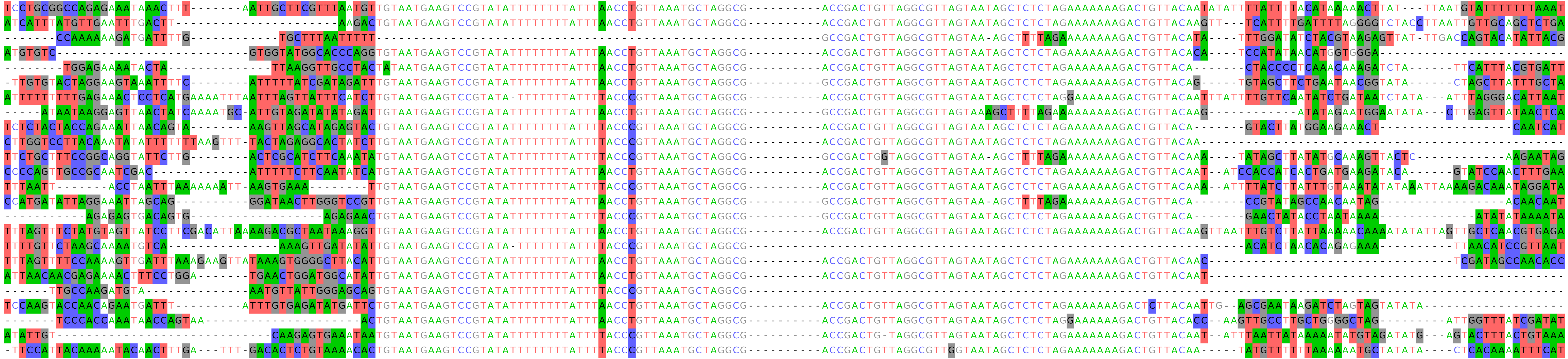


MSA length = 7253

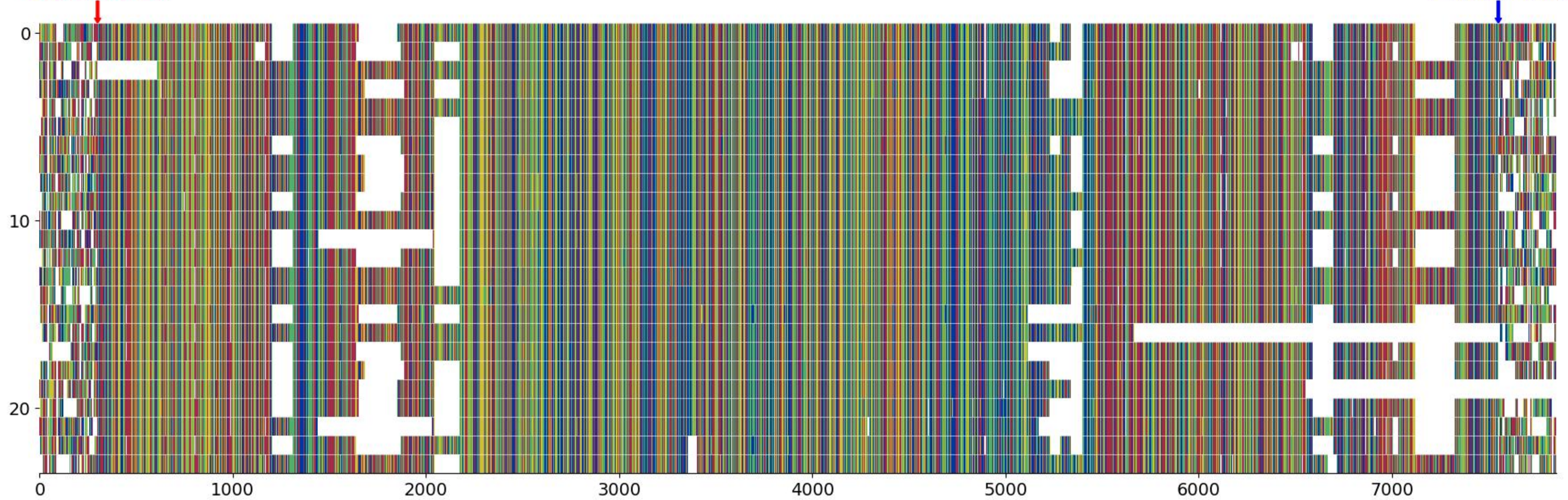
Start crop Point

End crop Point



