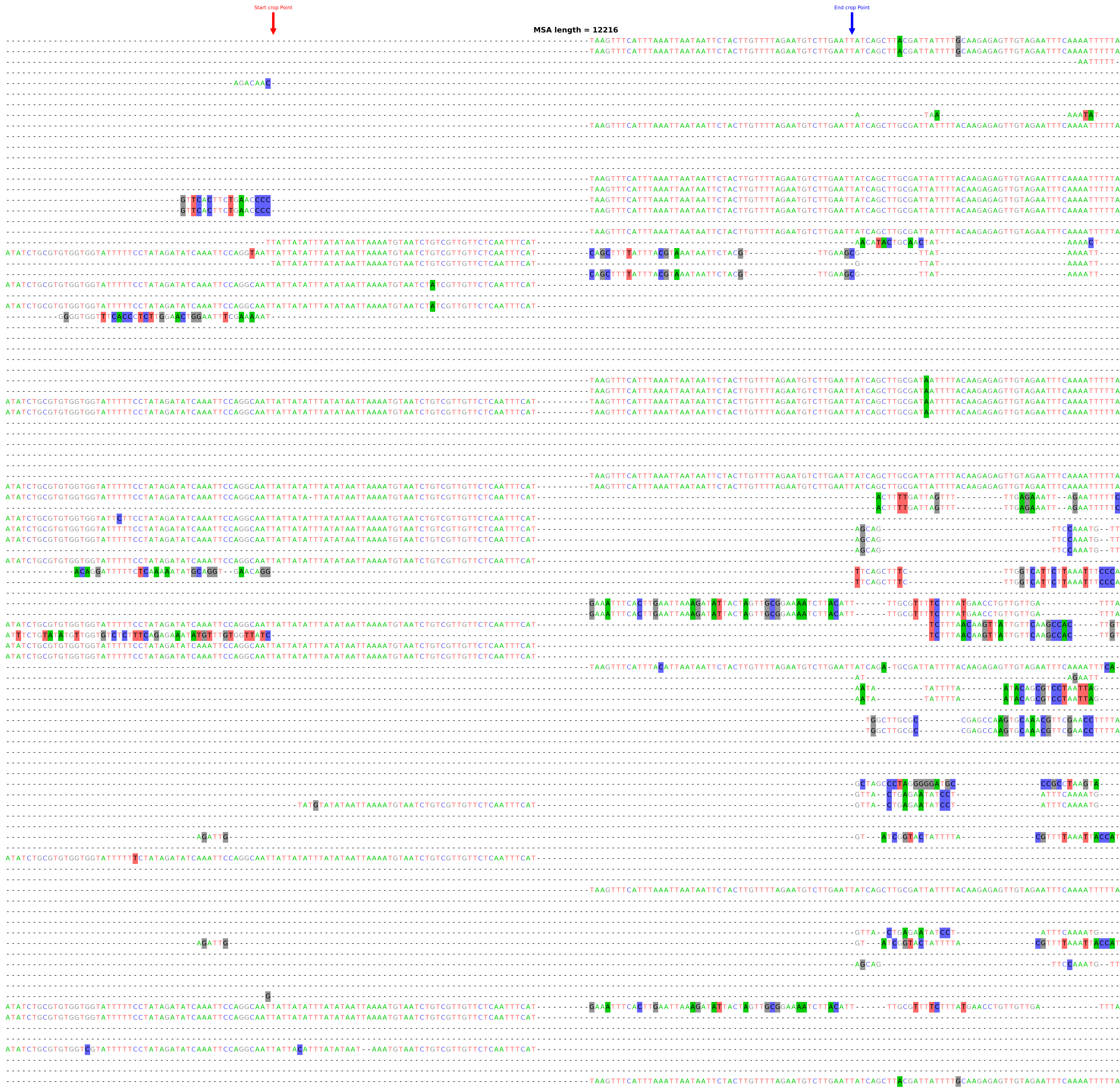
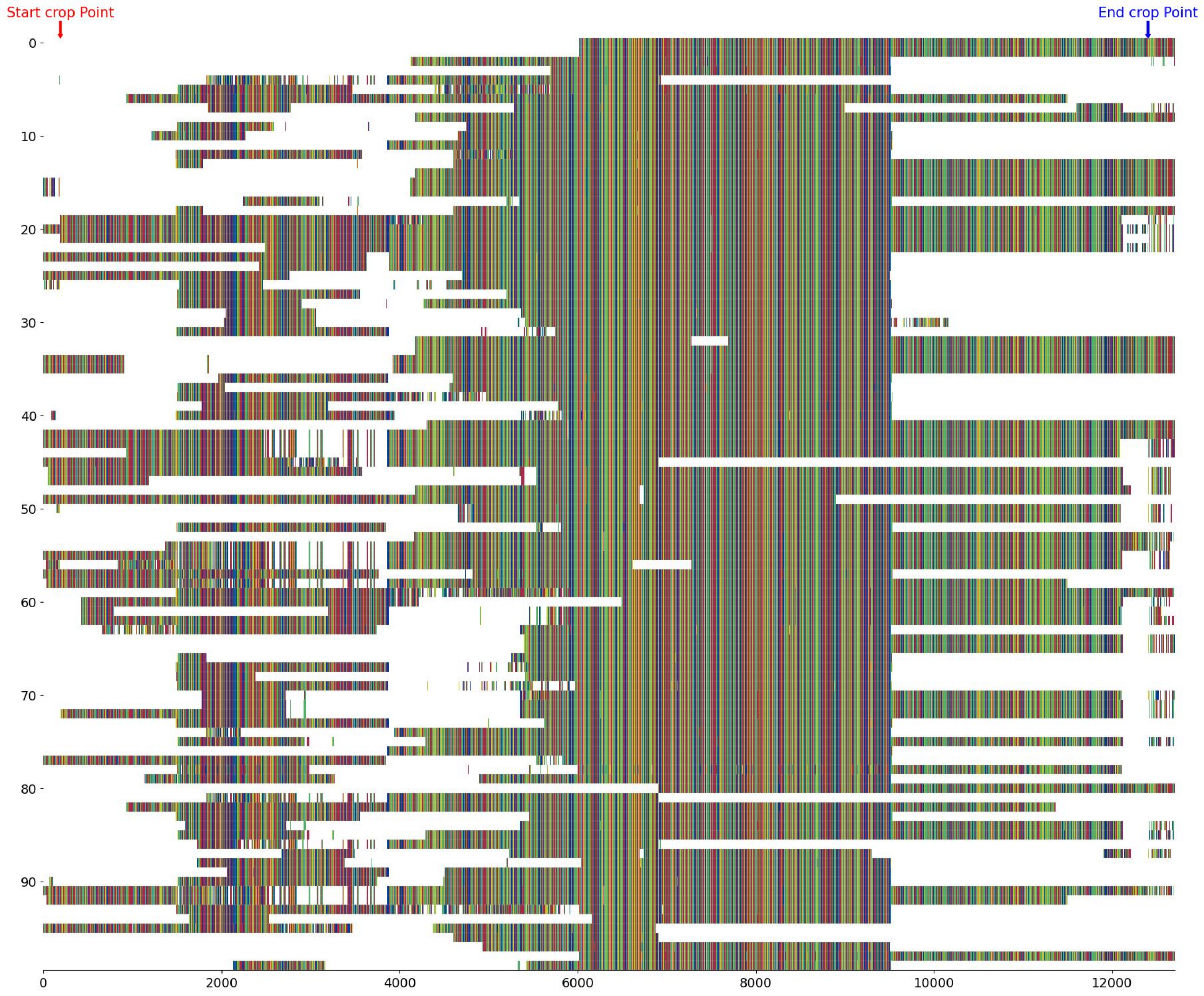
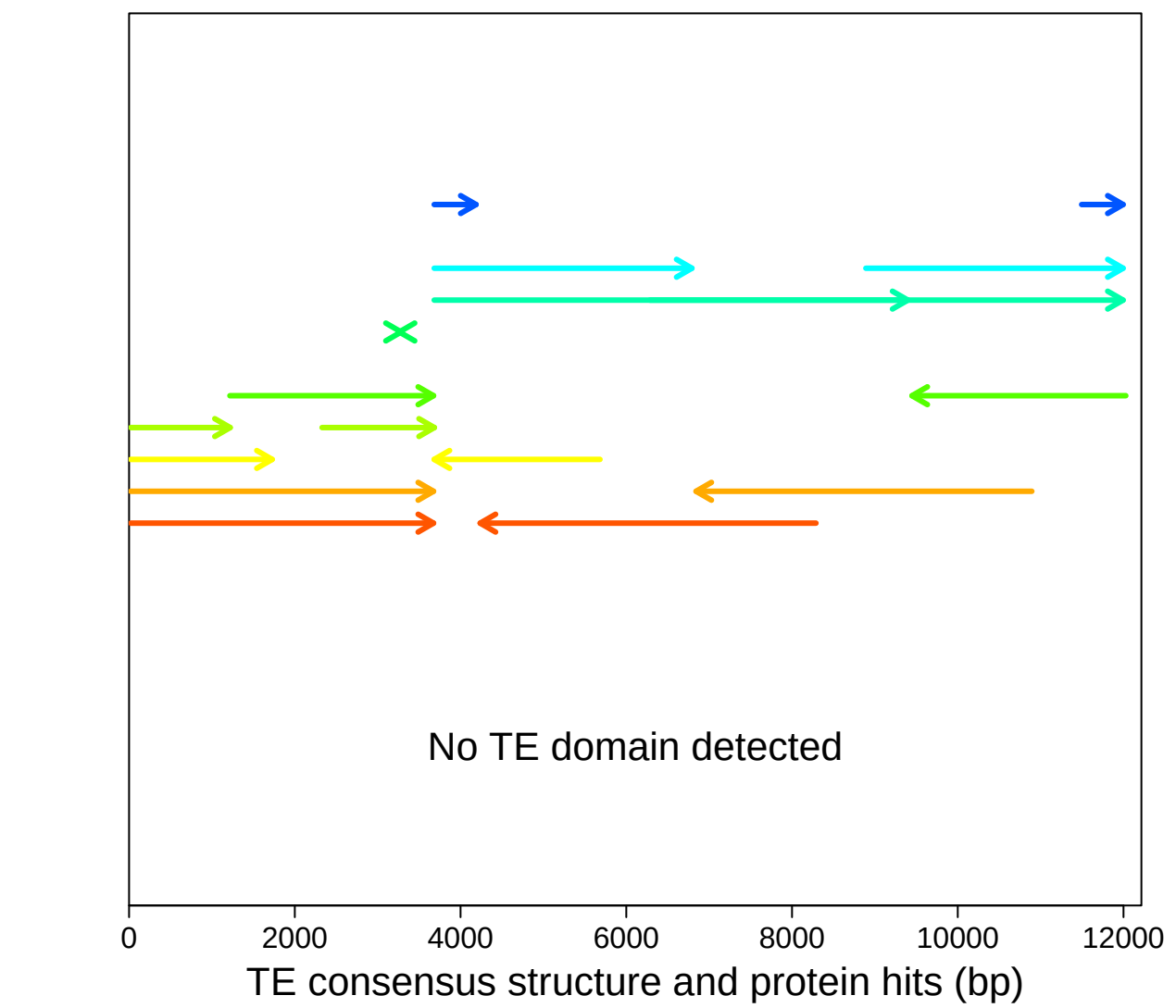
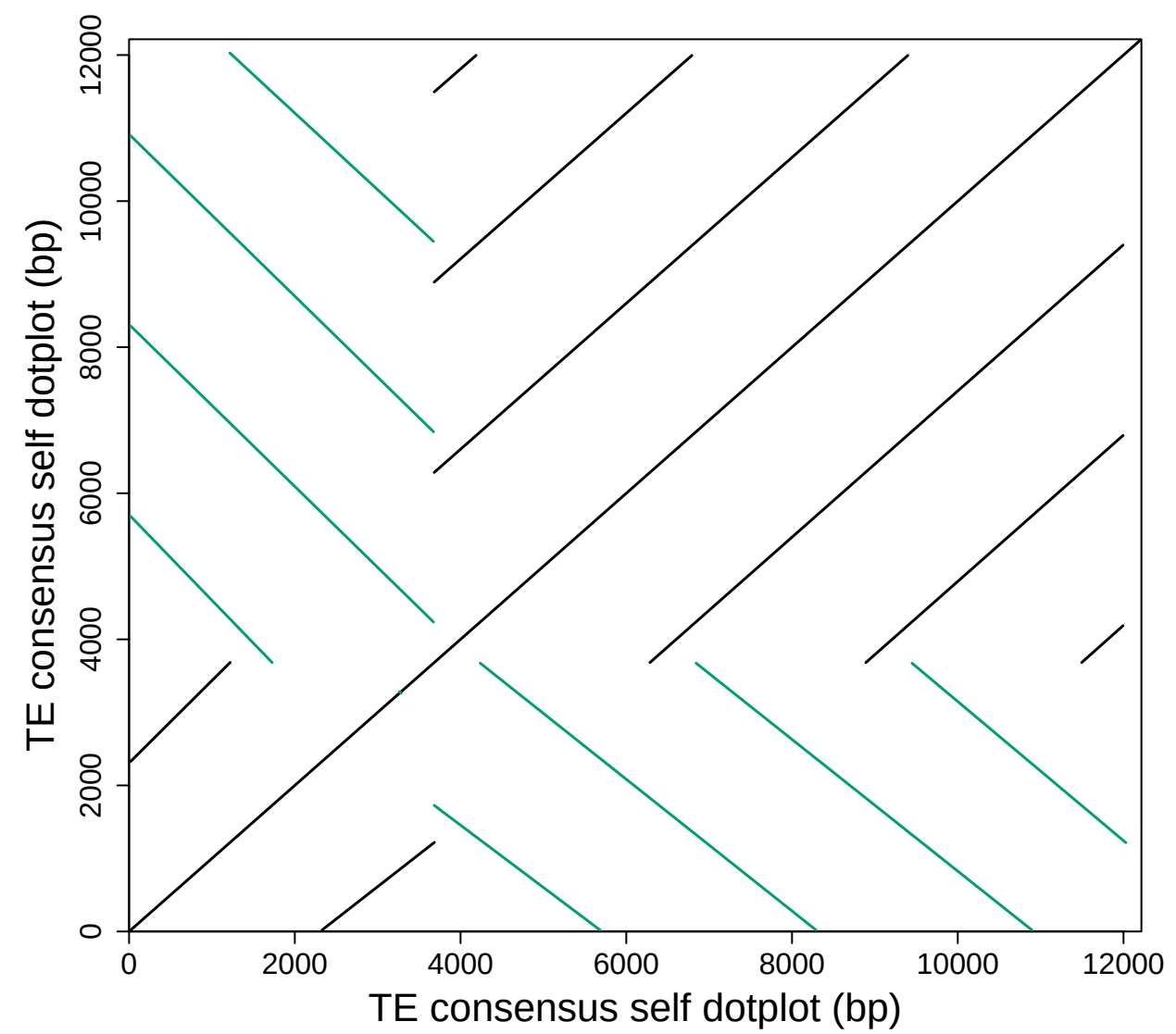
