

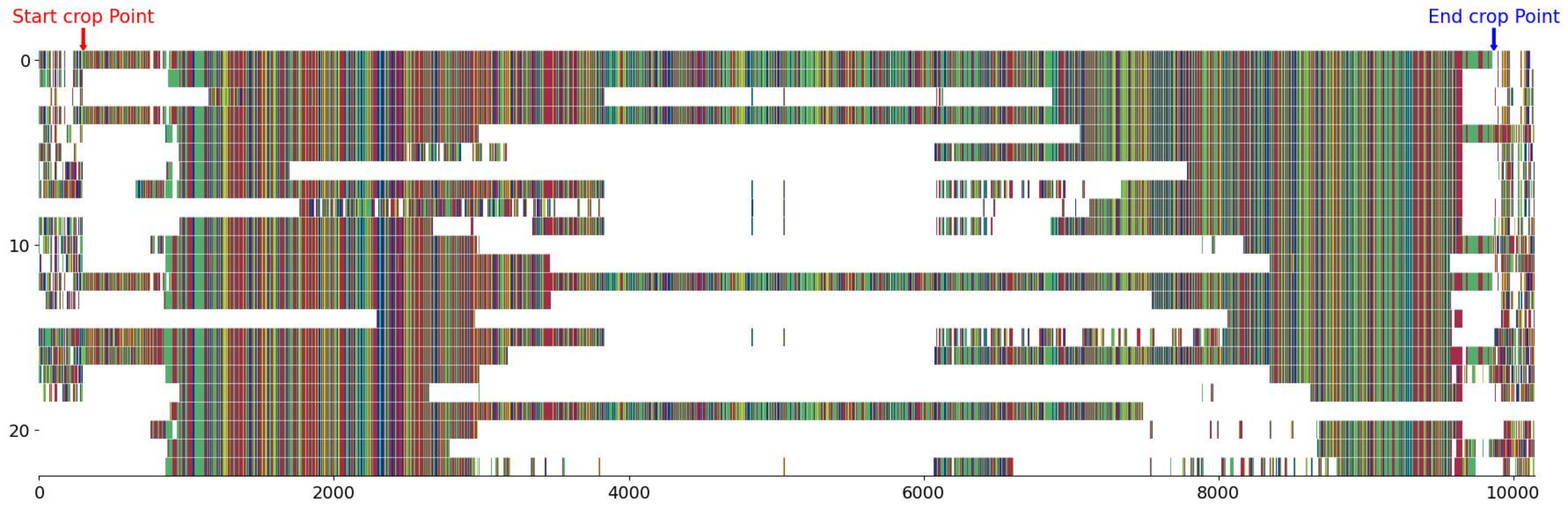
MSA length = 9570

Start crop Point



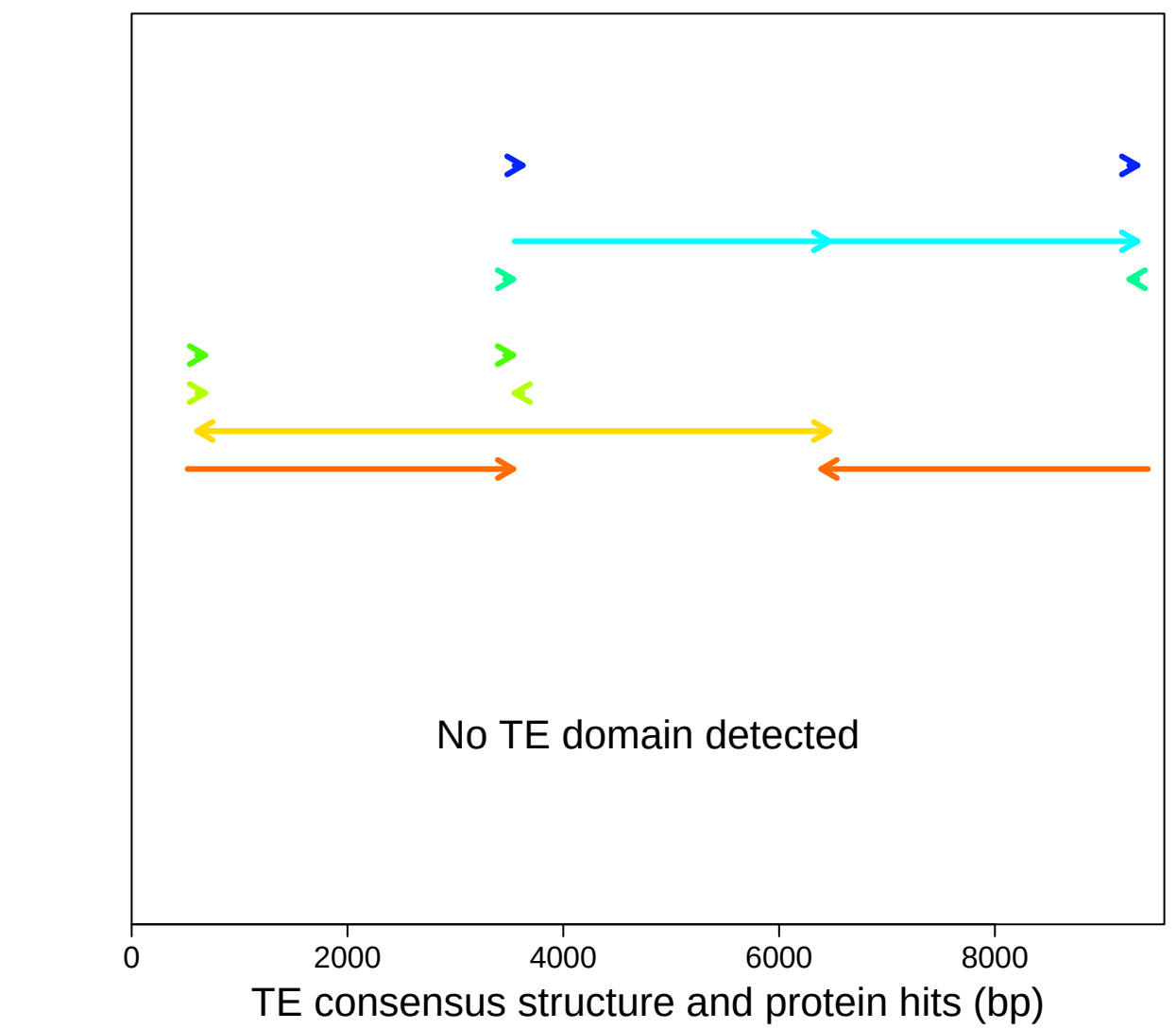
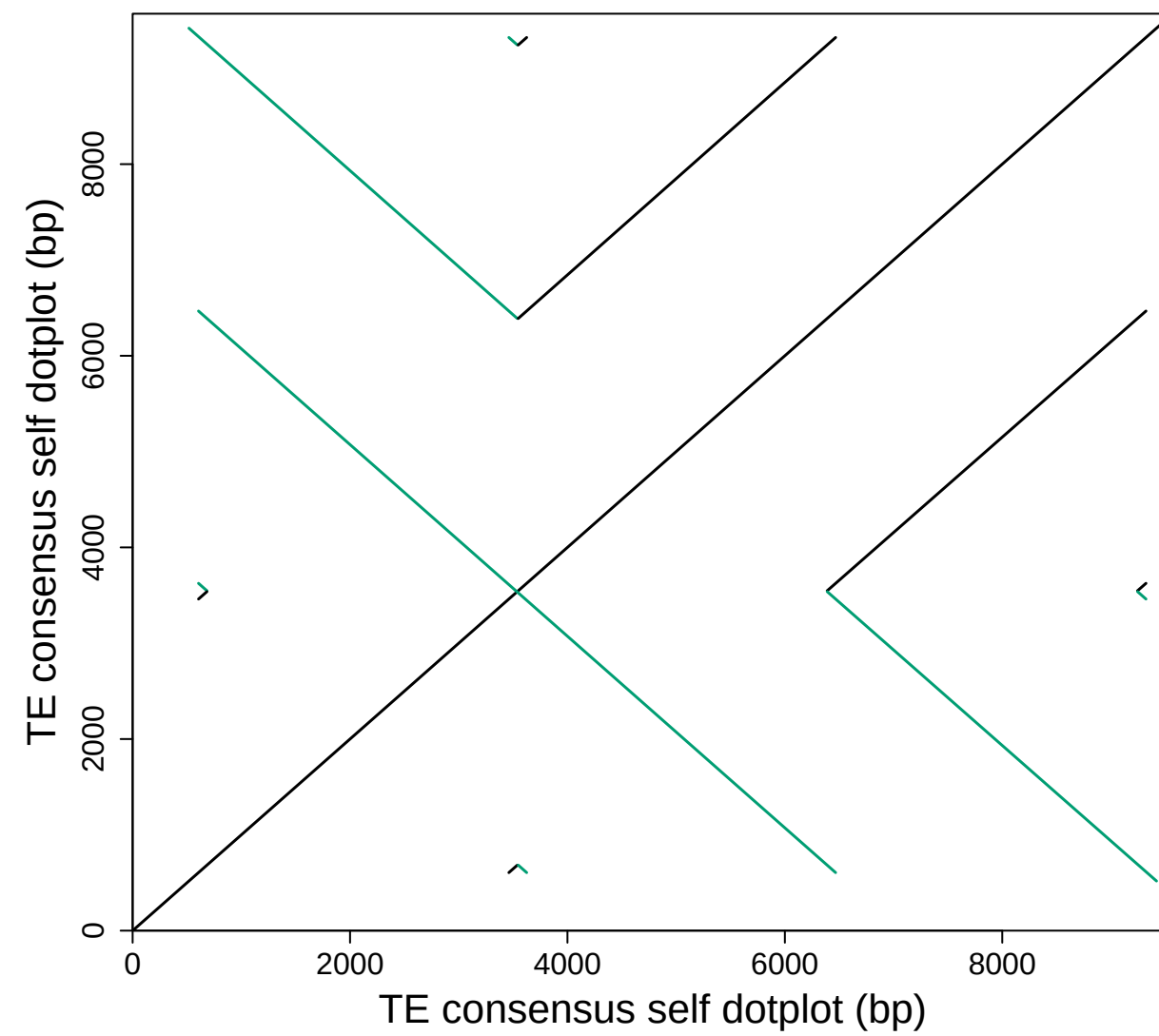
