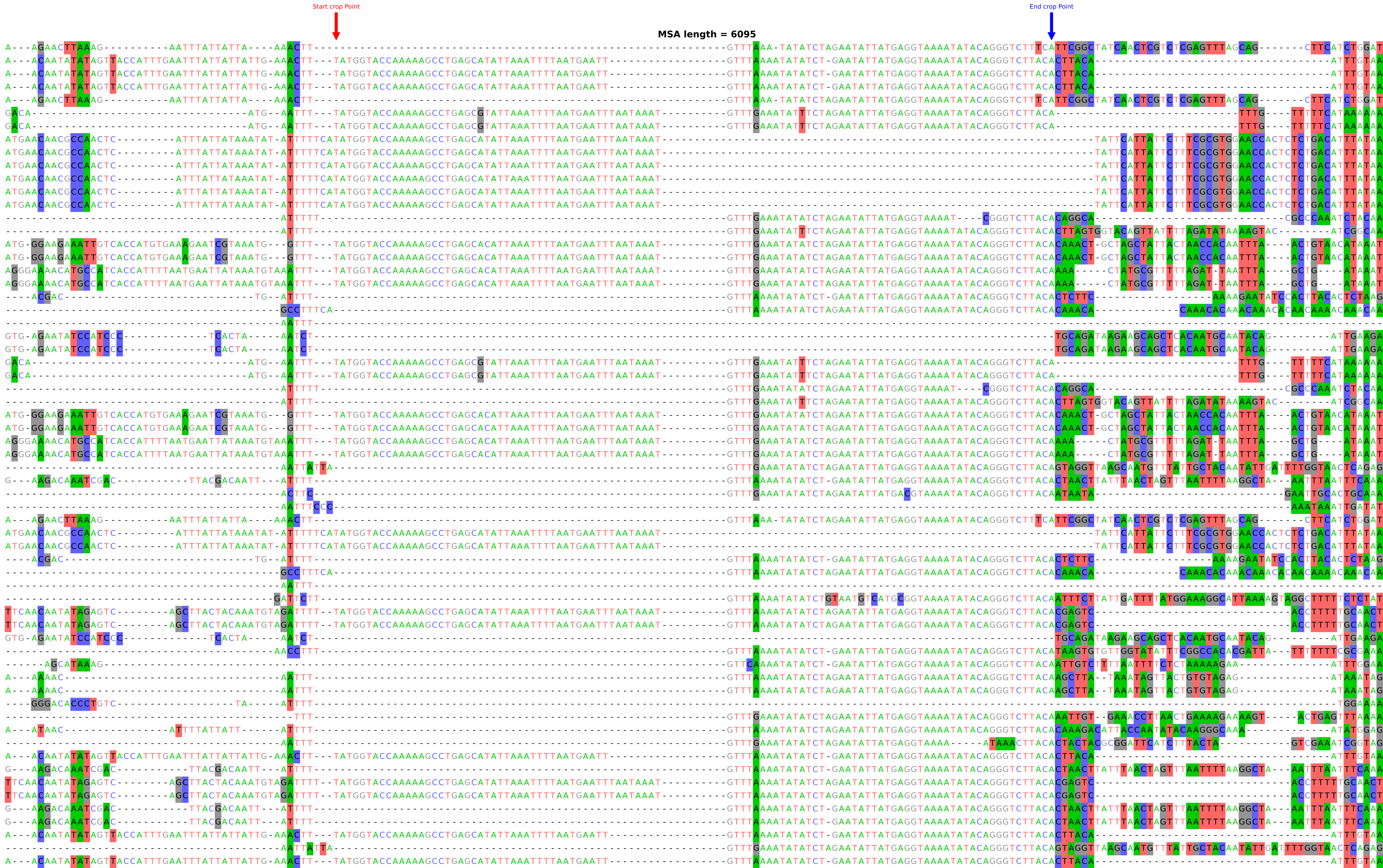
