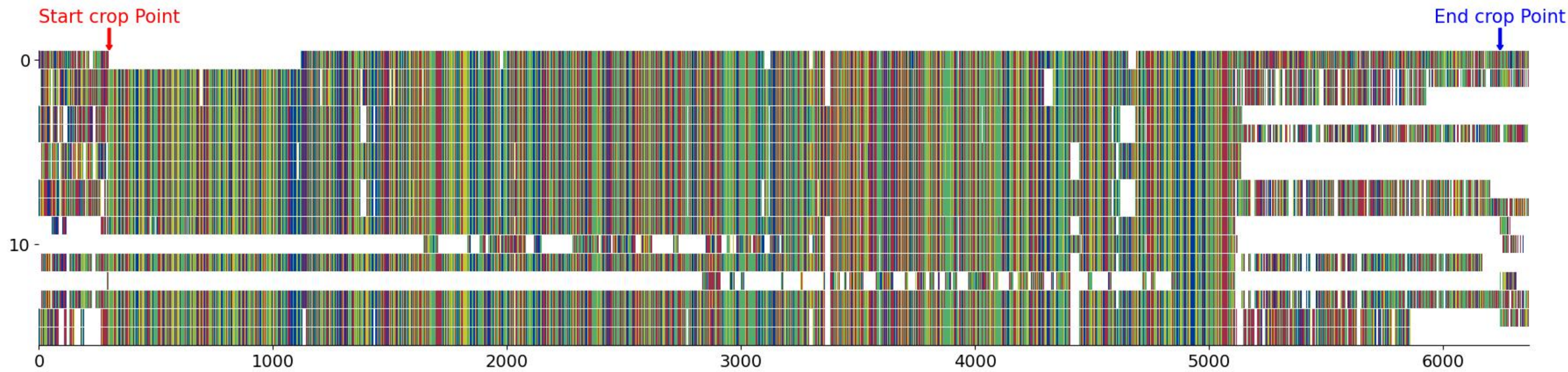


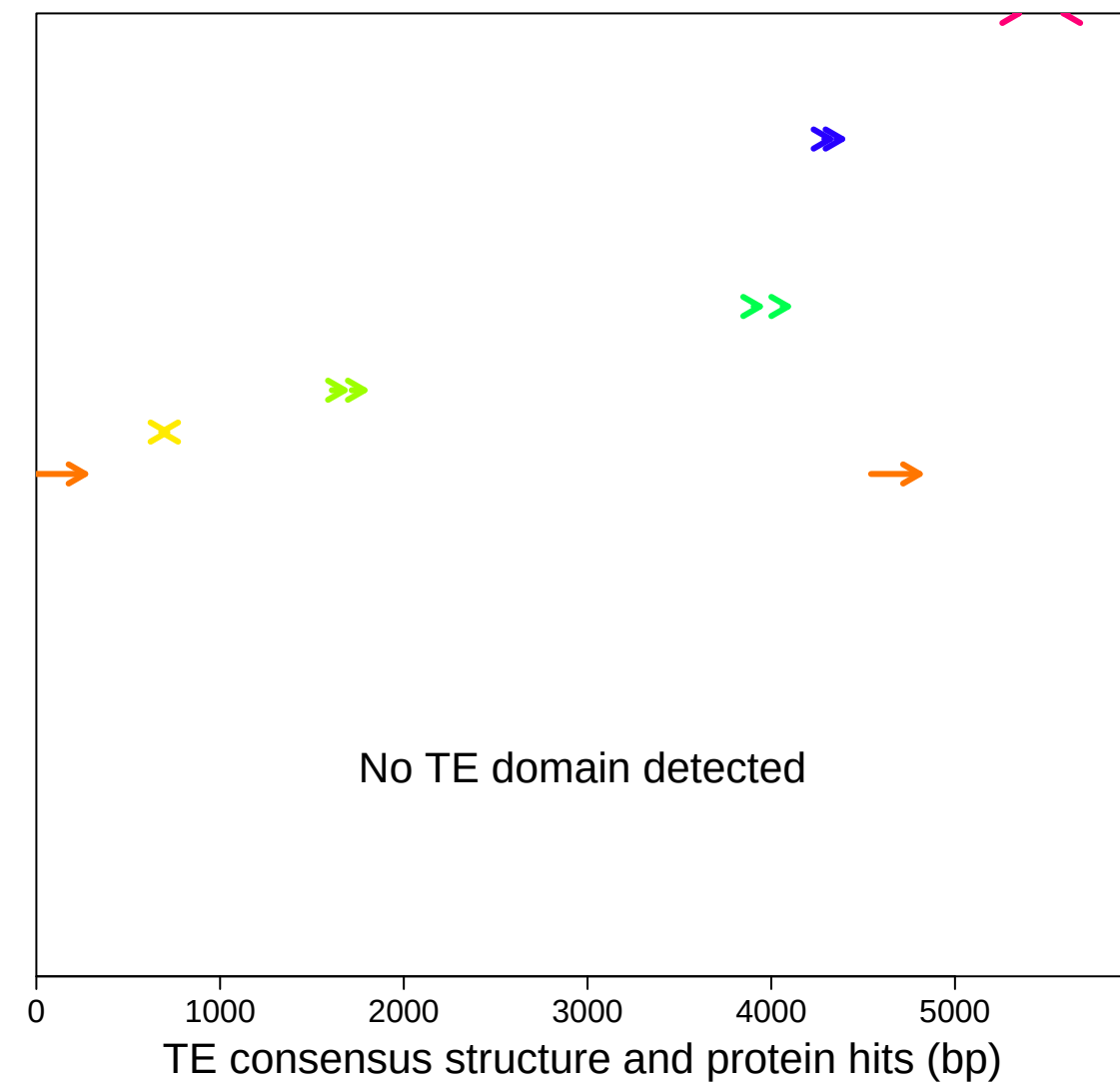
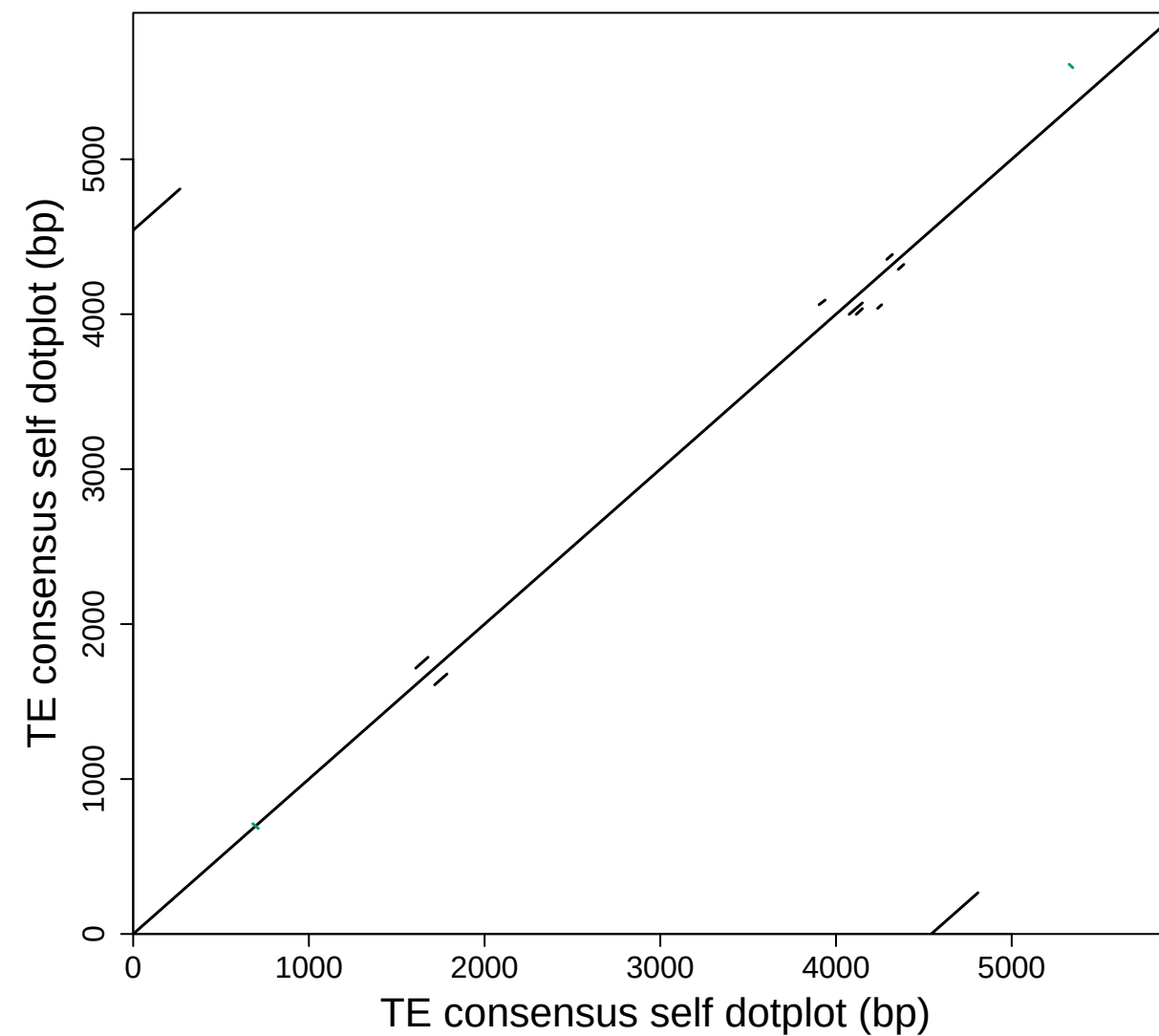
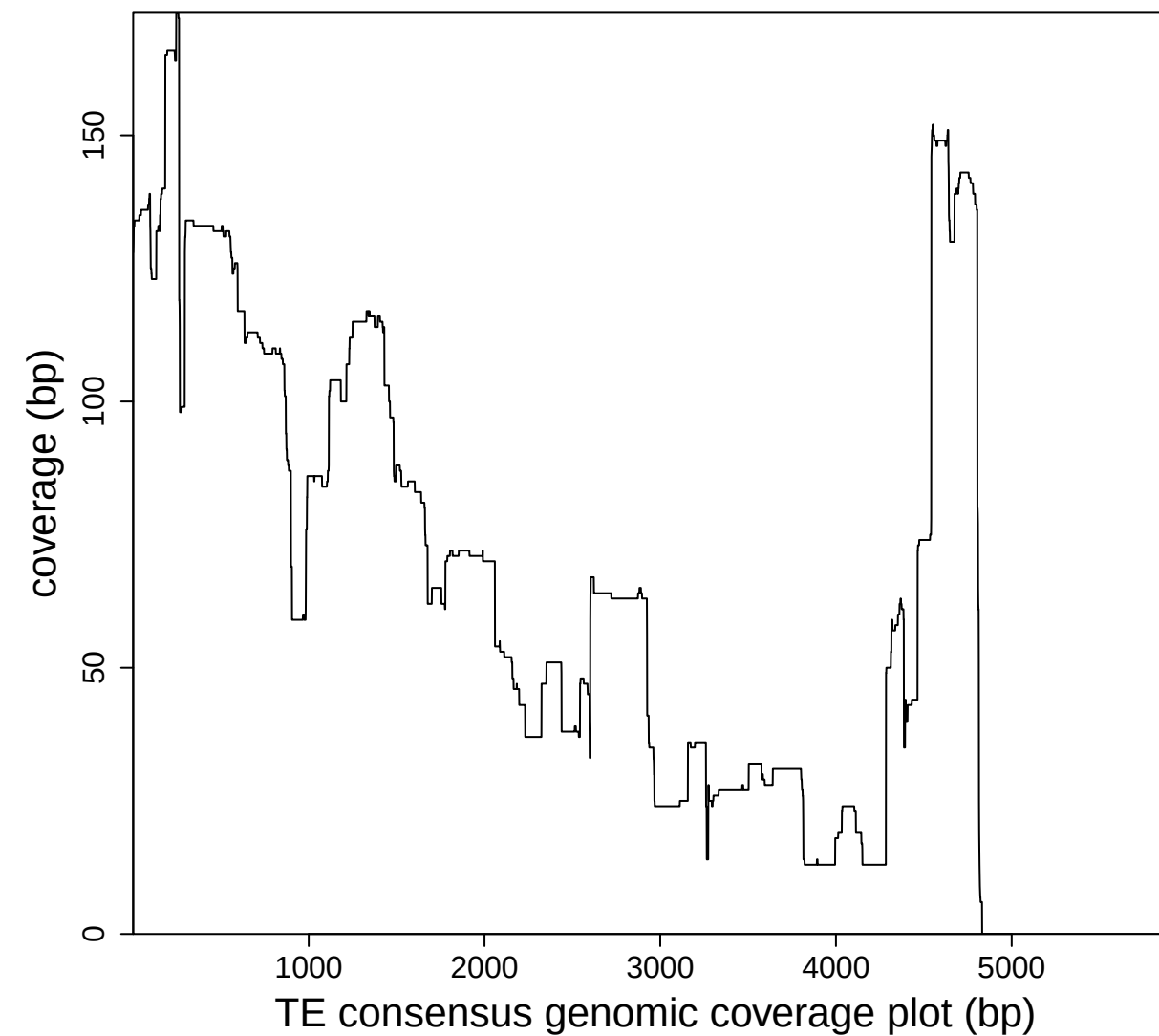
