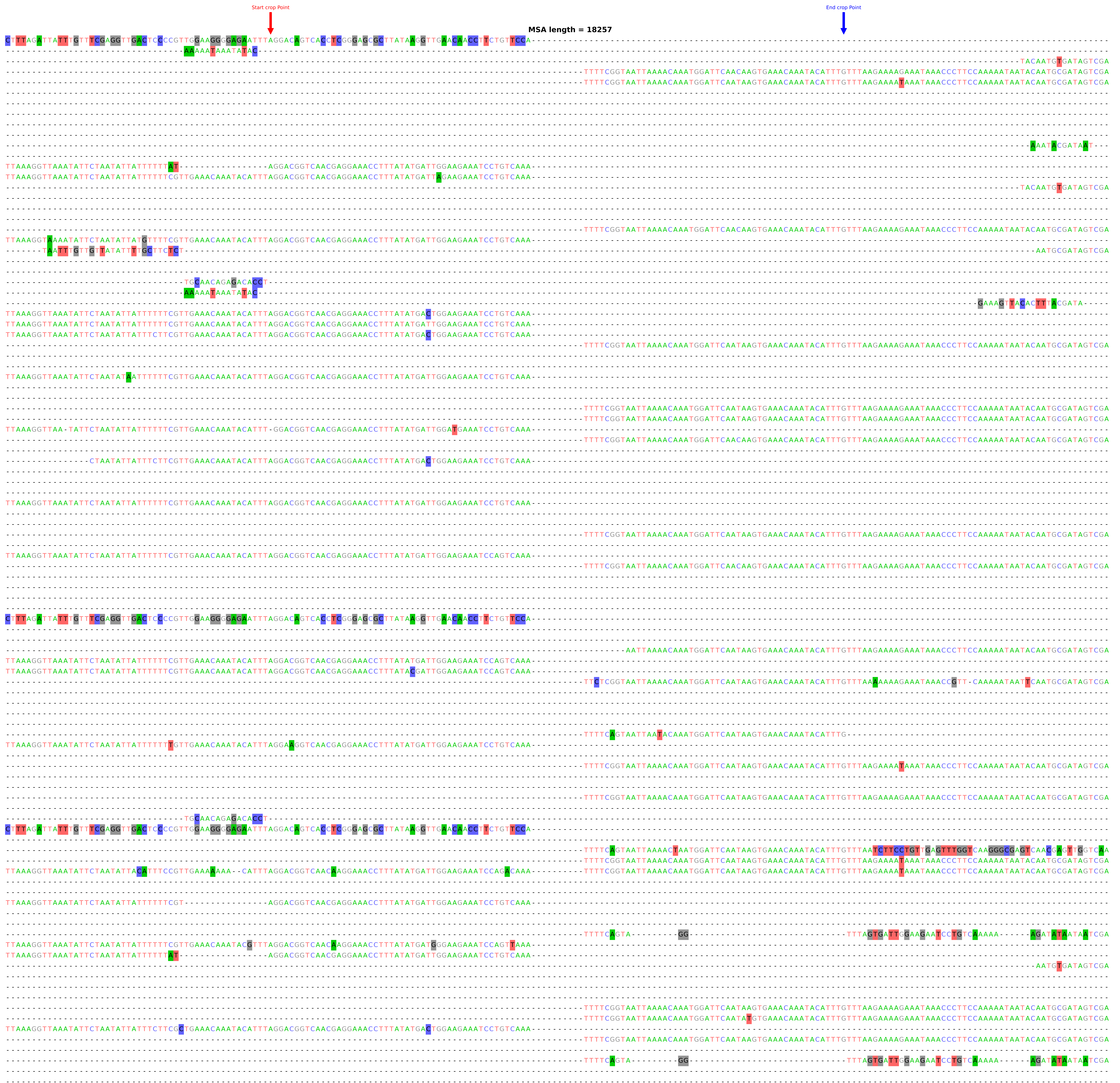
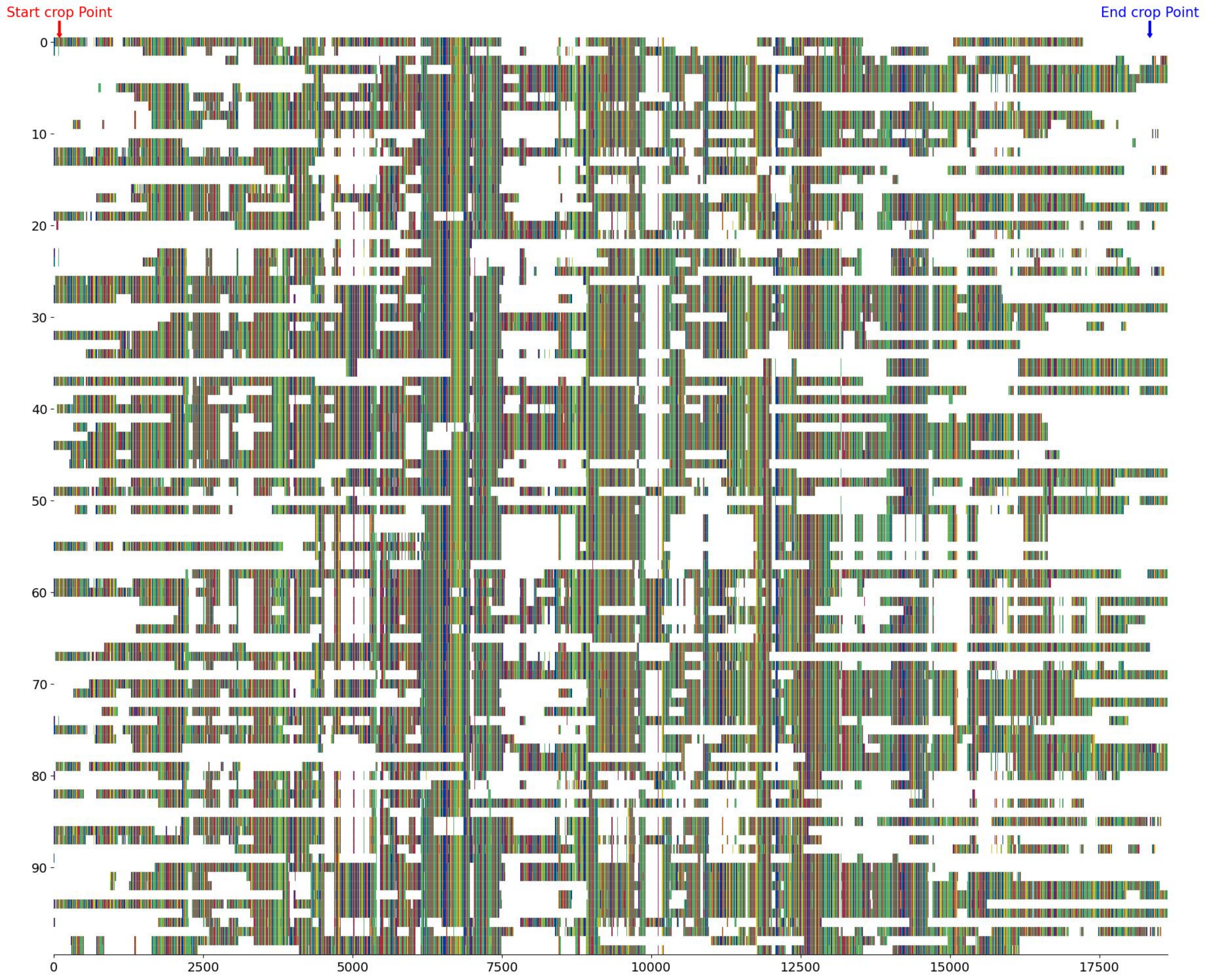
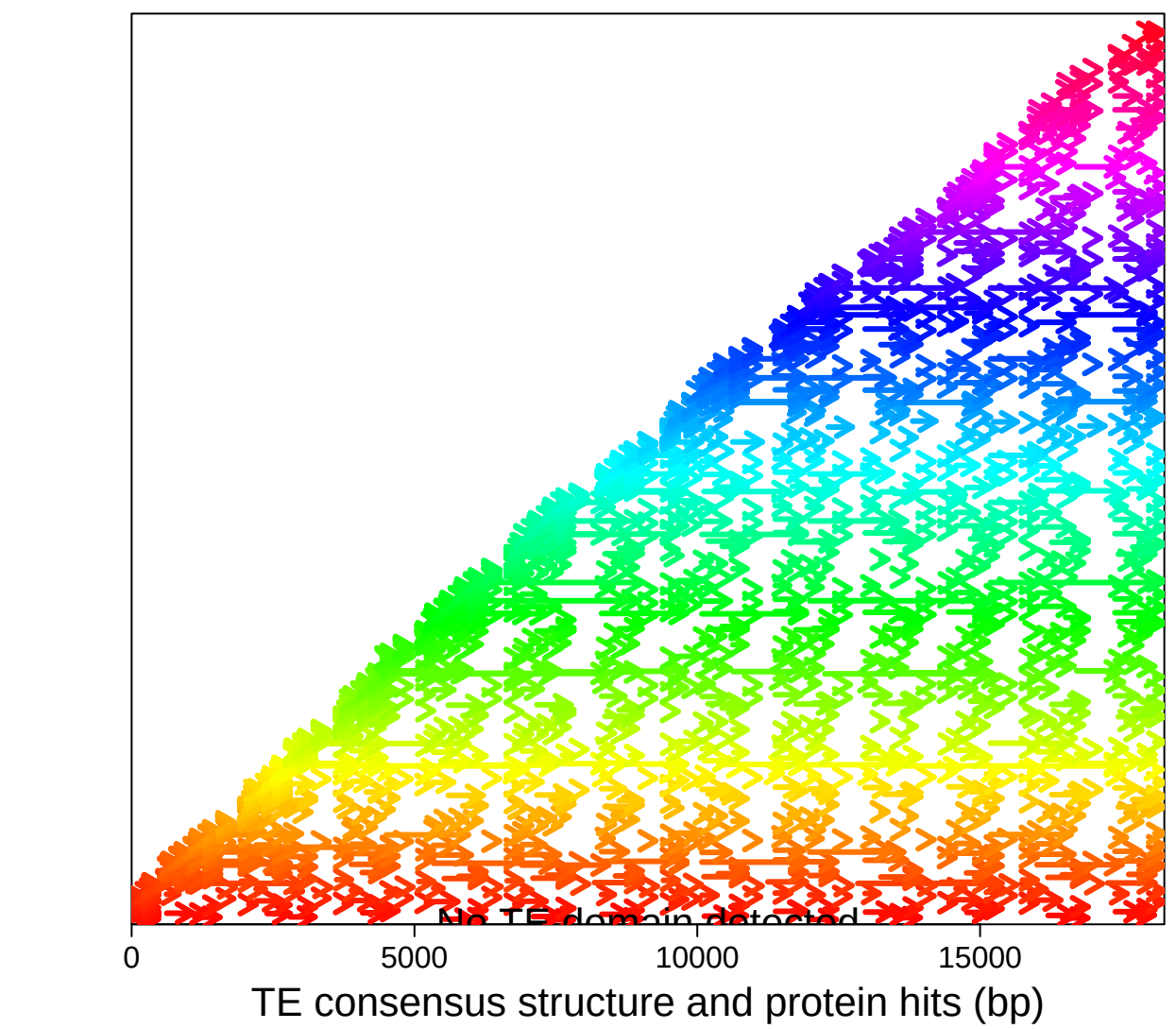
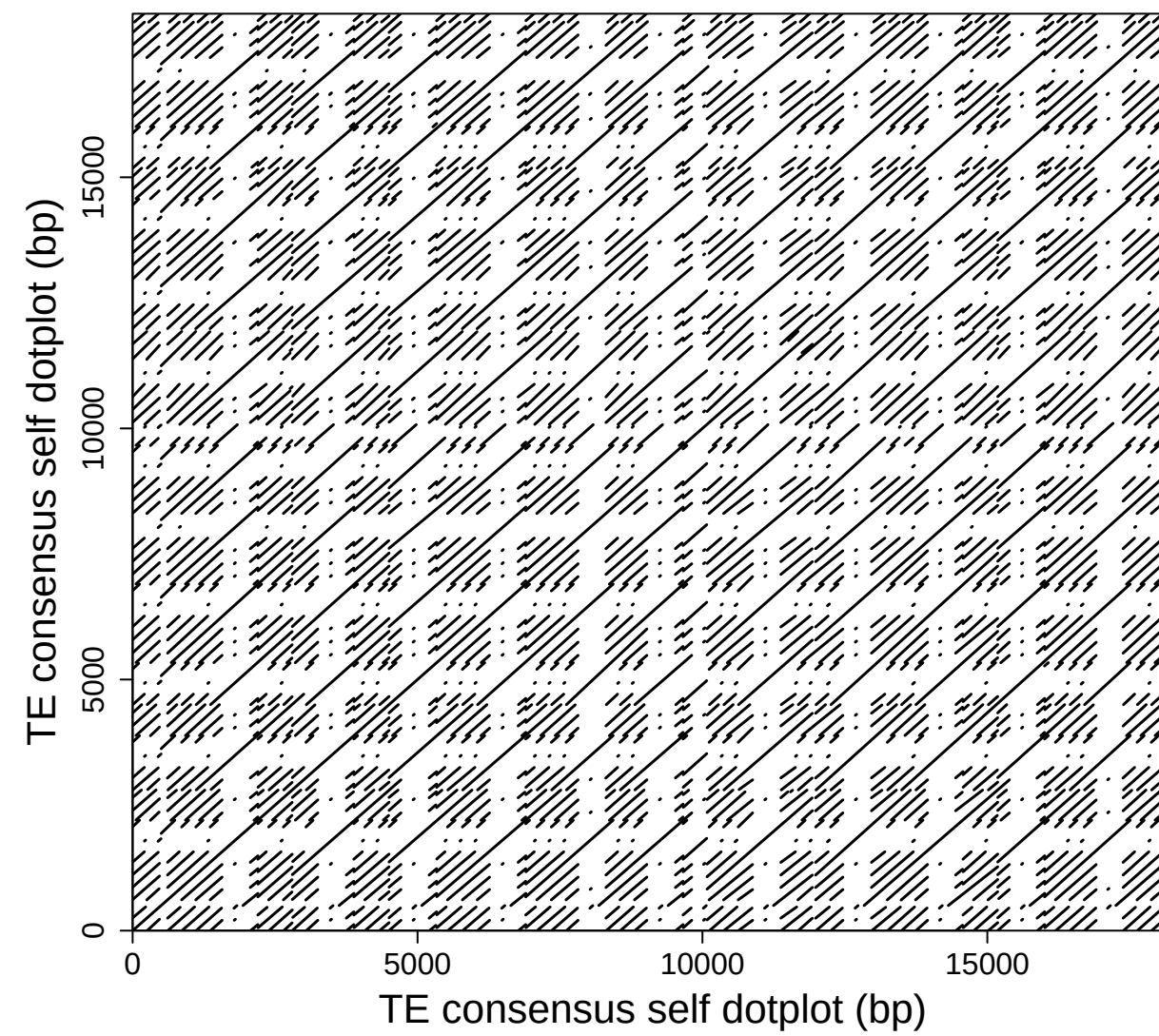