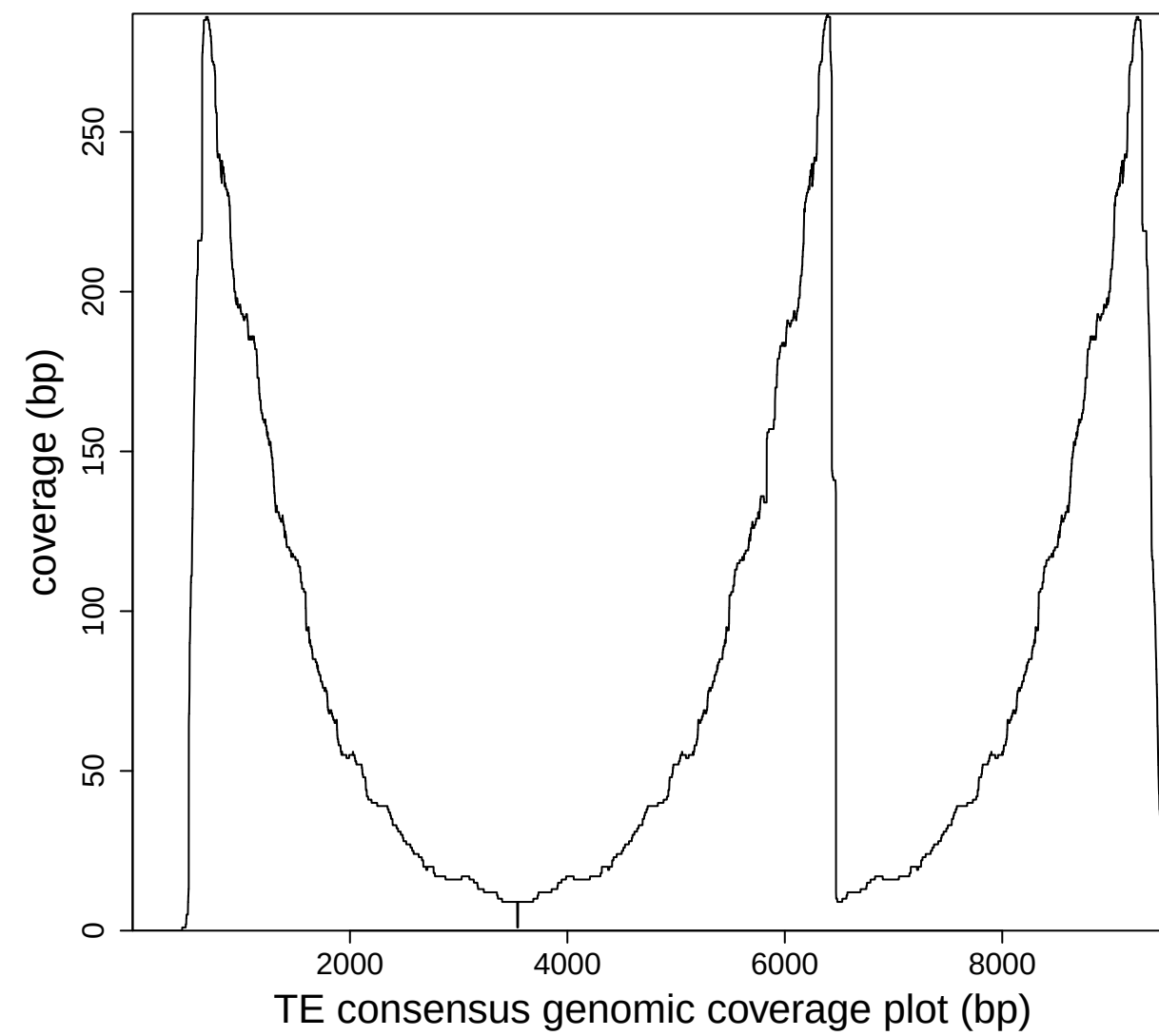
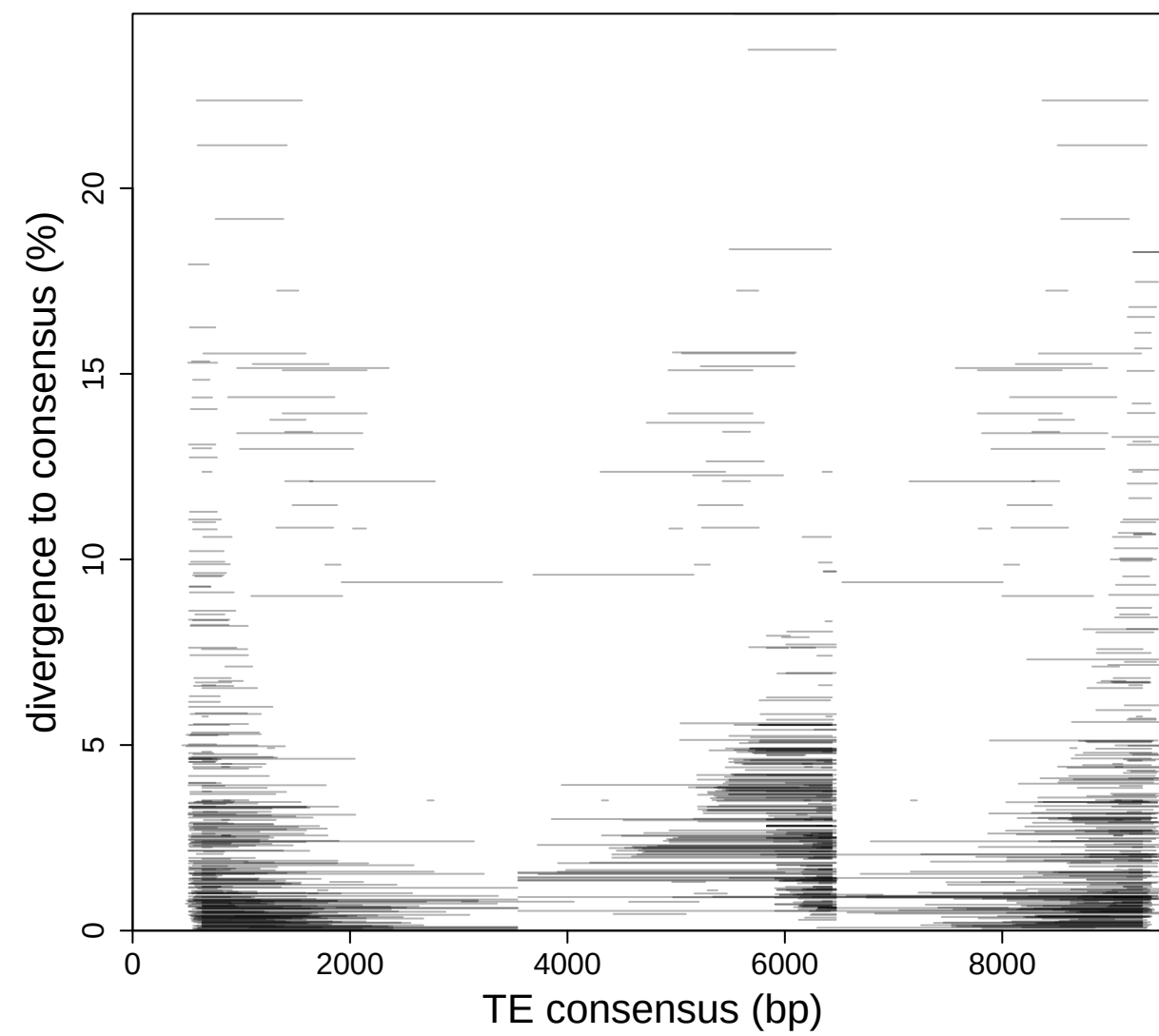
End crop Point





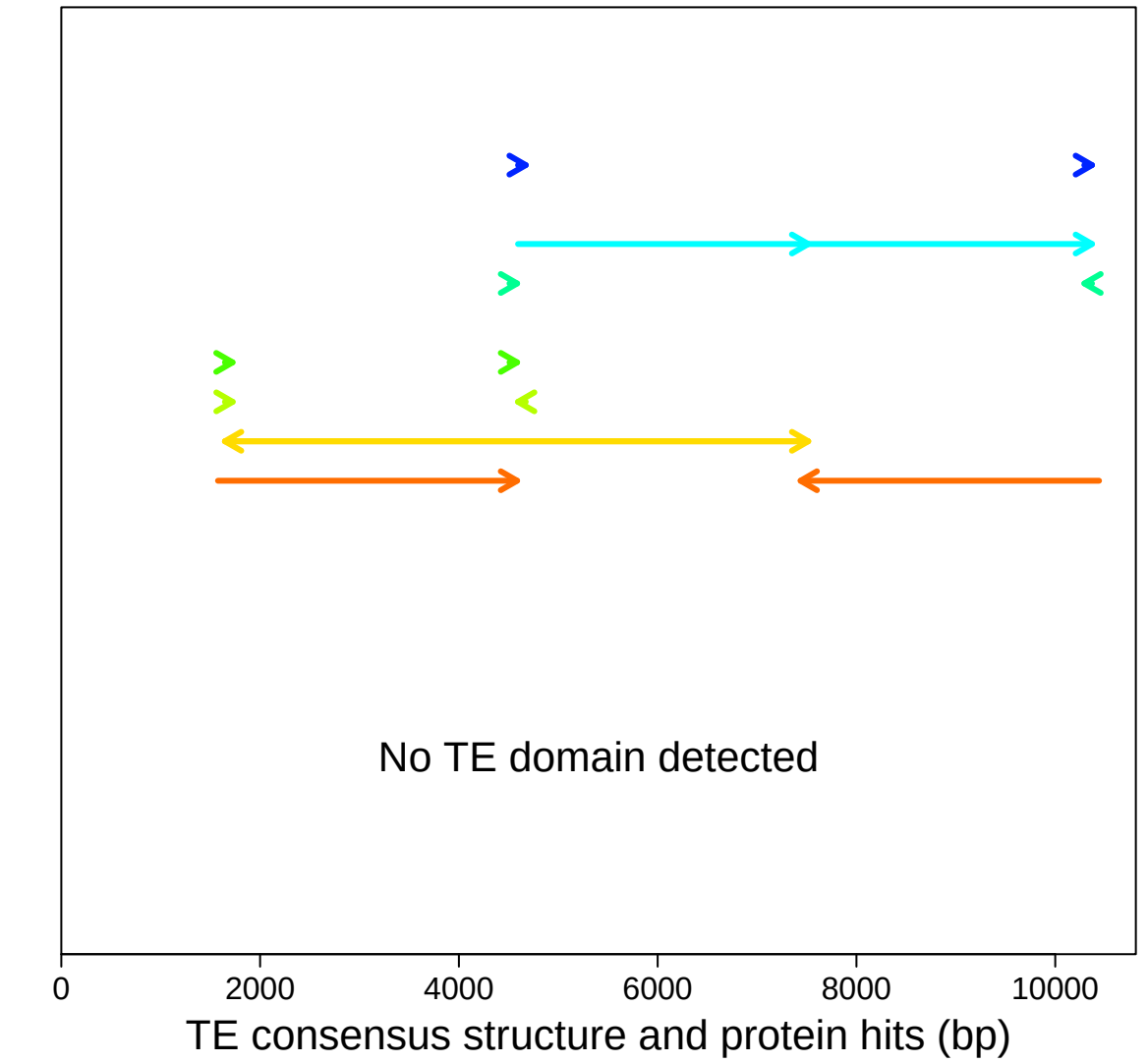
