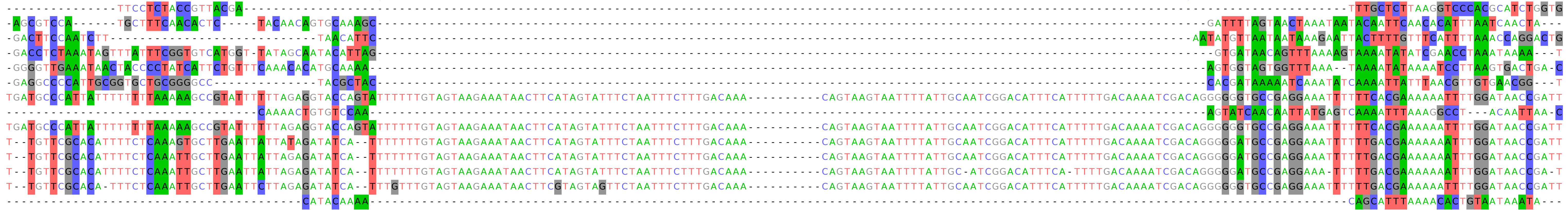
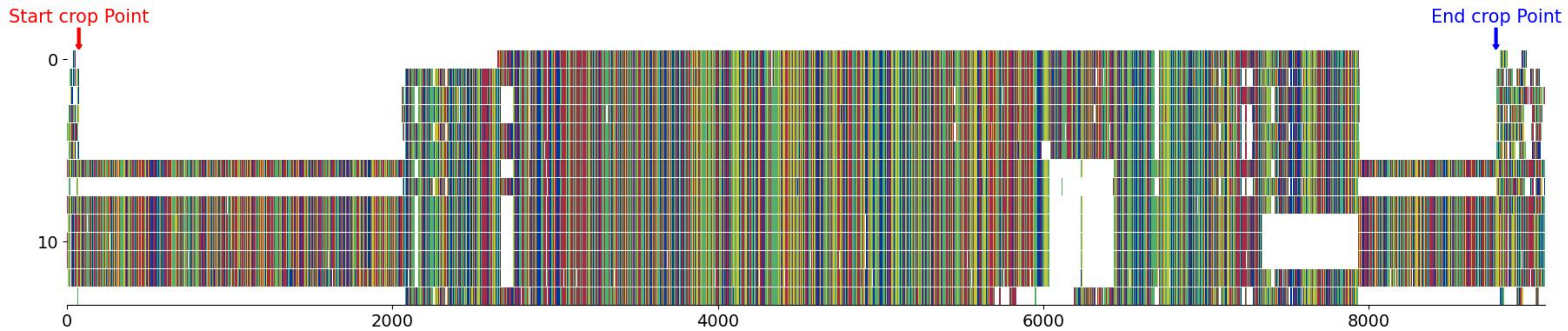


Start crop Point

End crop Point

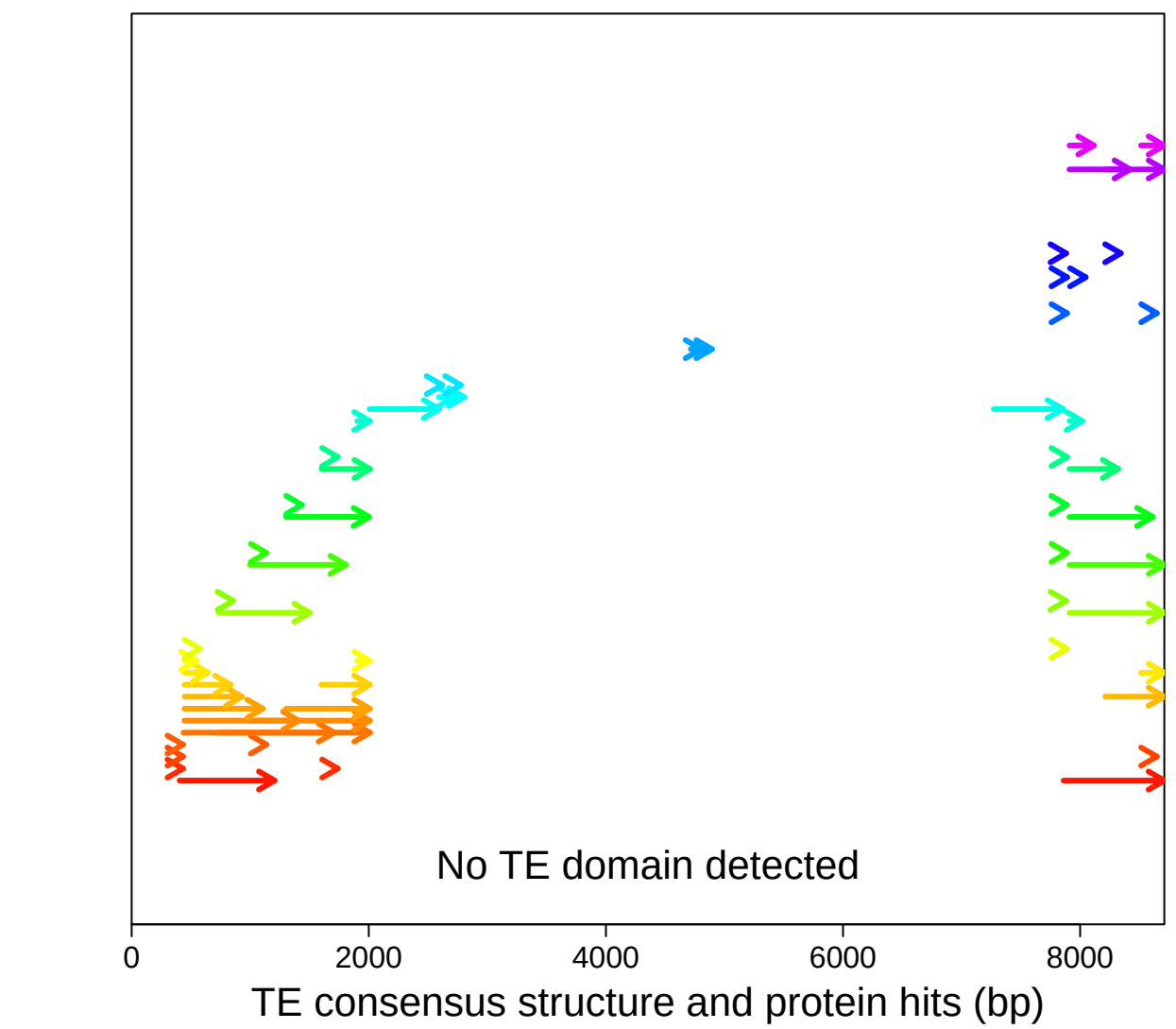