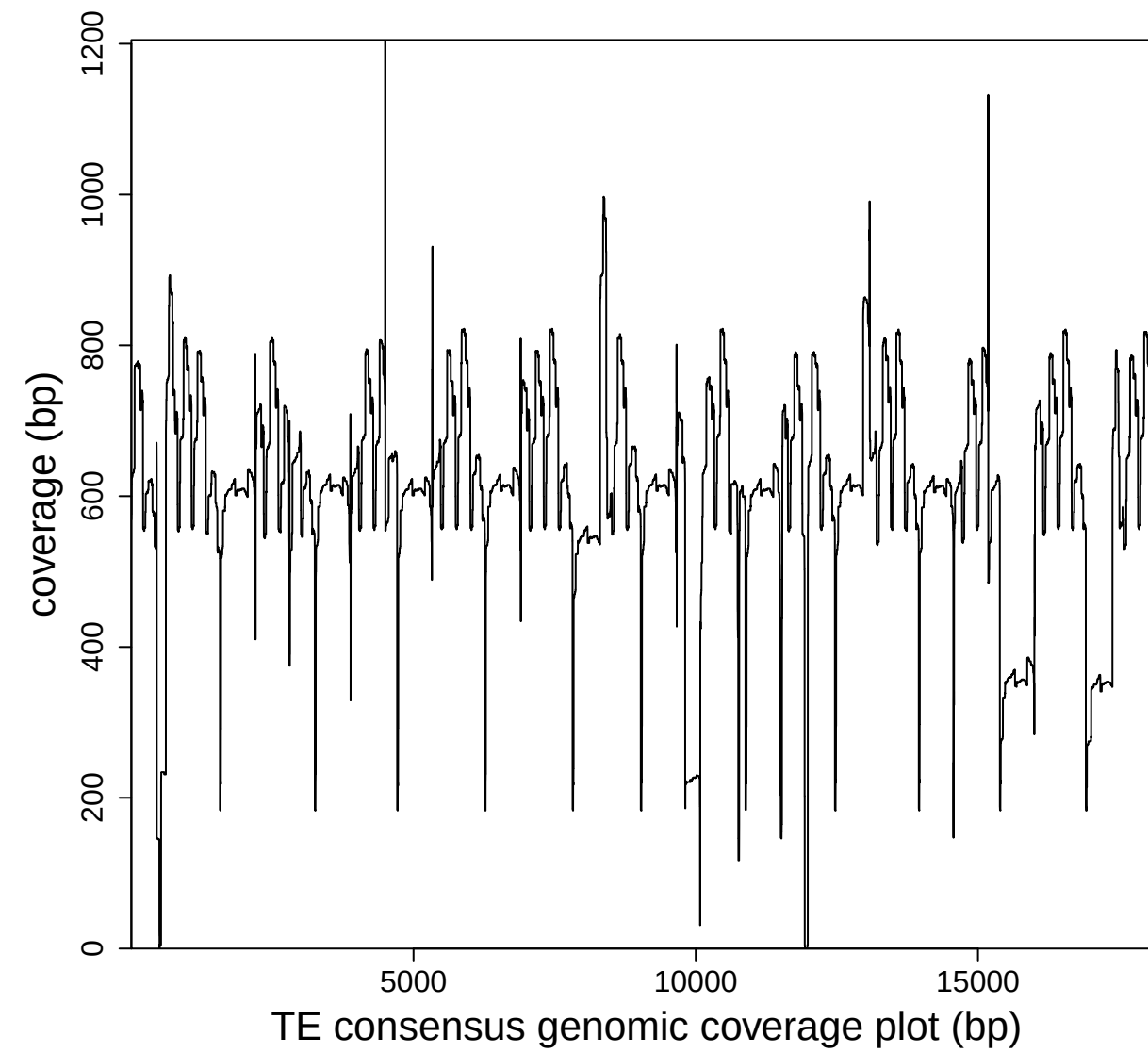
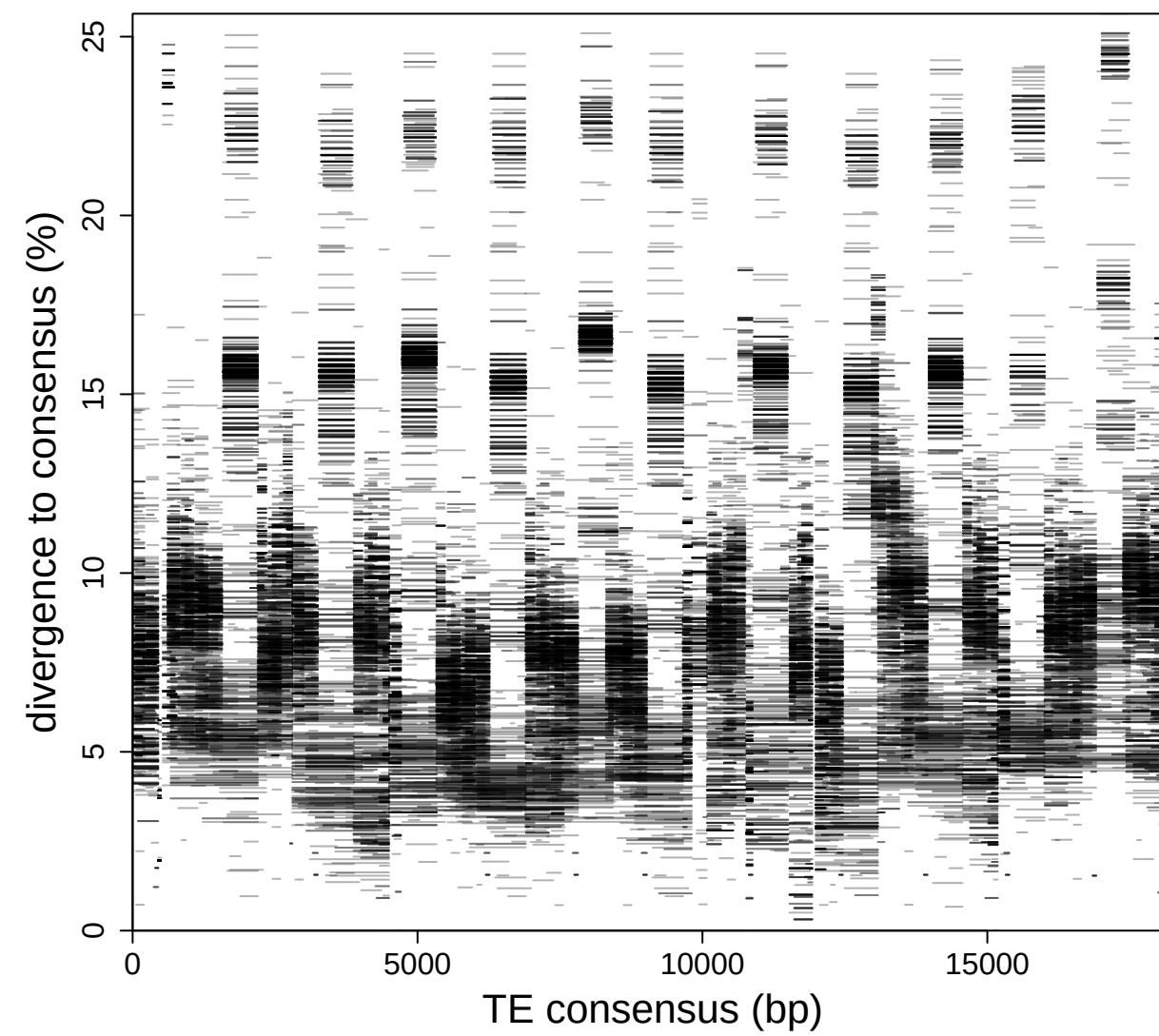






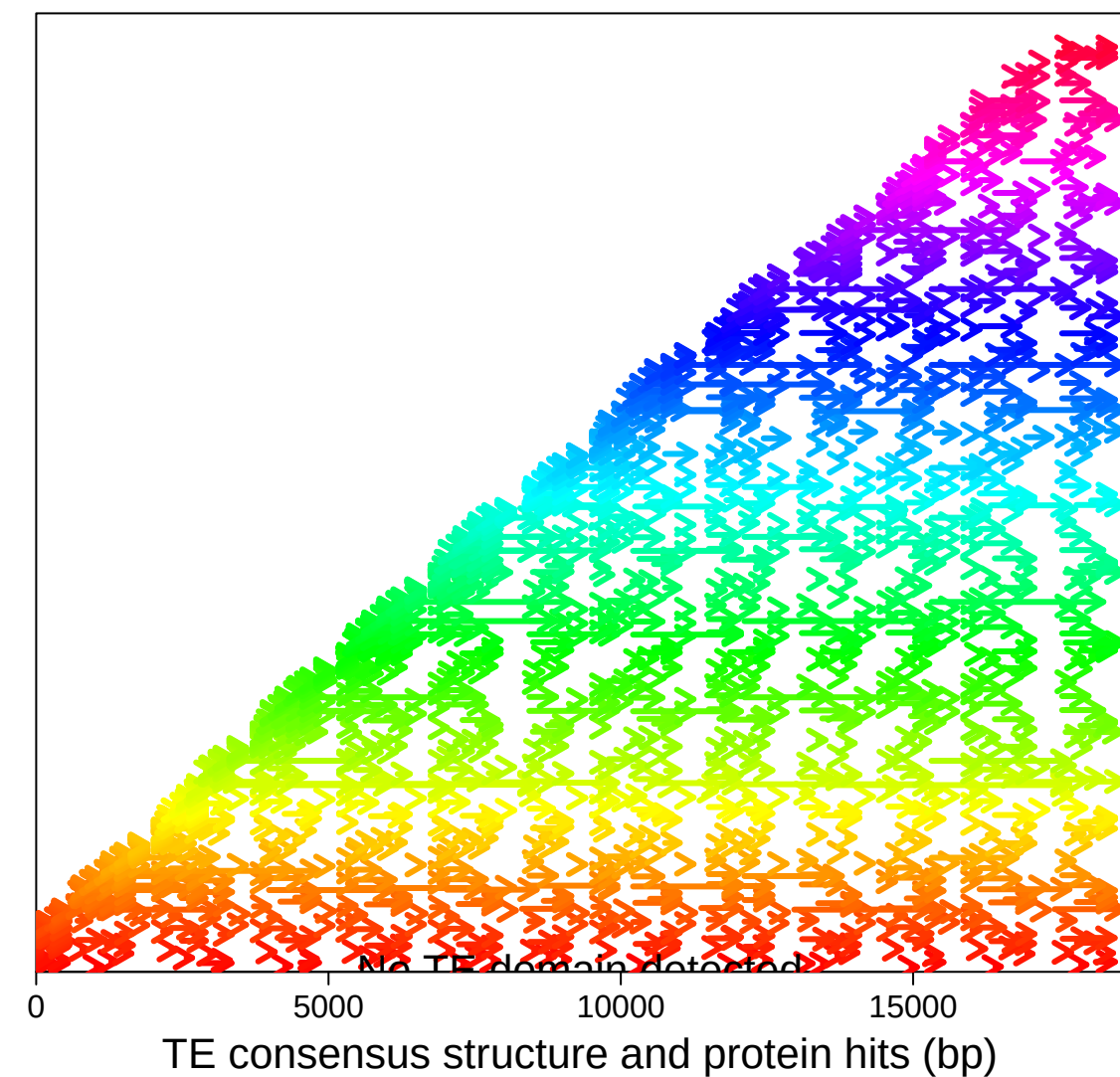