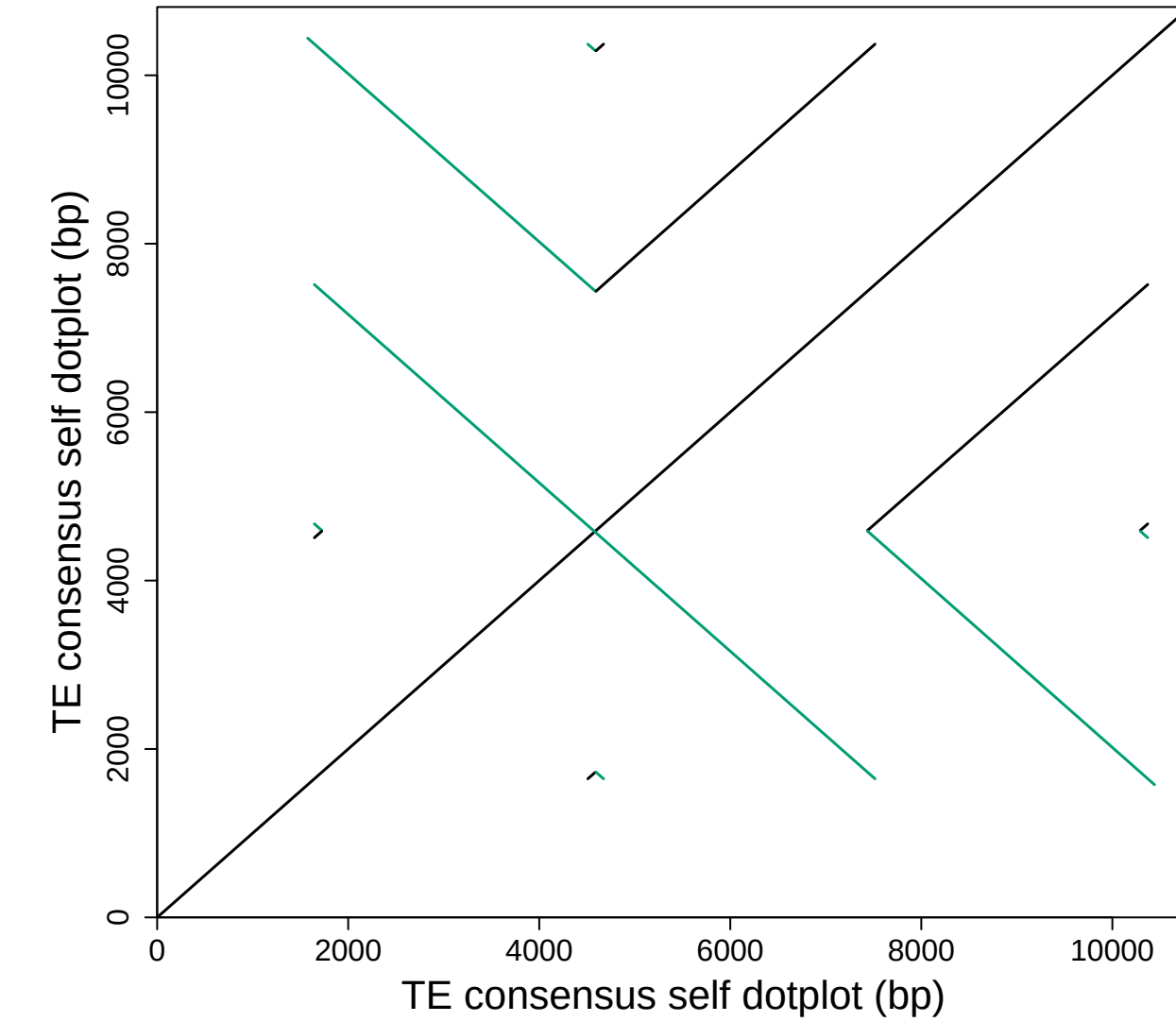
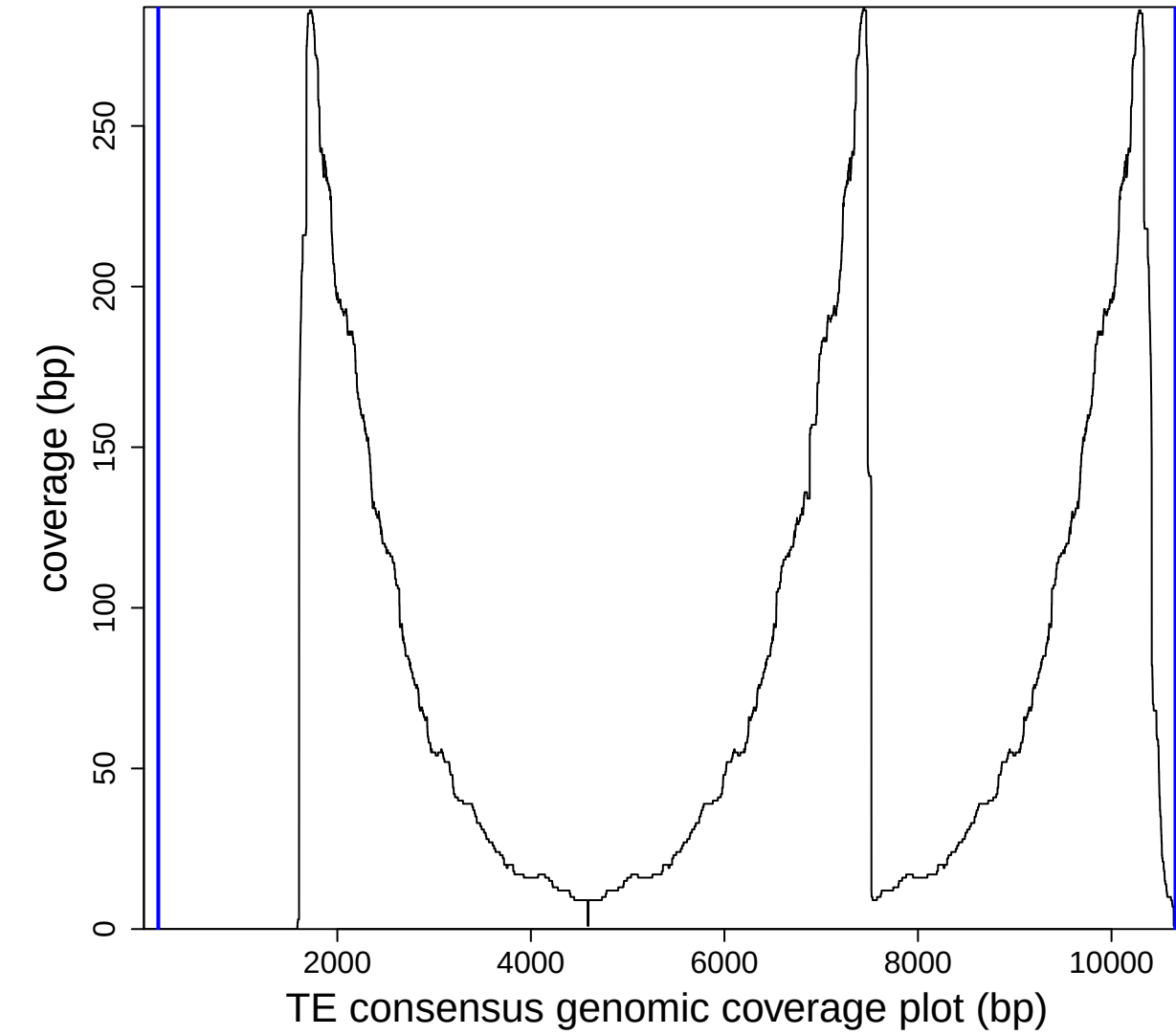
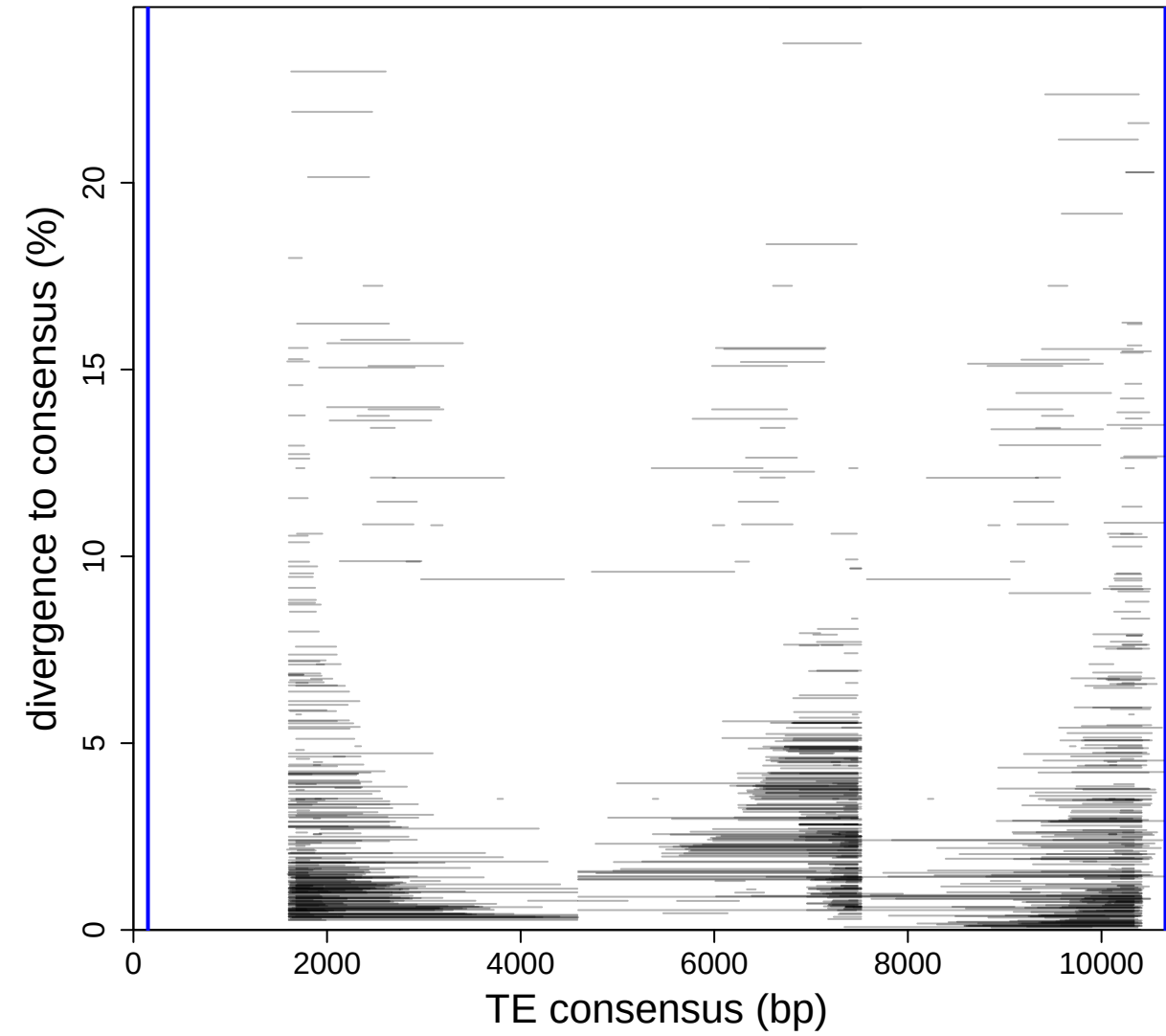
bed_fm_1.bed 0 0 n.bed g 1.bed fm_2.bed 0 0 bcln.fa aln.fa