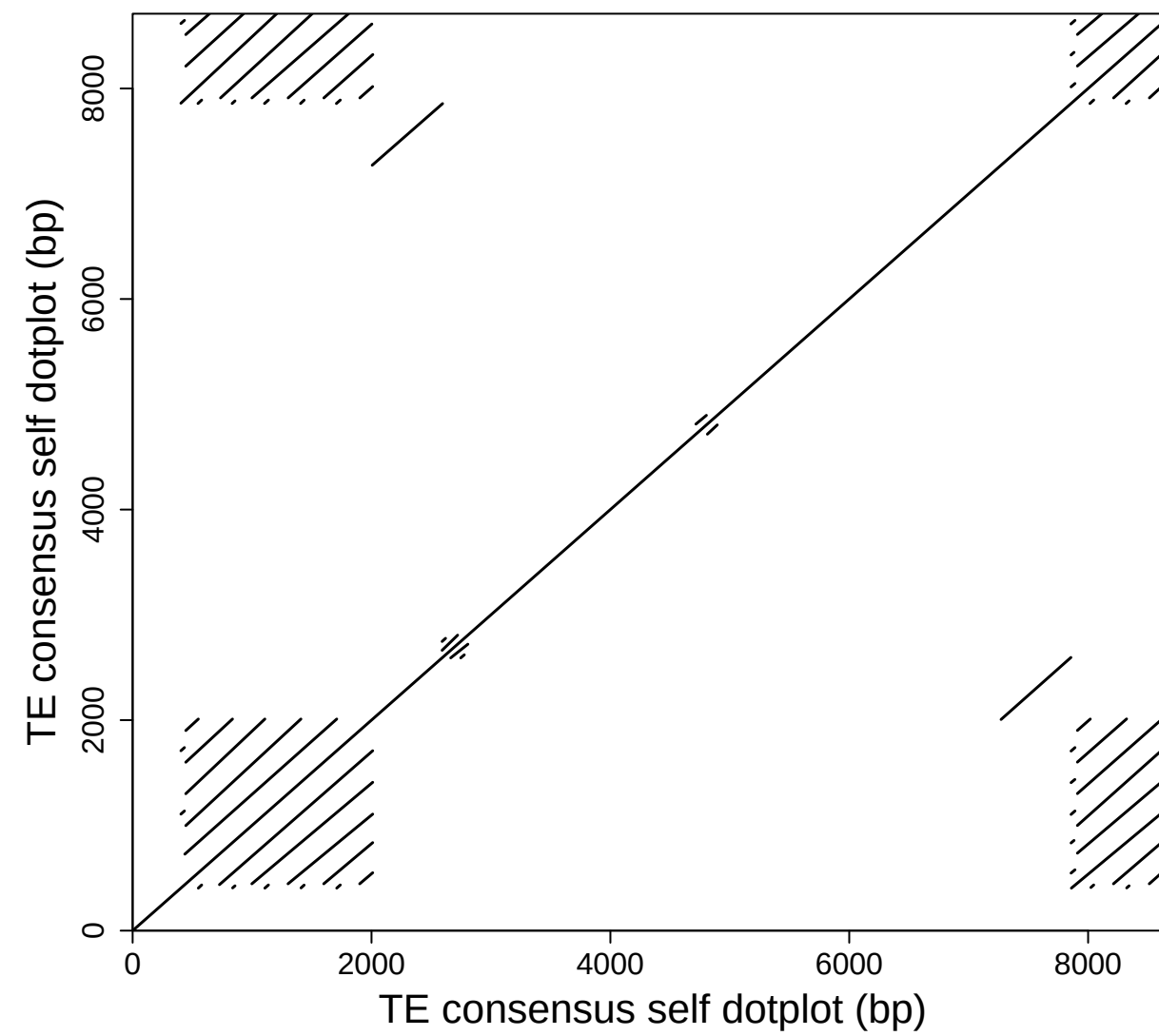
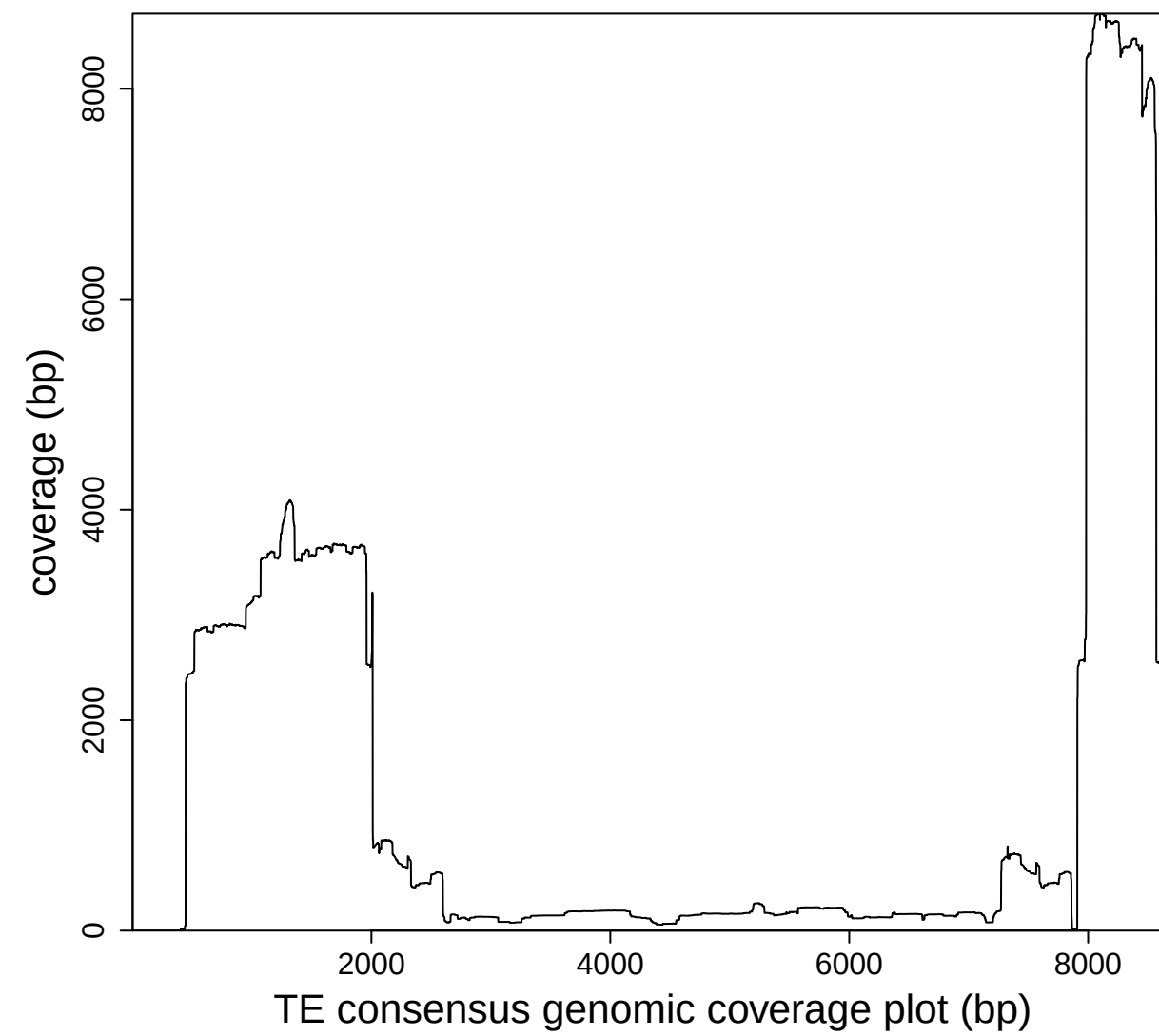
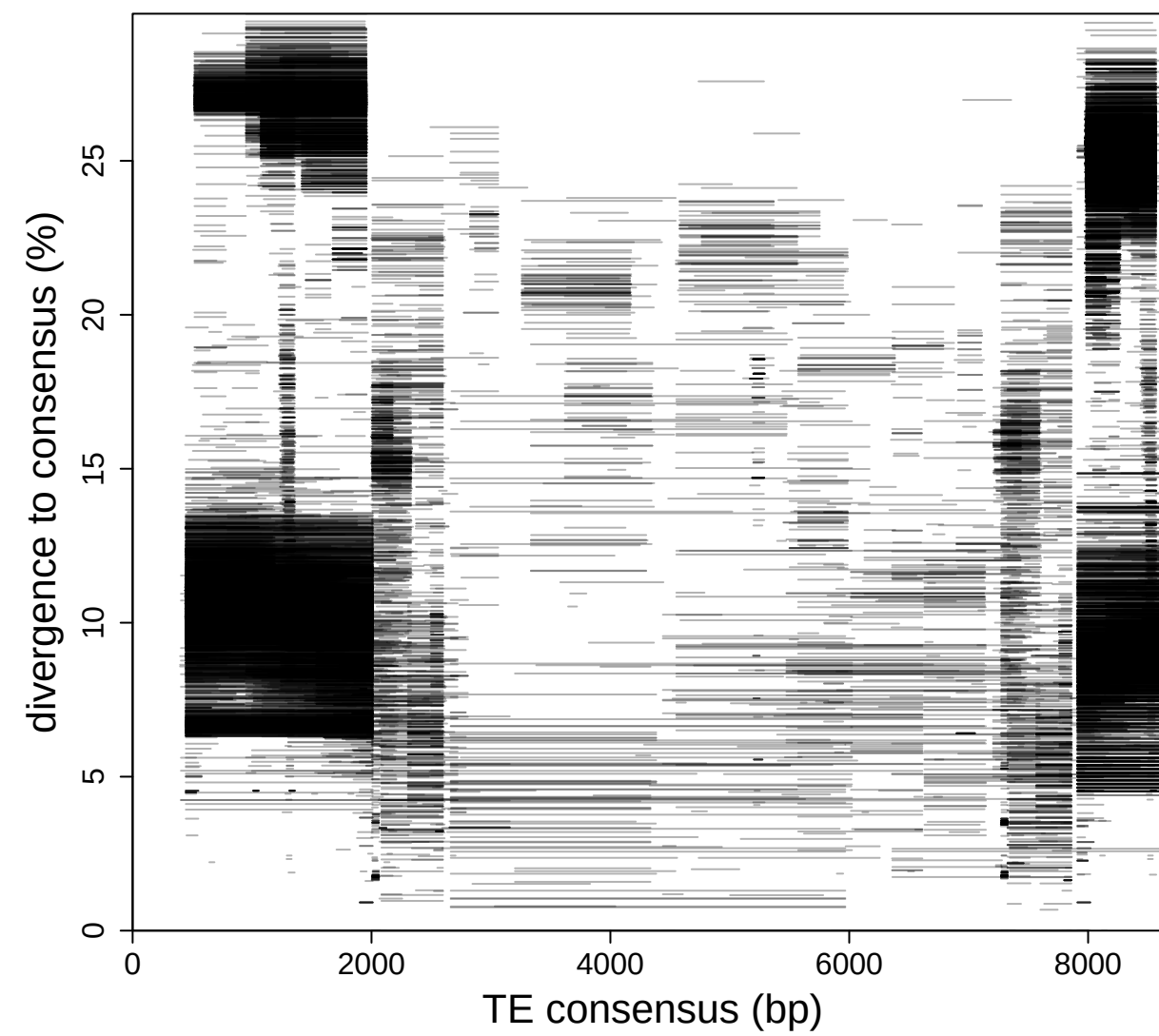
MSA length = 8710