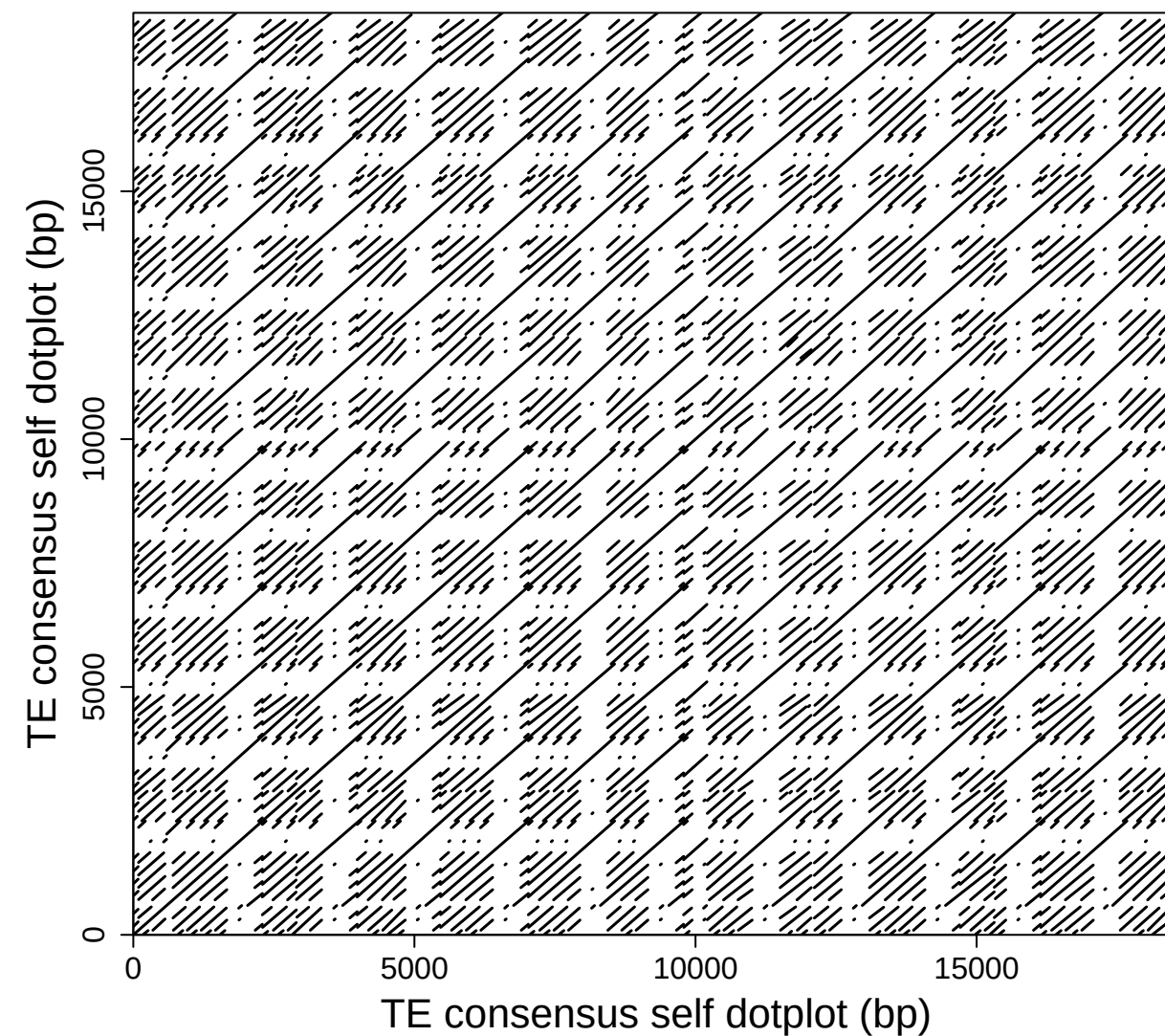
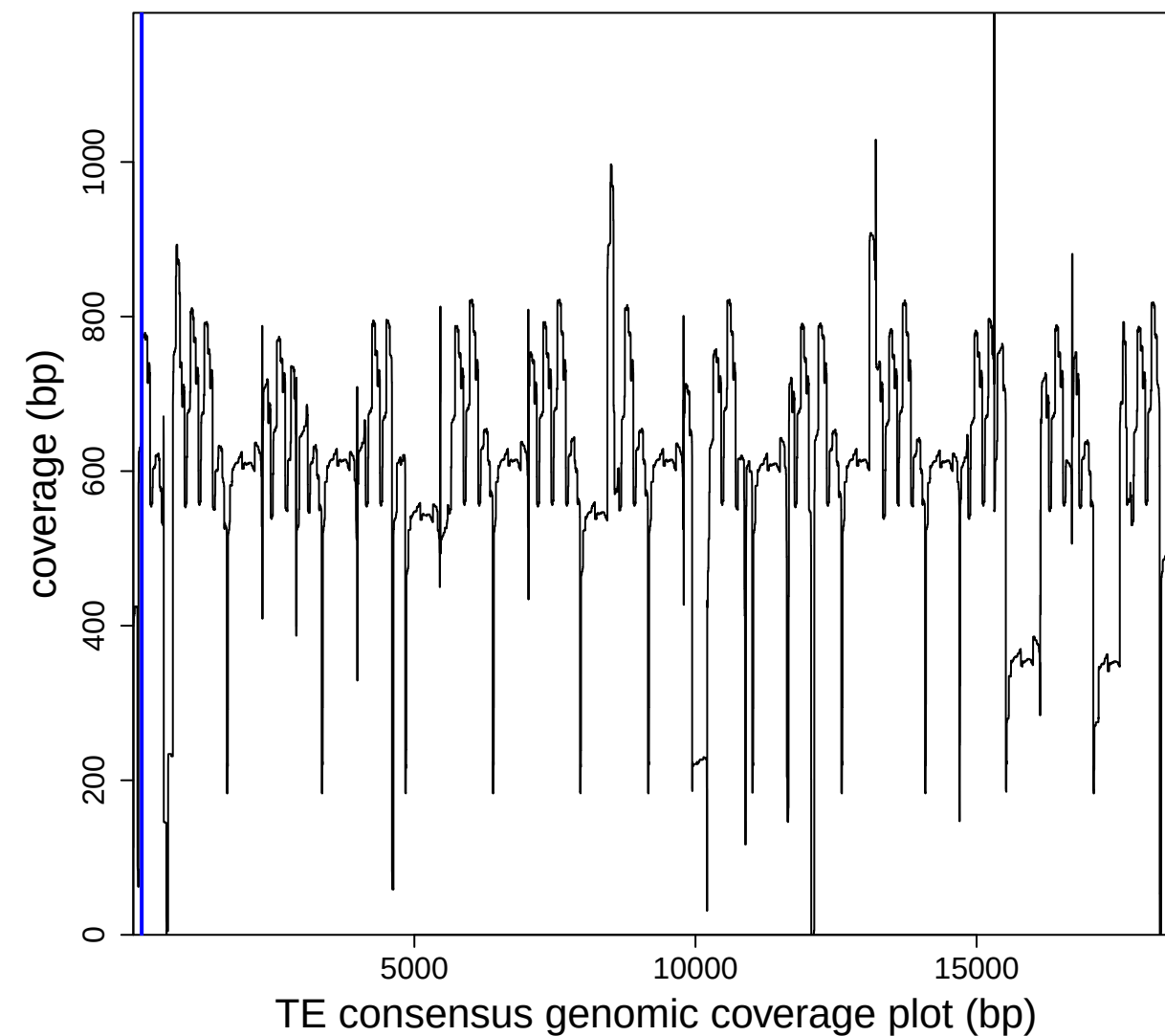
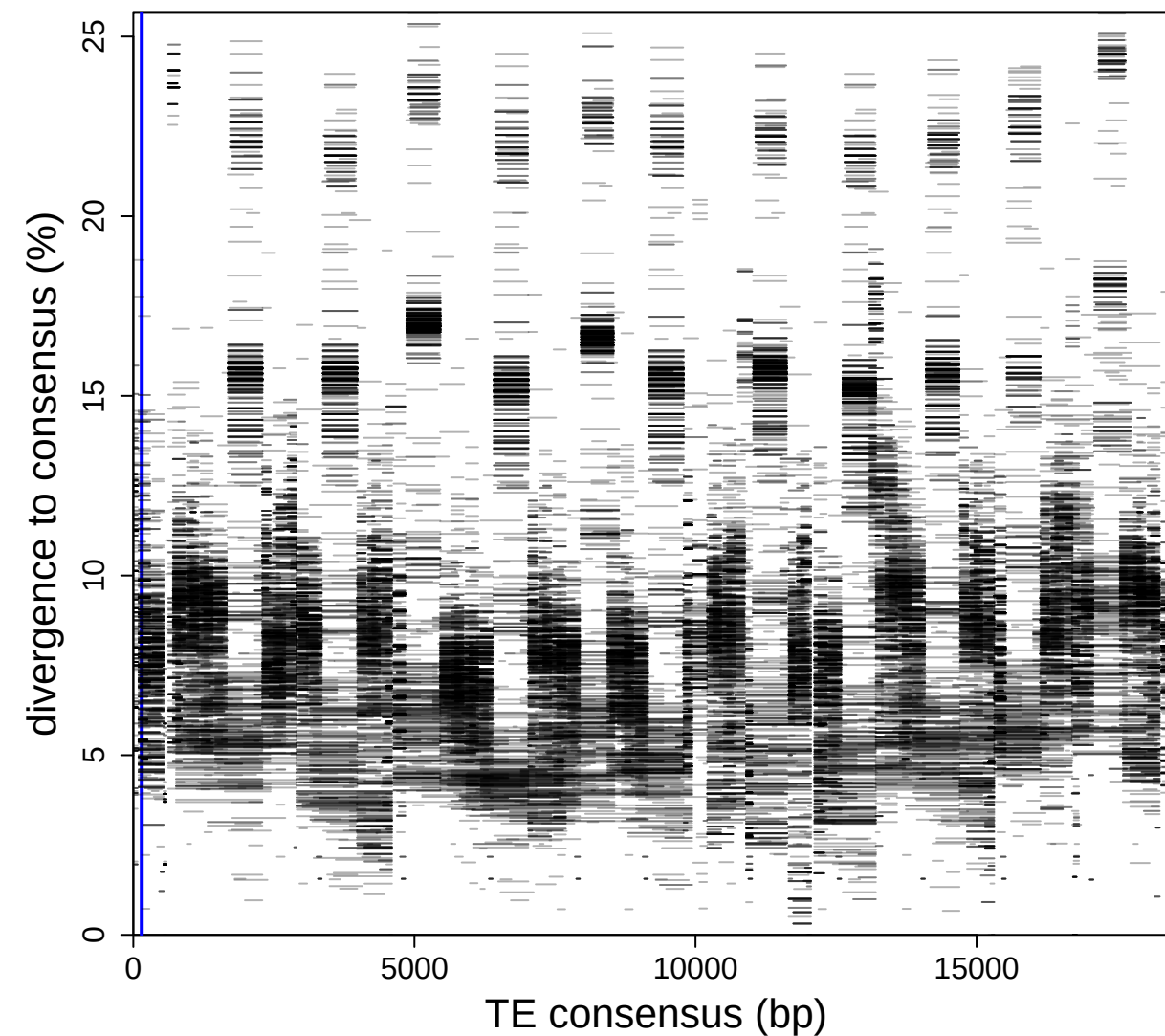
bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa c