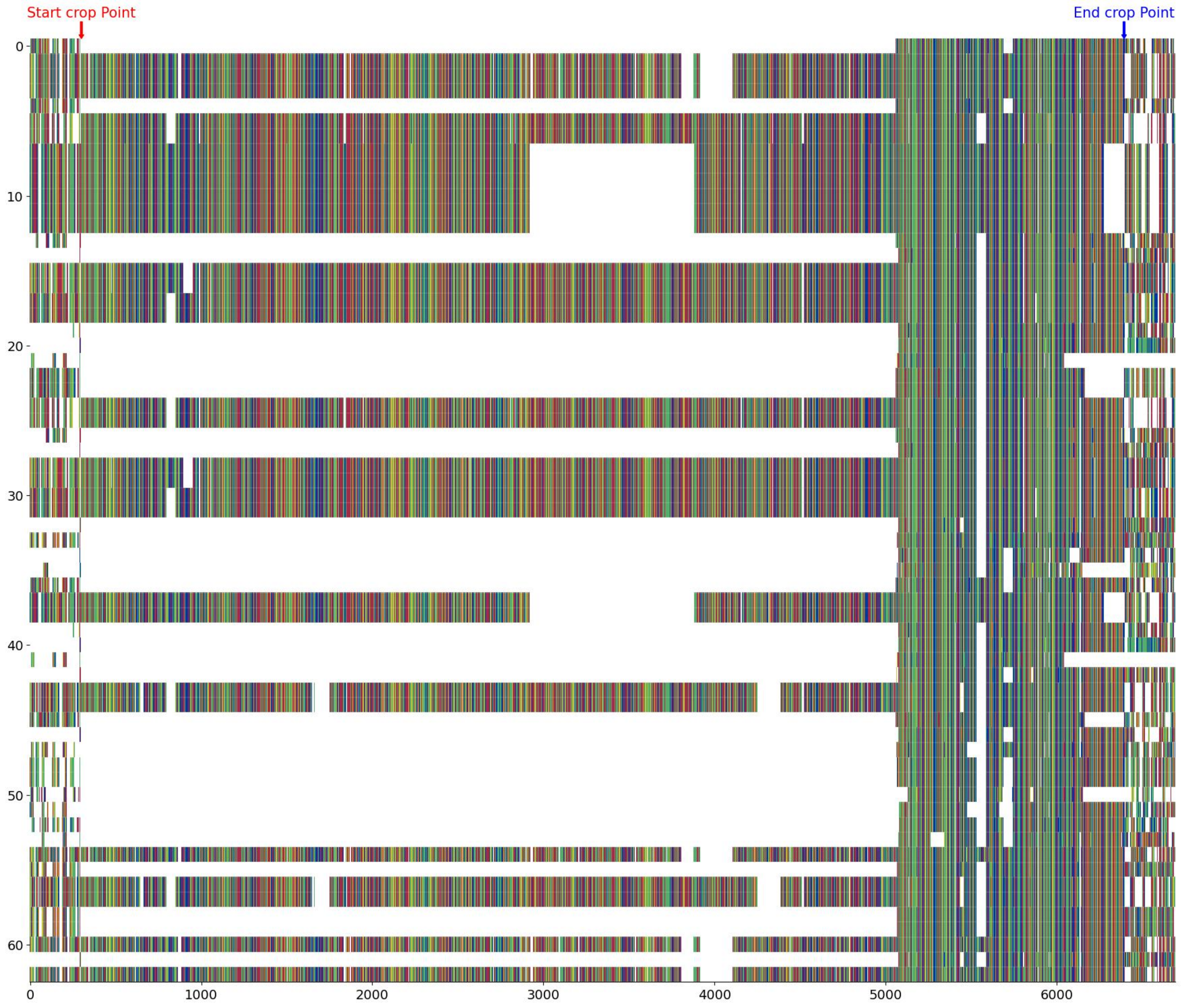