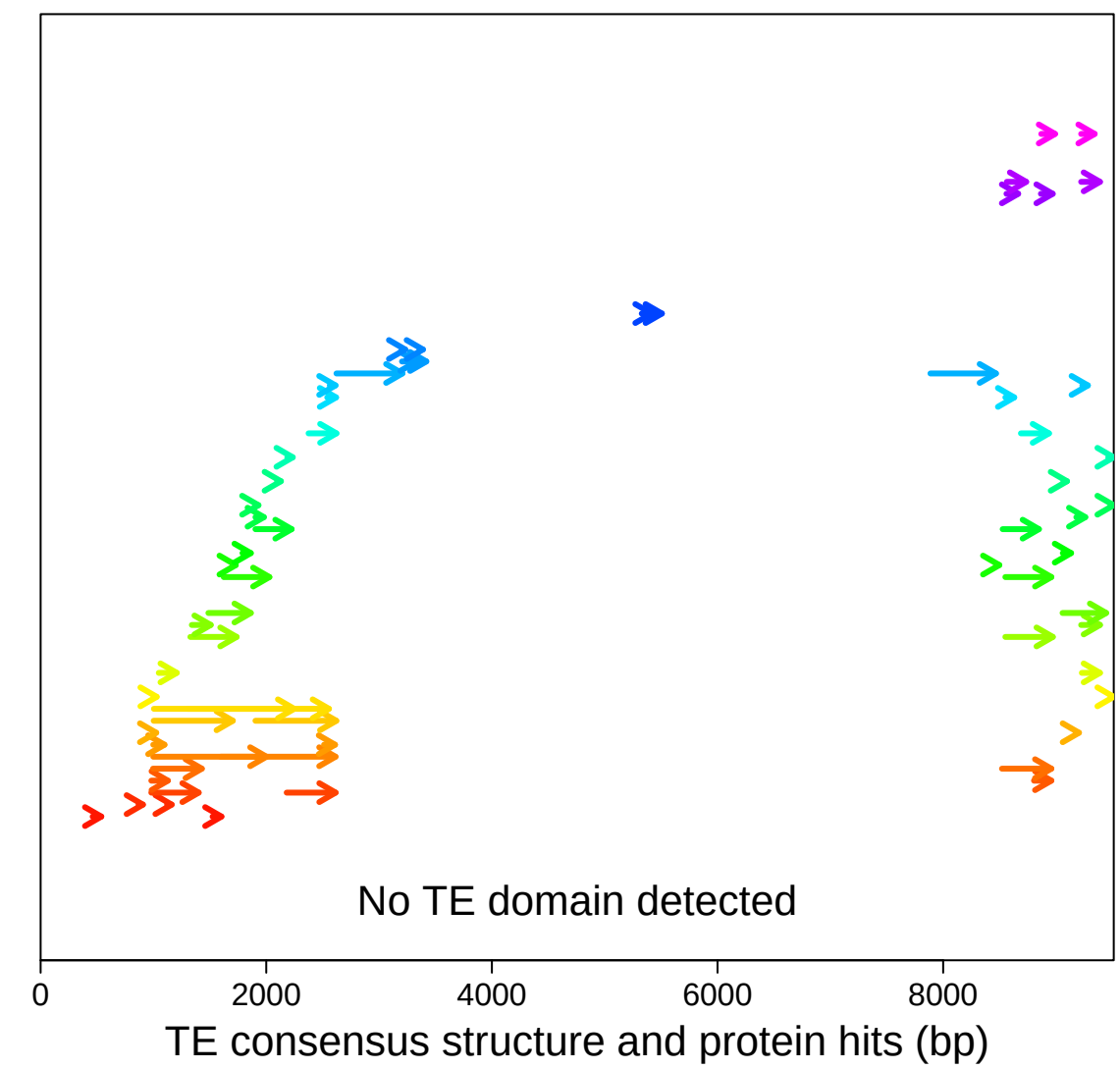
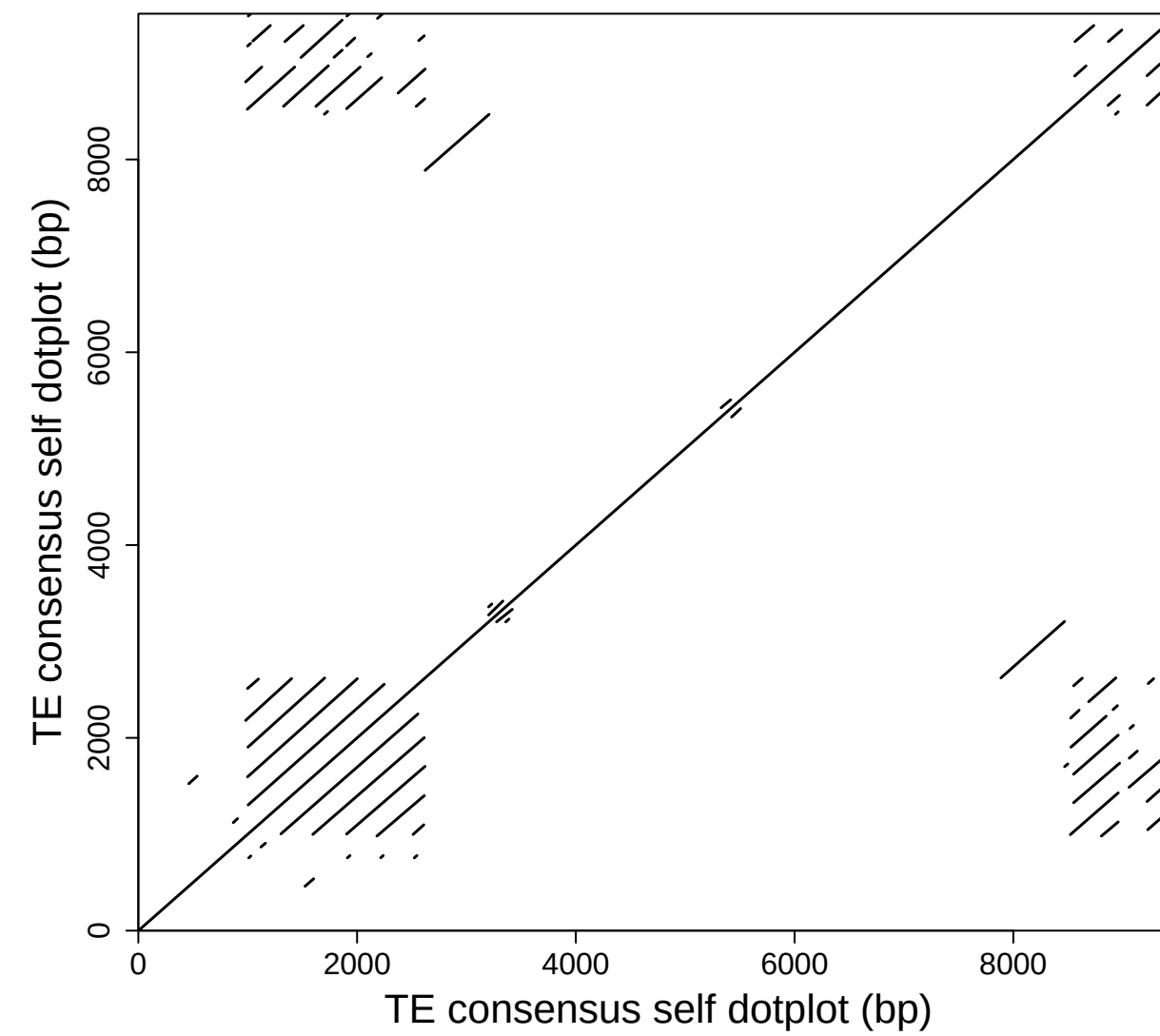
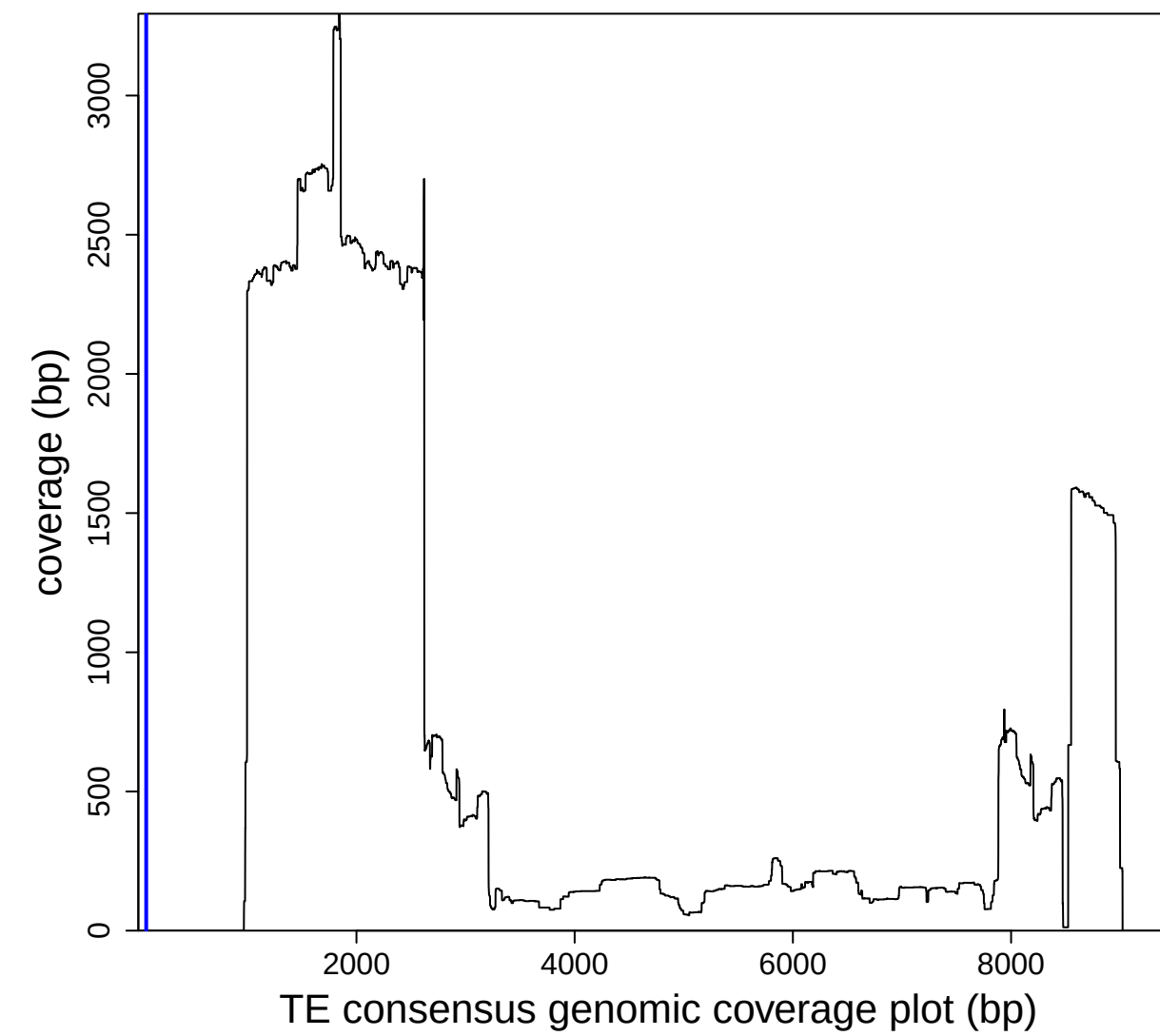
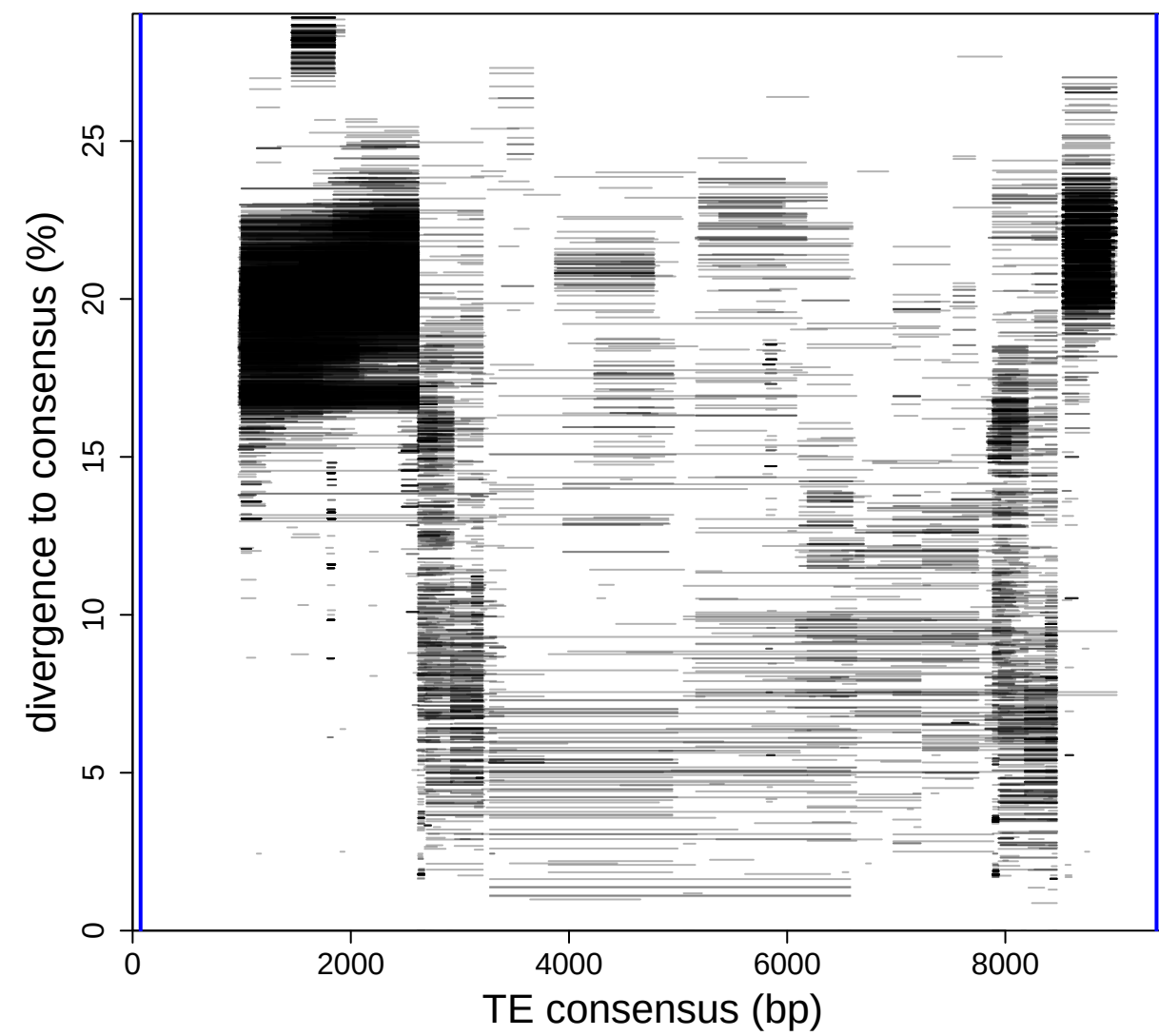