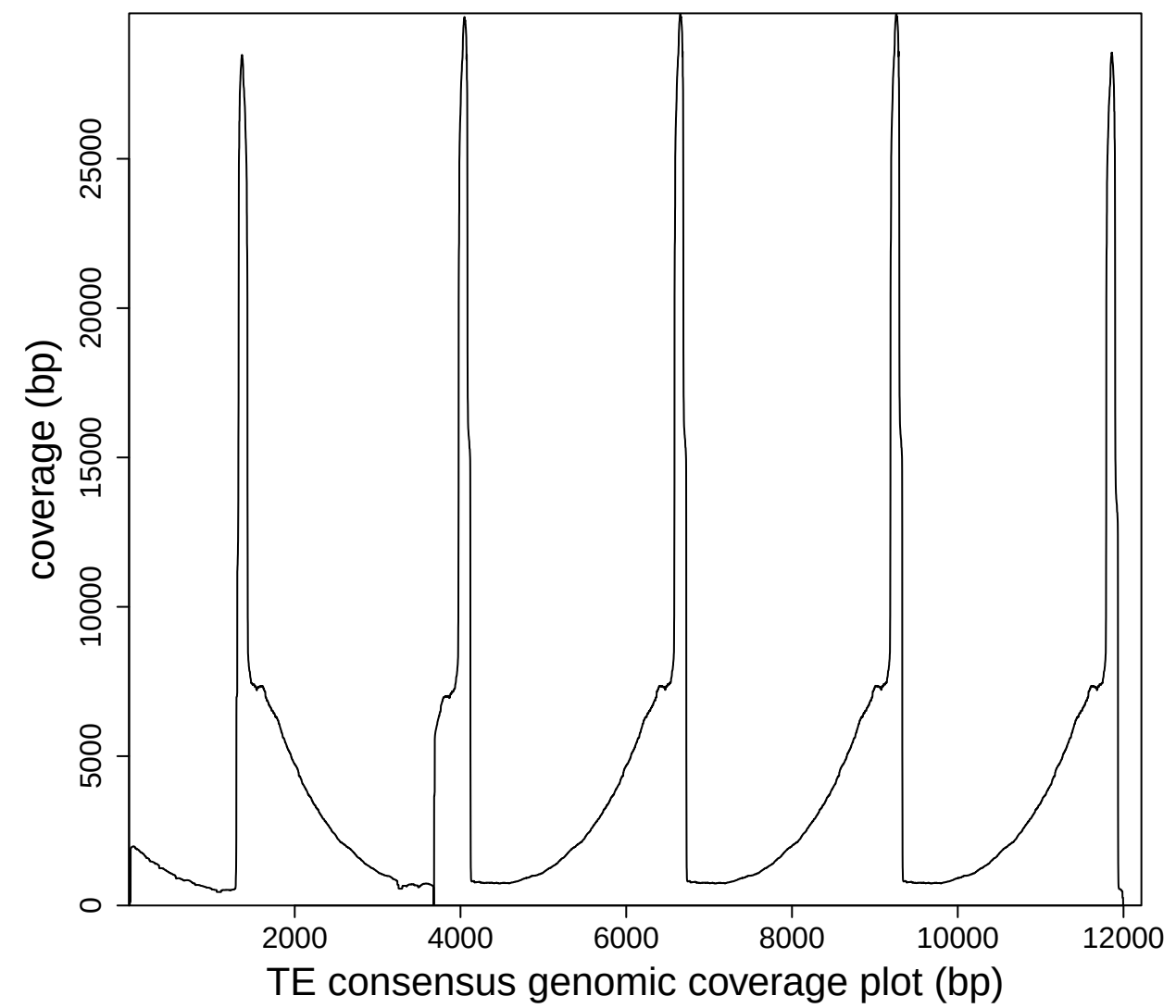
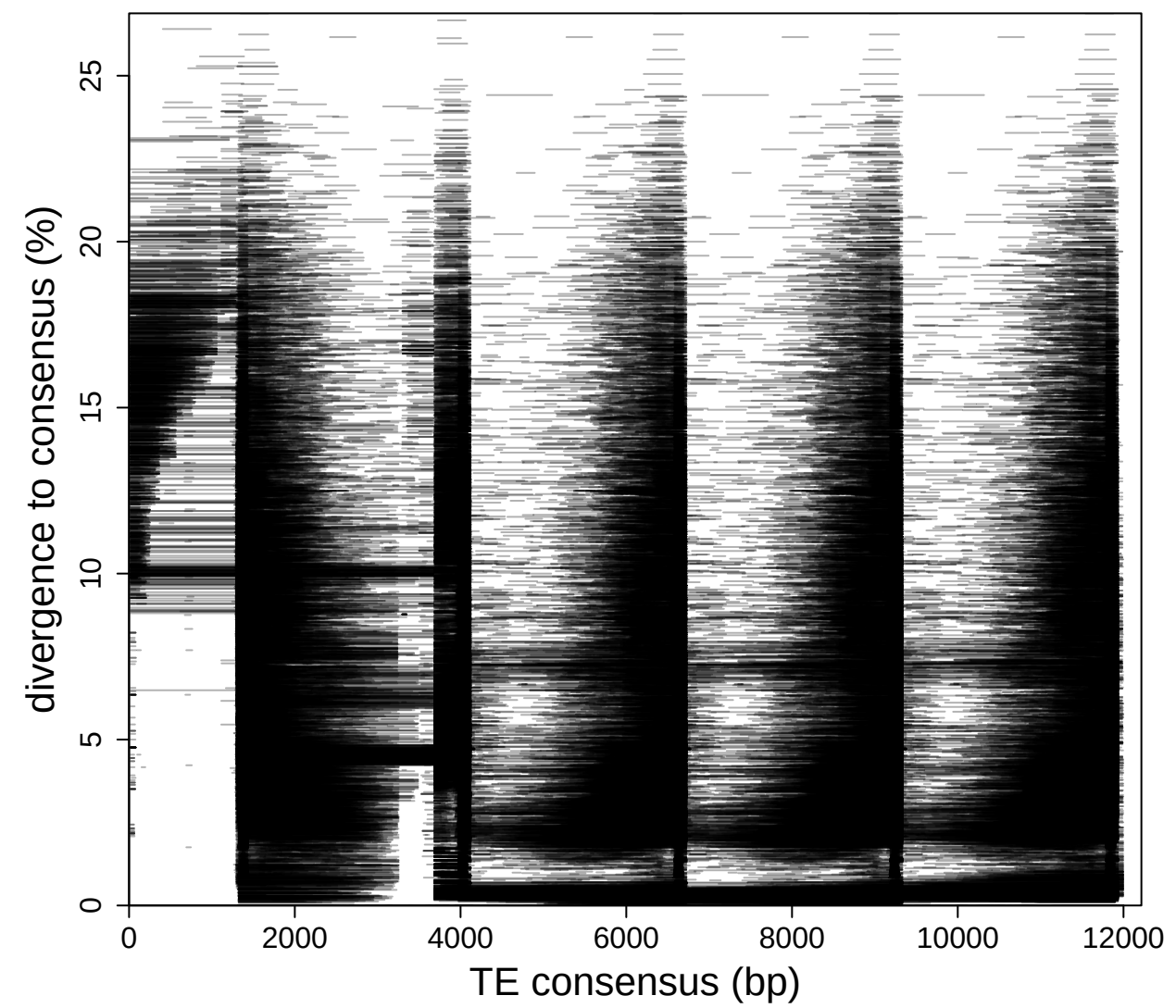






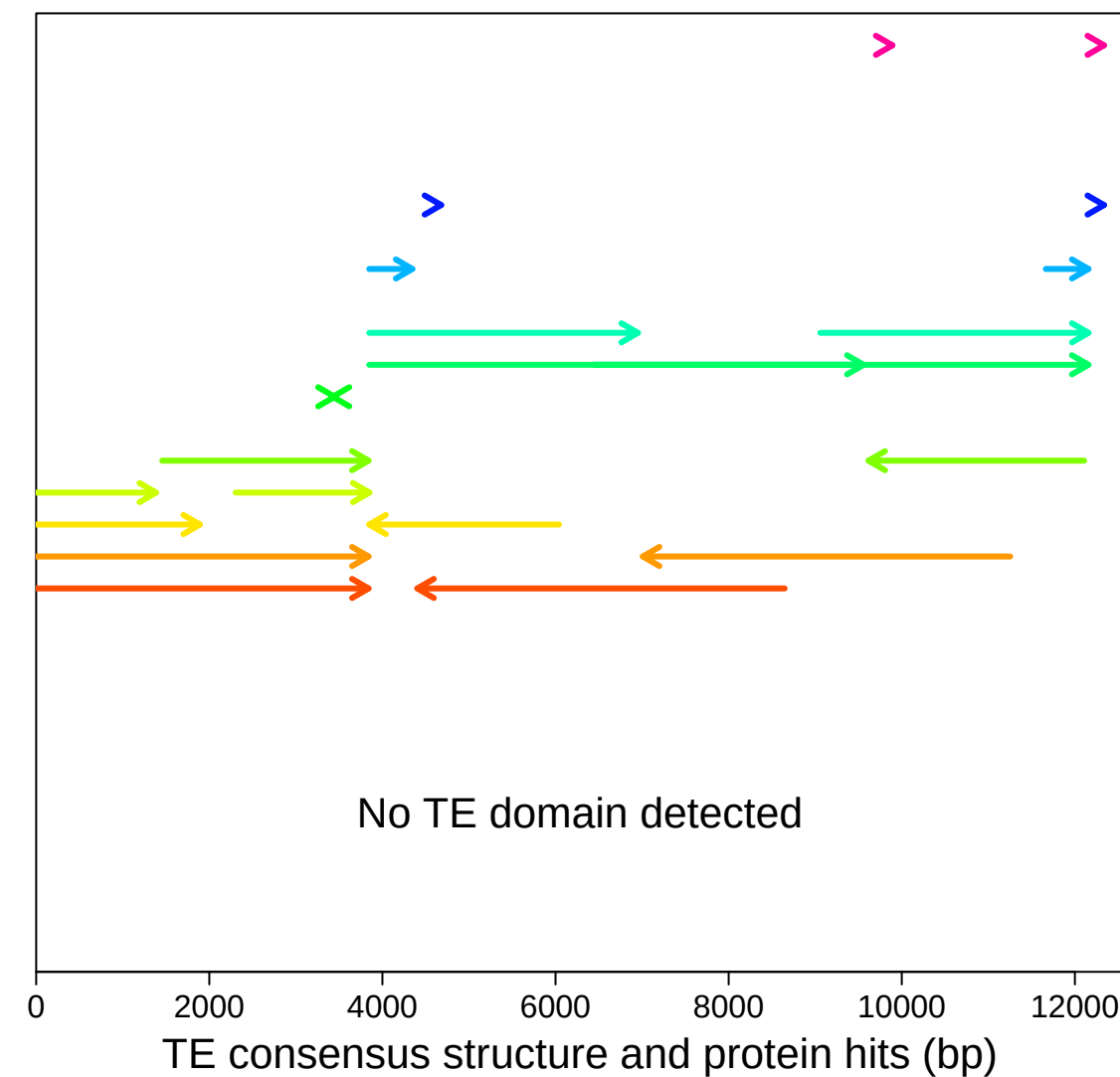
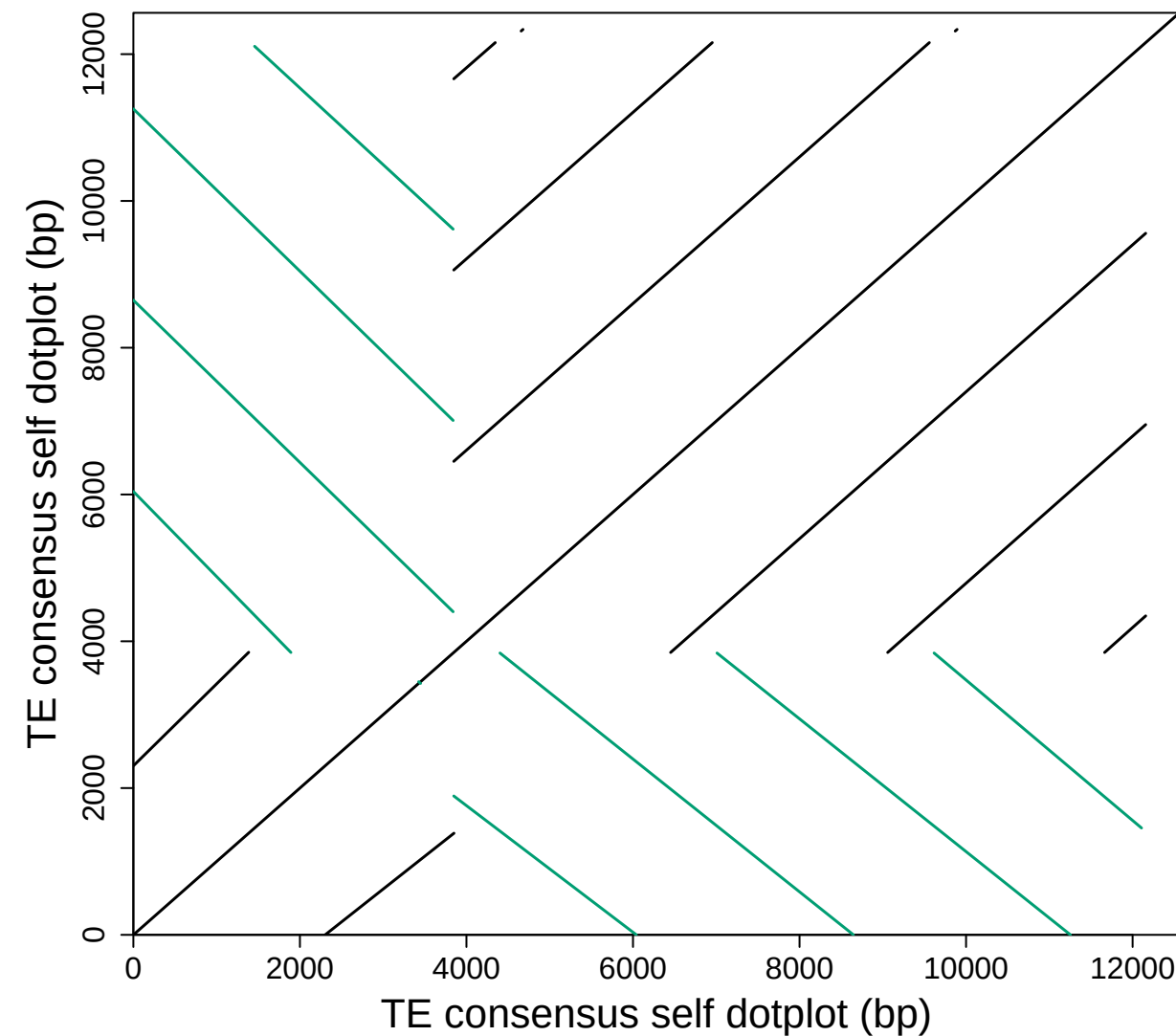