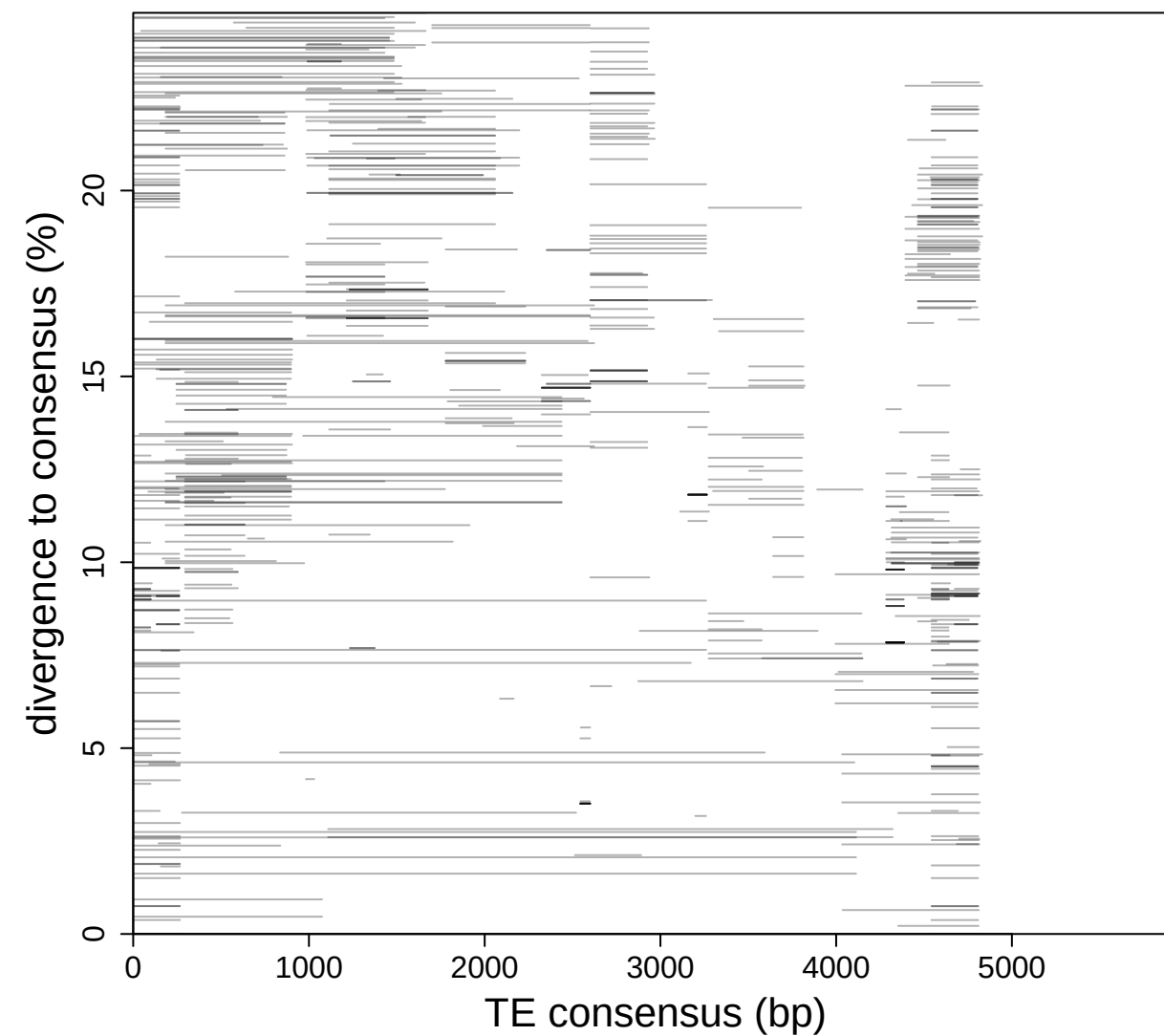
MSA length = 5945





_3.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_al