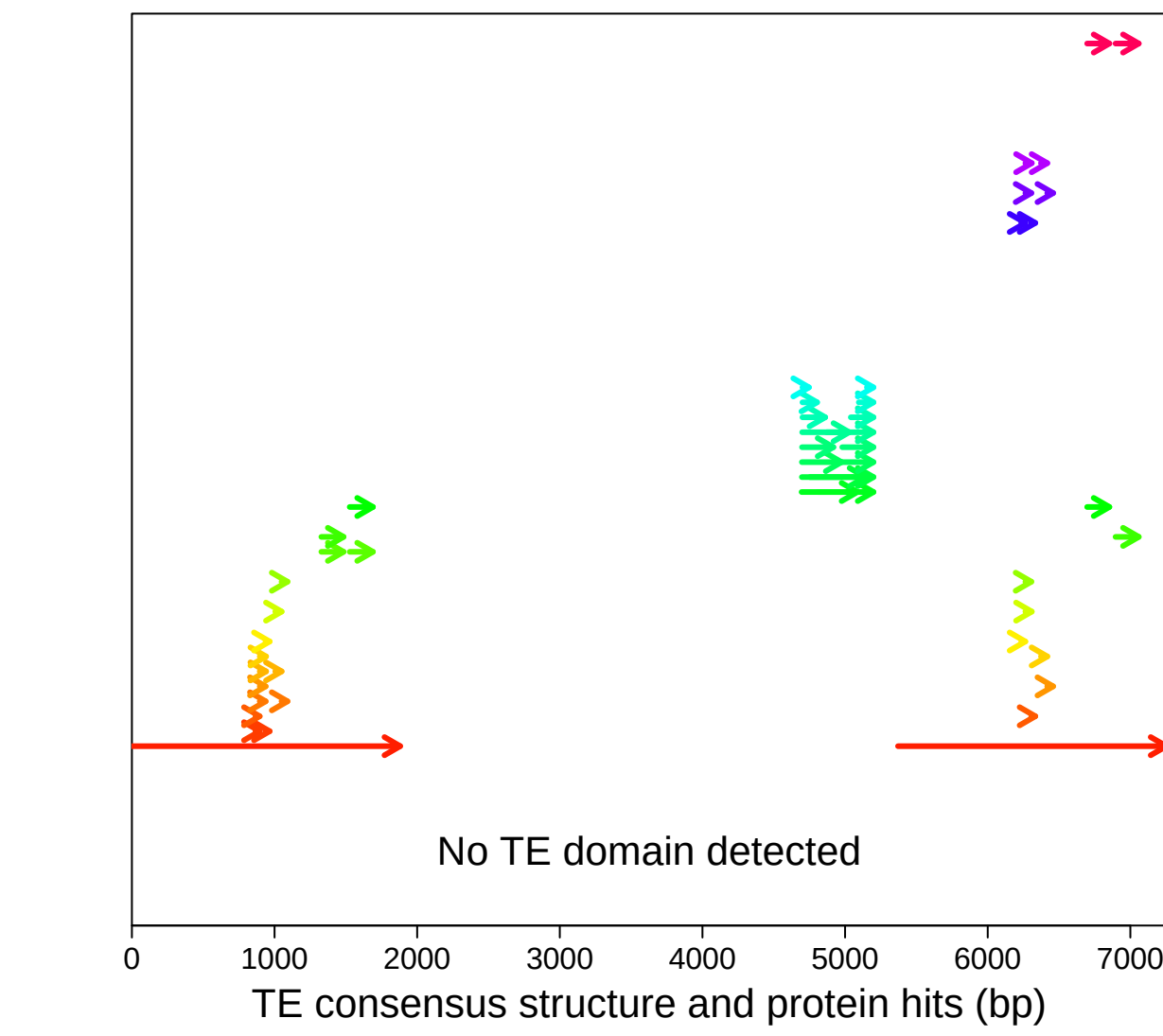
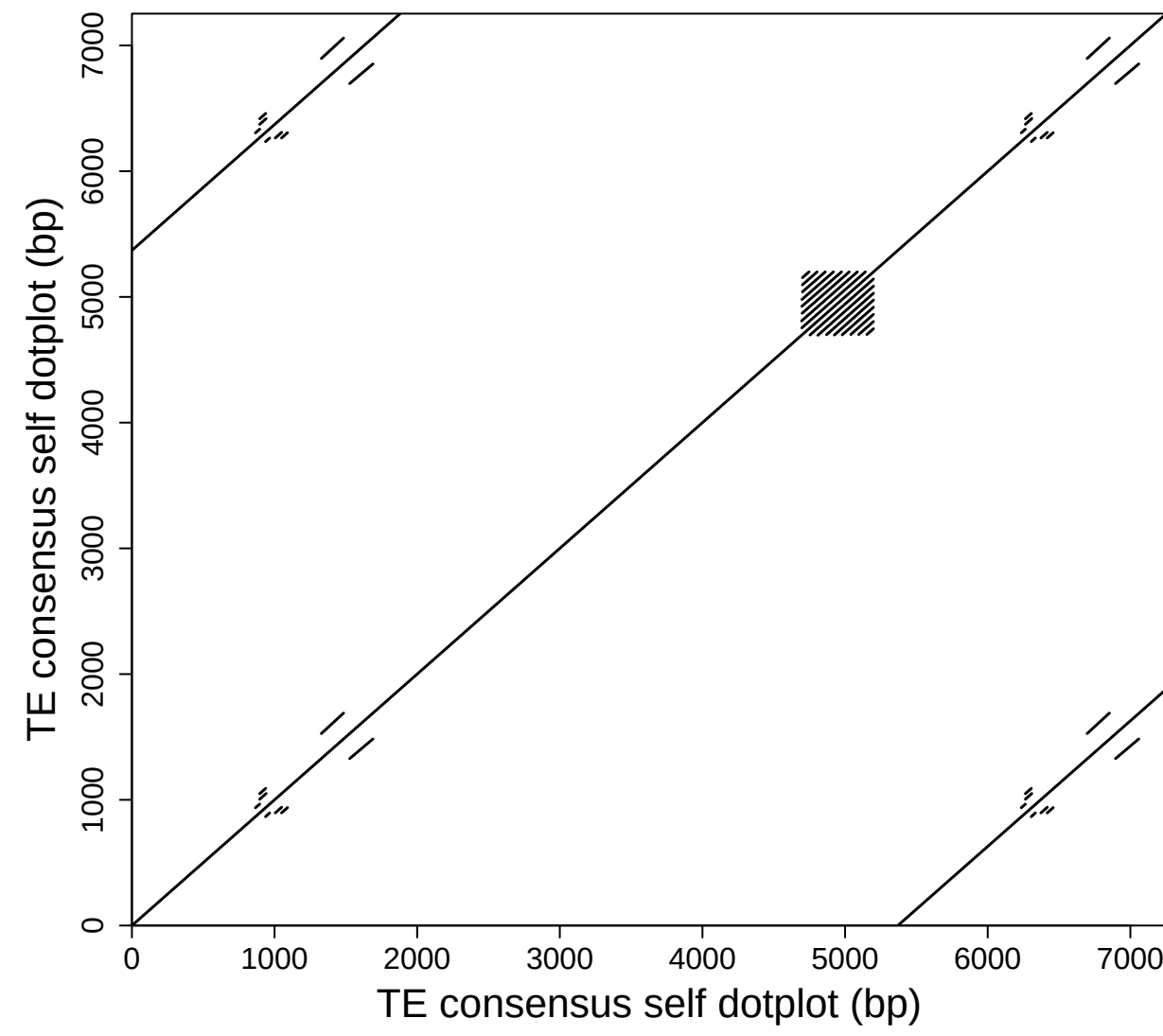
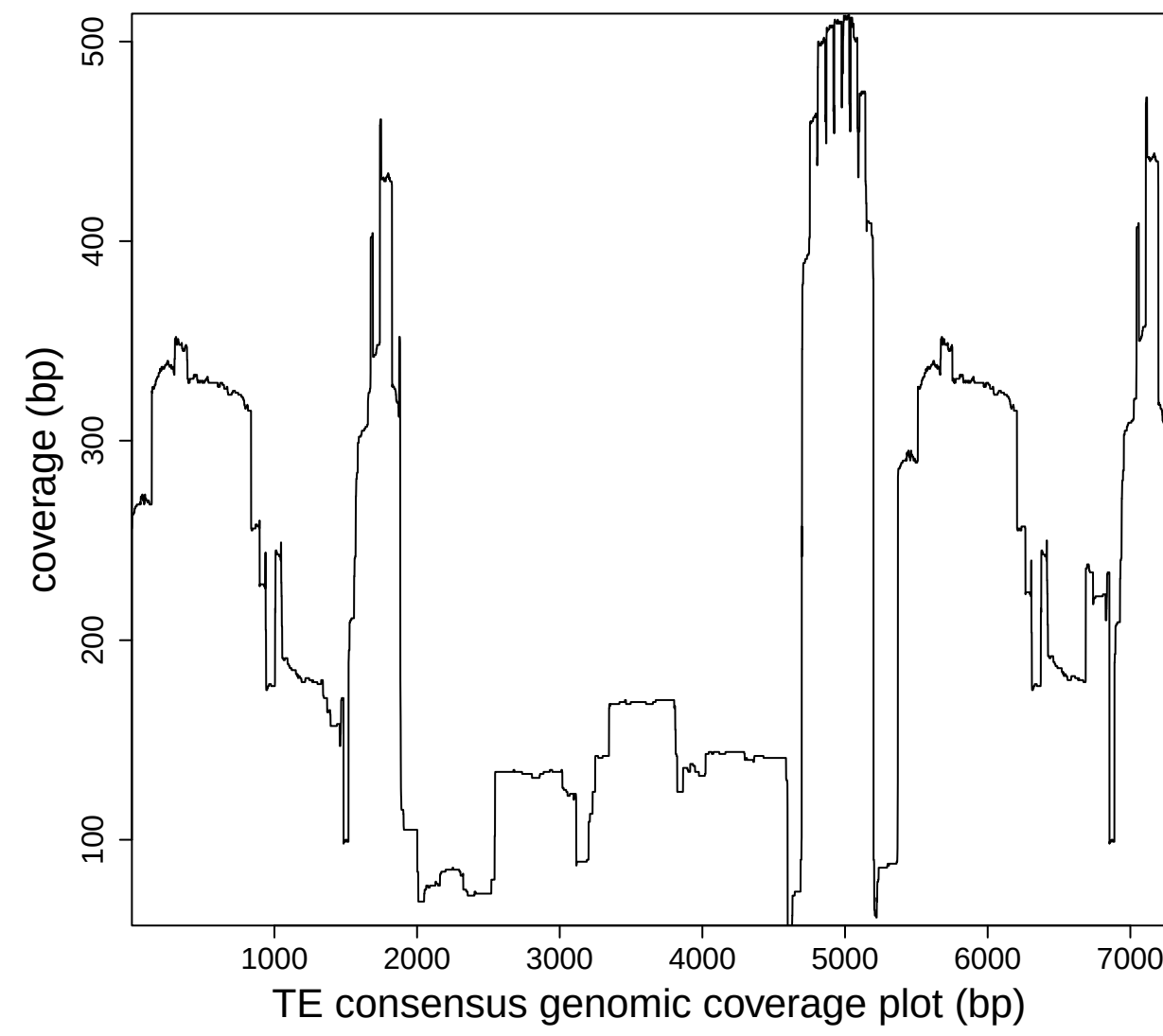
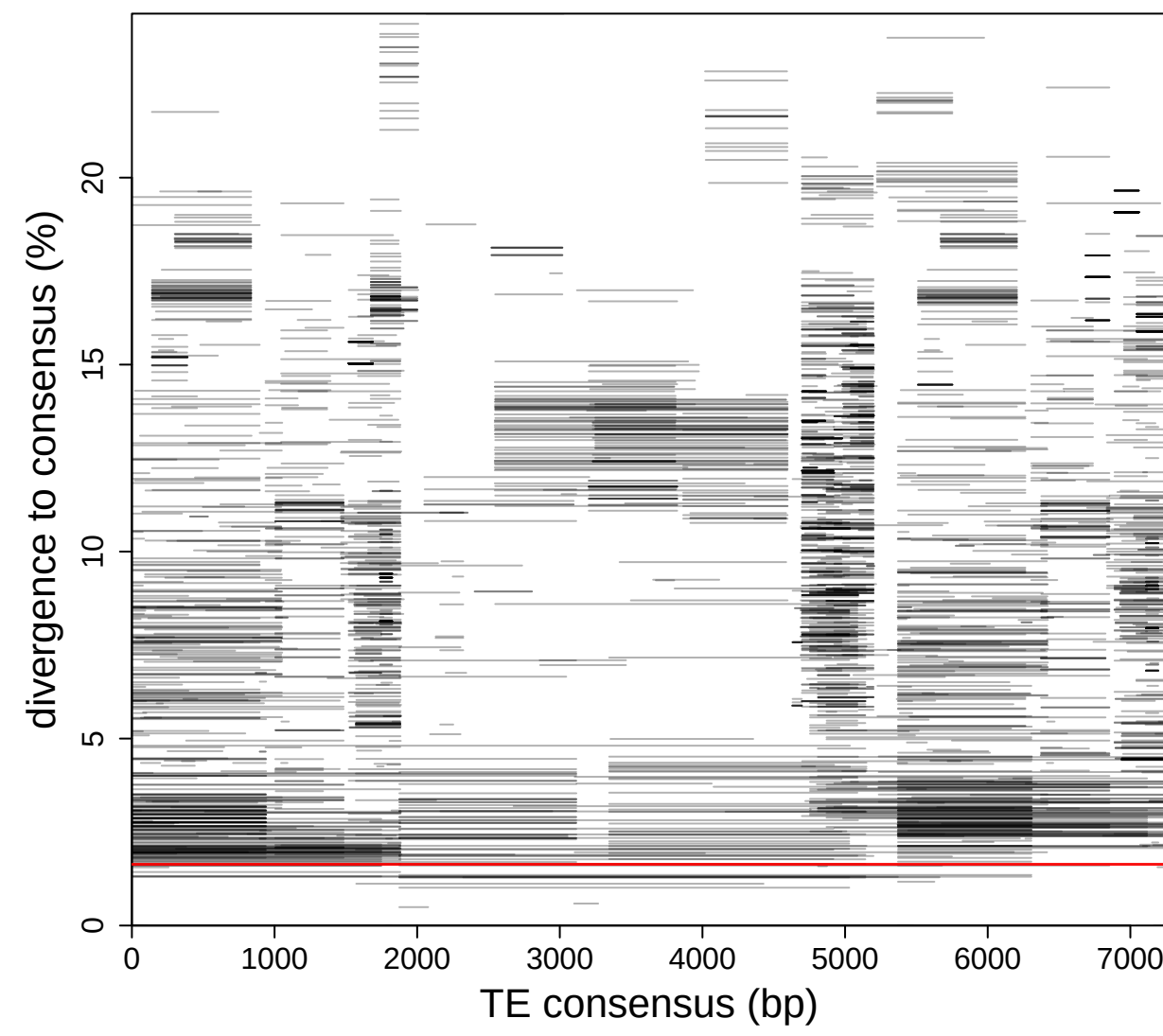