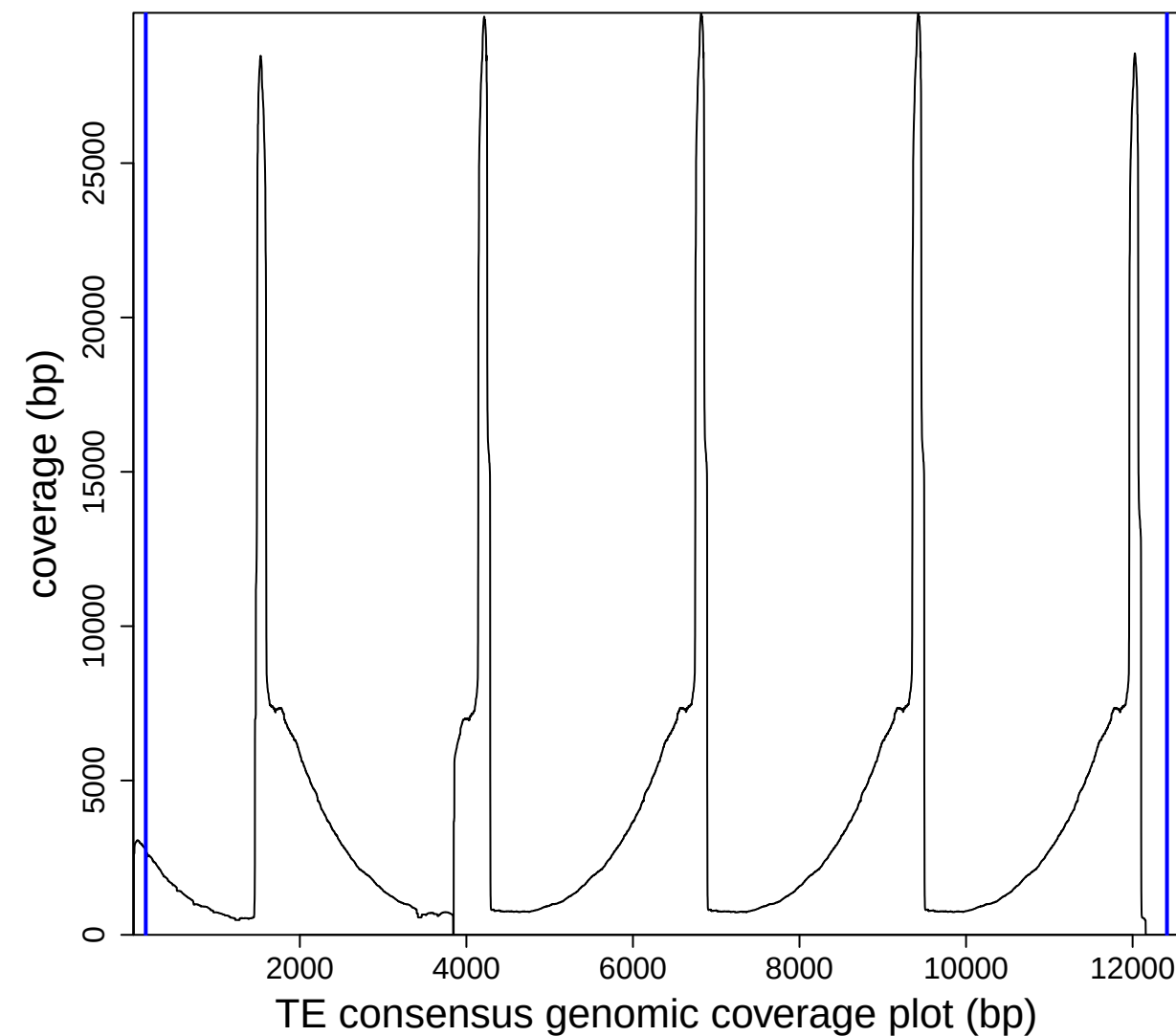
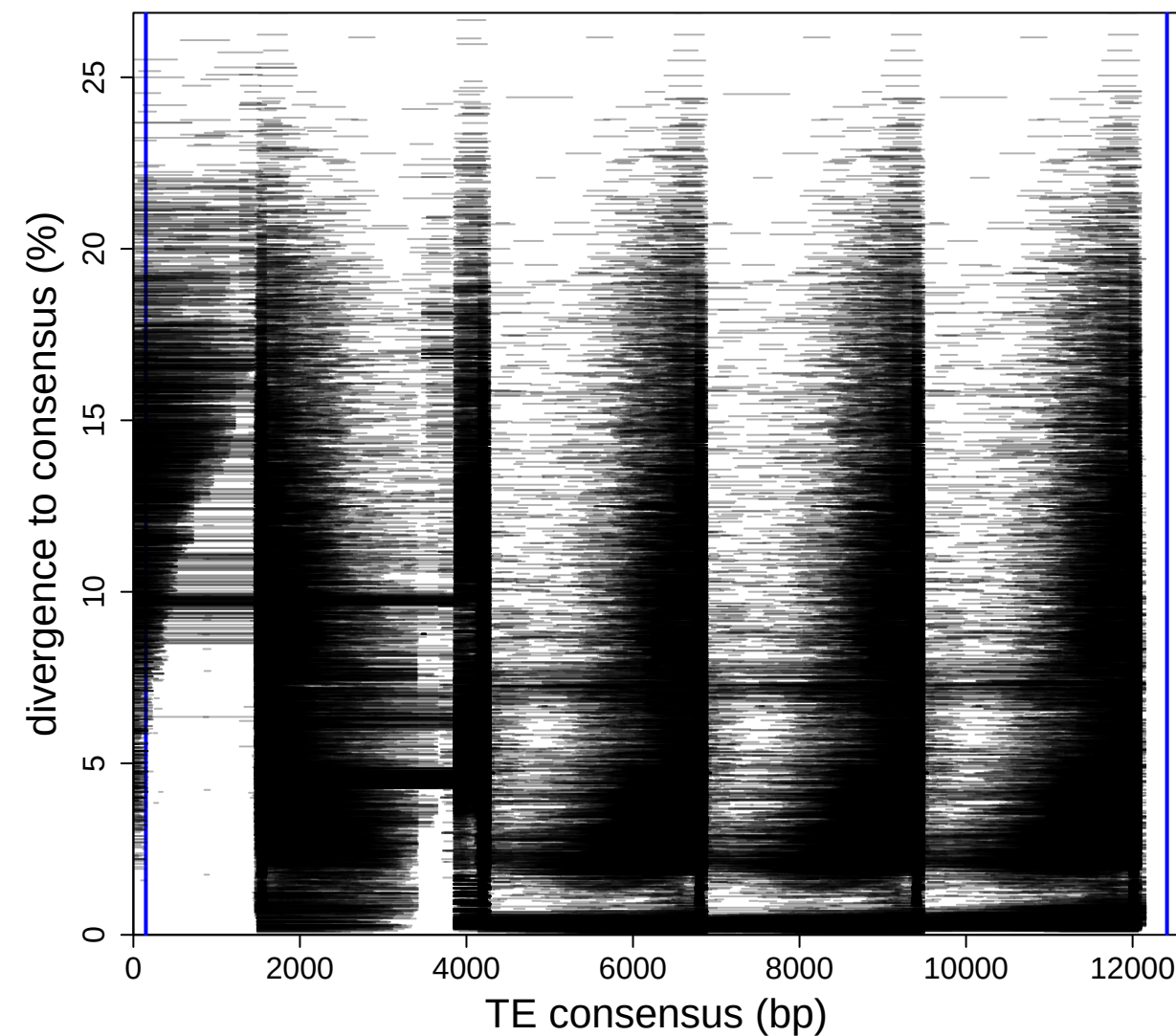
bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.fa cl.fa gs.fa c