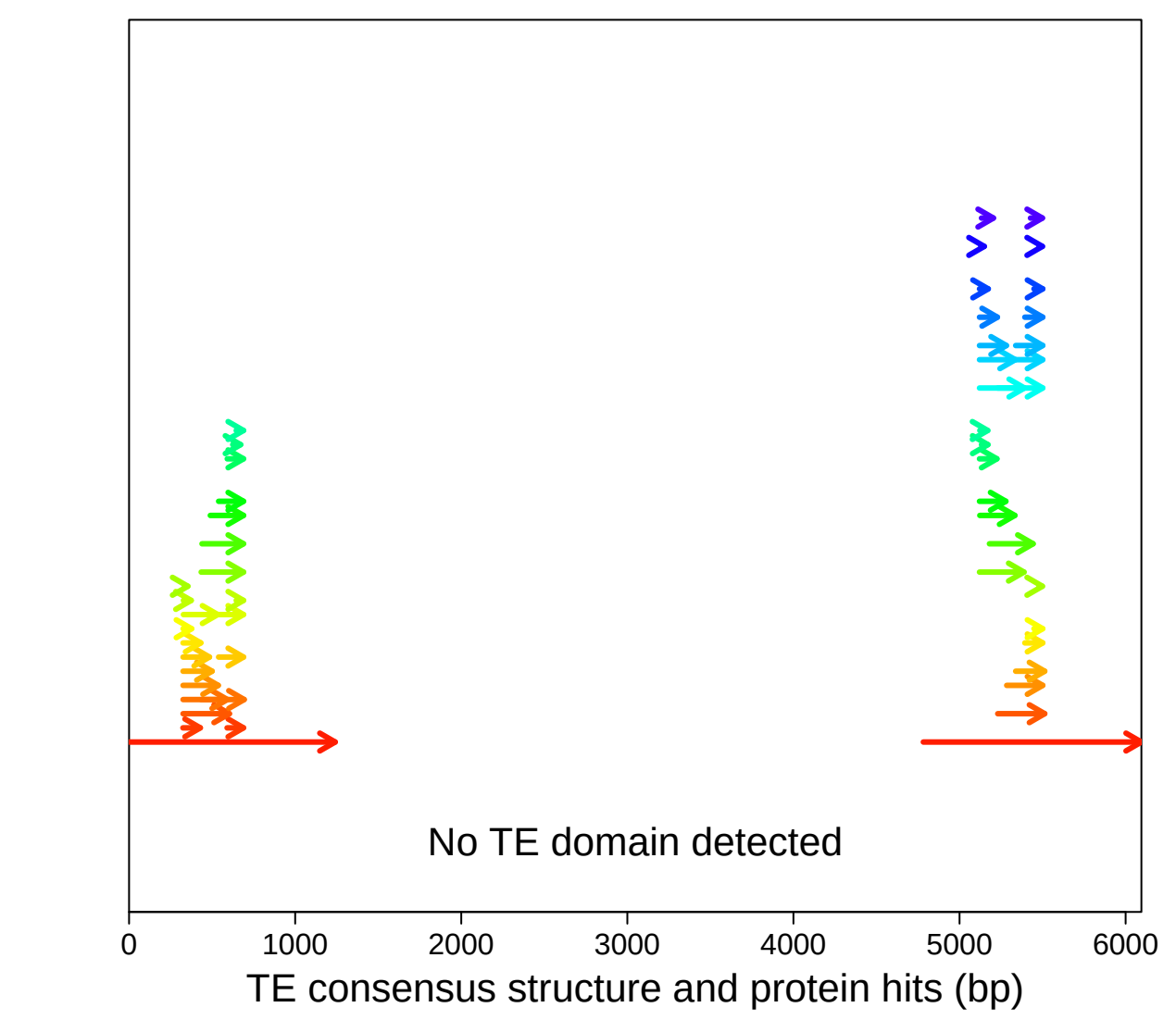