Start crop Point

End crop Point



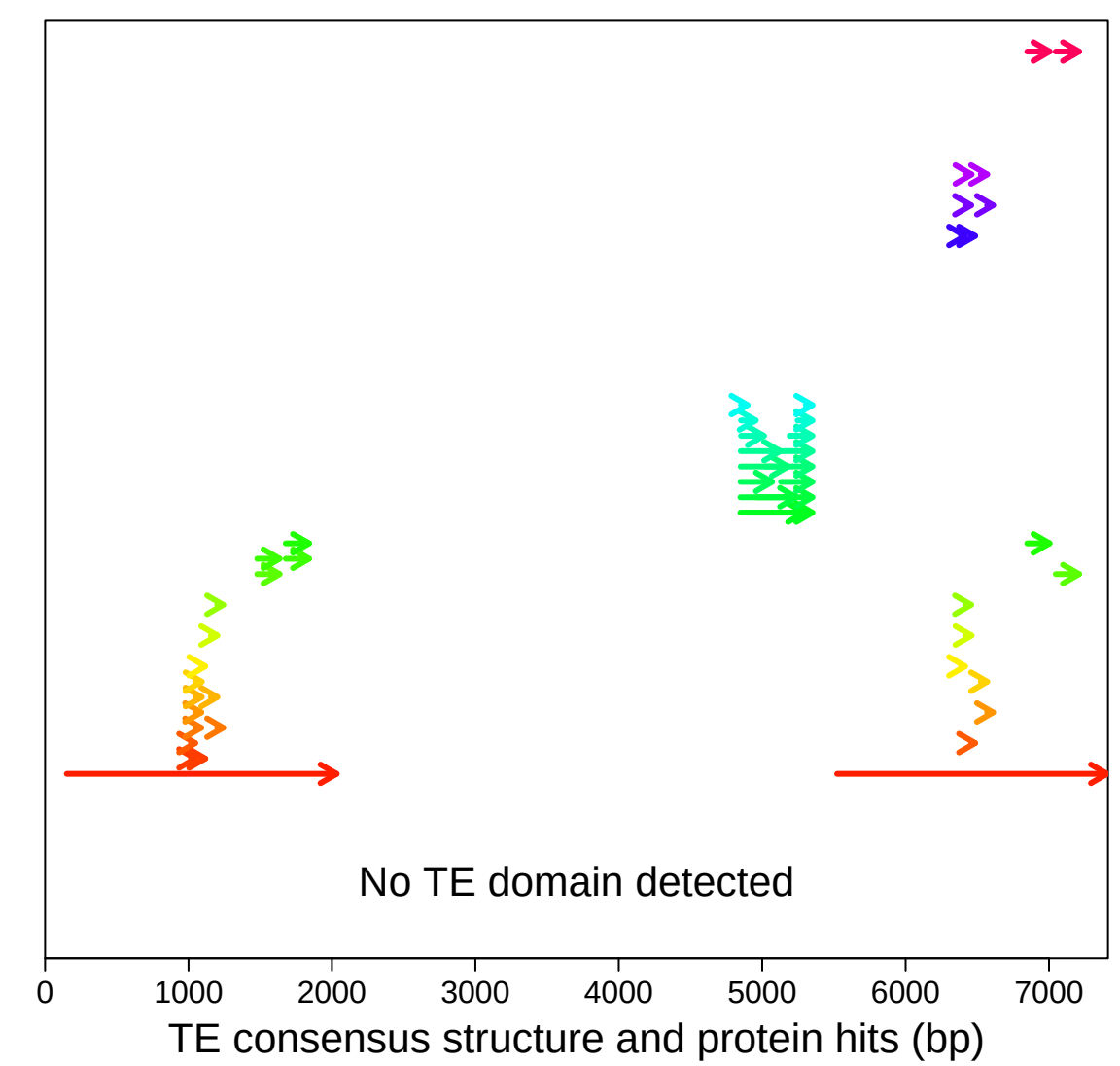
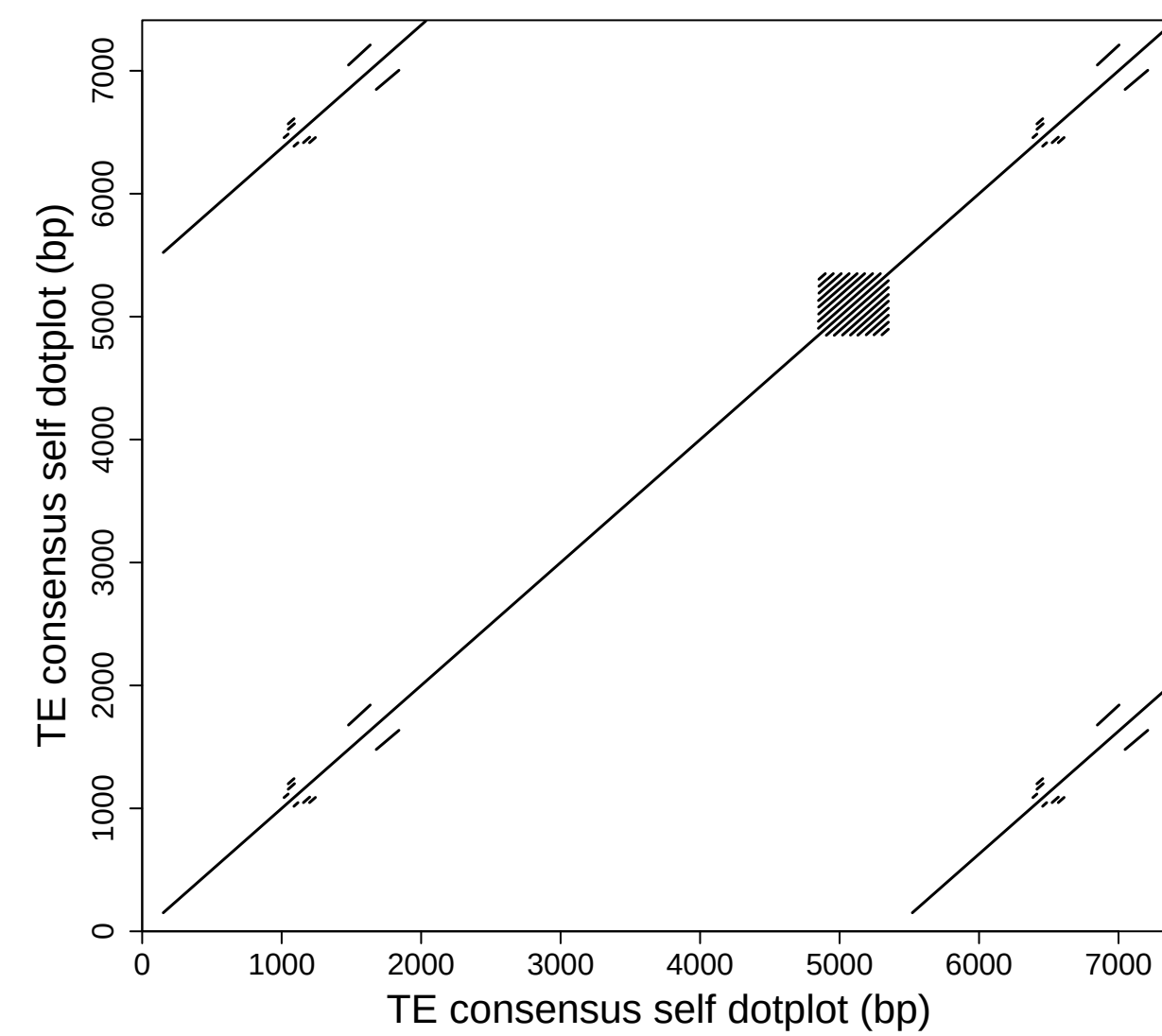
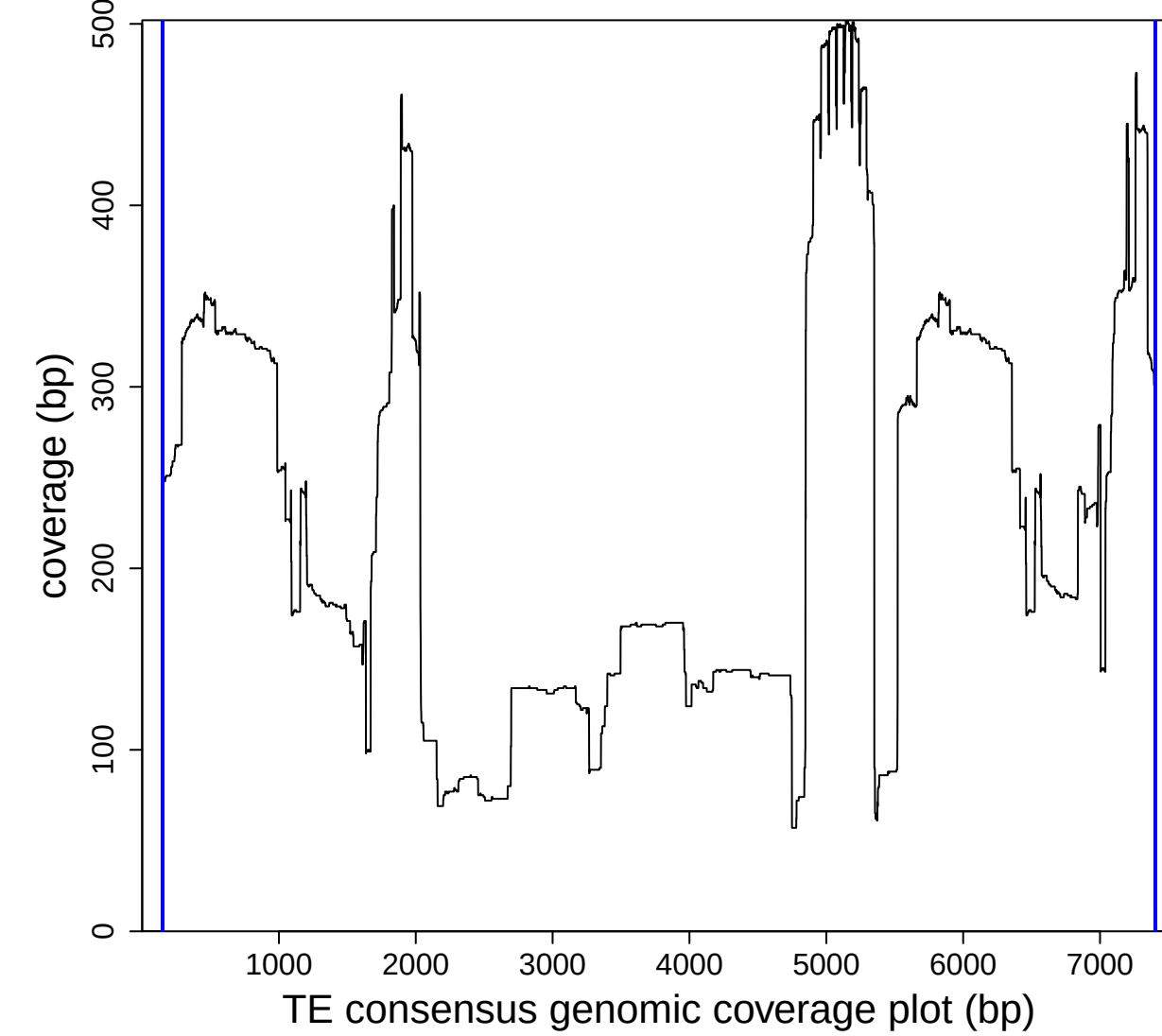
