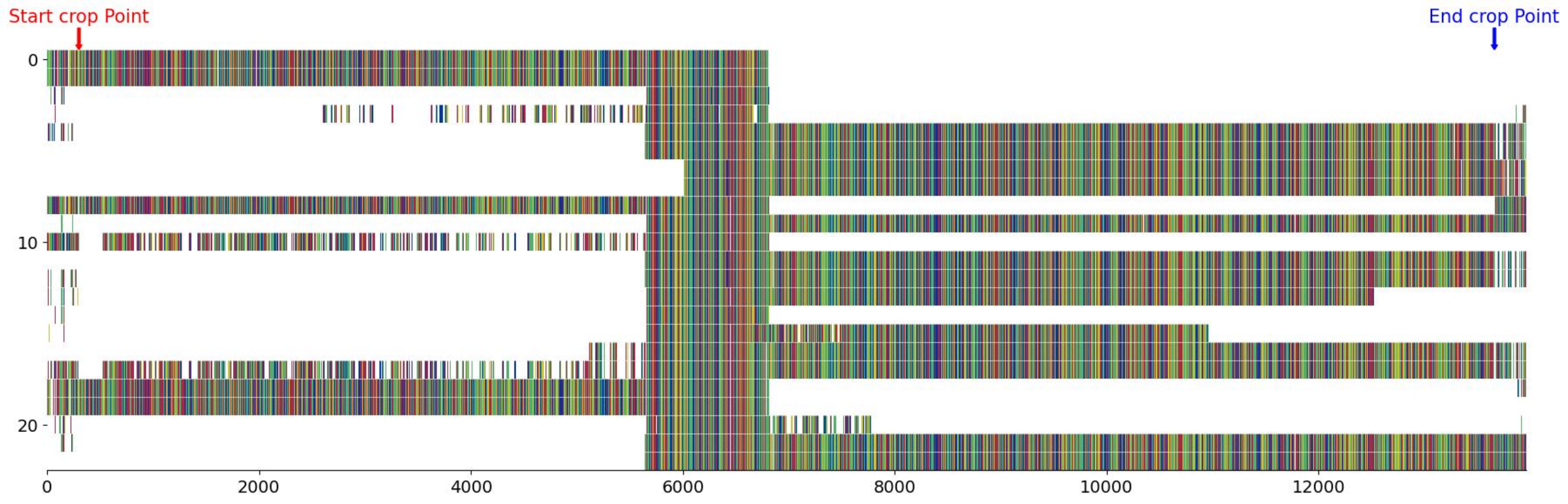


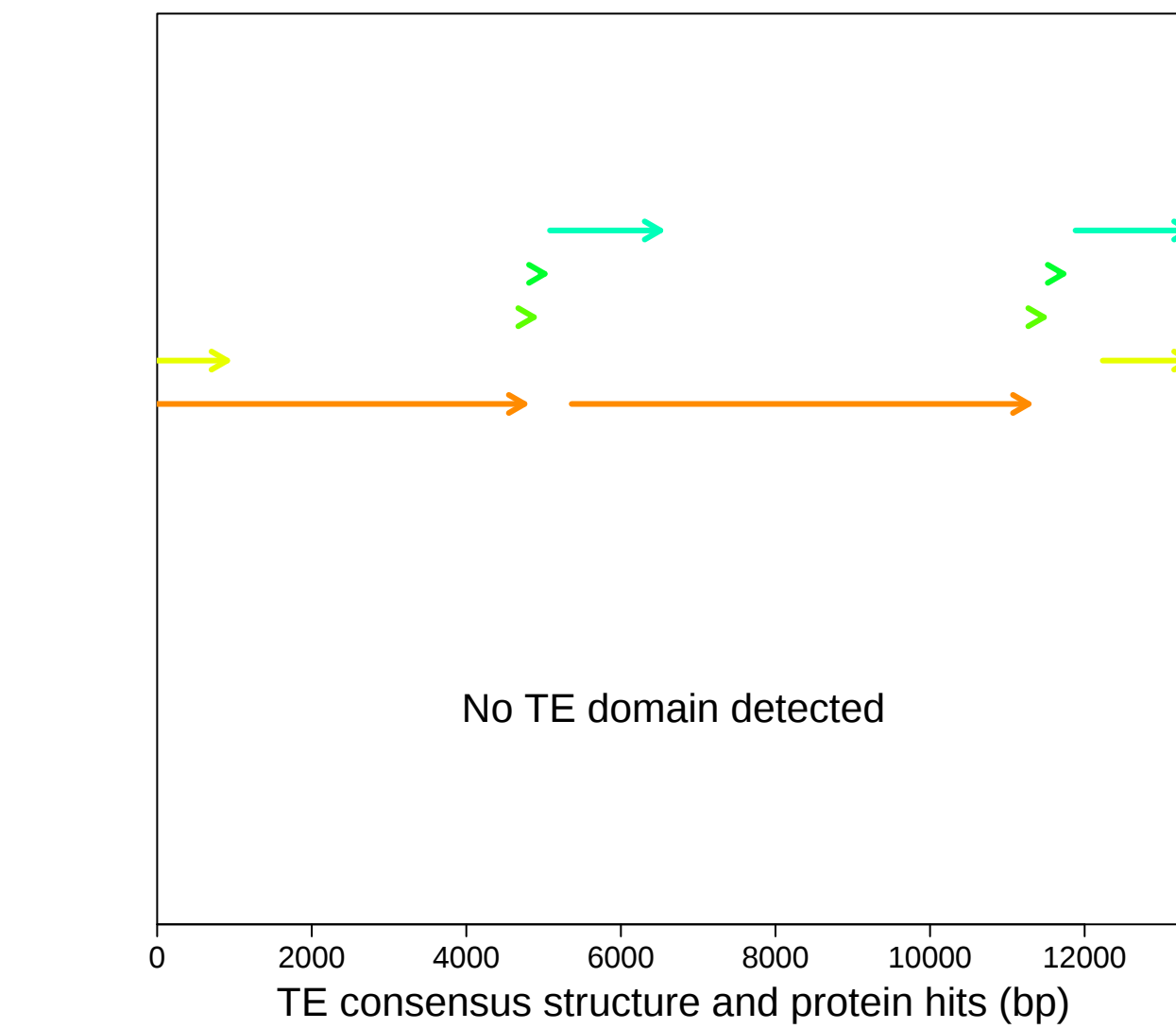
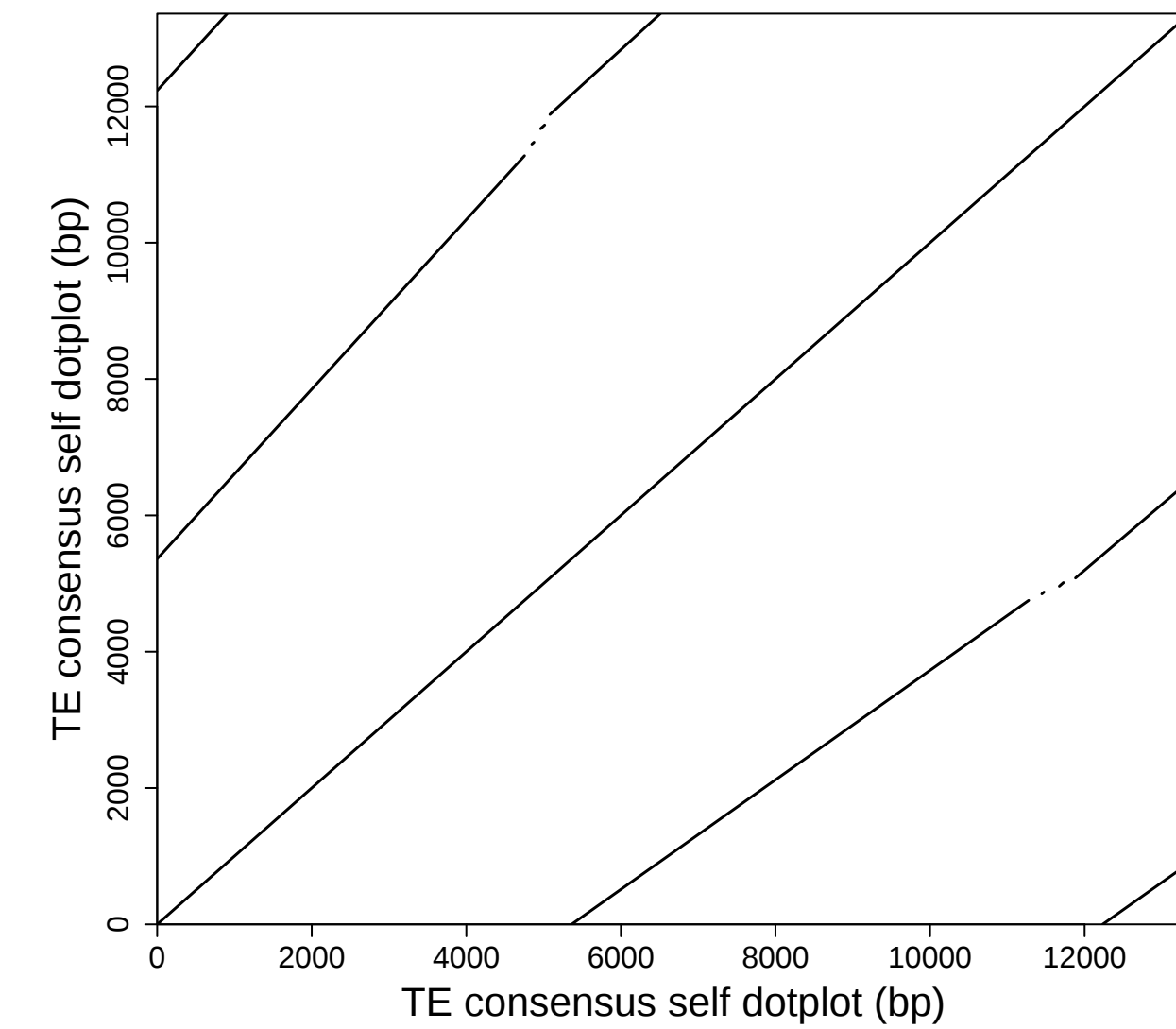
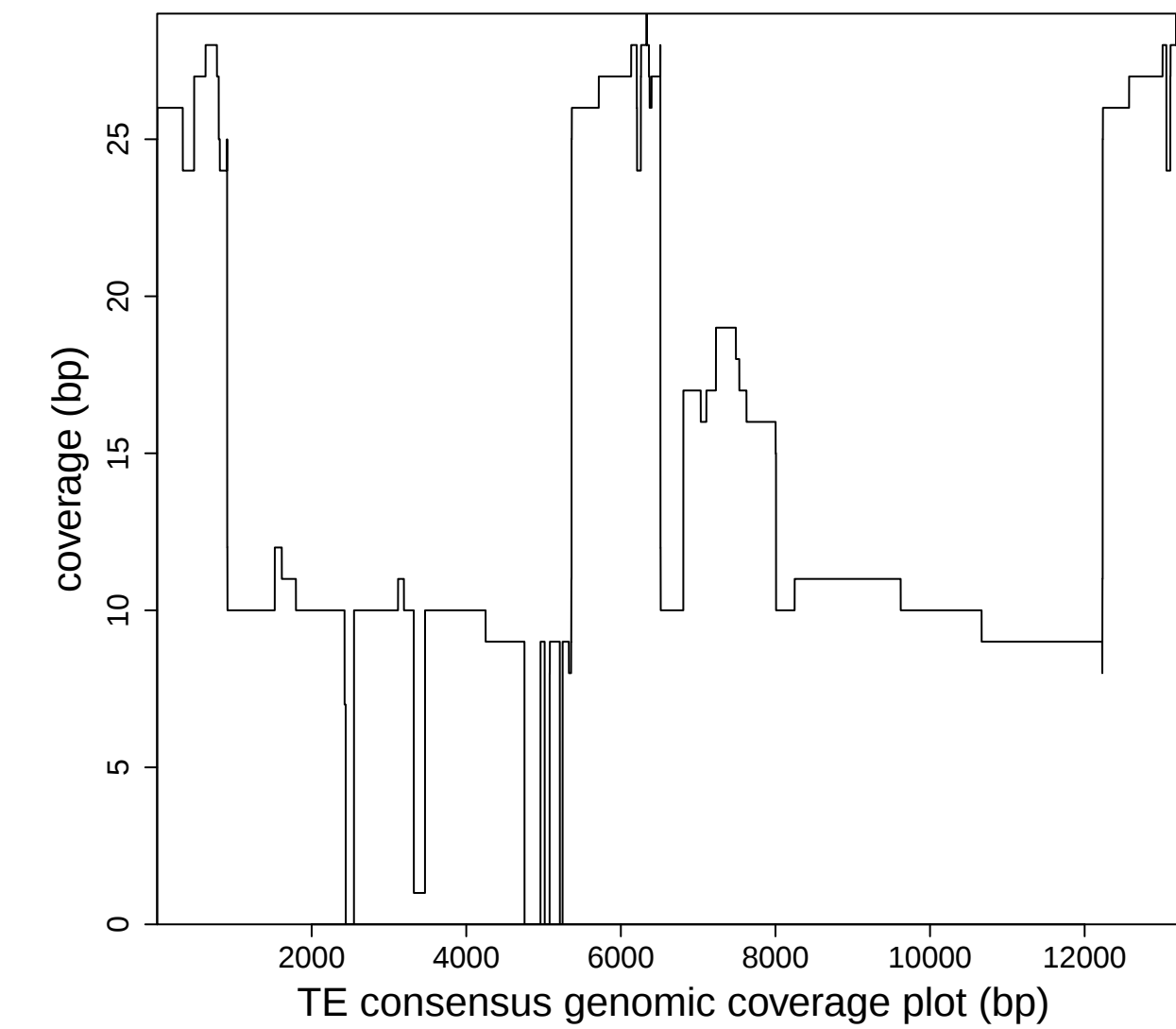