bed_uf.bed g_2.bed fm_1.bed 0_0_bcln.fa aln.fa cl.fa gs.fa c
size: 8710bp; fragments: 18448; full length: 0 (>=7839bp)

After TEtrimmer 8710 bp



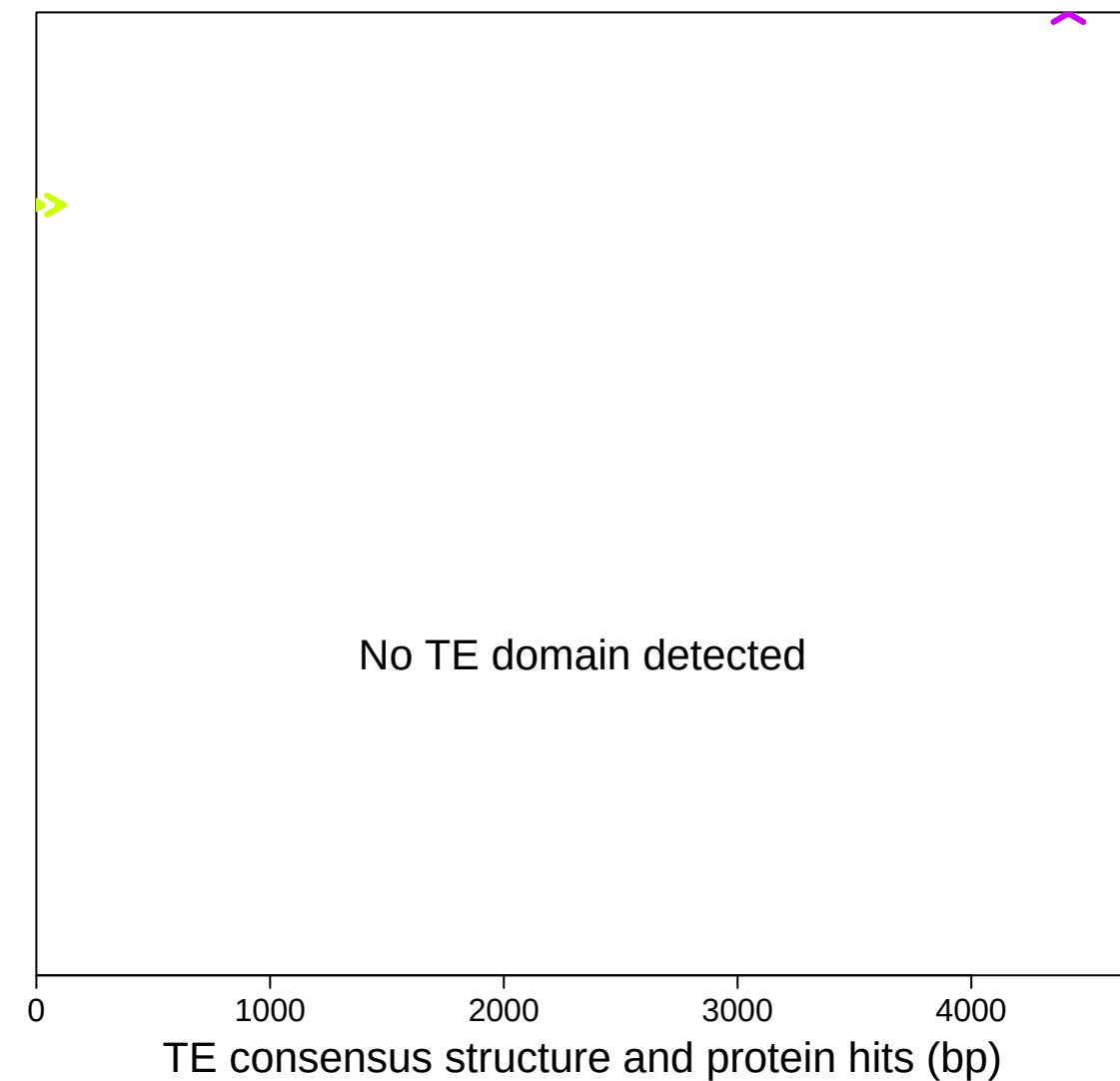