size: 9570bp; fragments: 987; full length: 0 (>=8613bp)

After TETrimmer 9570 bp

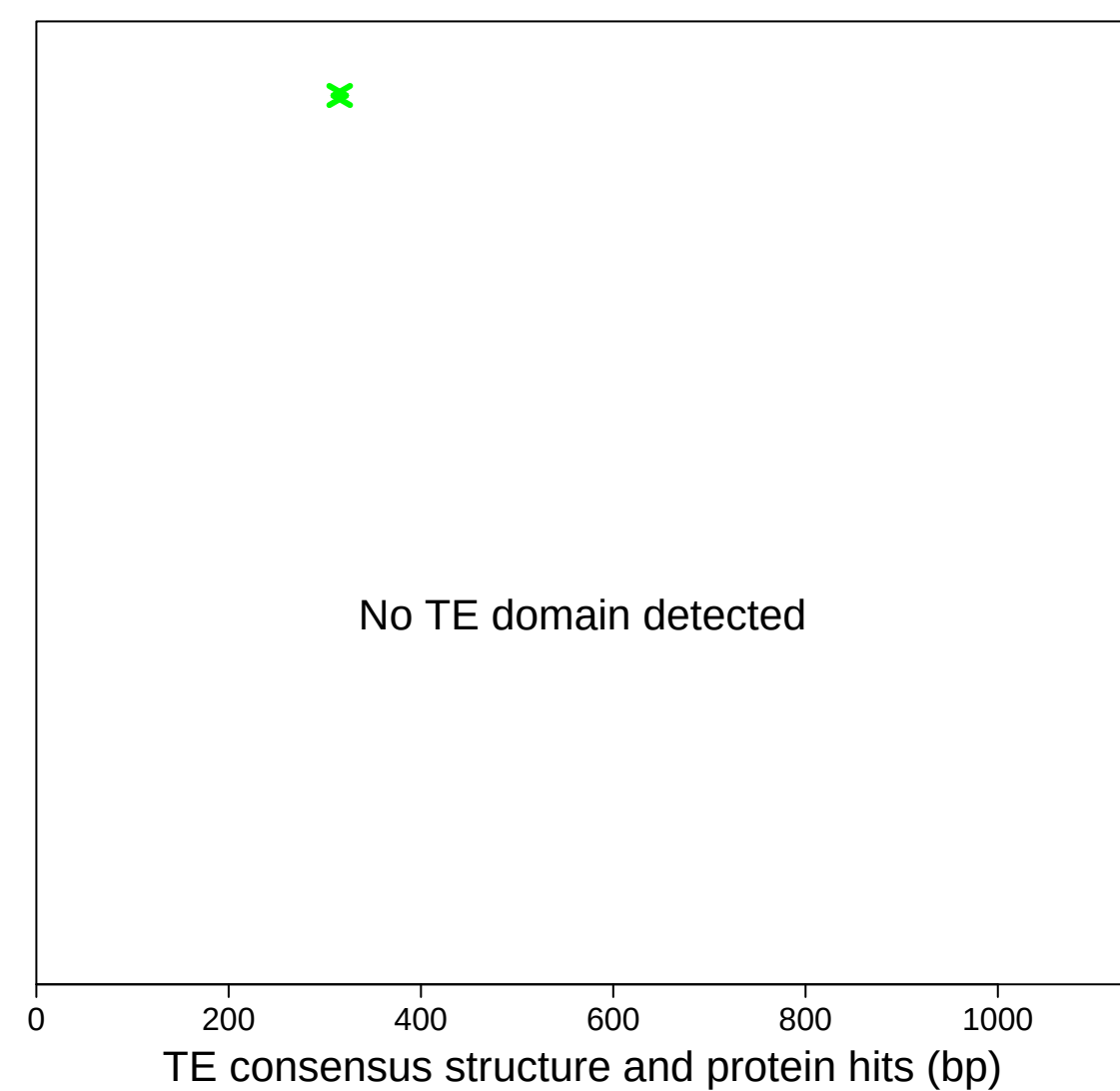
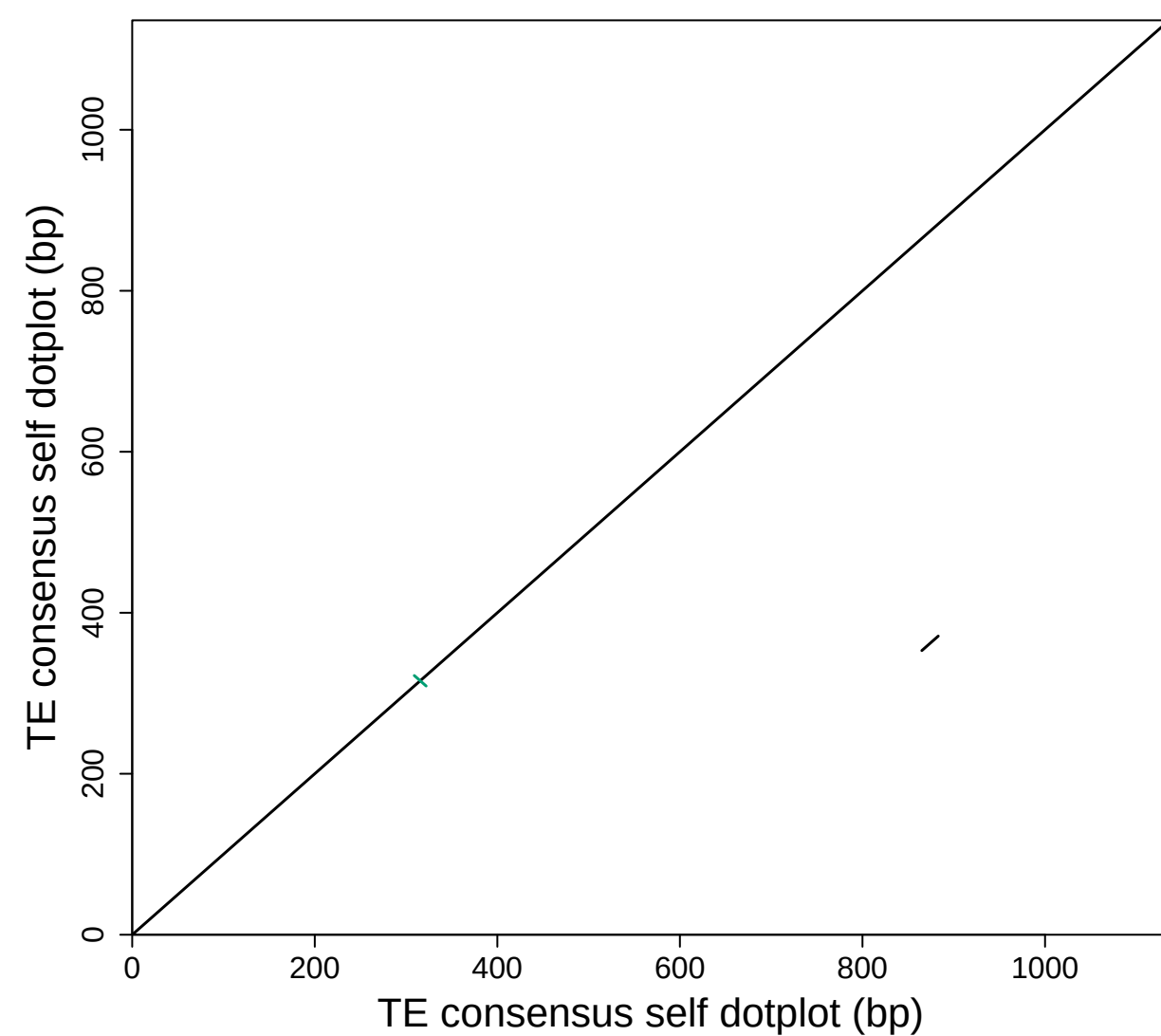