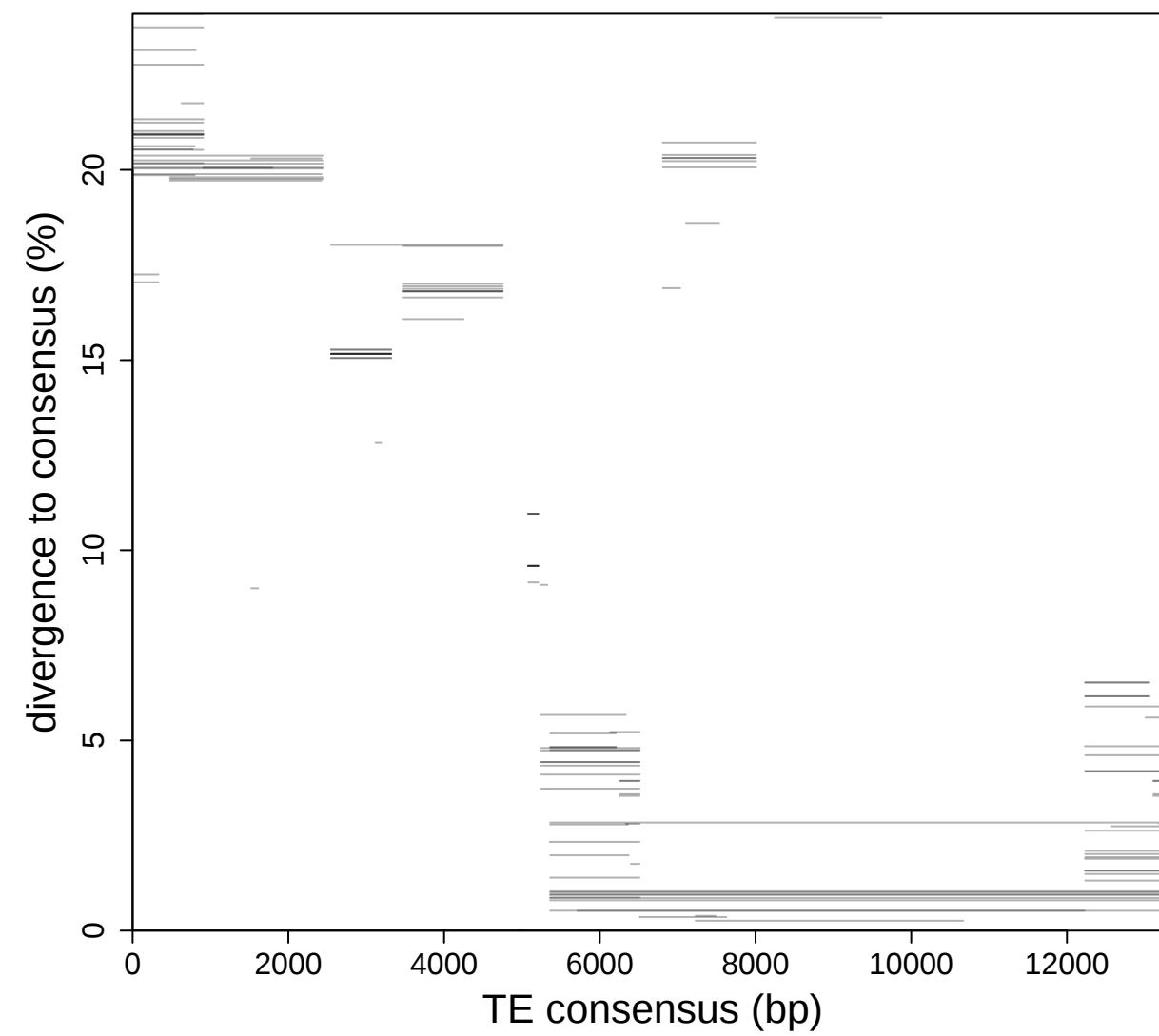
MSA length = 13361

[illegible]



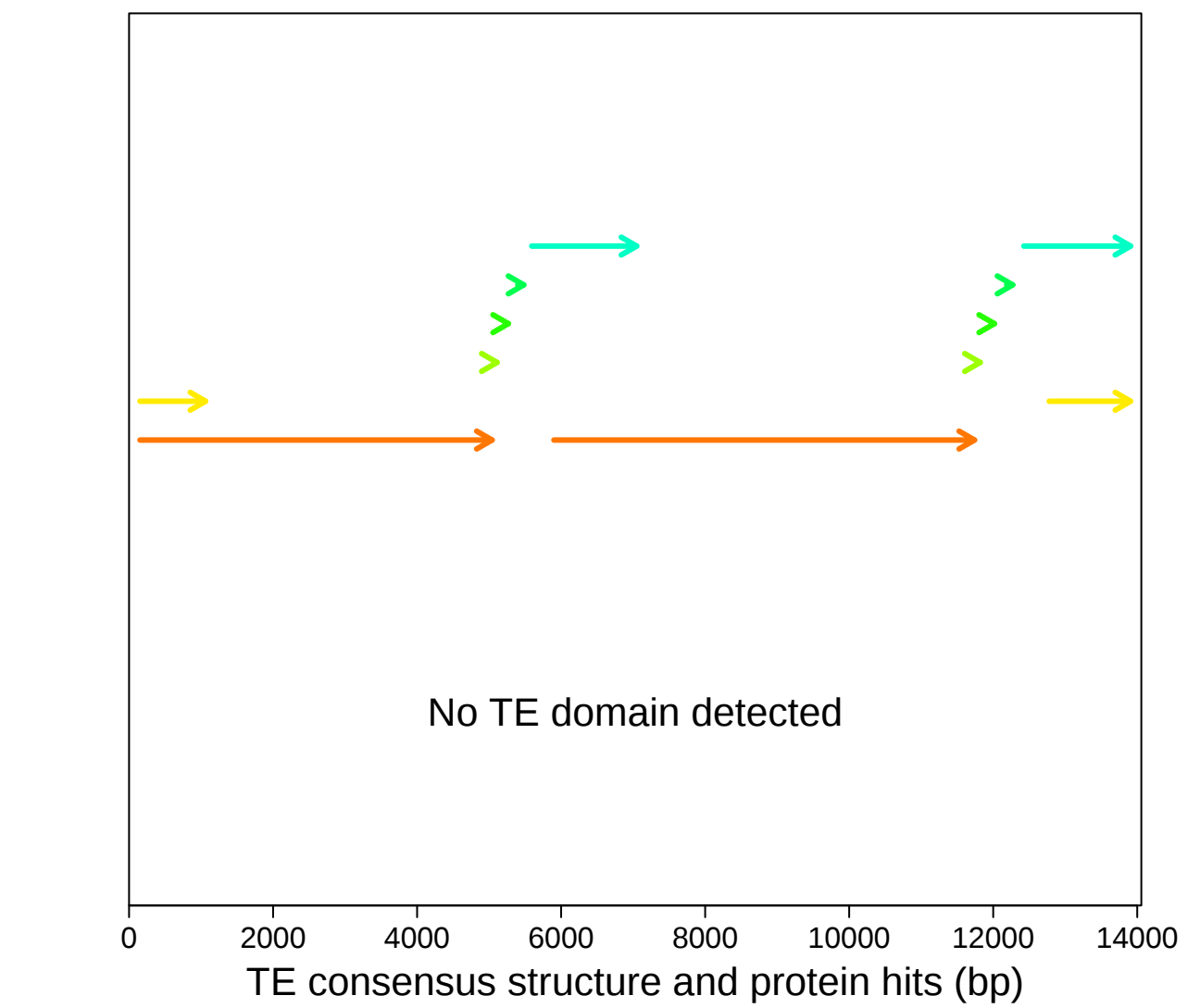
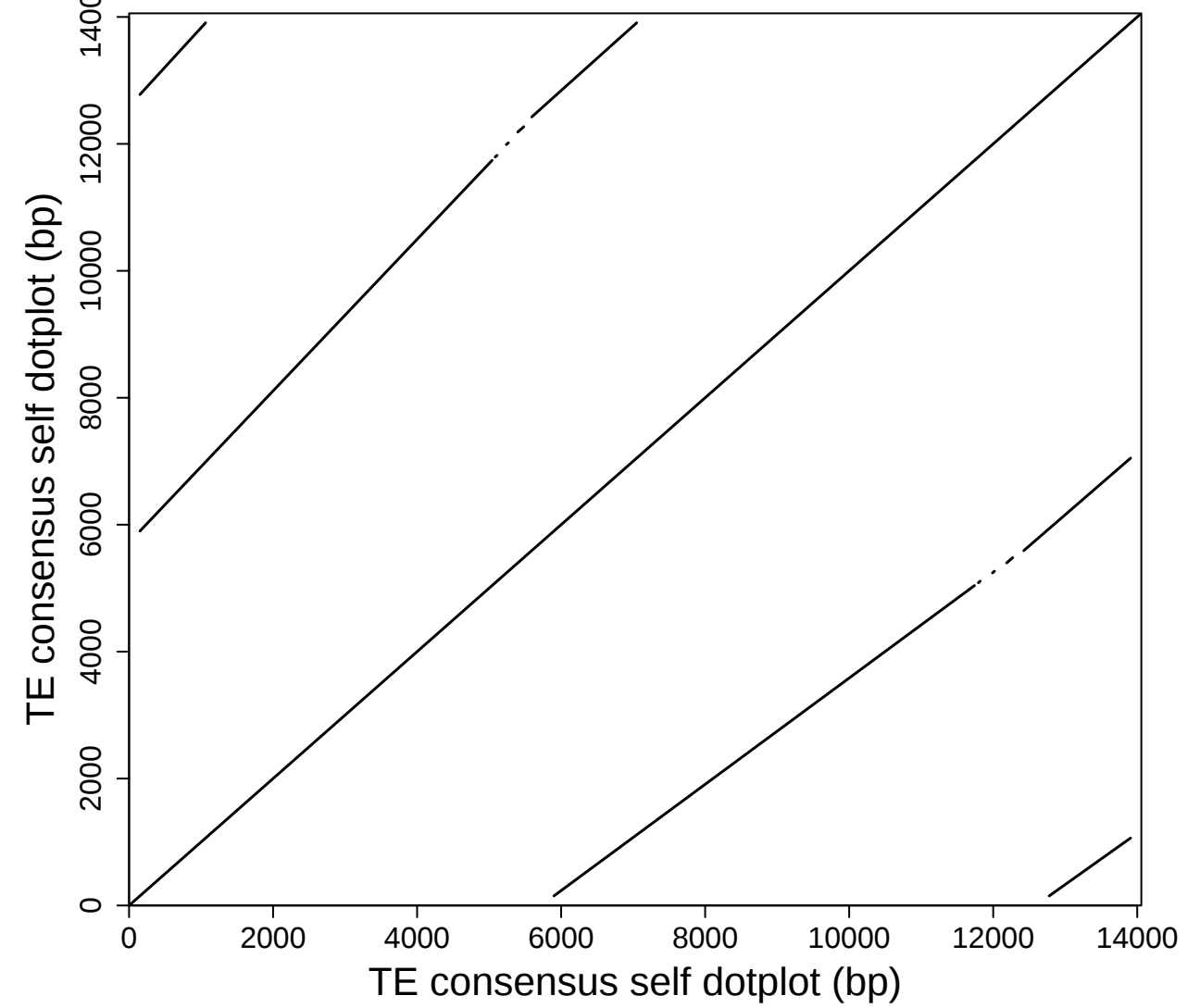