size: 12216bp; fragments: 174669; full length: 0 (>=10994.4bp)

After TEtrimmer 12216 bp



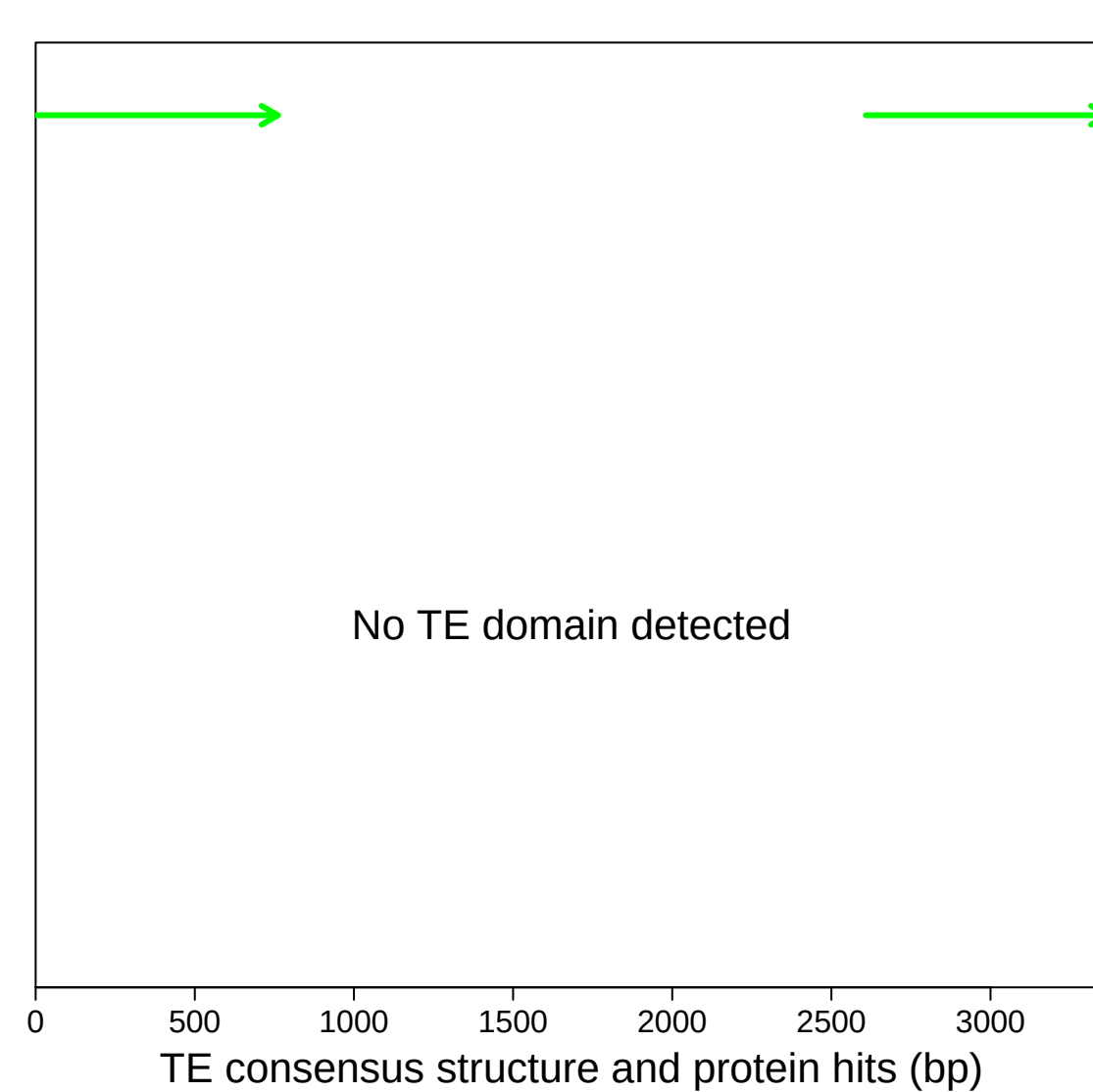