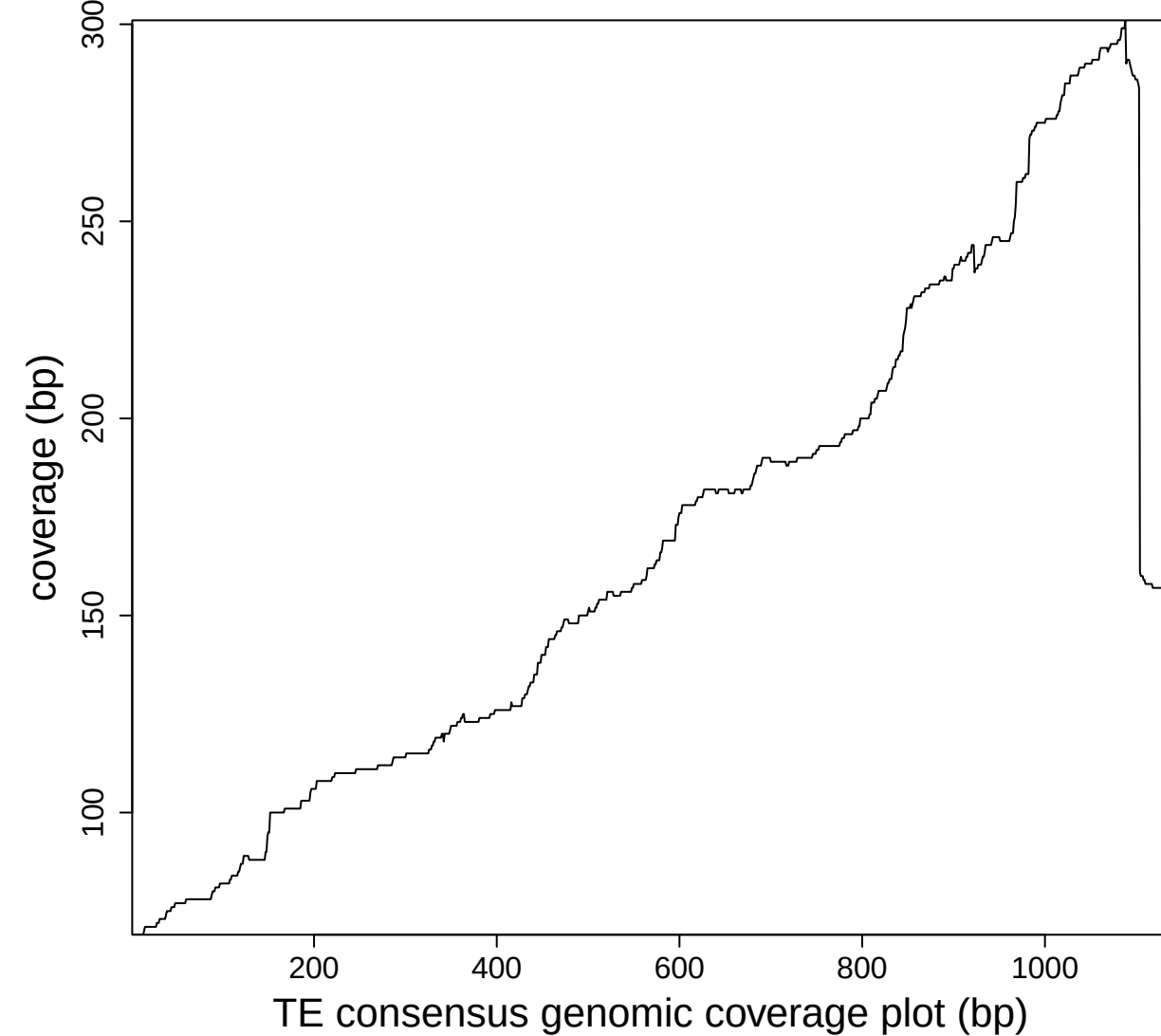
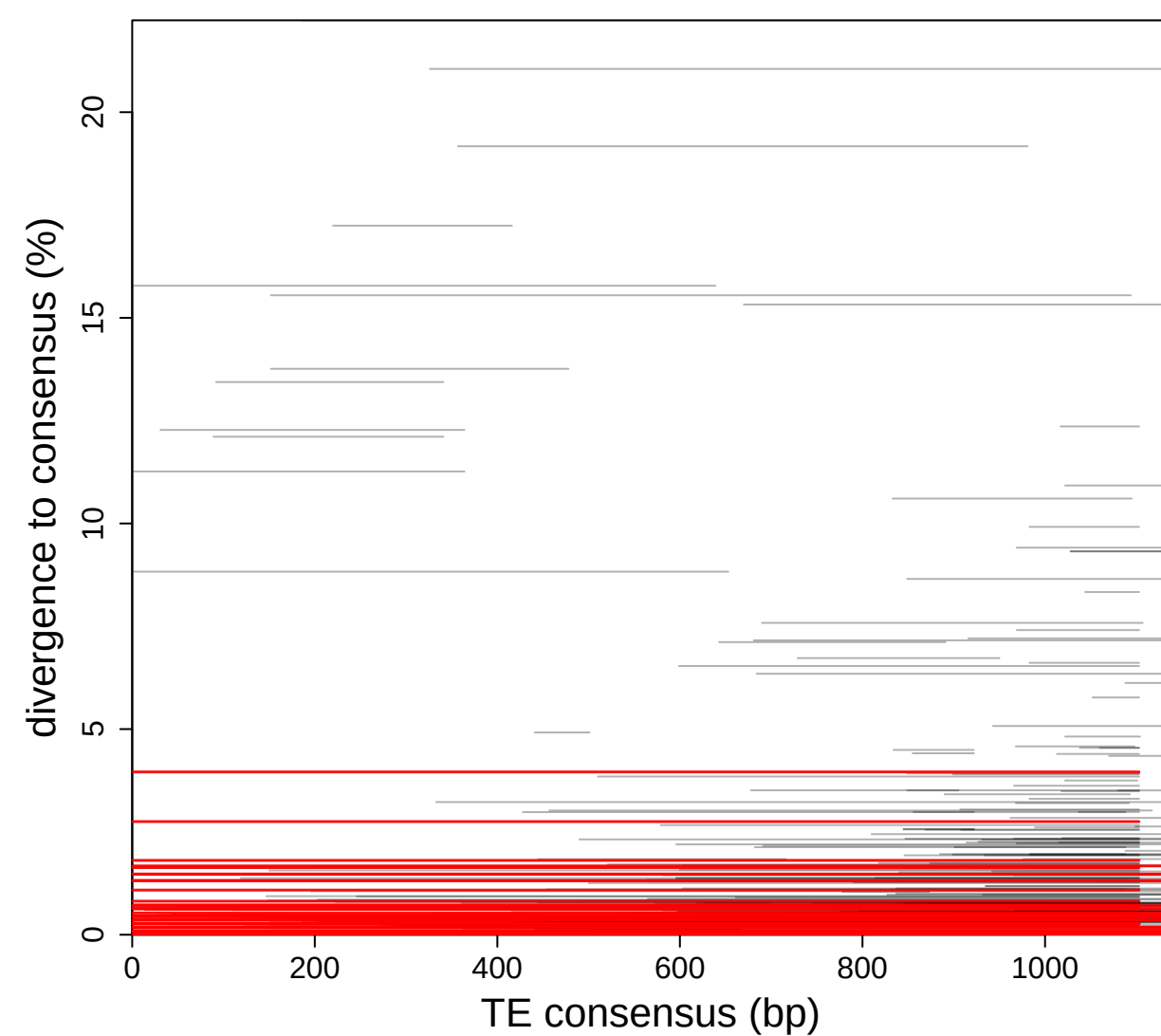


f.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcl