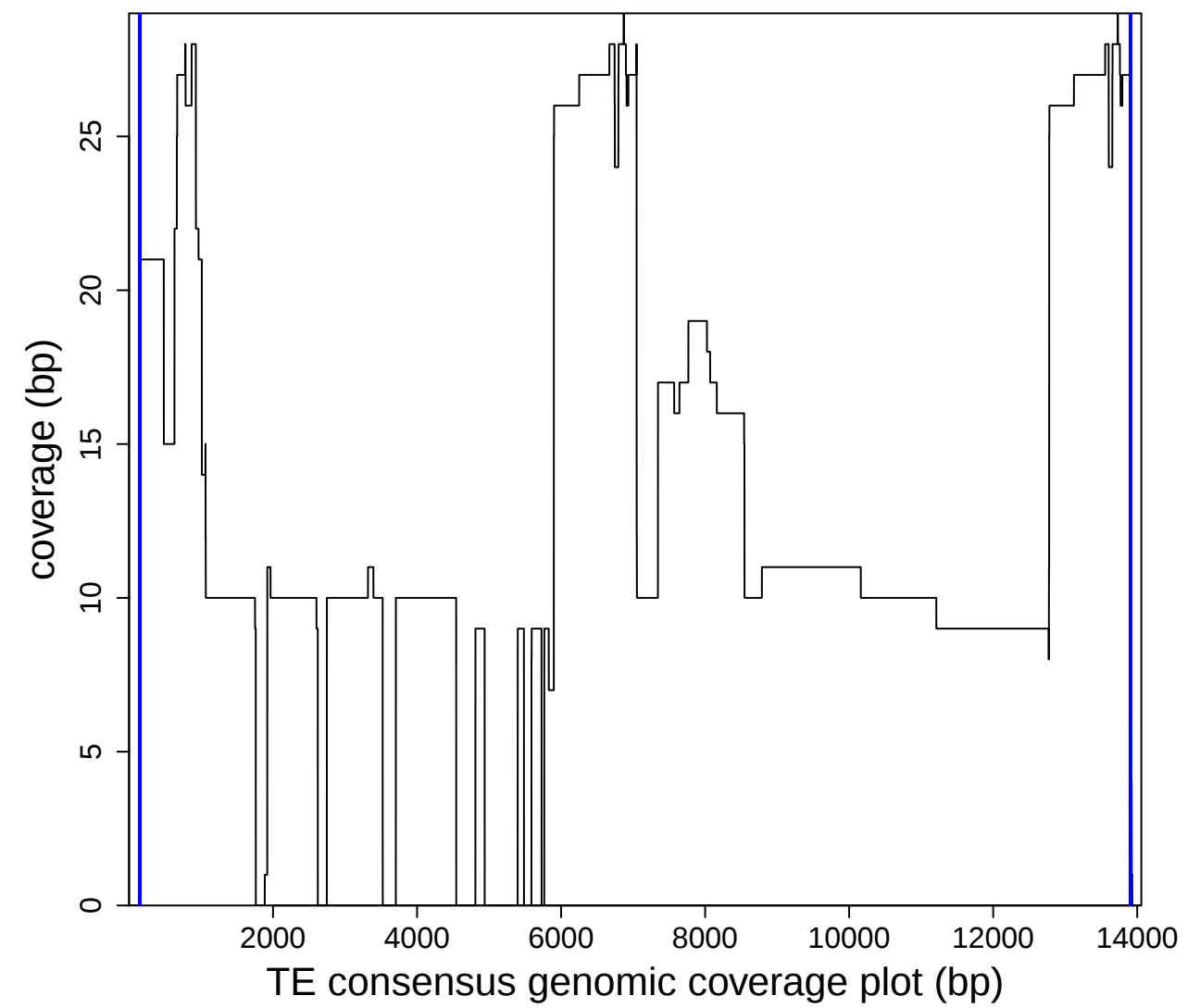
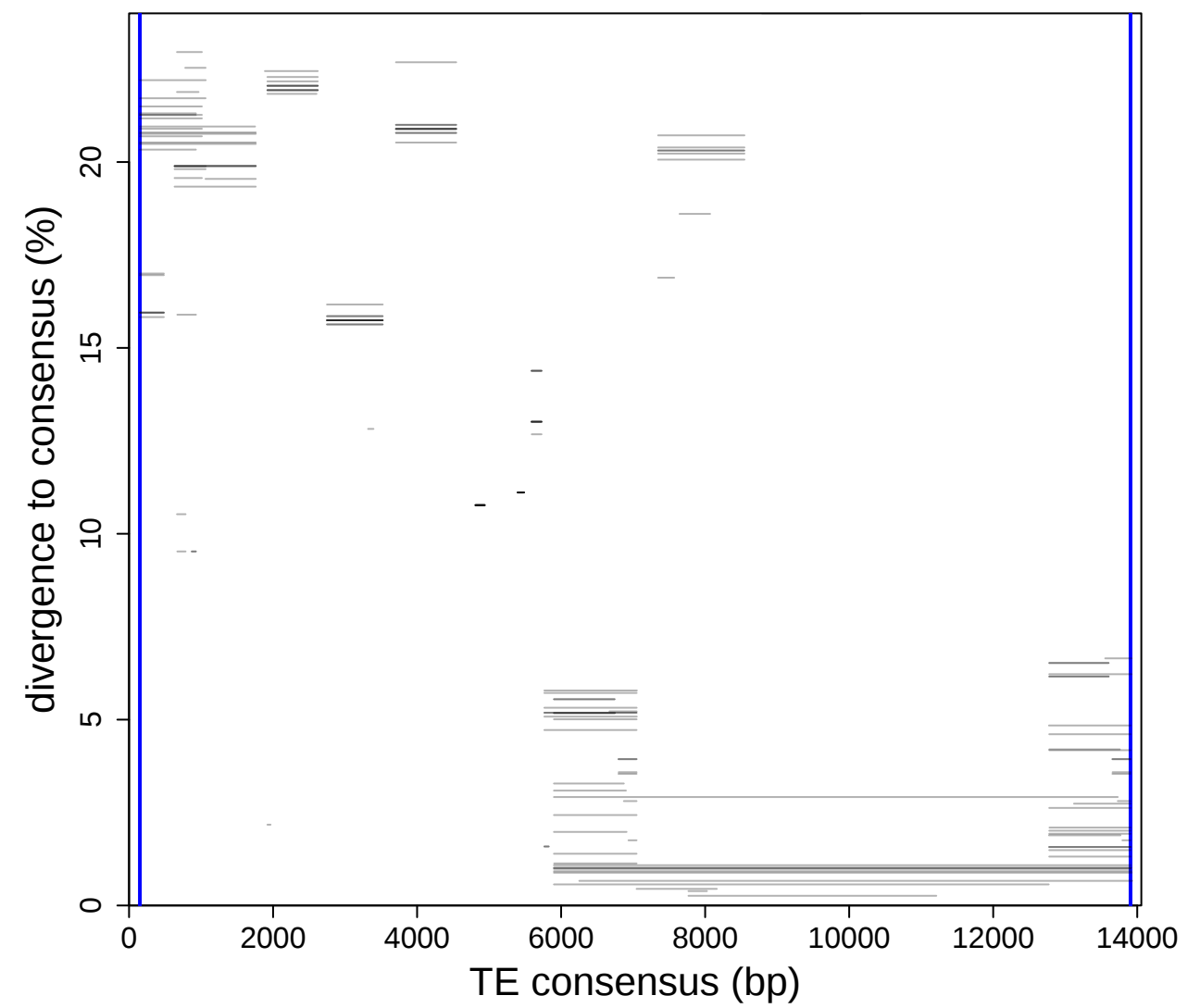
bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0_bcln.fa_aln.fa