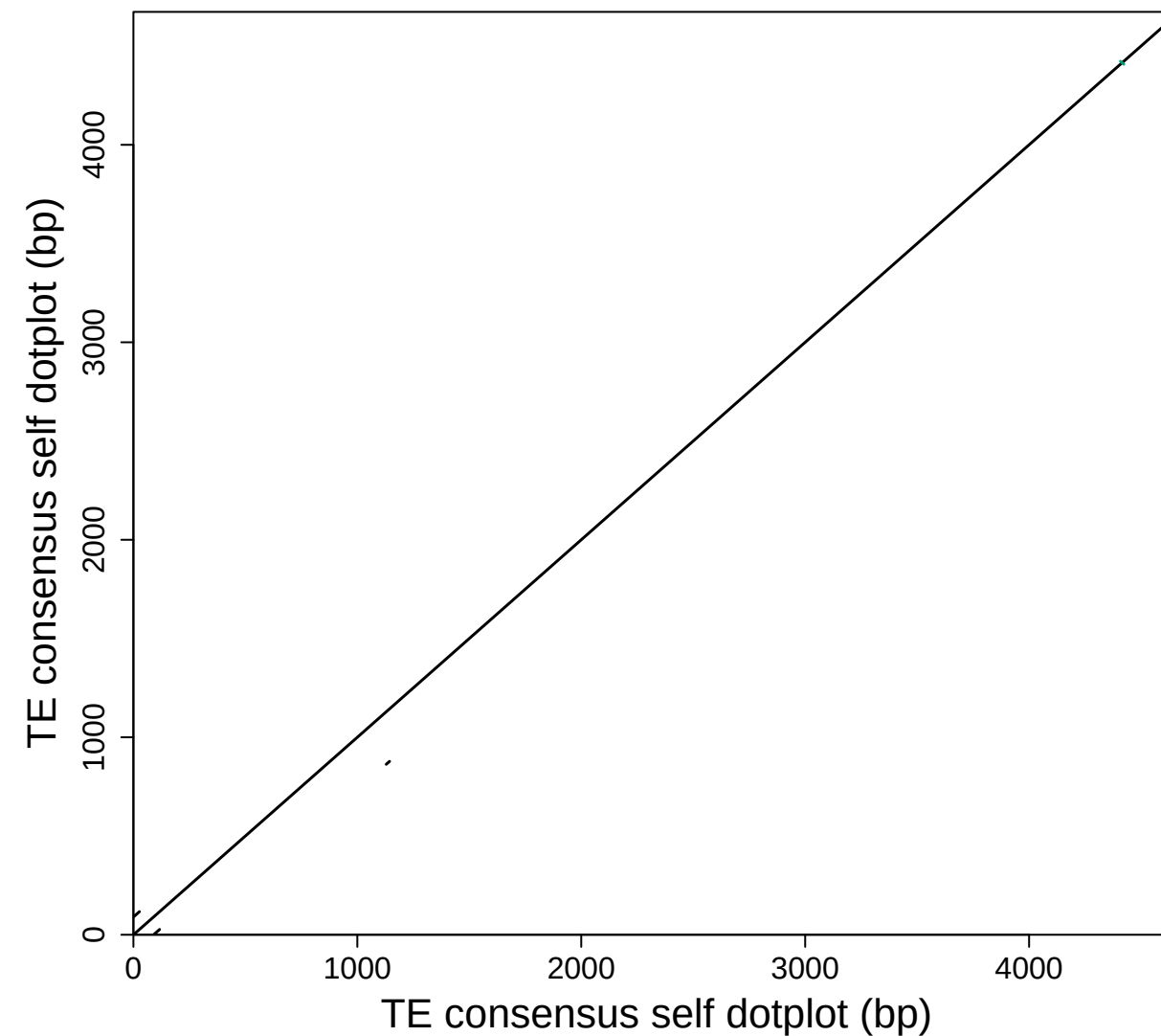
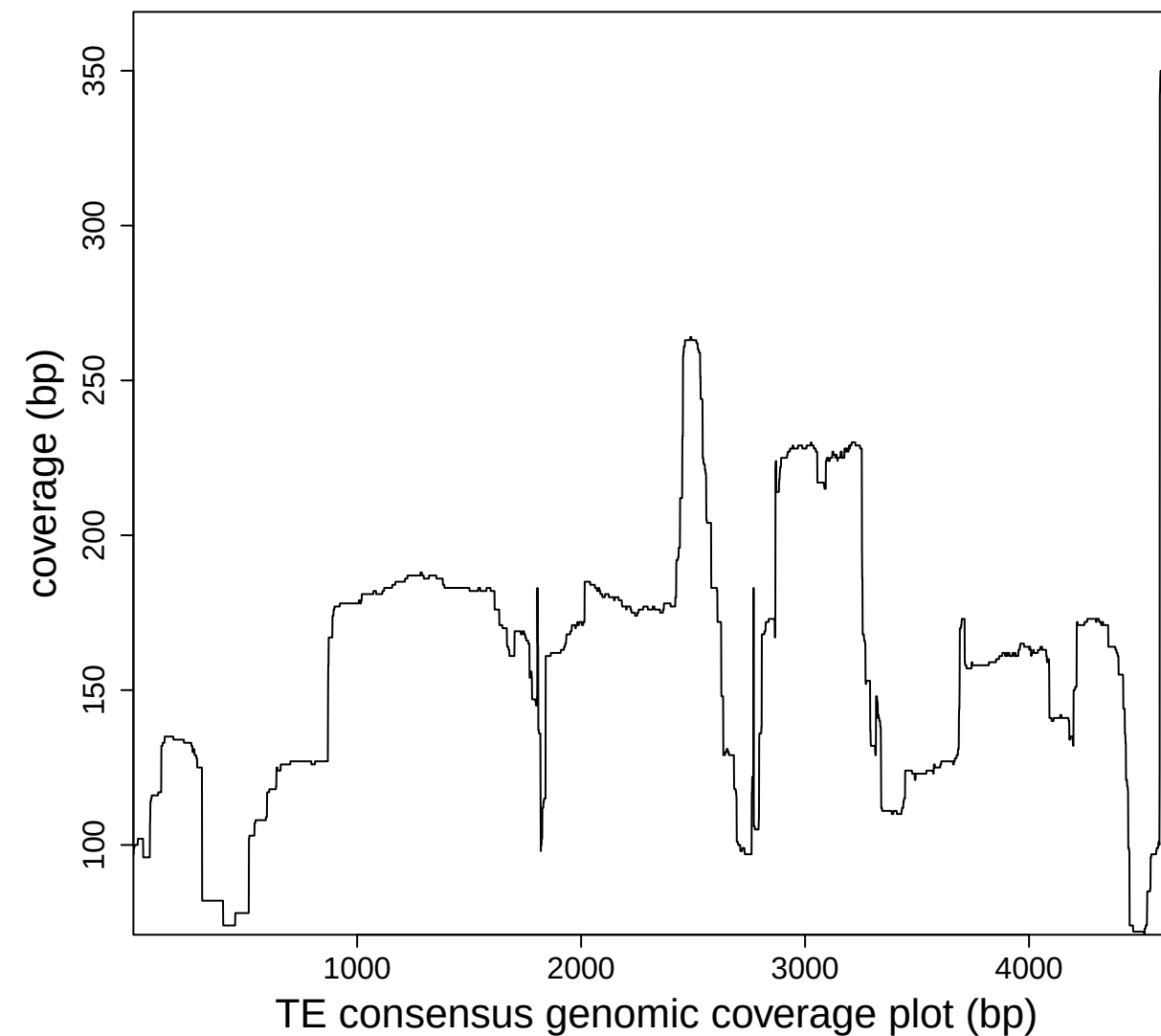
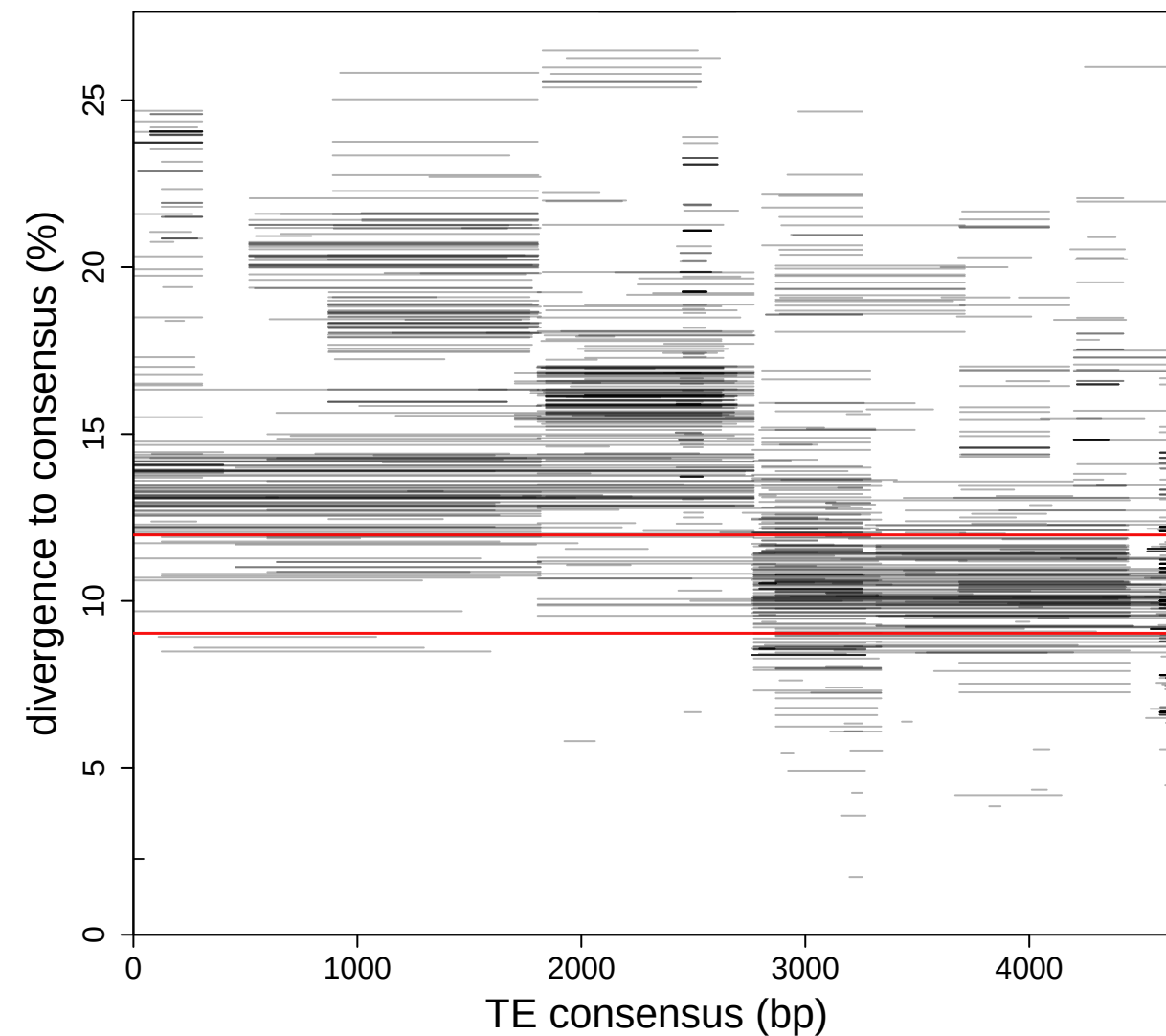
ly_1143.fasta.b.bed uf.bed g 