size: 18257bp; fragments: 24806; full length: 0 (>=16431.3bp)

After TEtrimmer 18257 bp



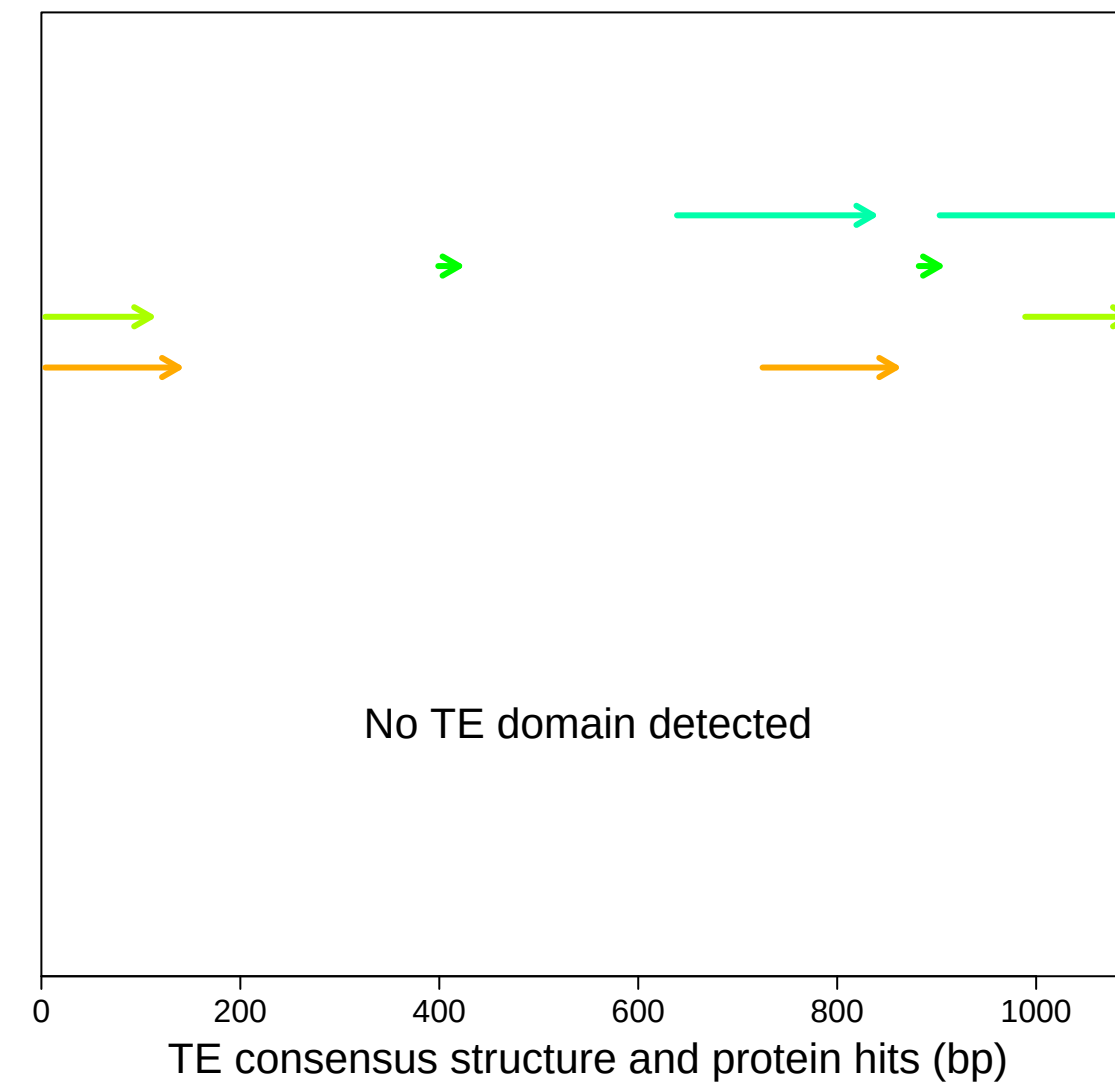