size: 13361bp; fragments: 144; full length: 0 (>=12024.9bp)

After TEtrimmer 13361 bp



ed_uf.bed_g_1.bed_fm_1.bed_0_0_n.bed_g_1.bed_fm_2.bed_0_0
size: 14057bp; fragments: 170; full length: 0 (>=12651.3bp)

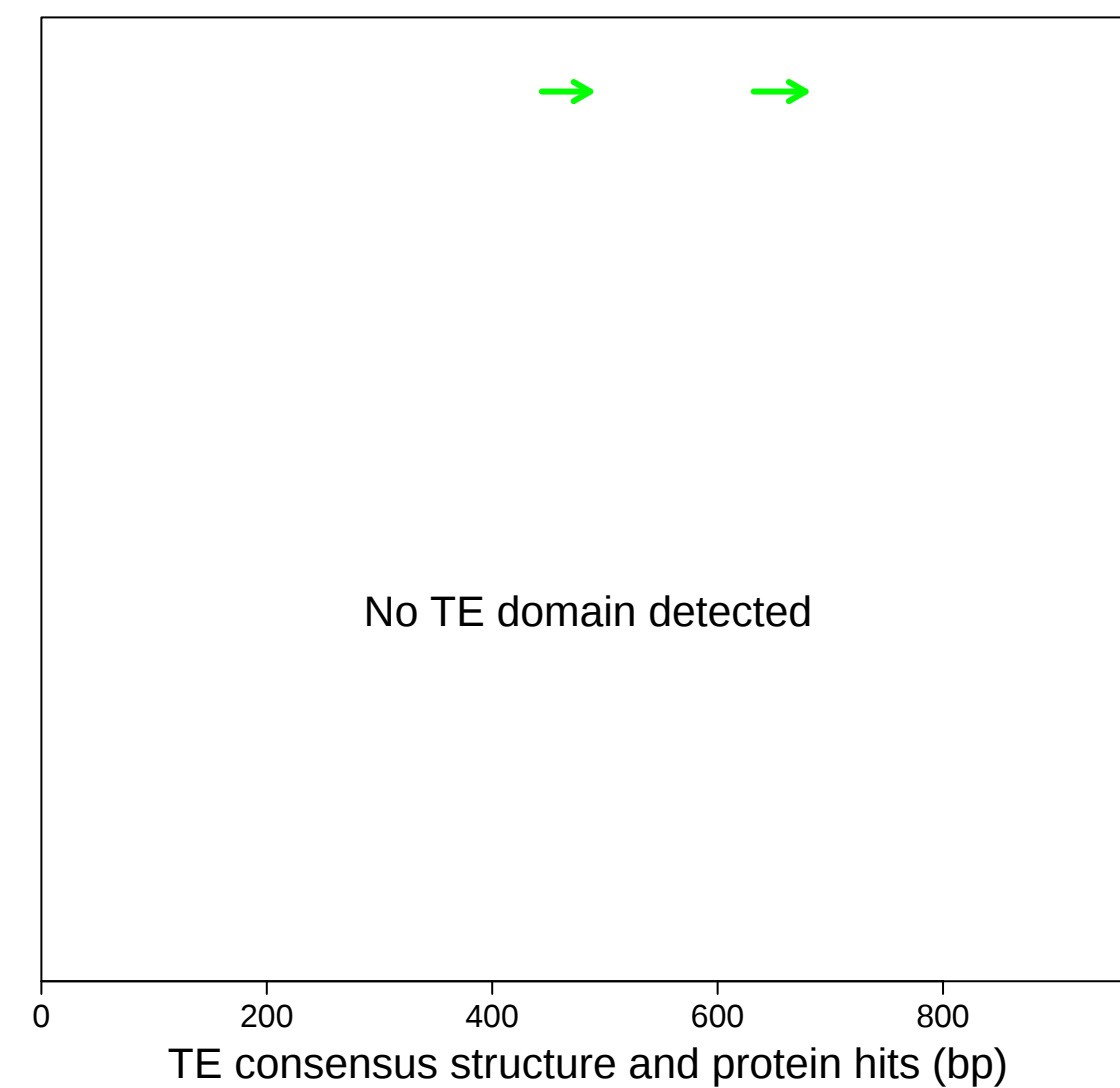
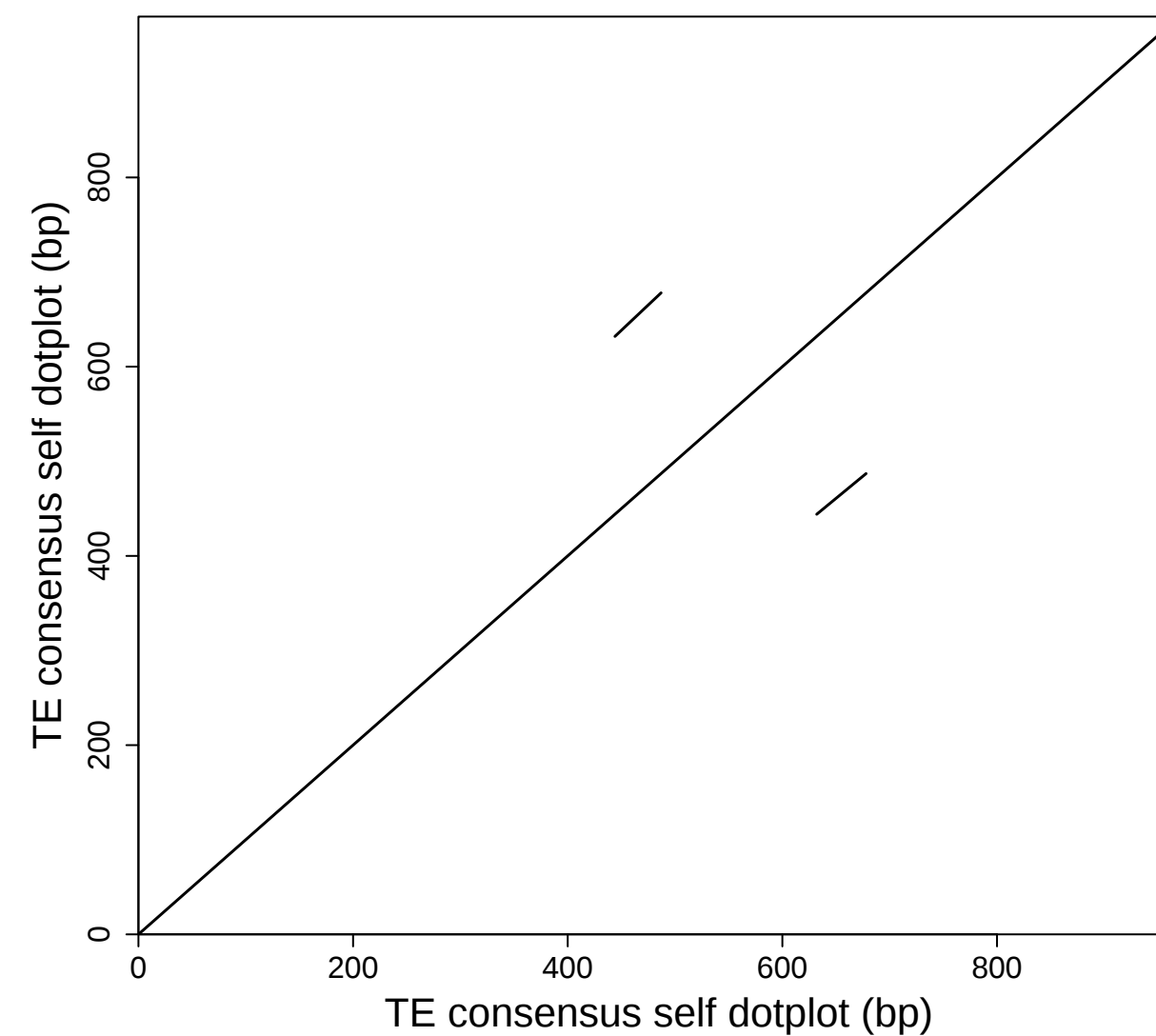