size: 5945bp; fragments: 651; full length: 0 (>=5350.5bp)

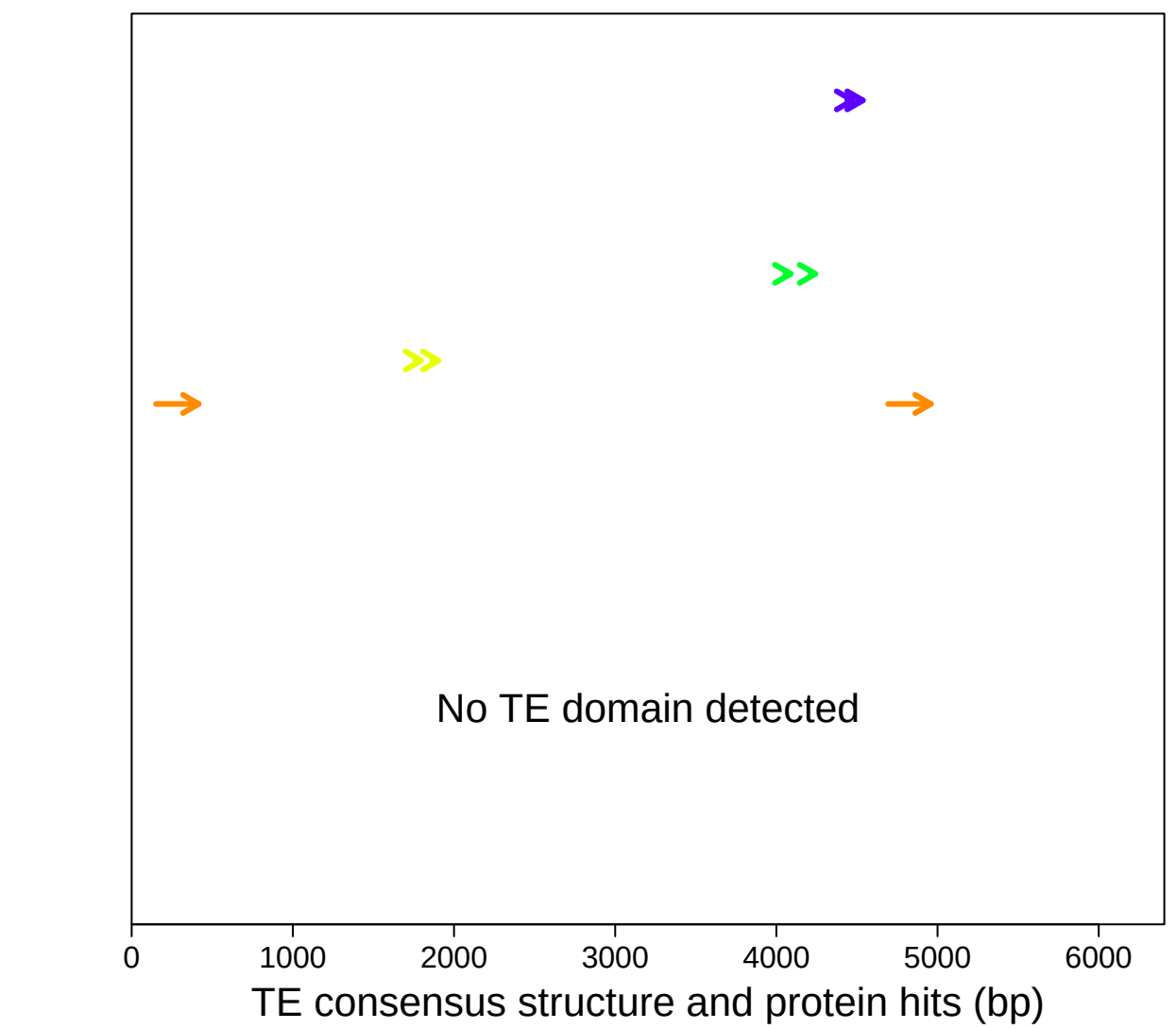
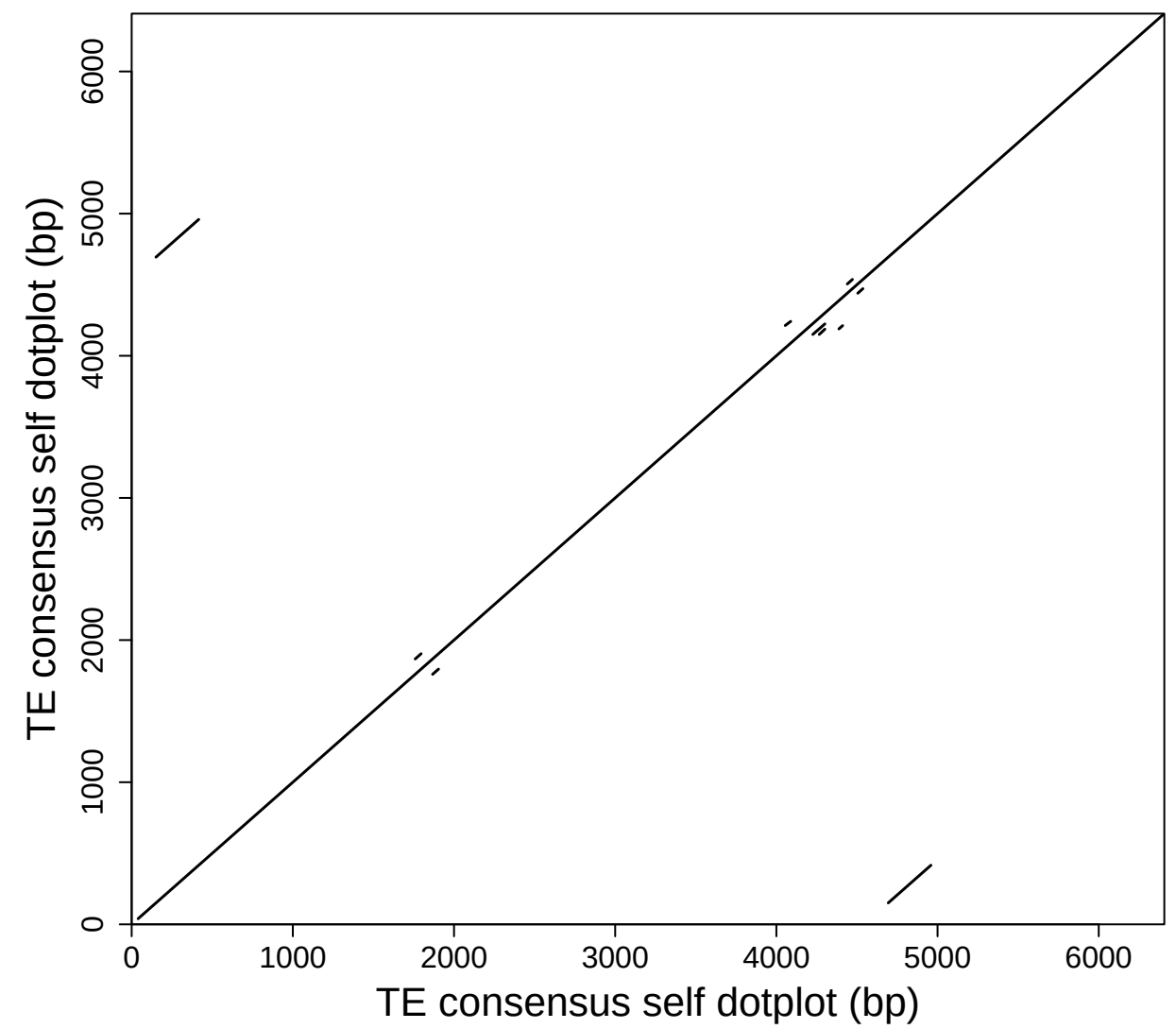
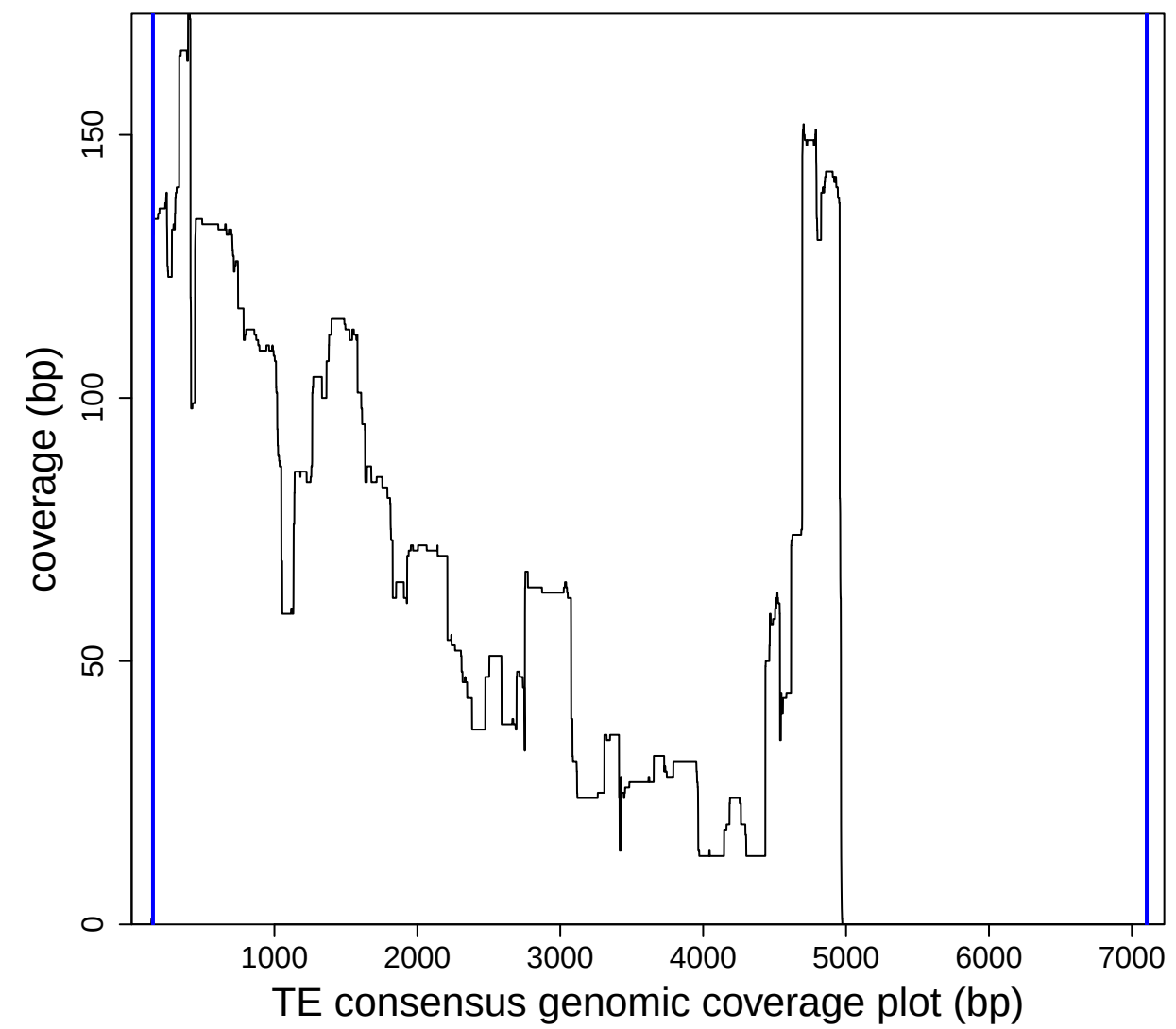