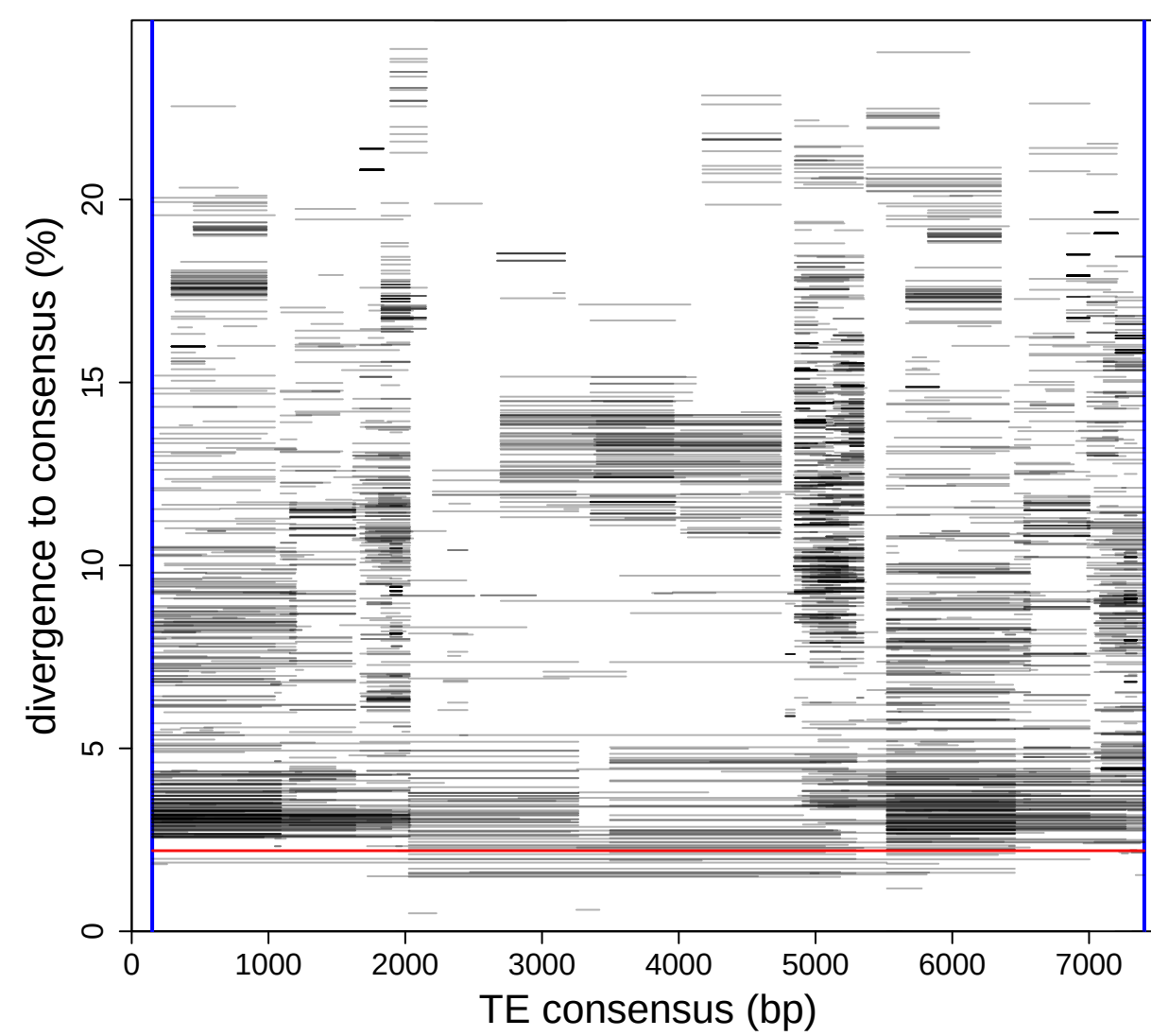
ed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 7253bp; fragments: 3113; full length: 1 (>=6527.7bp)