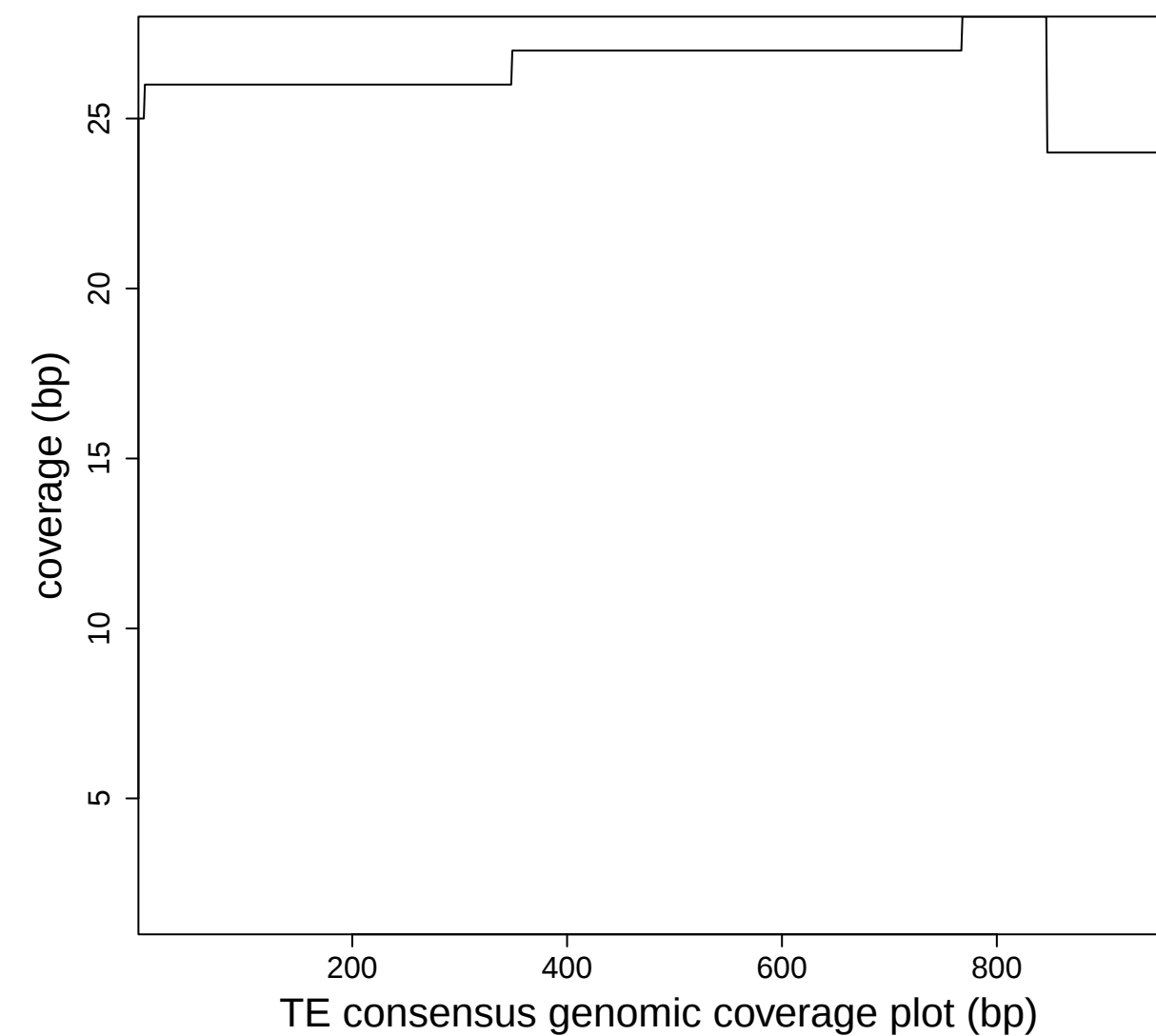
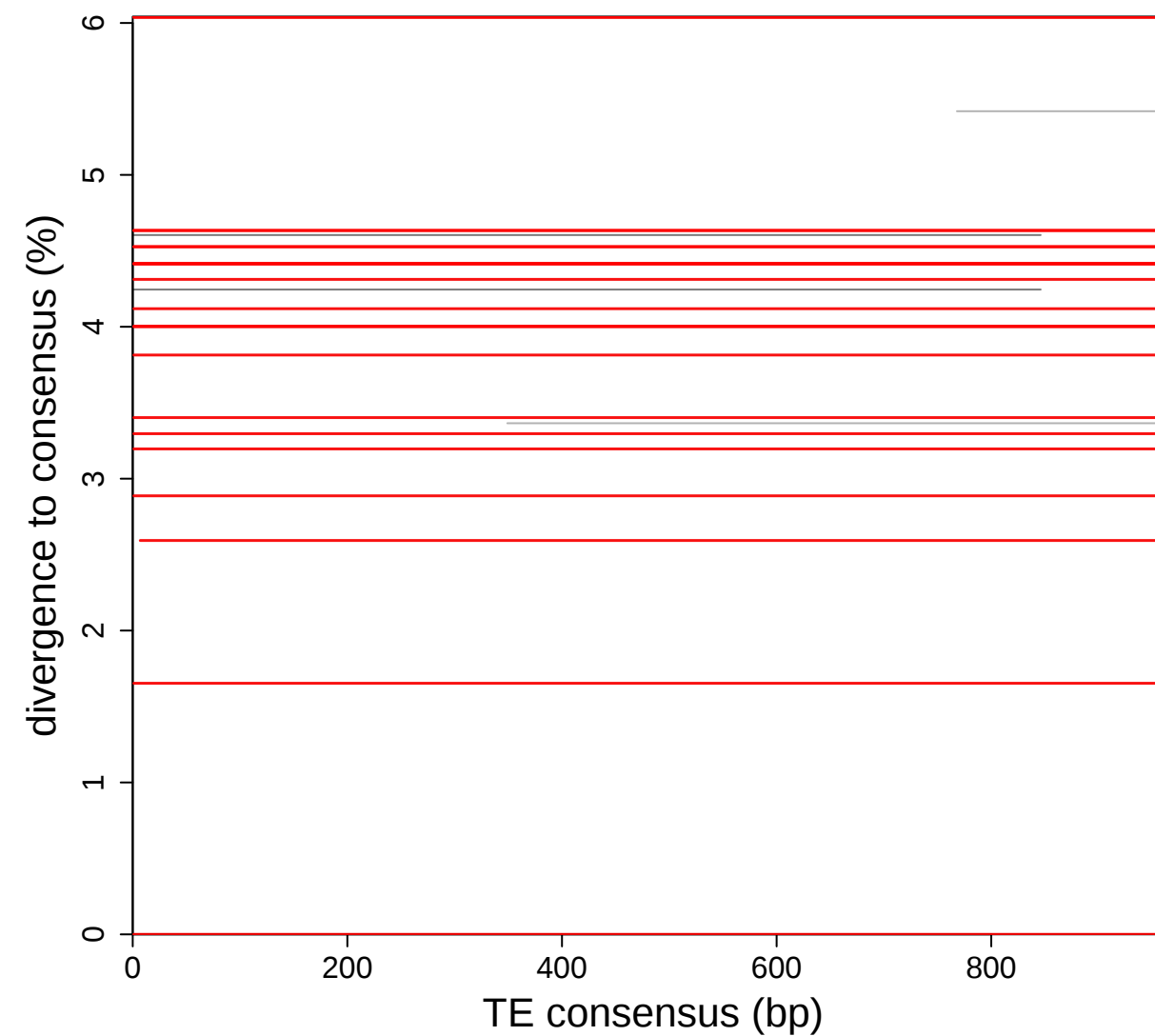
After TEtrimmer Extended plot Blue lines are boundaries