2.bed fm 1.bed 0 0 bcln.fa aln.f
size: 9536bp; fragments: 8836; full length: 0 (>=8582.4bp)

After TEtrimmer Extended plot Blue lines are boundaries

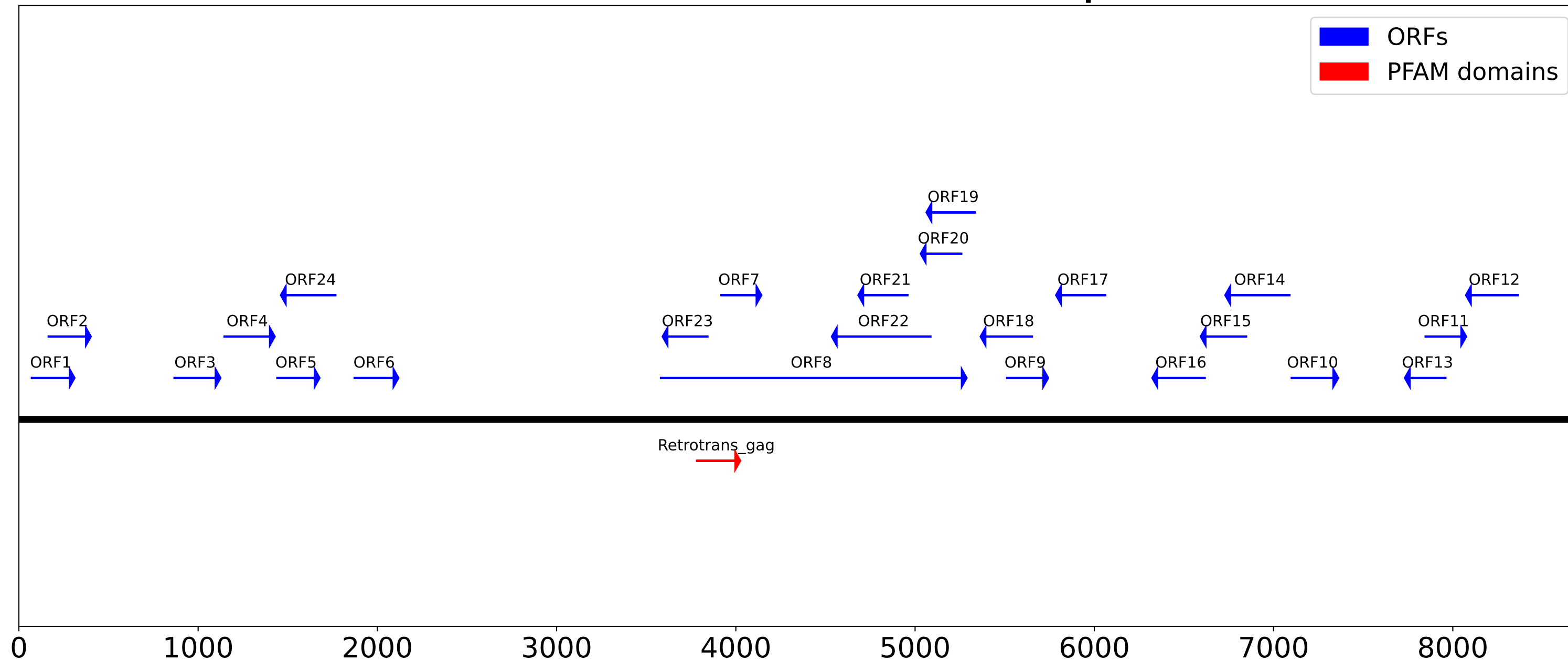


TE: rnd_4_family_1143
size: 4673bp; fragments: 1229; full length: 2 (≥ 4205.7 bp)

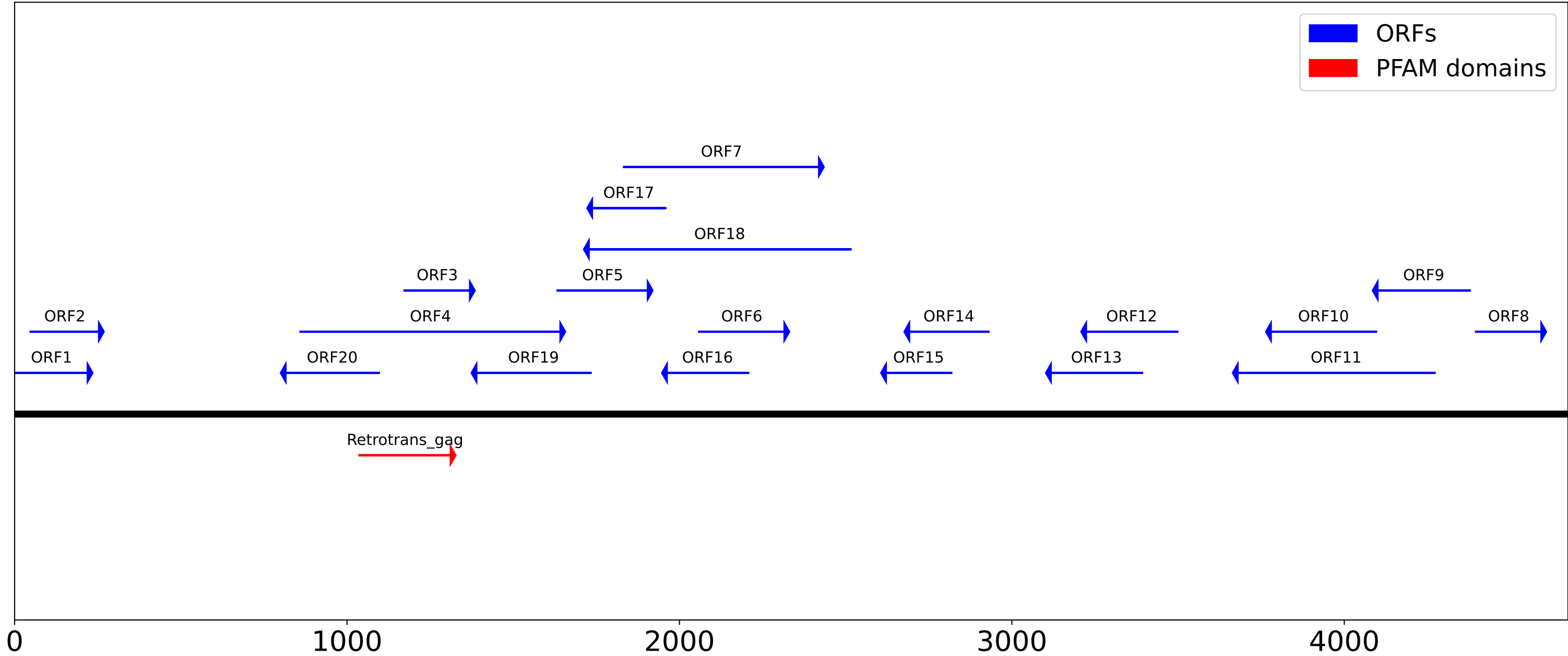