After TEtrimmer 7253 bp



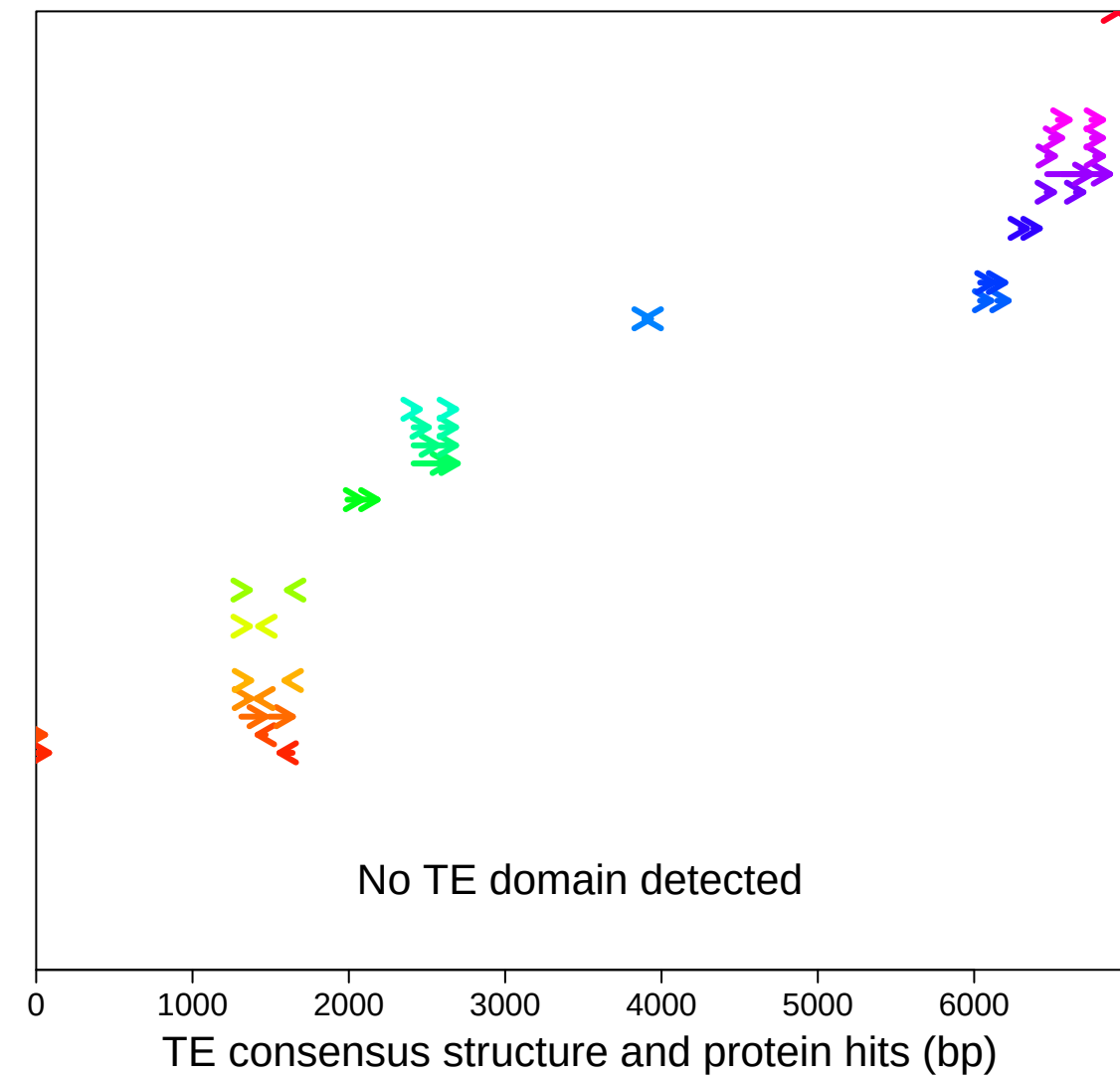
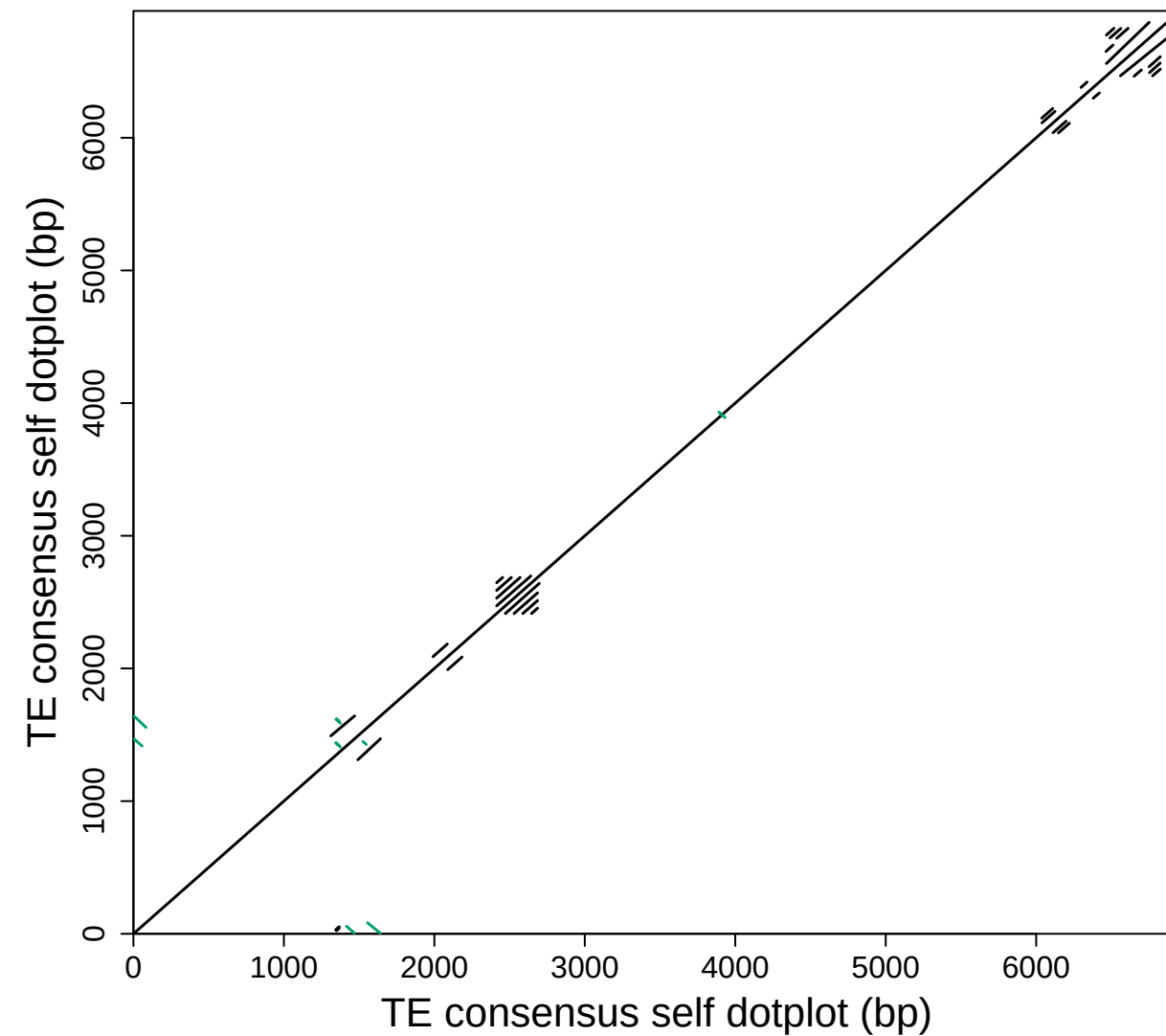
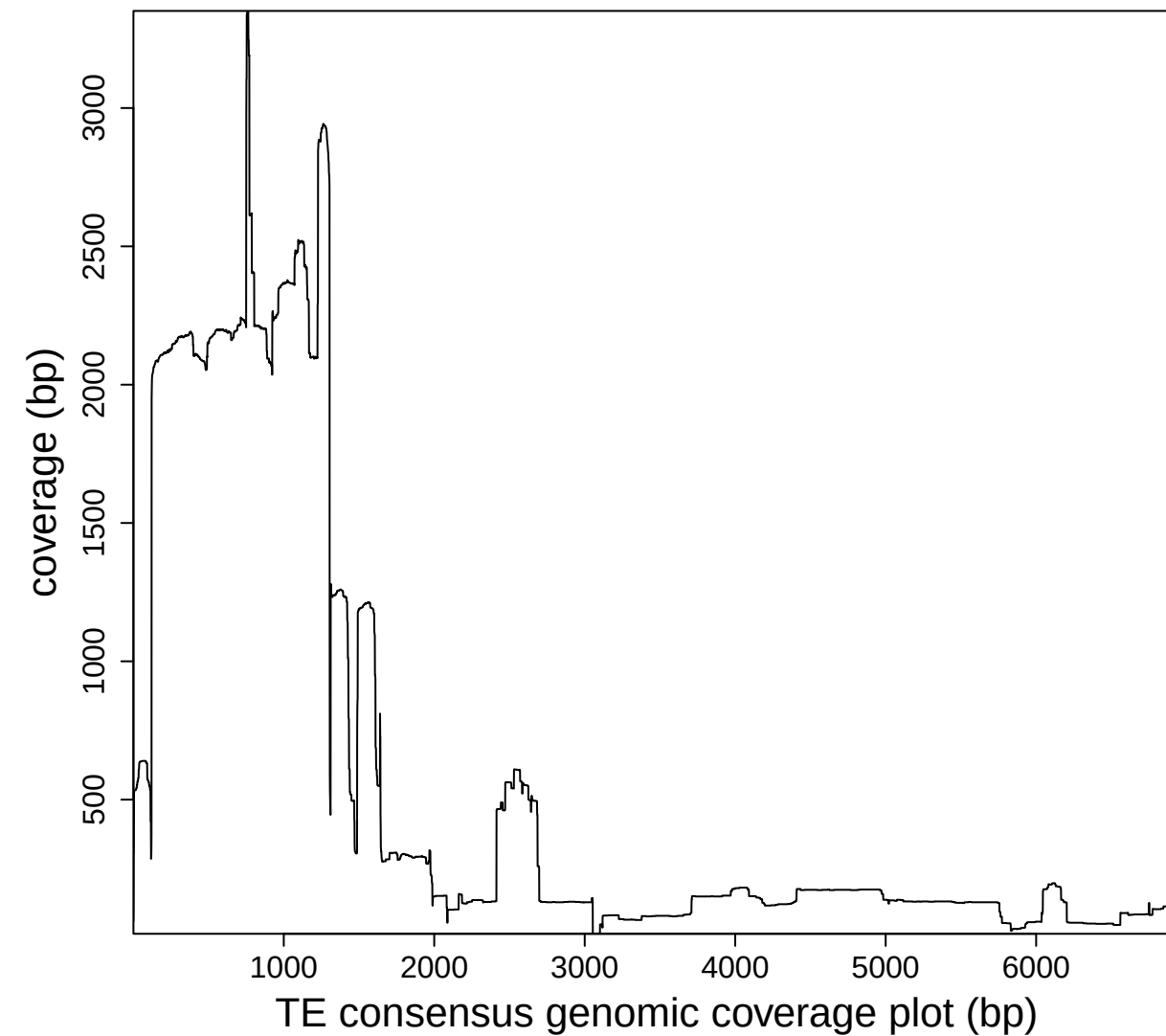