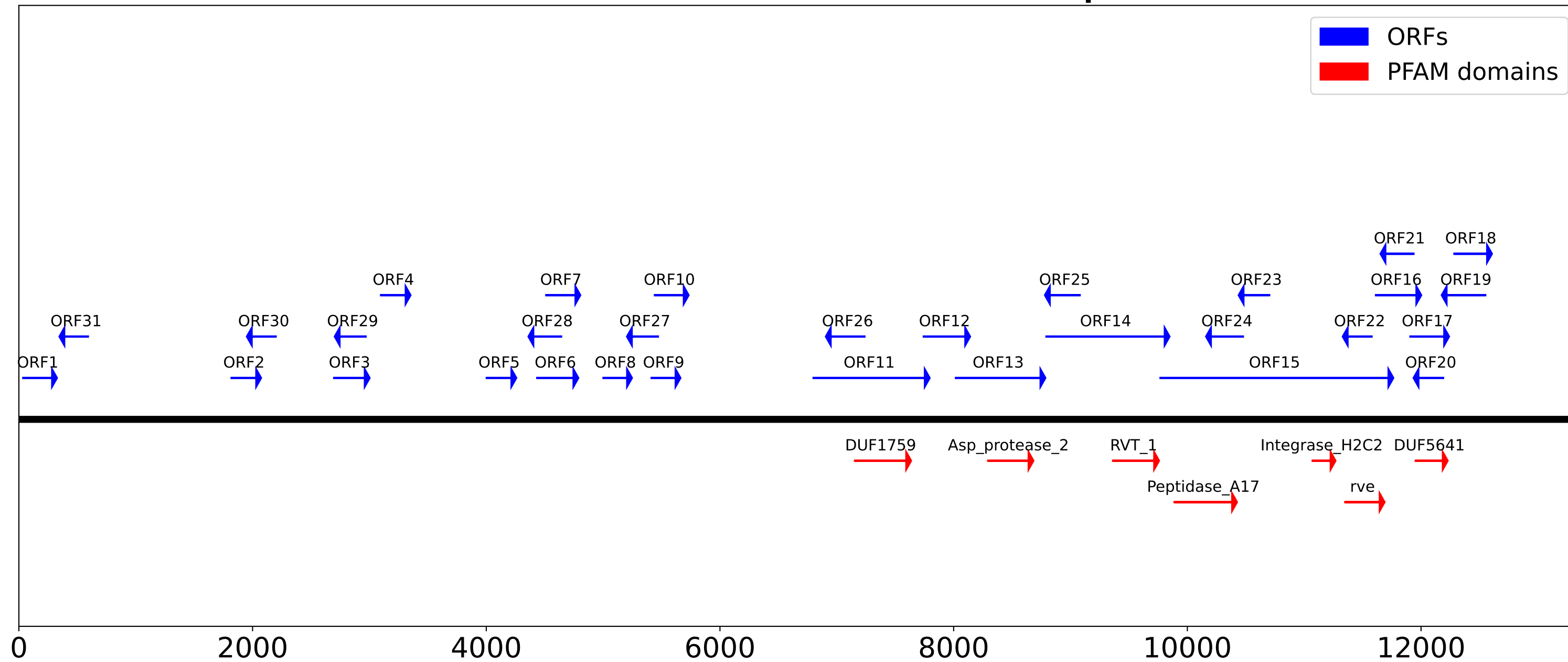
Before TEtrimmer 970 bp

TE: ltr_1_family_378

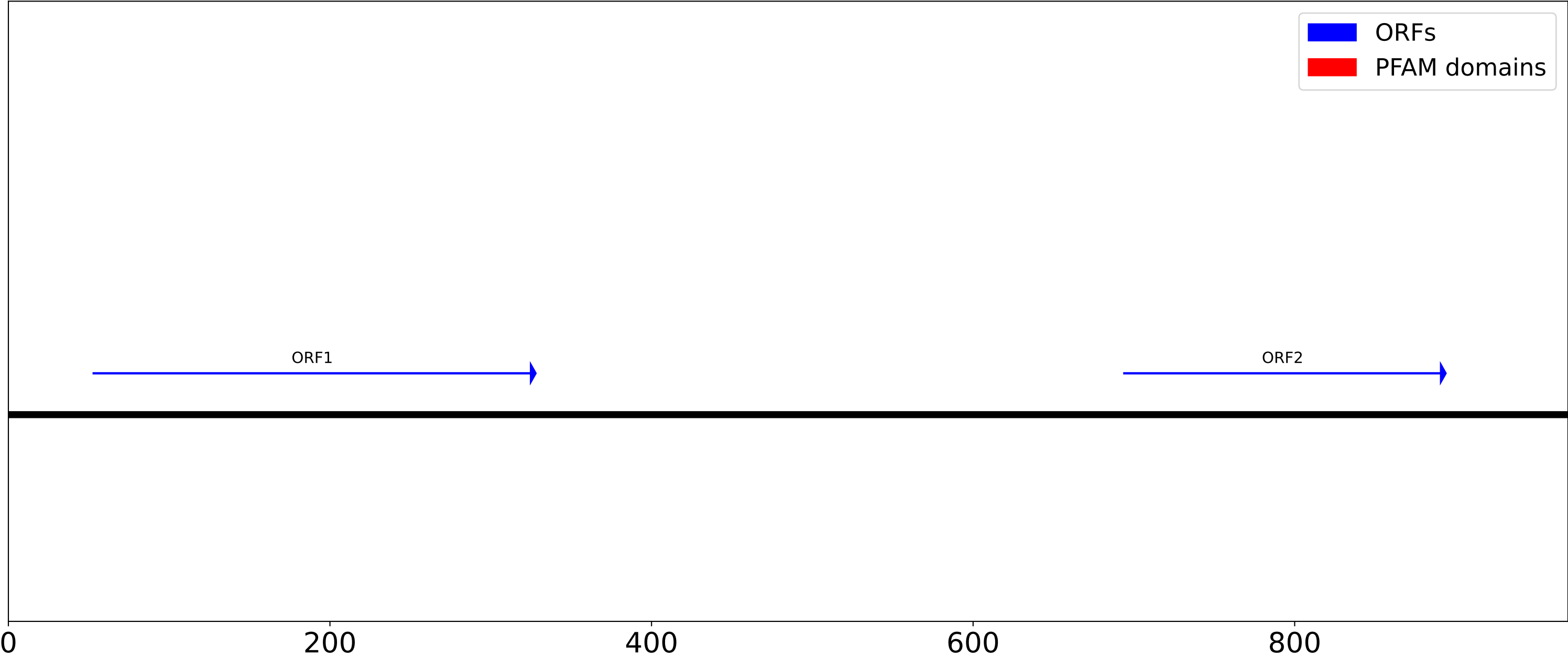
size: 970bp; fragments: 28; full length: 22 (≥ 873 bp)



After TEtrimmer ORF and PFAM domain plot



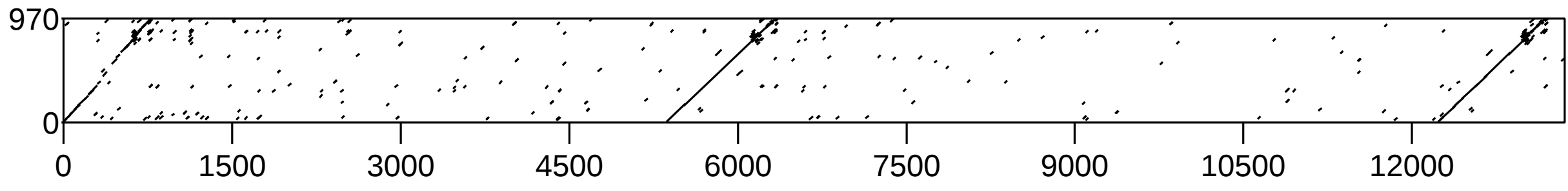
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)