size: 10811bp; fragments: 987; full length: 0 (>=9729.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

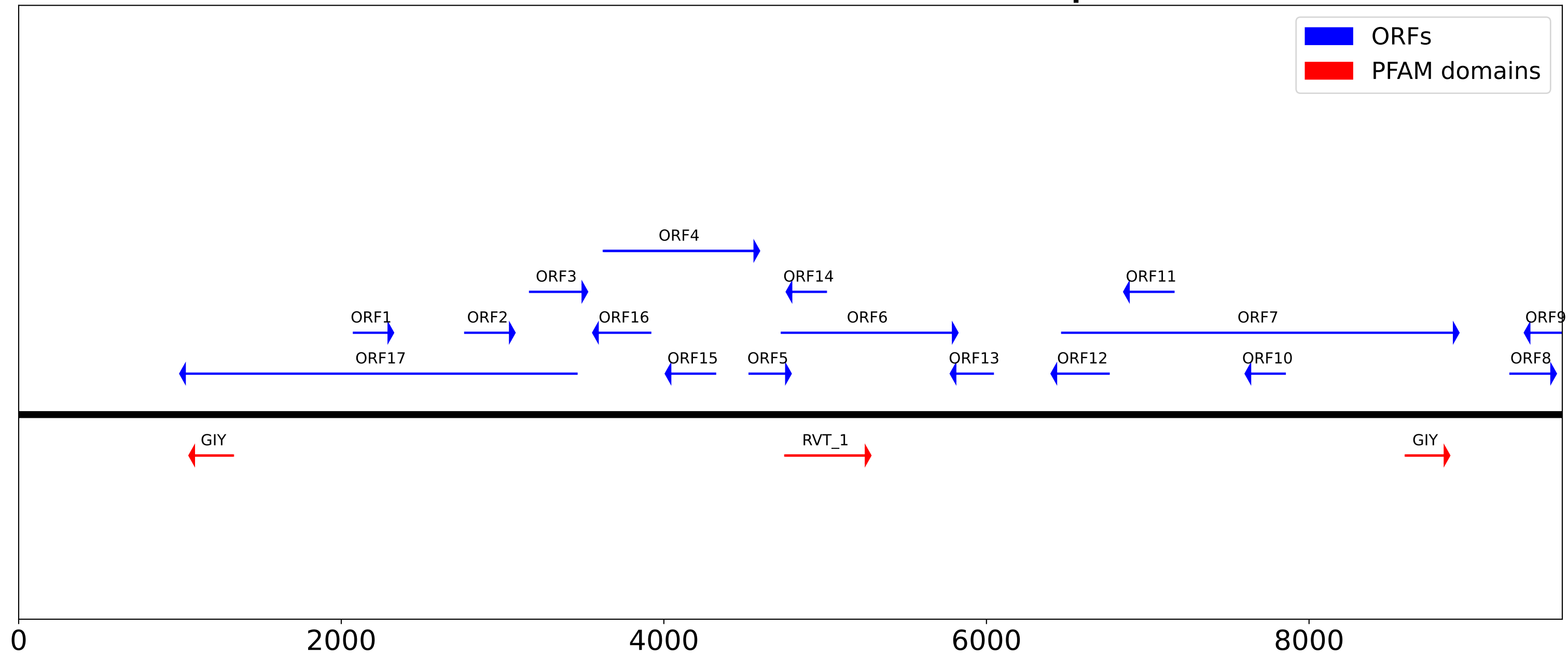


TE: rnd_1_family_462