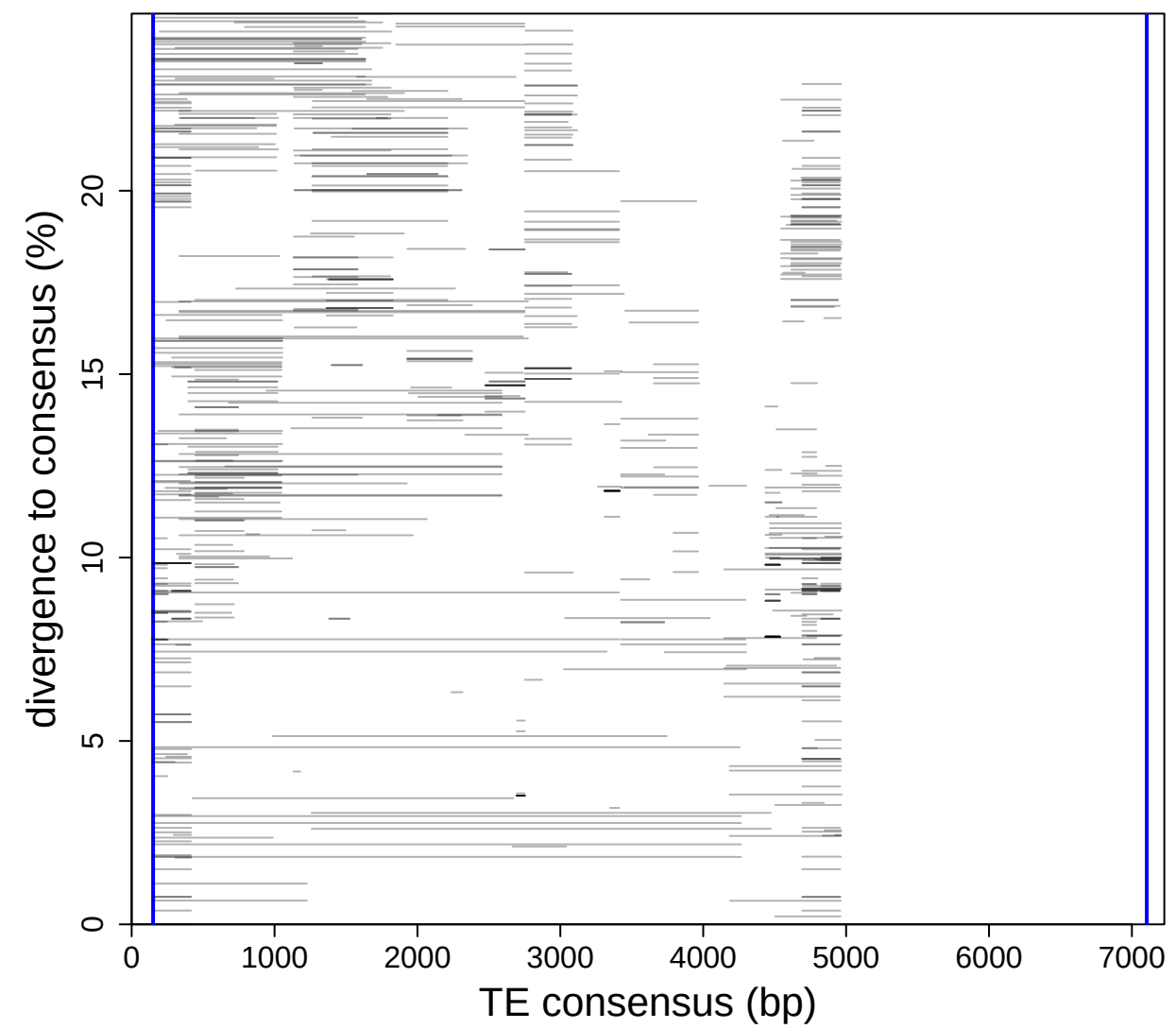
After TEtrimmer 5945 bp



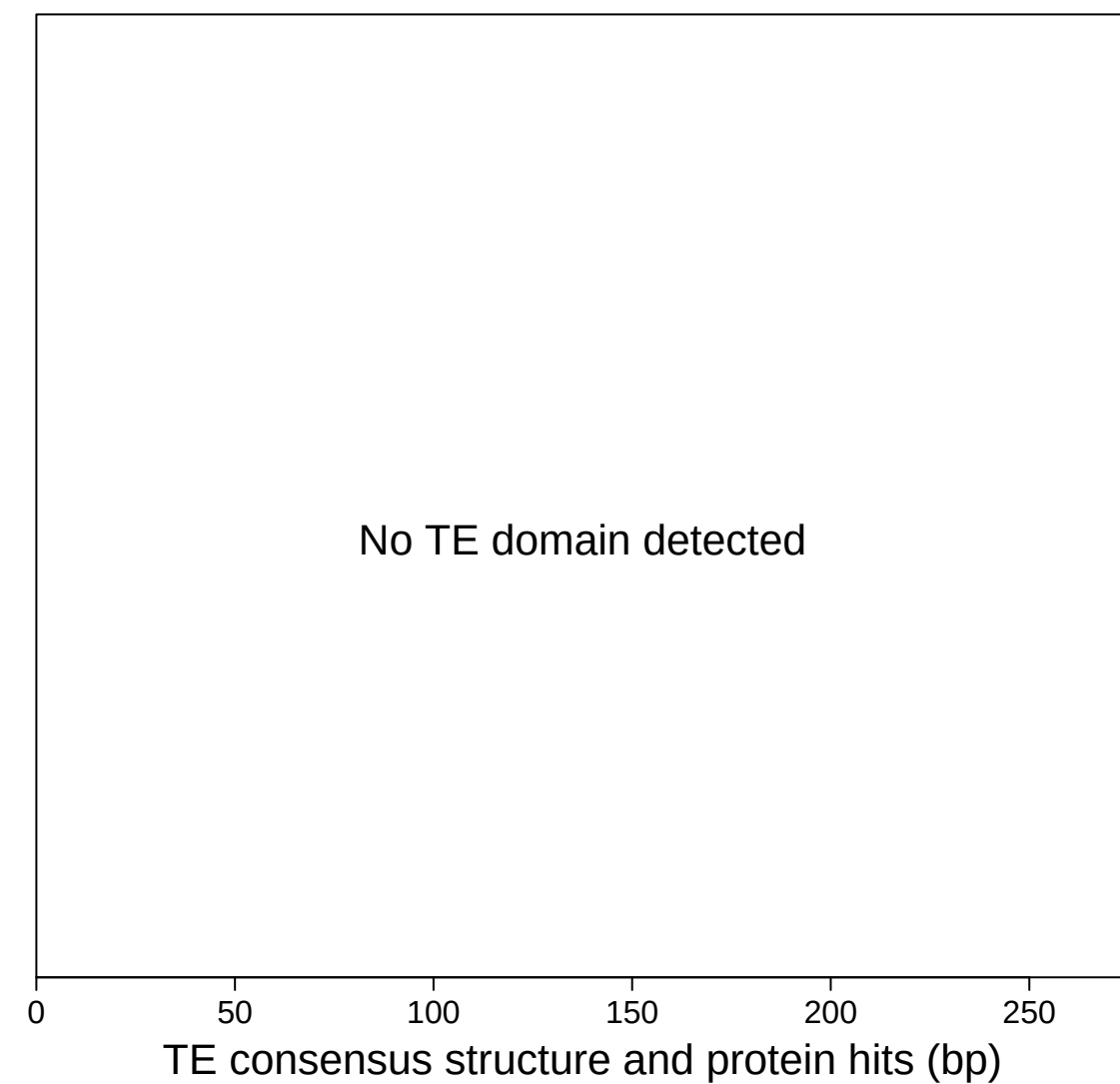
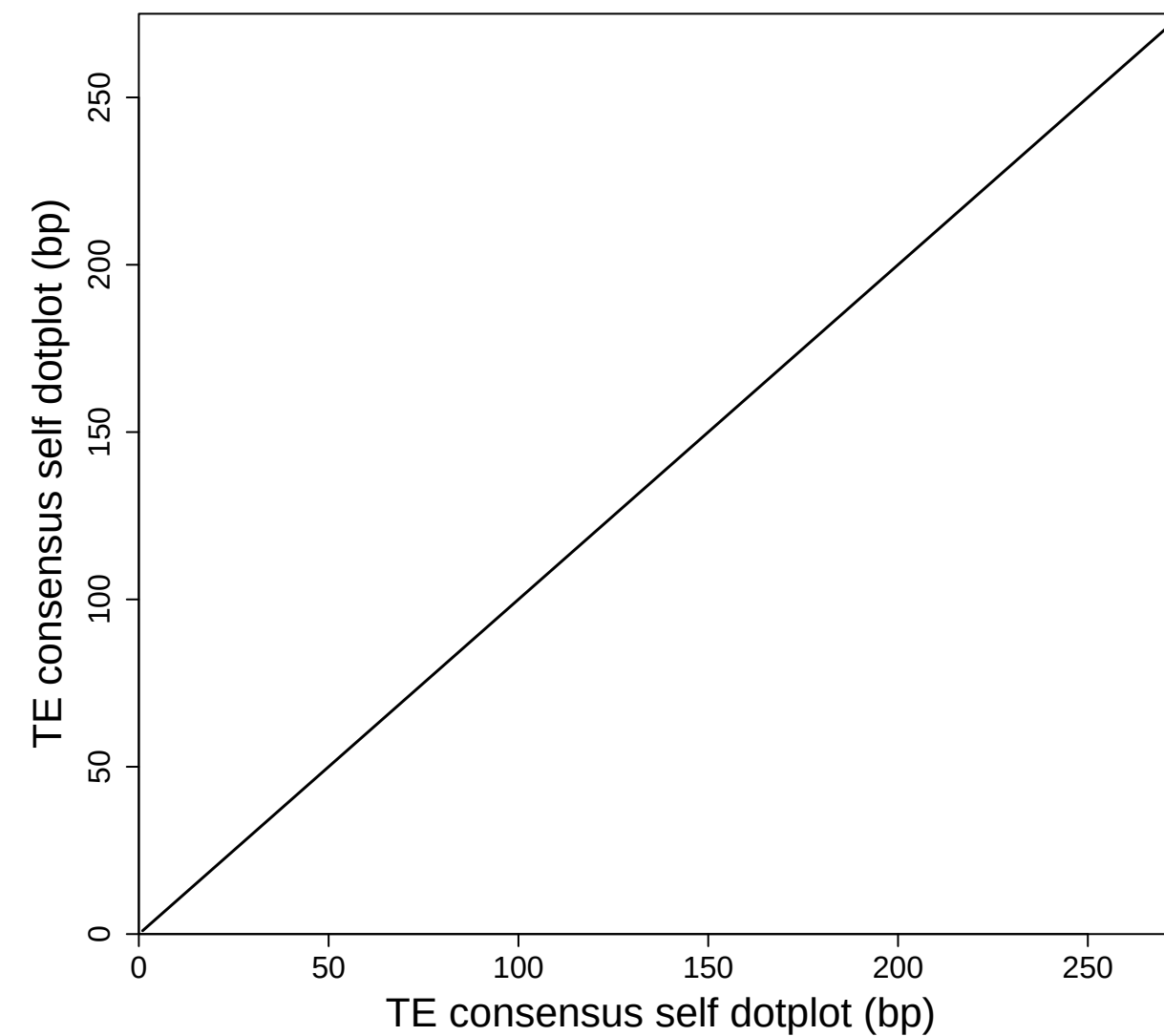