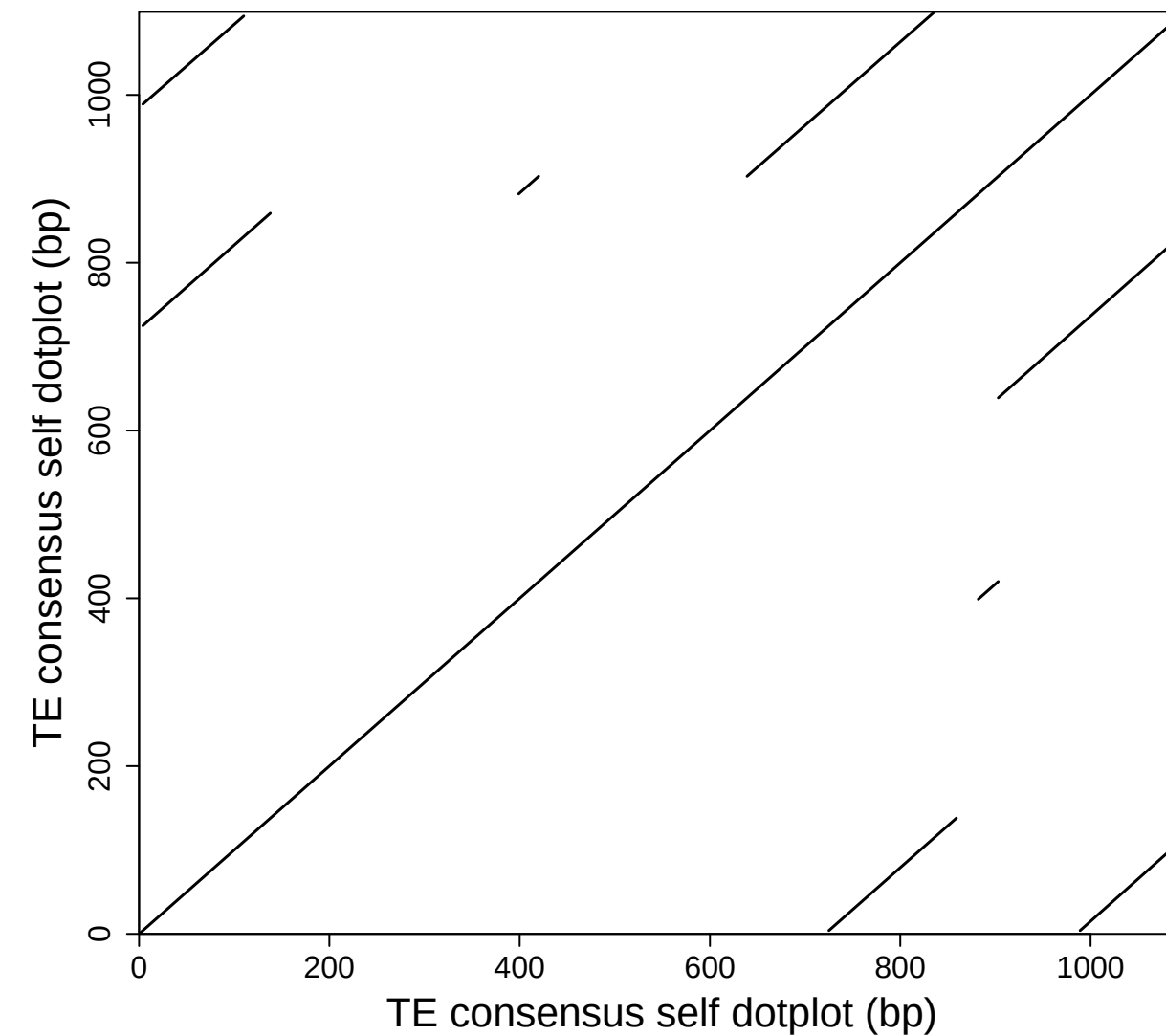
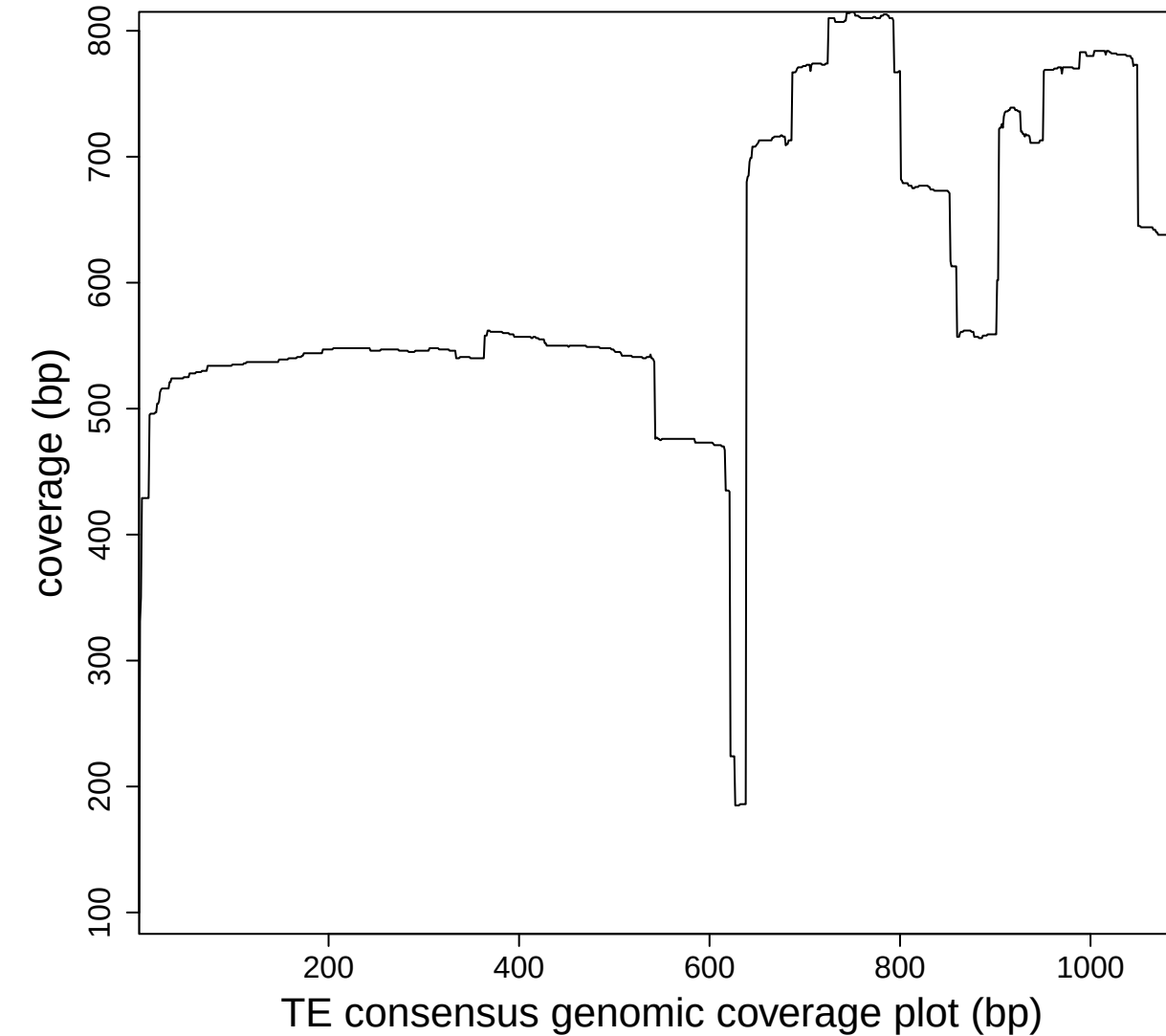
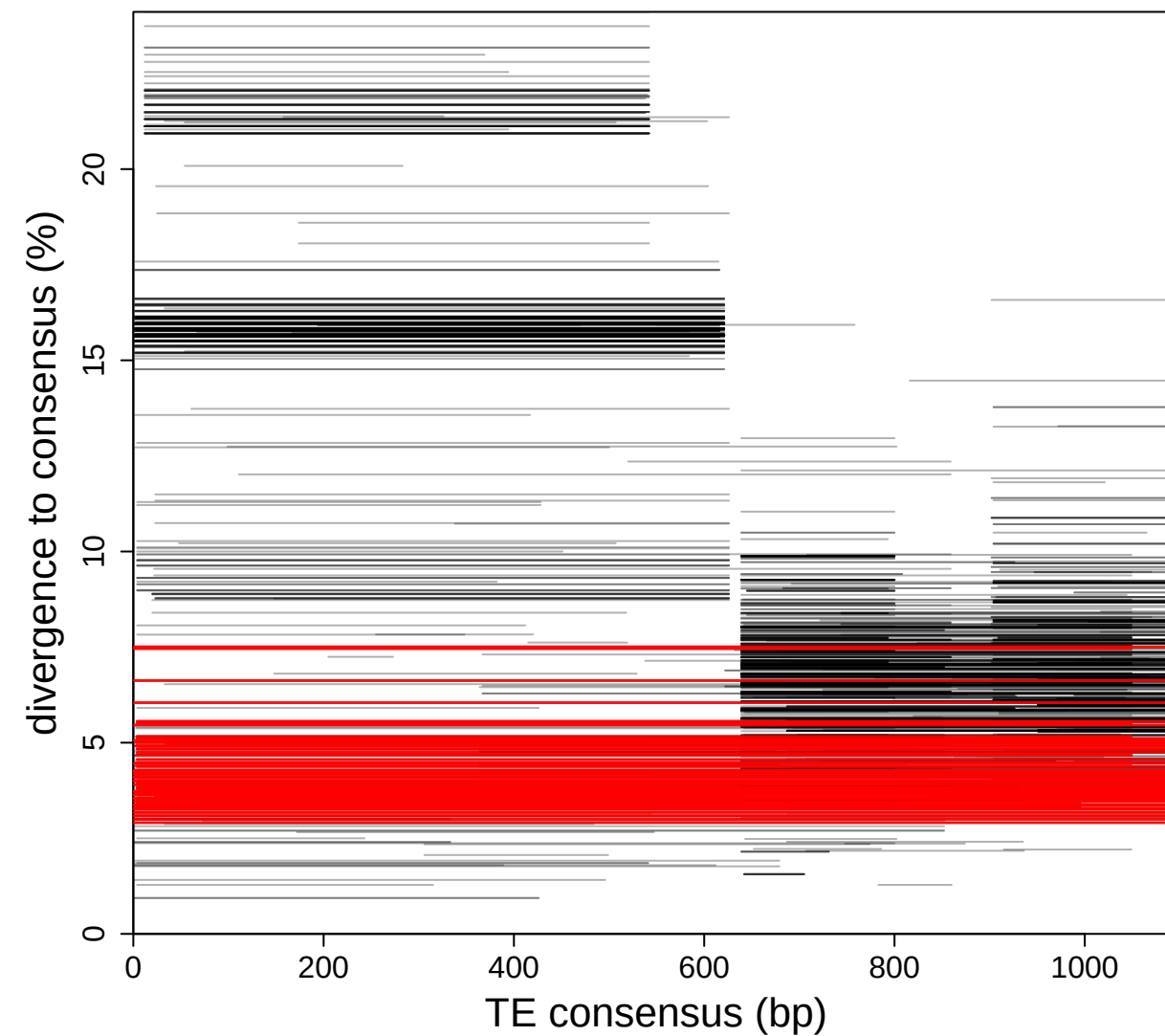
L0158.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 