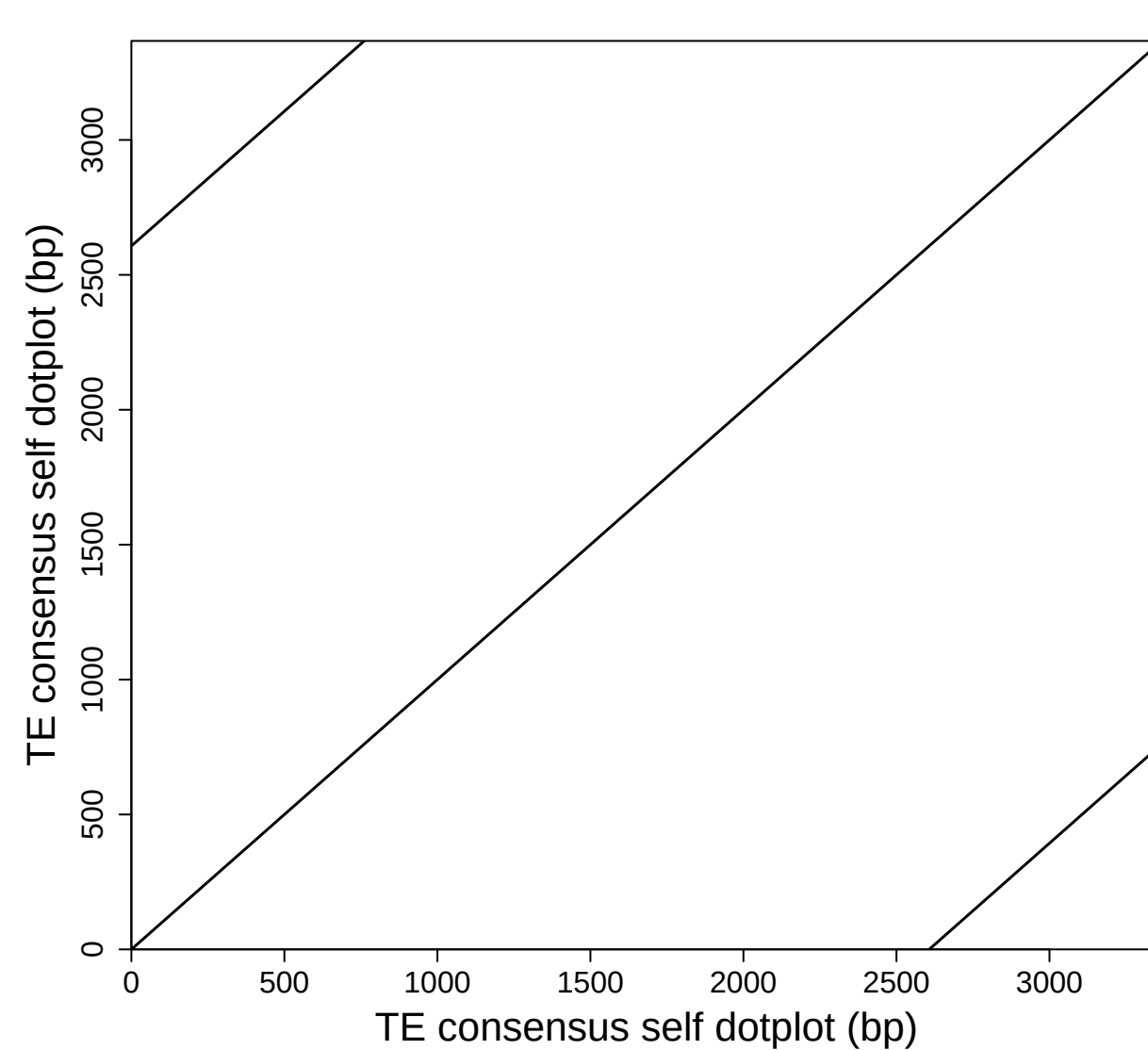
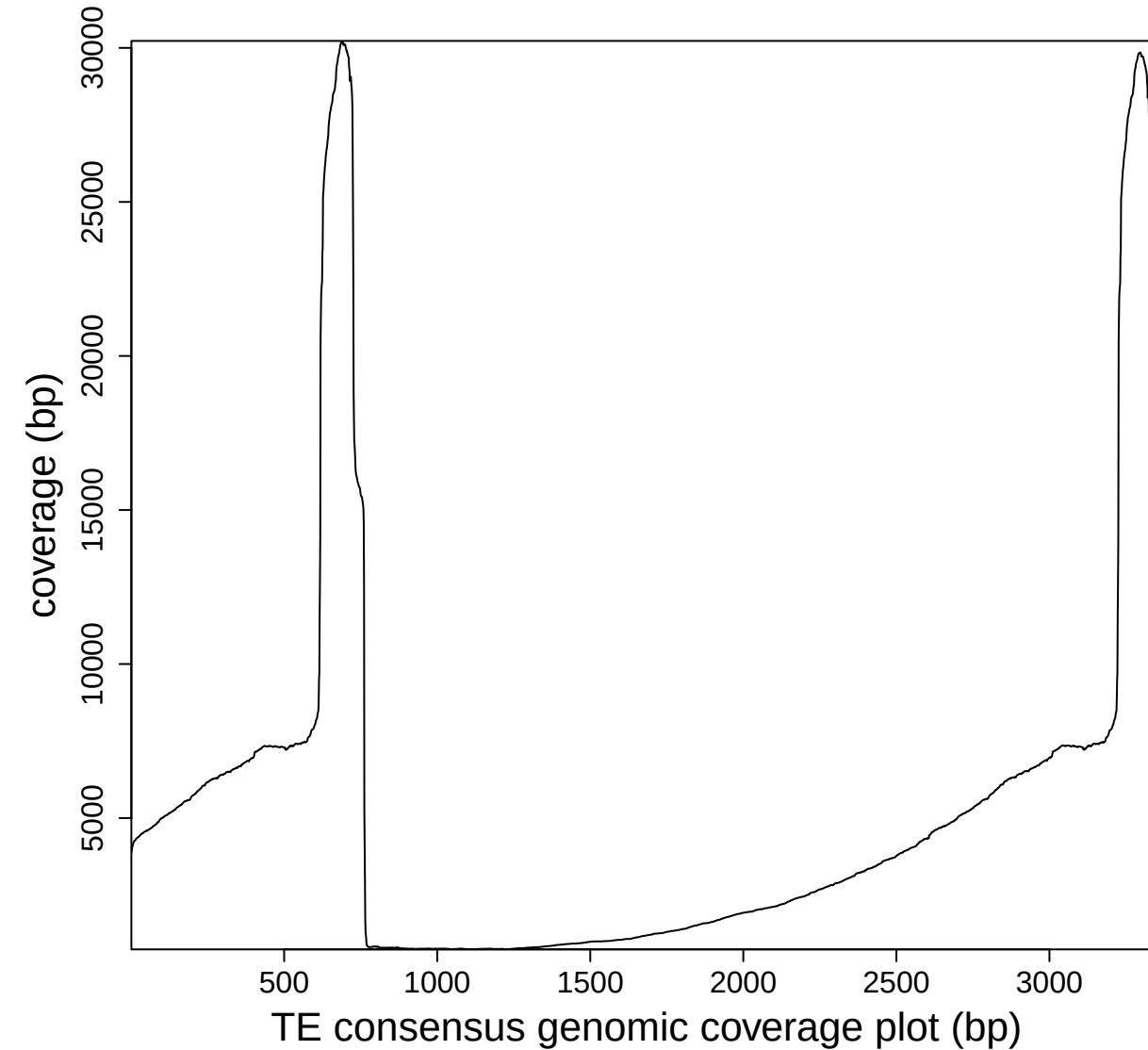
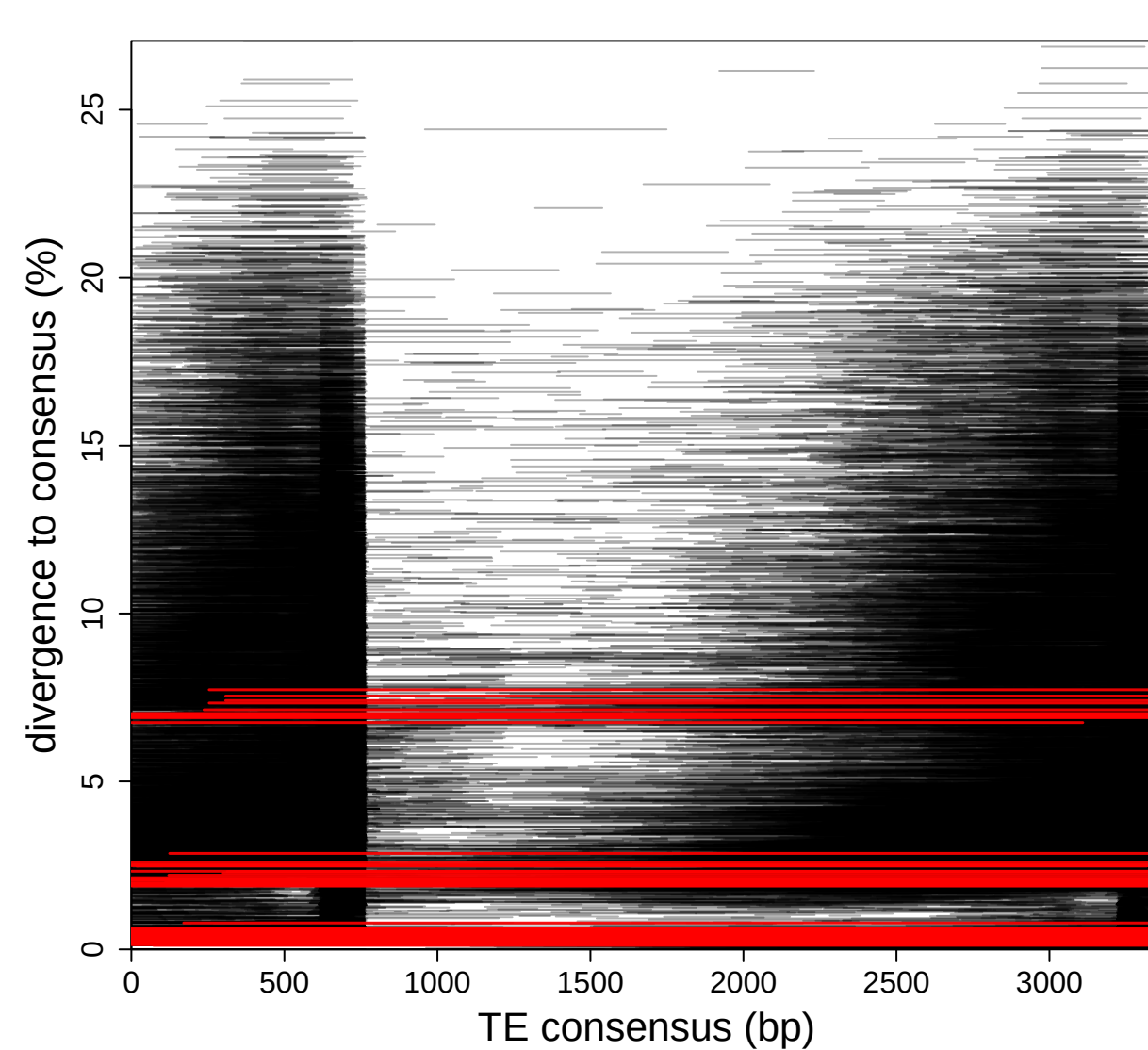
9.fasta.b.bed uf.bed g_1.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