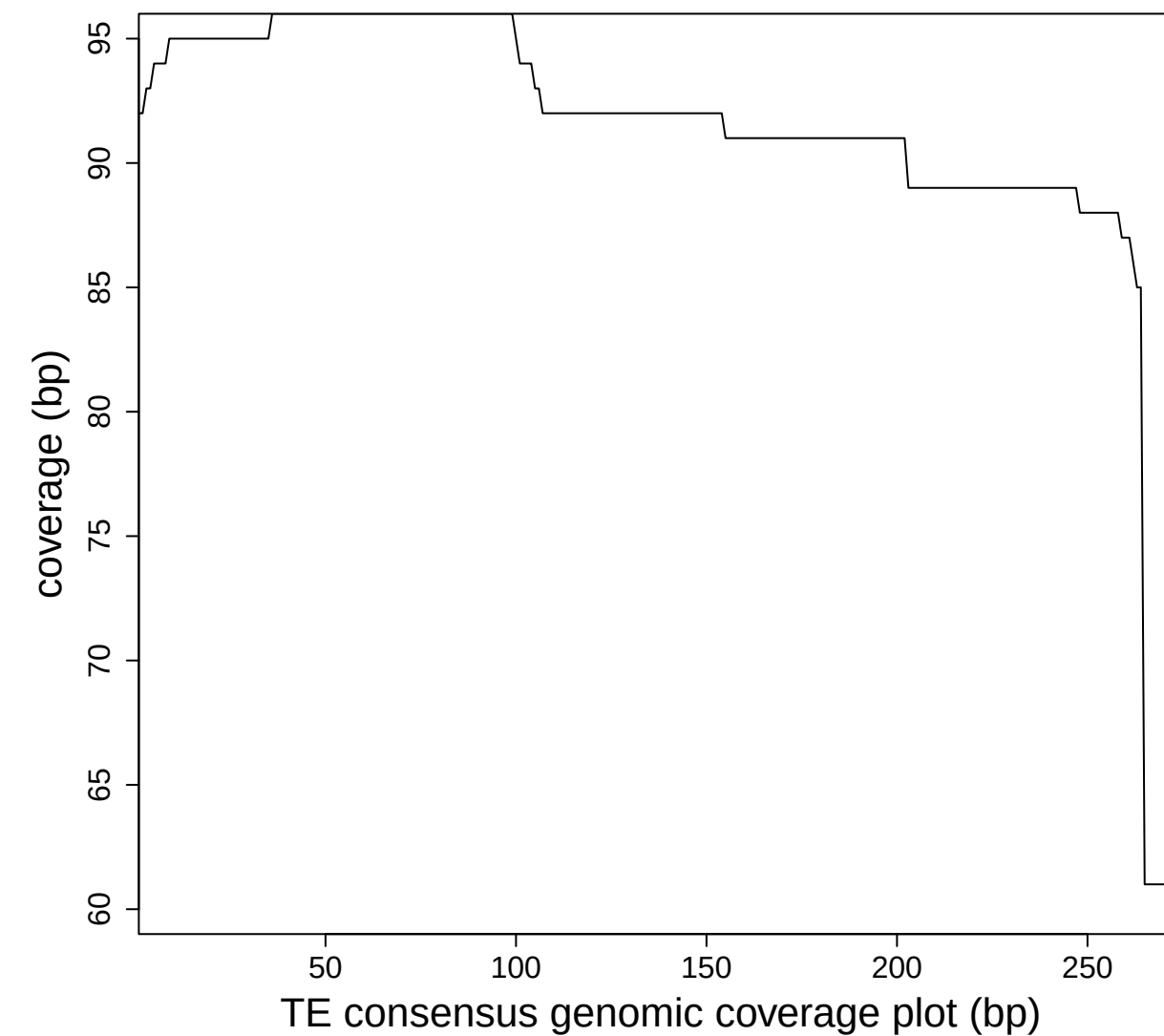
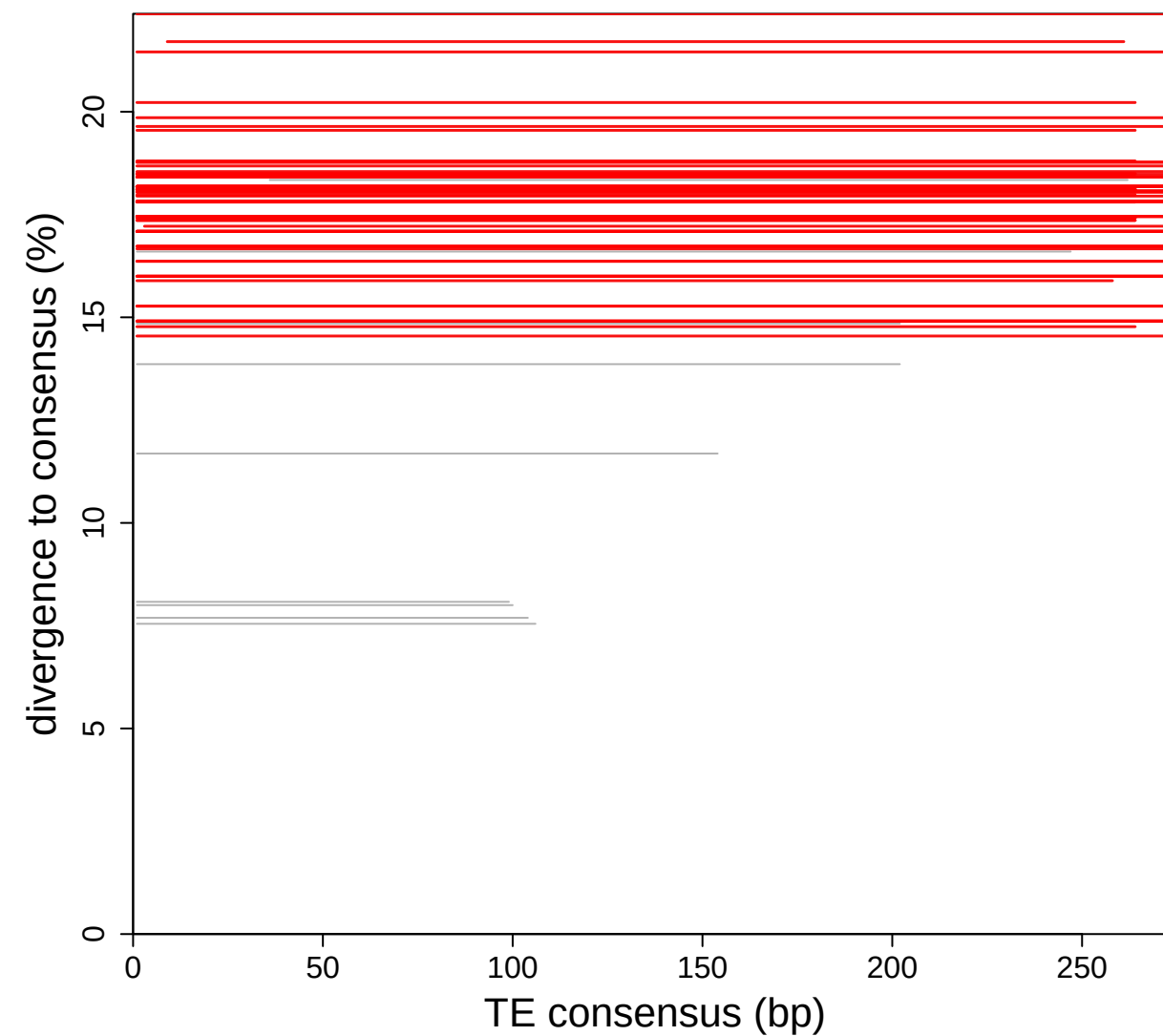
bed_g_3.bed fm_1.bed 0 0 n.bed_g_1.bed fm_2.bed 0 0 bclr