18600bp; fragments: 25502; full length: 0 (>=16740bp)

After TEtrimmer Extended plot Blue lines are boundaries

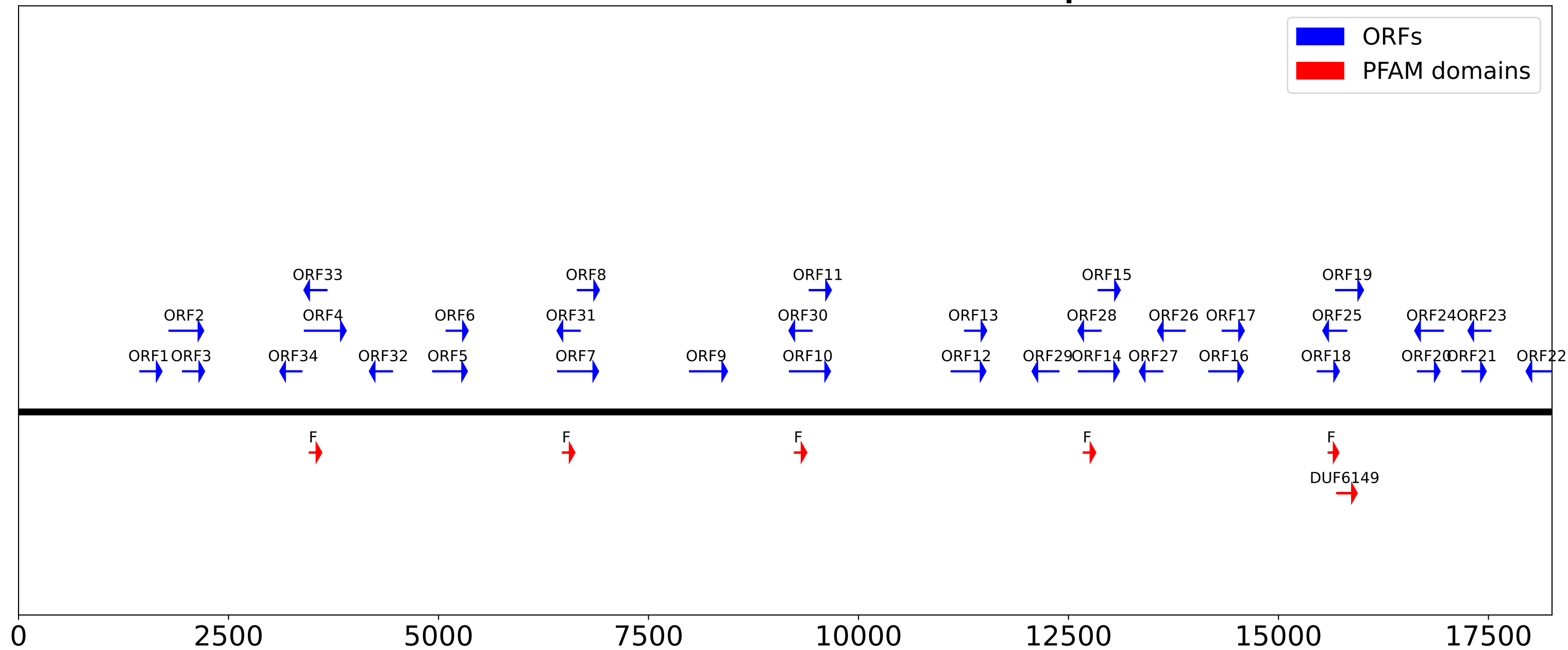