size: 1136bp; fragments: 335; full length: 73 (≥ 1022.4 bp)

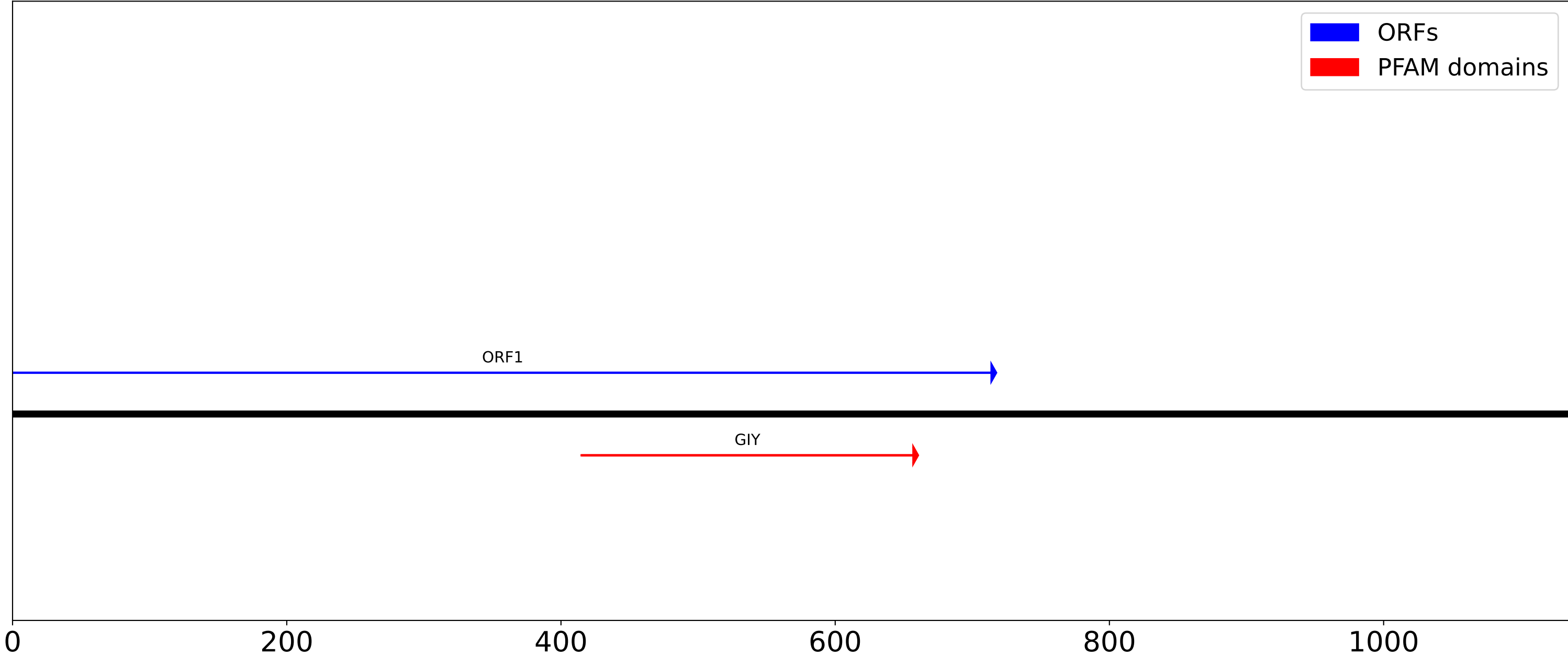


Before TETrimmer 1136 bp

After TEtrimmer ORF and PFAM domain plot