Before TEtrimmer 4673 bp



After TEtrimmer ORF and PFAM domain plot



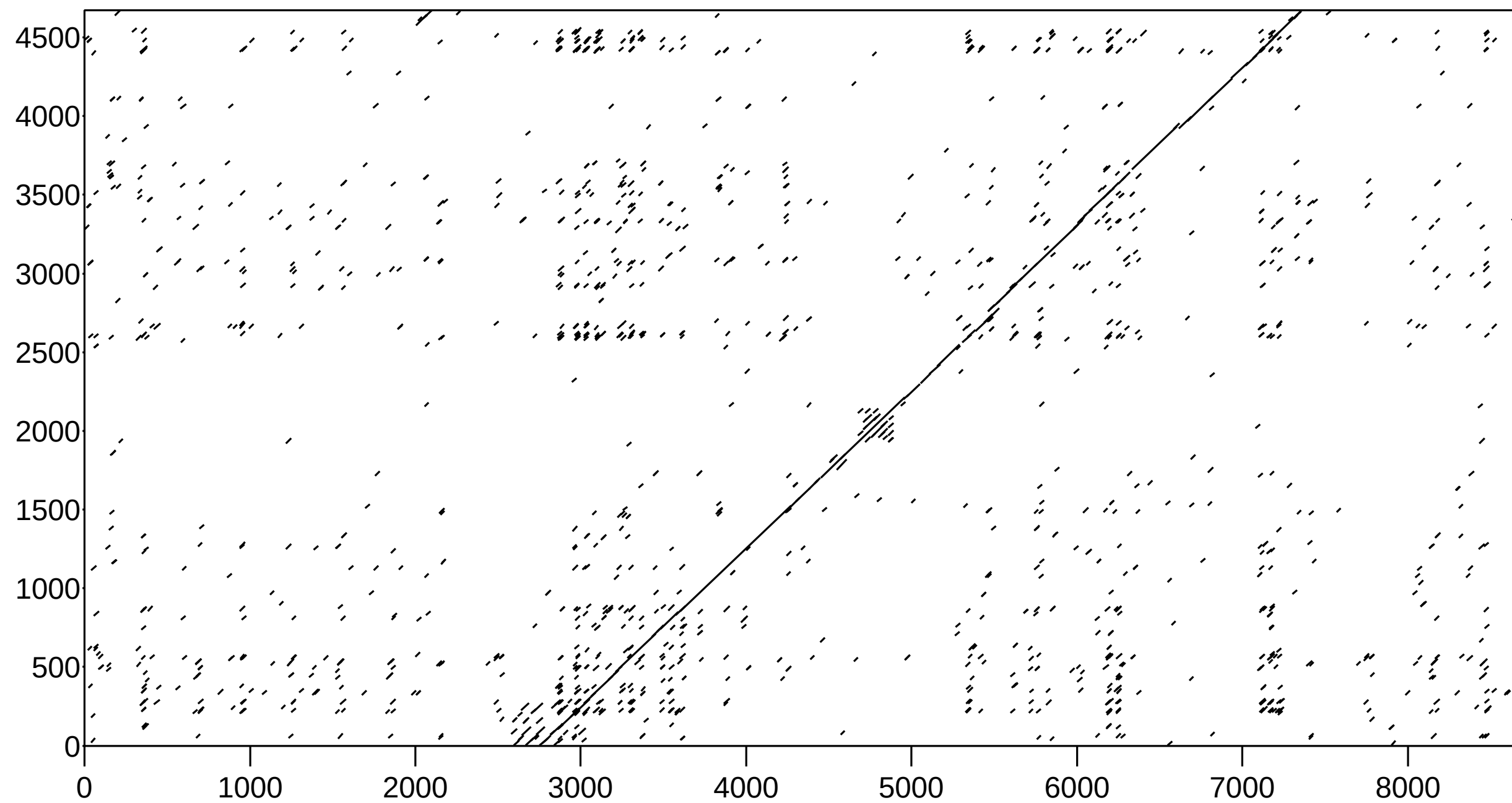
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)