size: 7227bp; fragments: 649; full length: 0 (>=6504.3bp)

After TEtrimmer Extended plot

Blue lines are boundaries

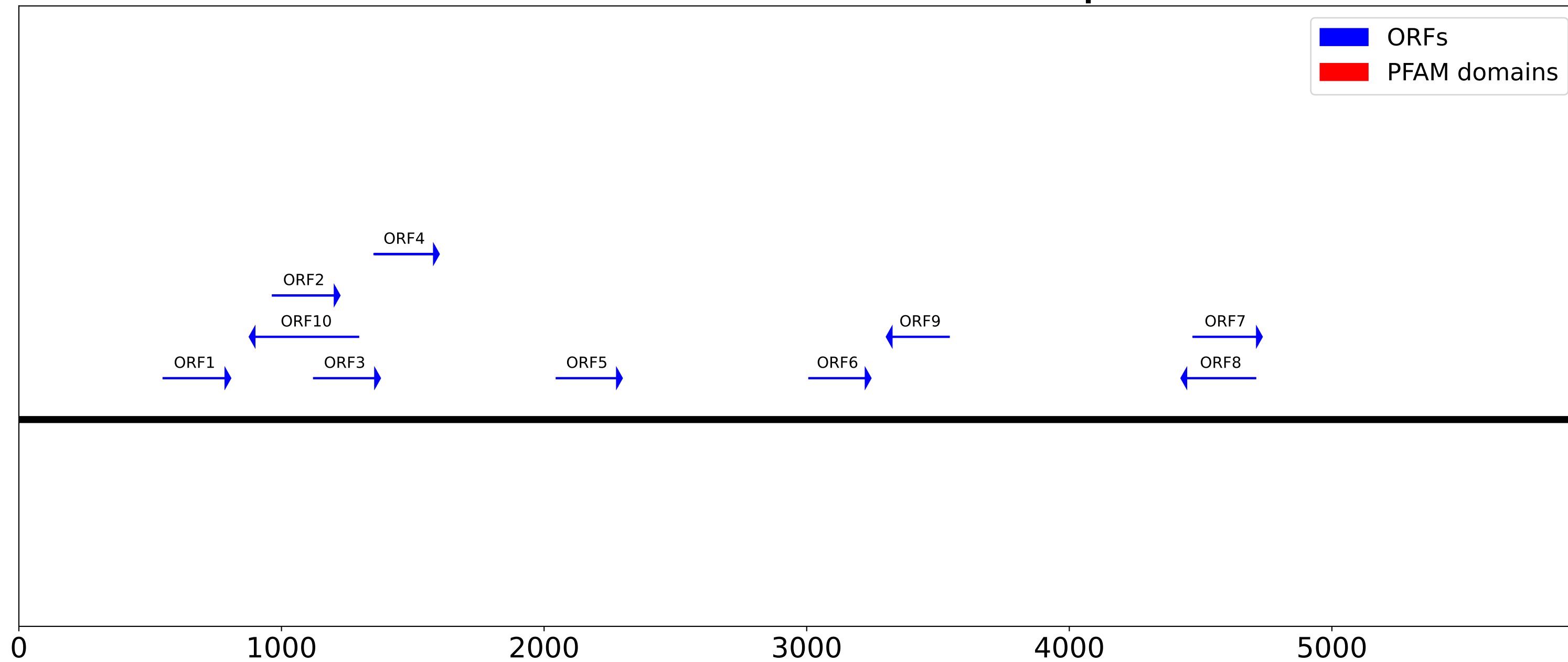