 12563bp; fragments: 175958; full length: 0 (>=11306.7bp)

After TEtrimmer Extended plot Blue lines are boundaries

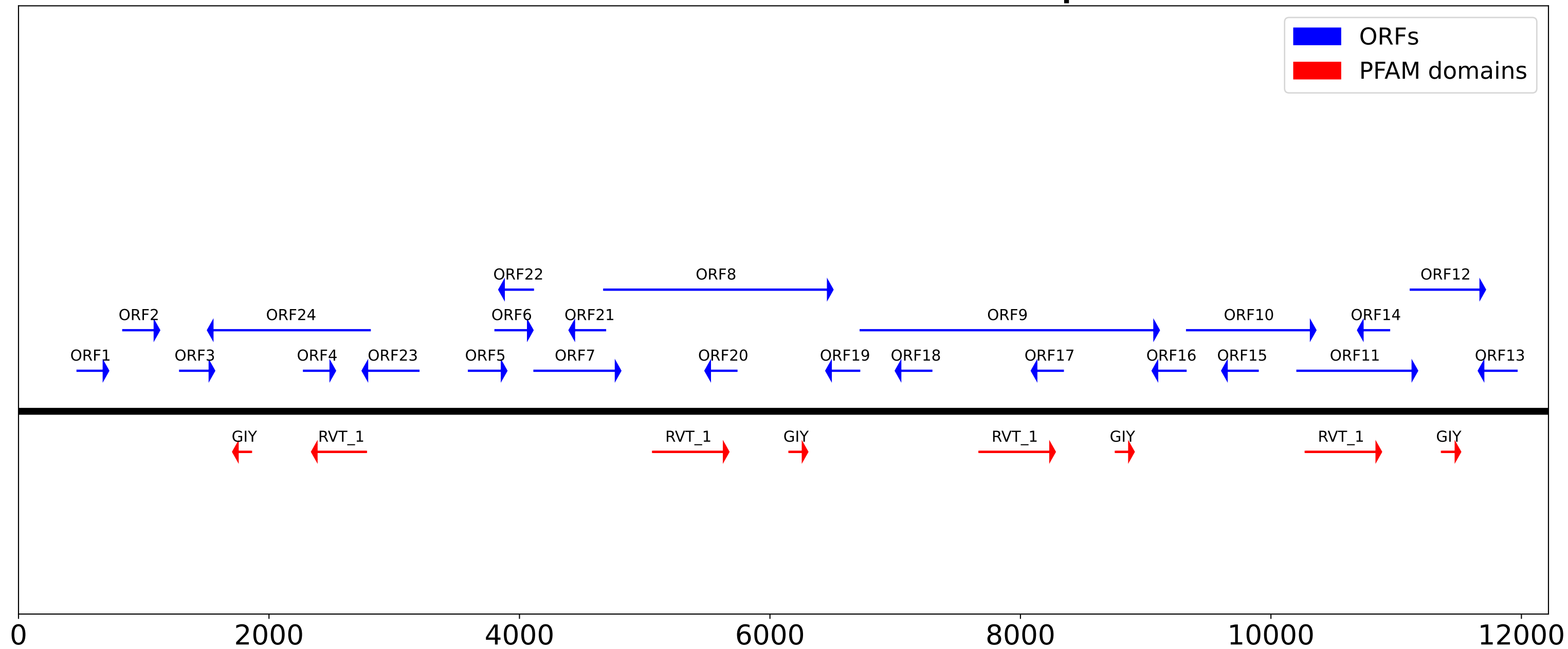


TE: rnd_4_family_1439
size: 3367bp; fragments: 71006; full length: 203 (≥ 3030.3 bp)

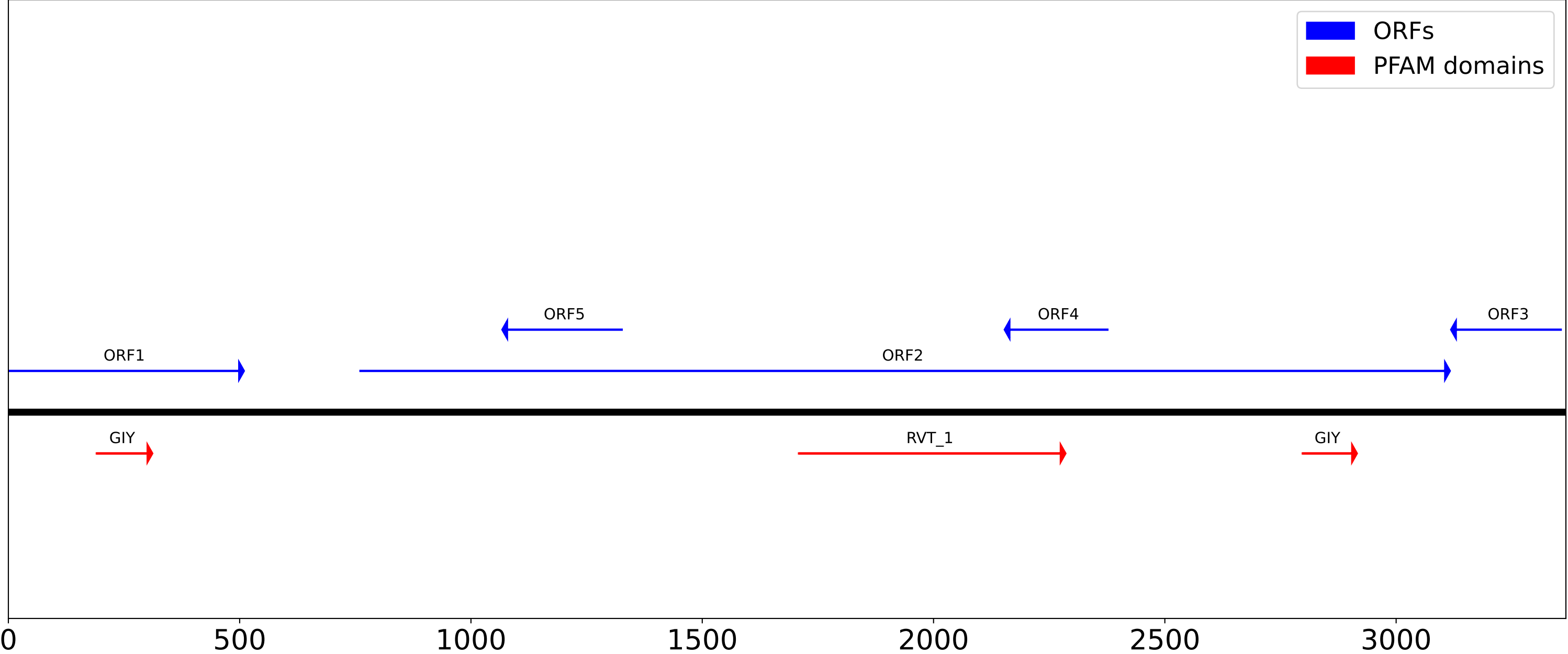