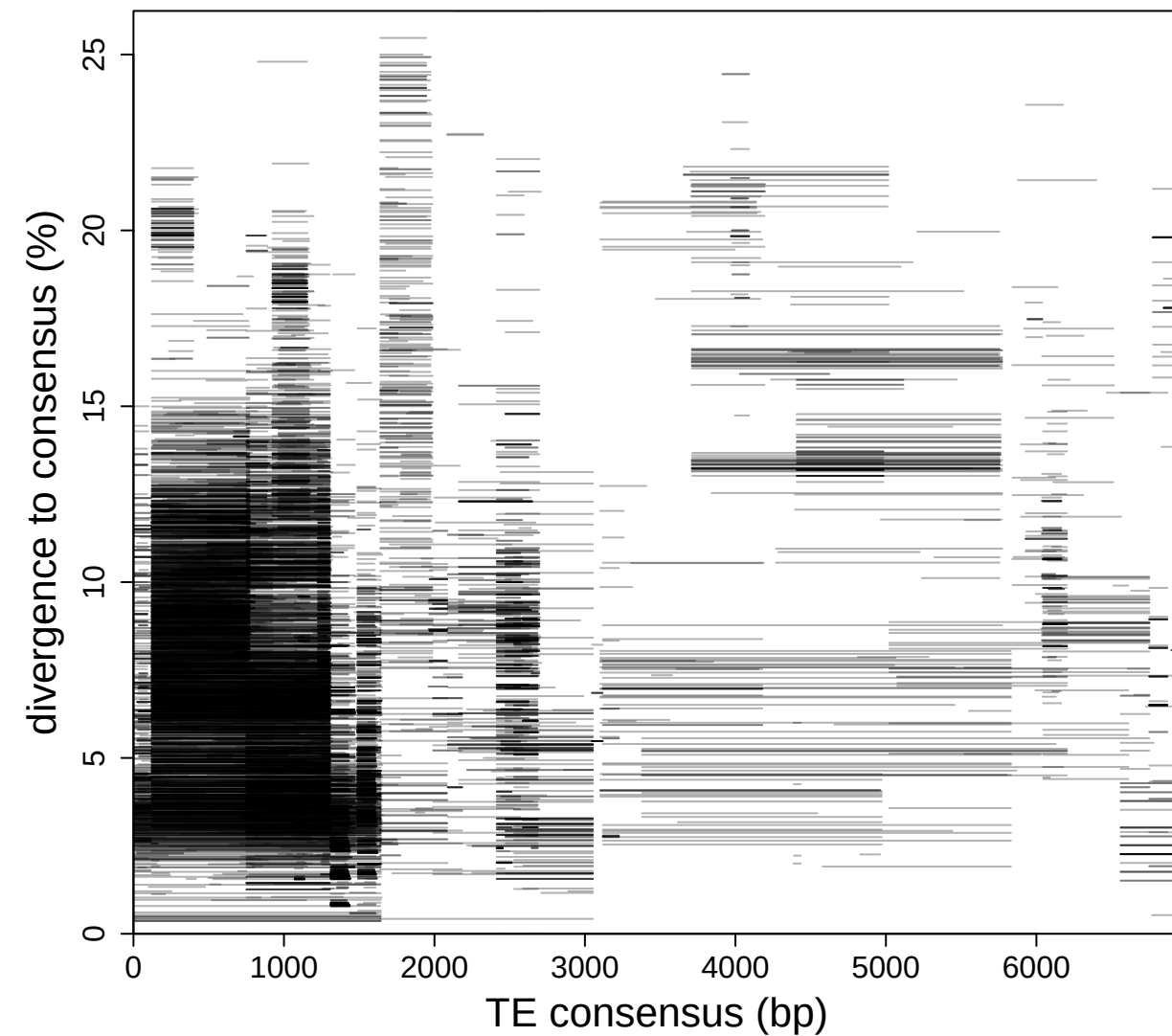
ly_903.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 7554bp; fragments: 3121; full length: 1 (>=6798.