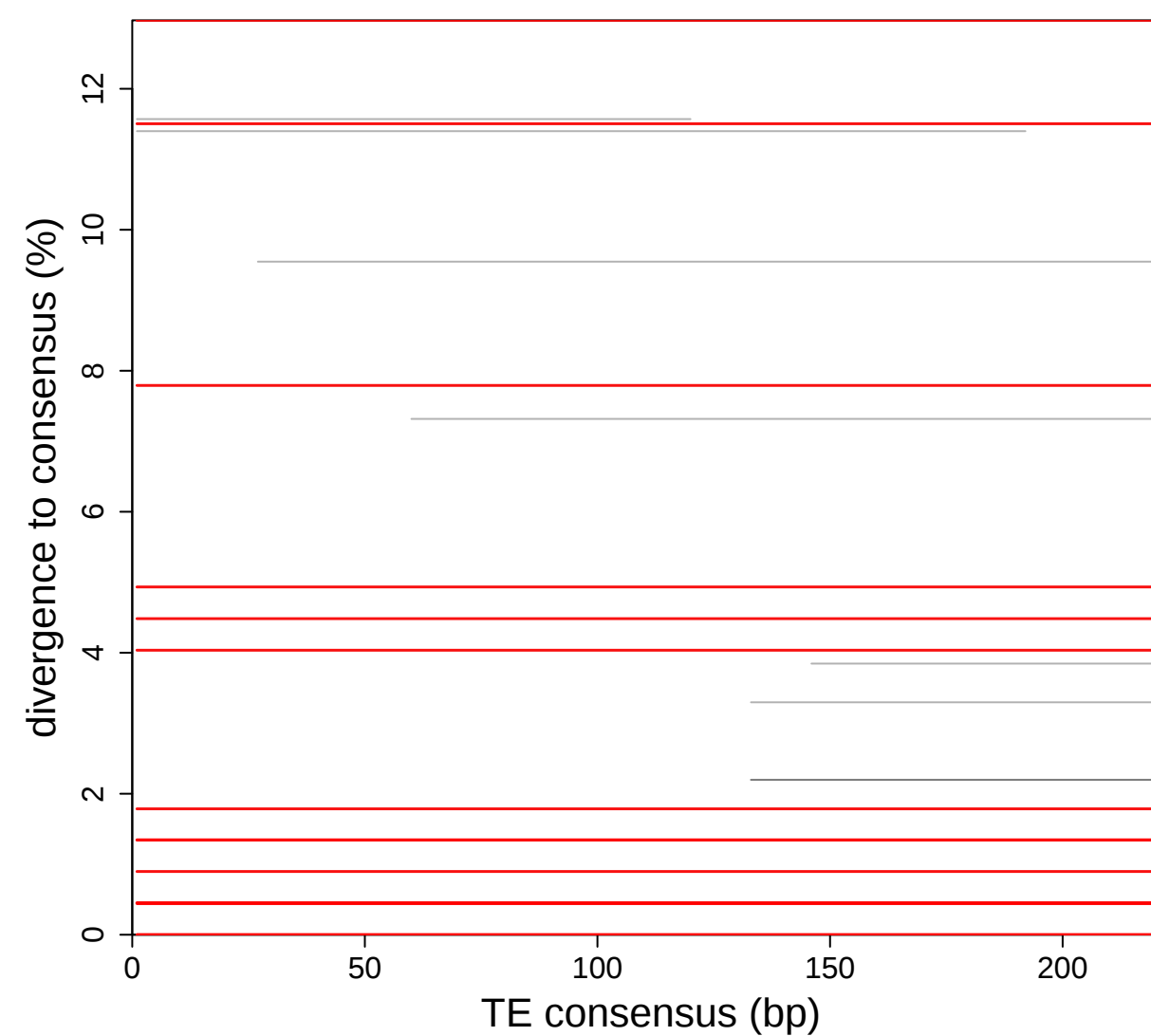
Blue lines are boundaries



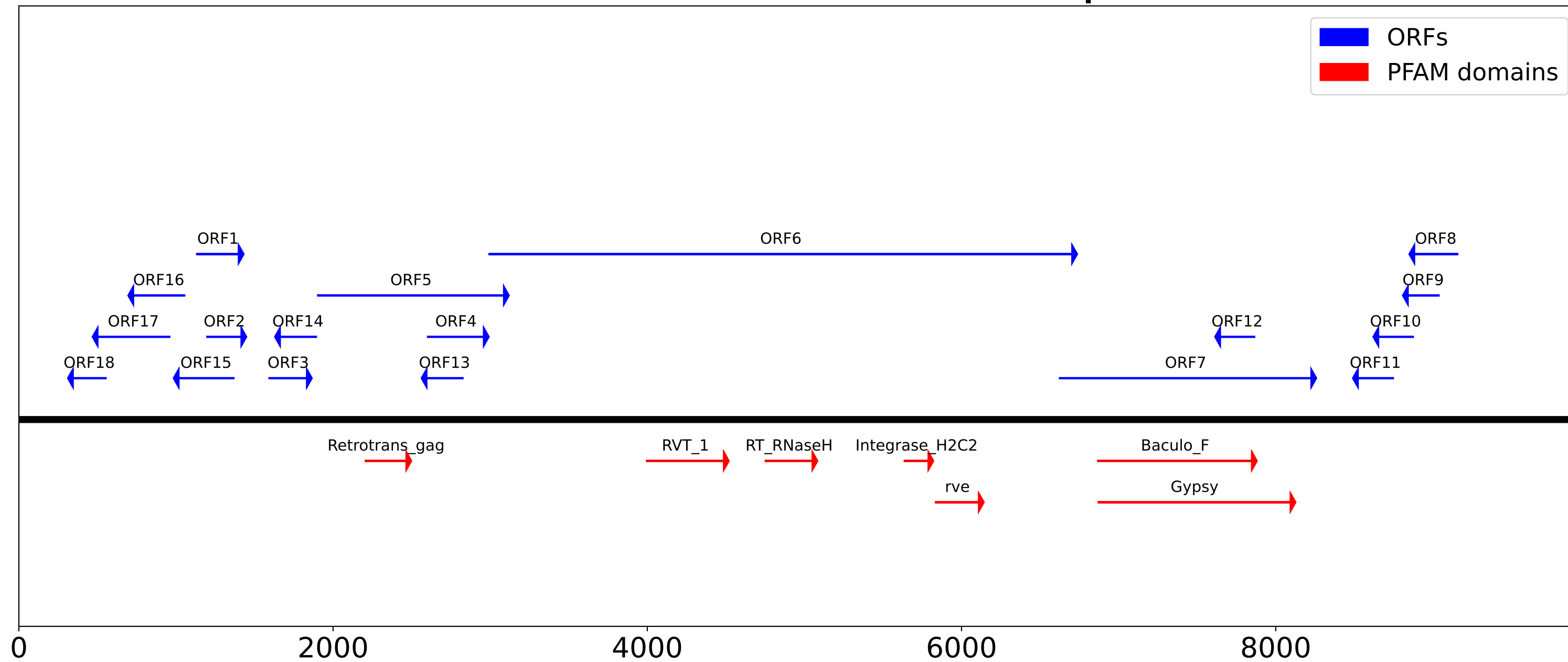
Before TEtrimmer 223 bp

TE: ltr_1_family_311

size: 223bp; fragments: 25; full length: 17 (≥ 200.7 bp)



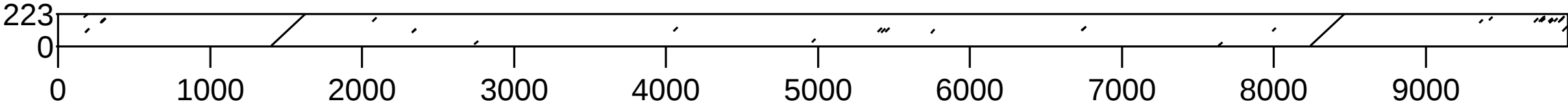
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)