TE: ltr_1_family_307
size: 275bp; fragments: 96; full length: 87 (≥ 247.5 bp)

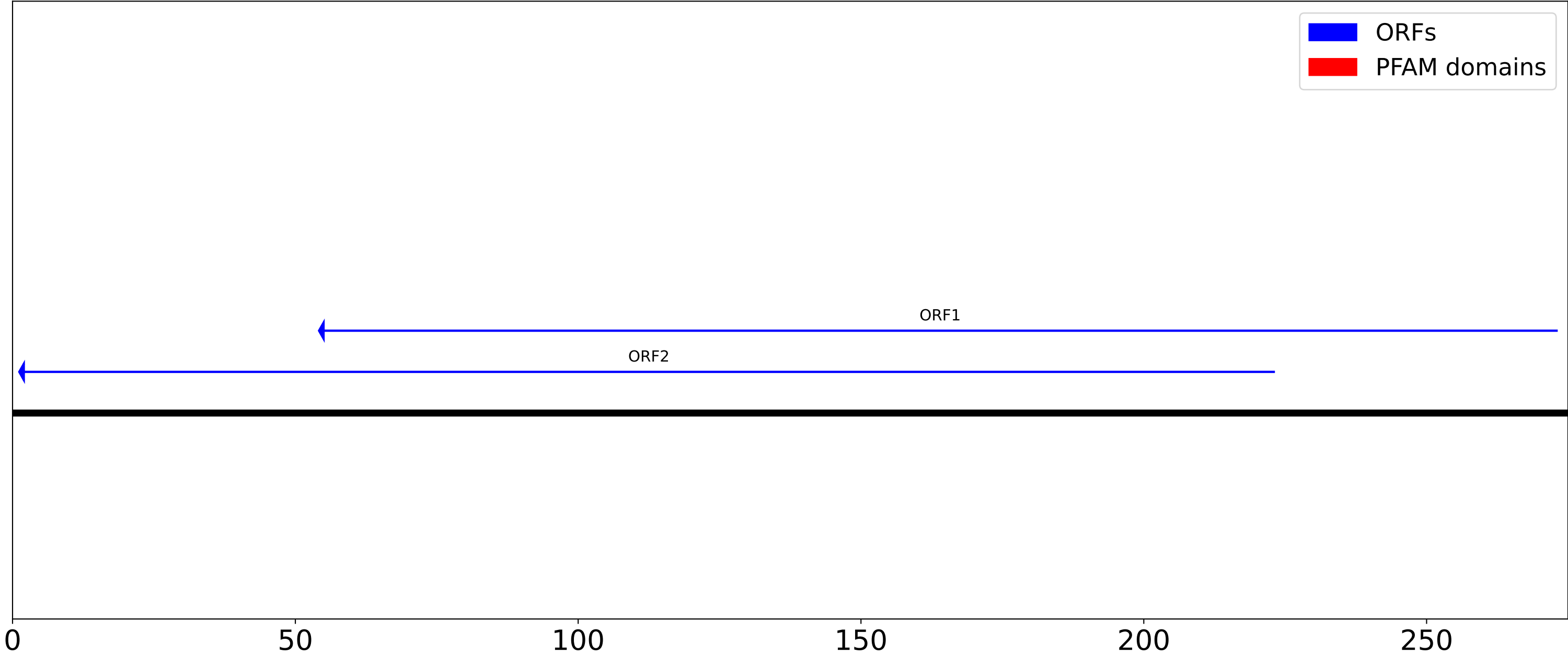


Before TEtrimmer 275 bp

After TEtrimmer ORF and PFAM domain plot



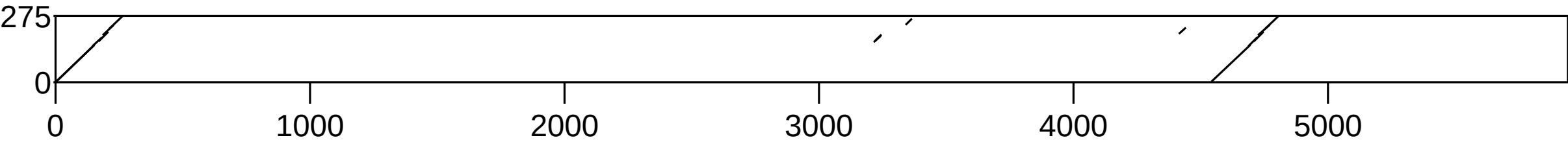
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)