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

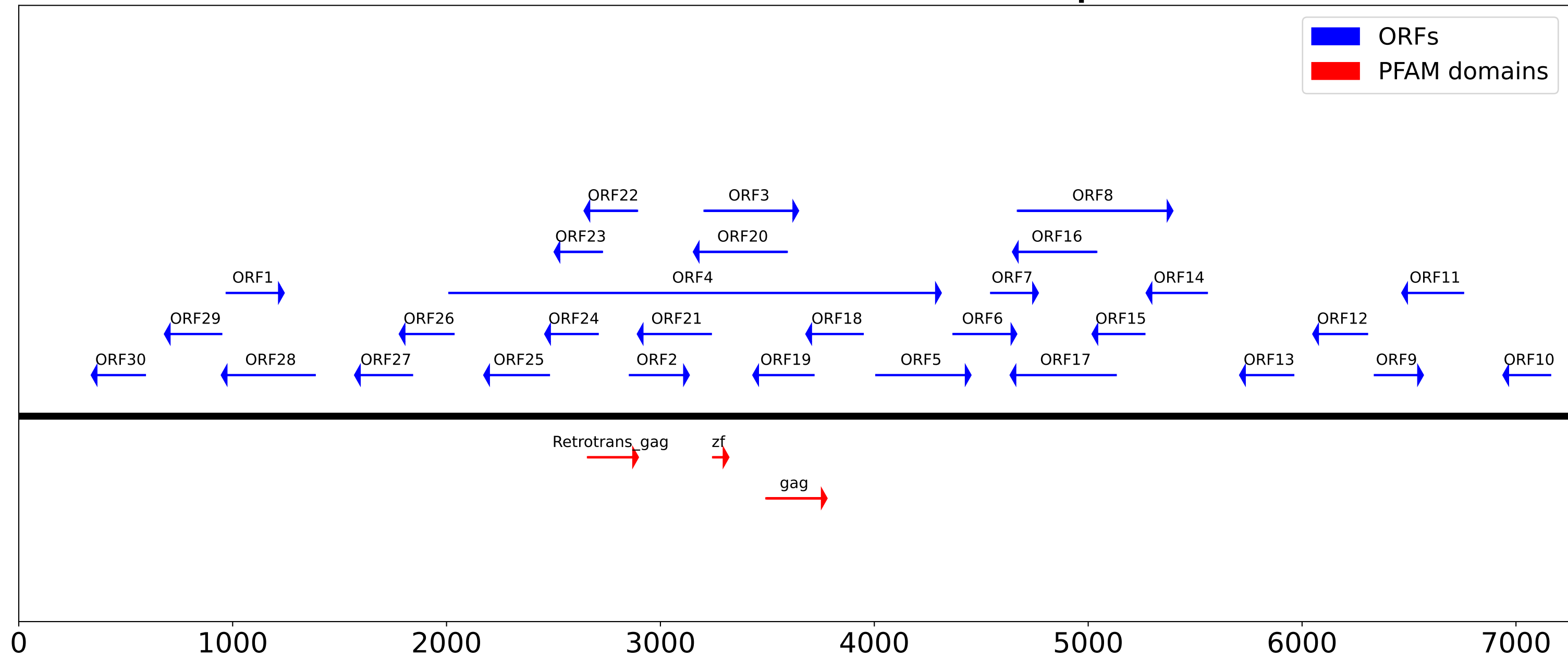


TE: ltr_1_family_903
size: 6956bp; fragments: 9712; full length: 0 (>=6260.4bp)