TE: rnd_5_family_10158
size: 1099bp; fragments: 1524; full length: 98 (≥ 989.1 bp)

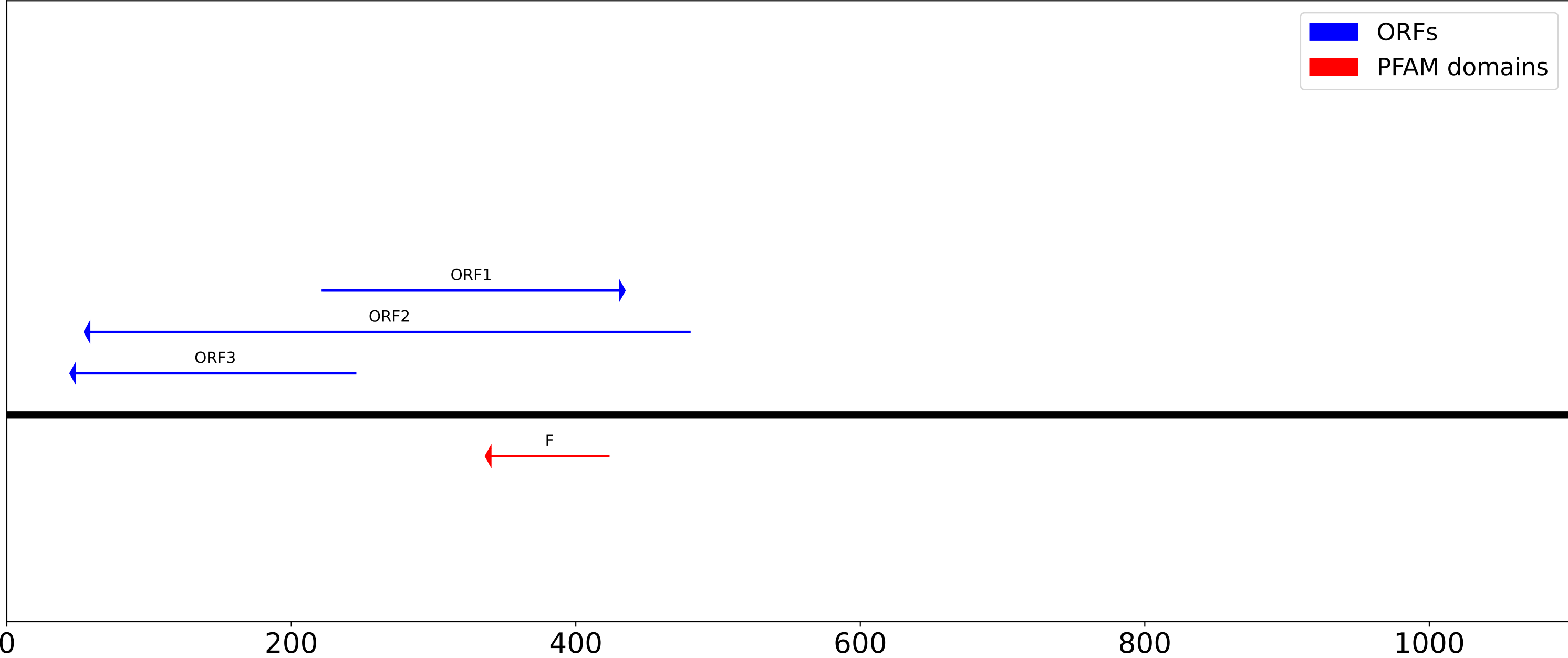
Before TEtrimmer 1099 bp



After TETrimmer ORF and PFAM domain plot



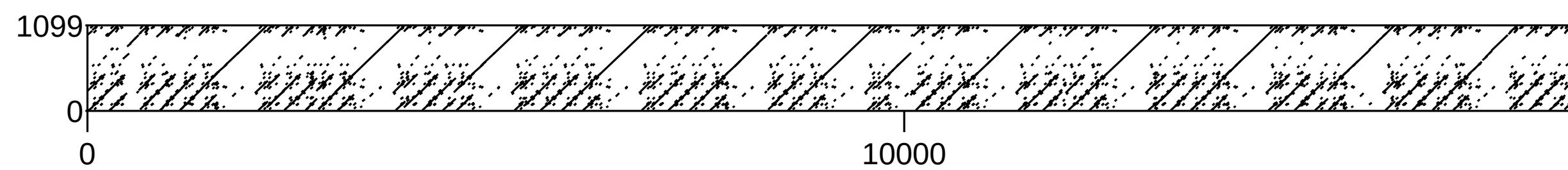
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)