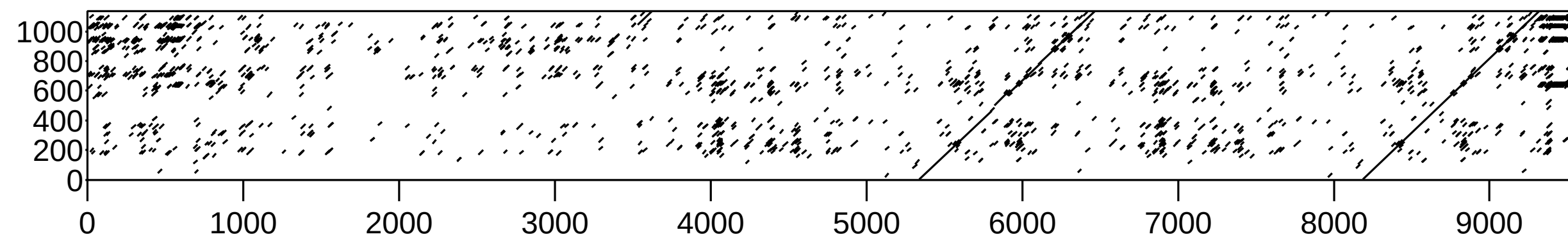
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

sensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)