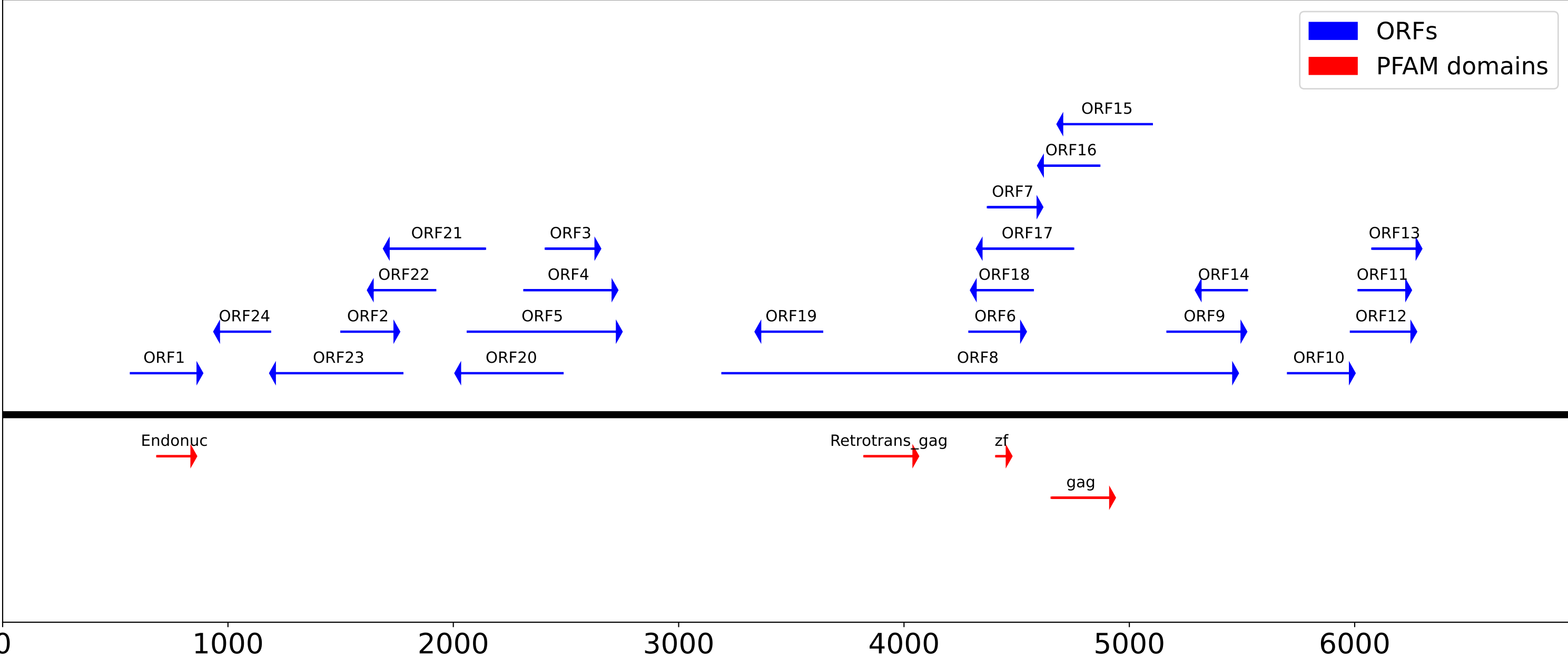
Before TEtrimmer 6956 bp



After TETrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