Before TEtrimmer 3367 bp



After TEtrimmer ORF and PFAM domain plot



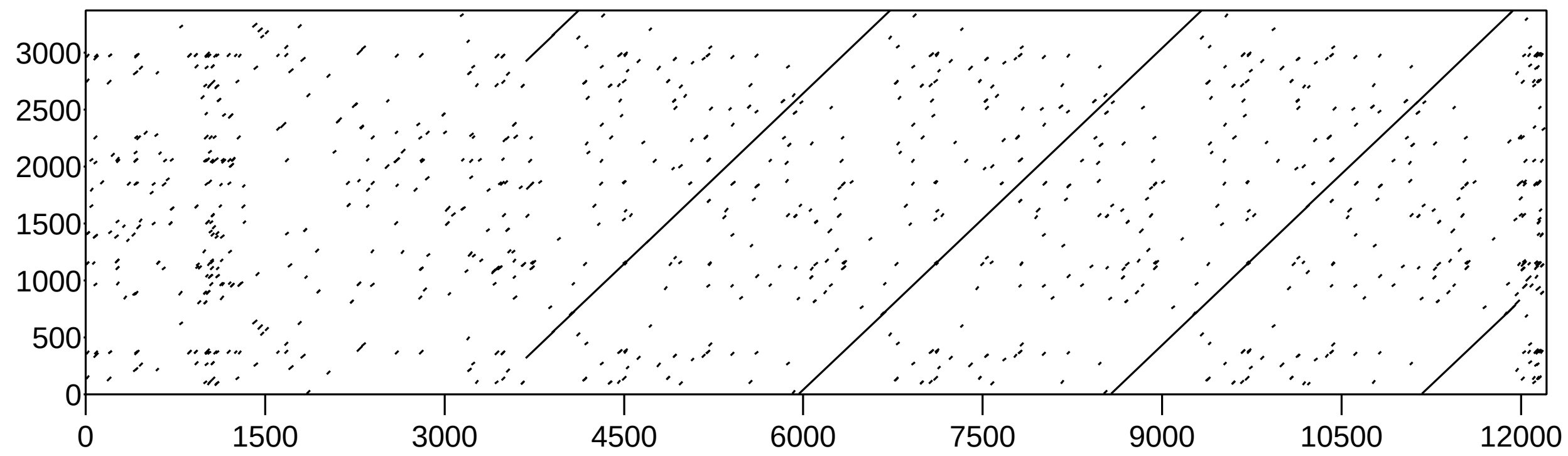
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)