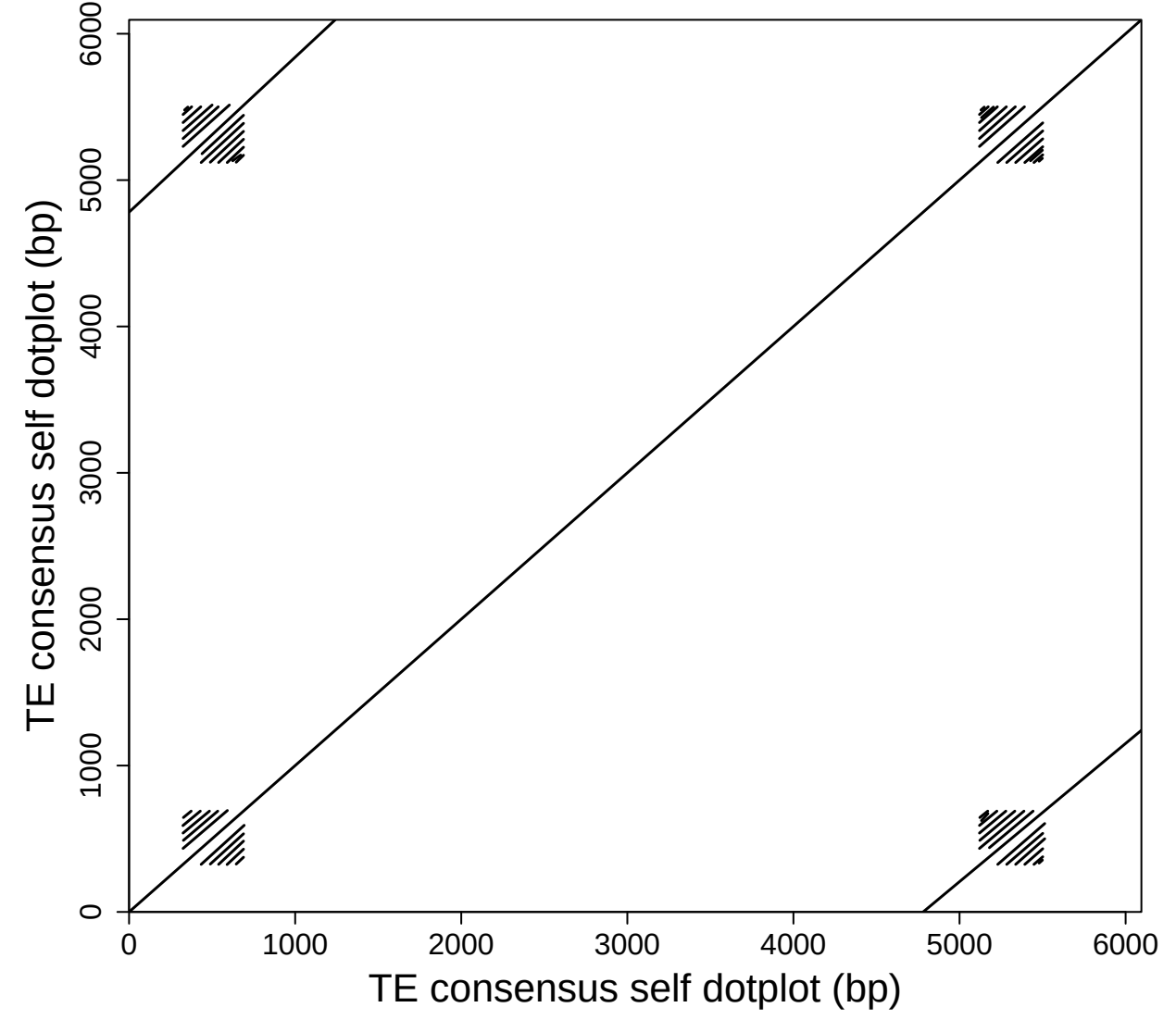
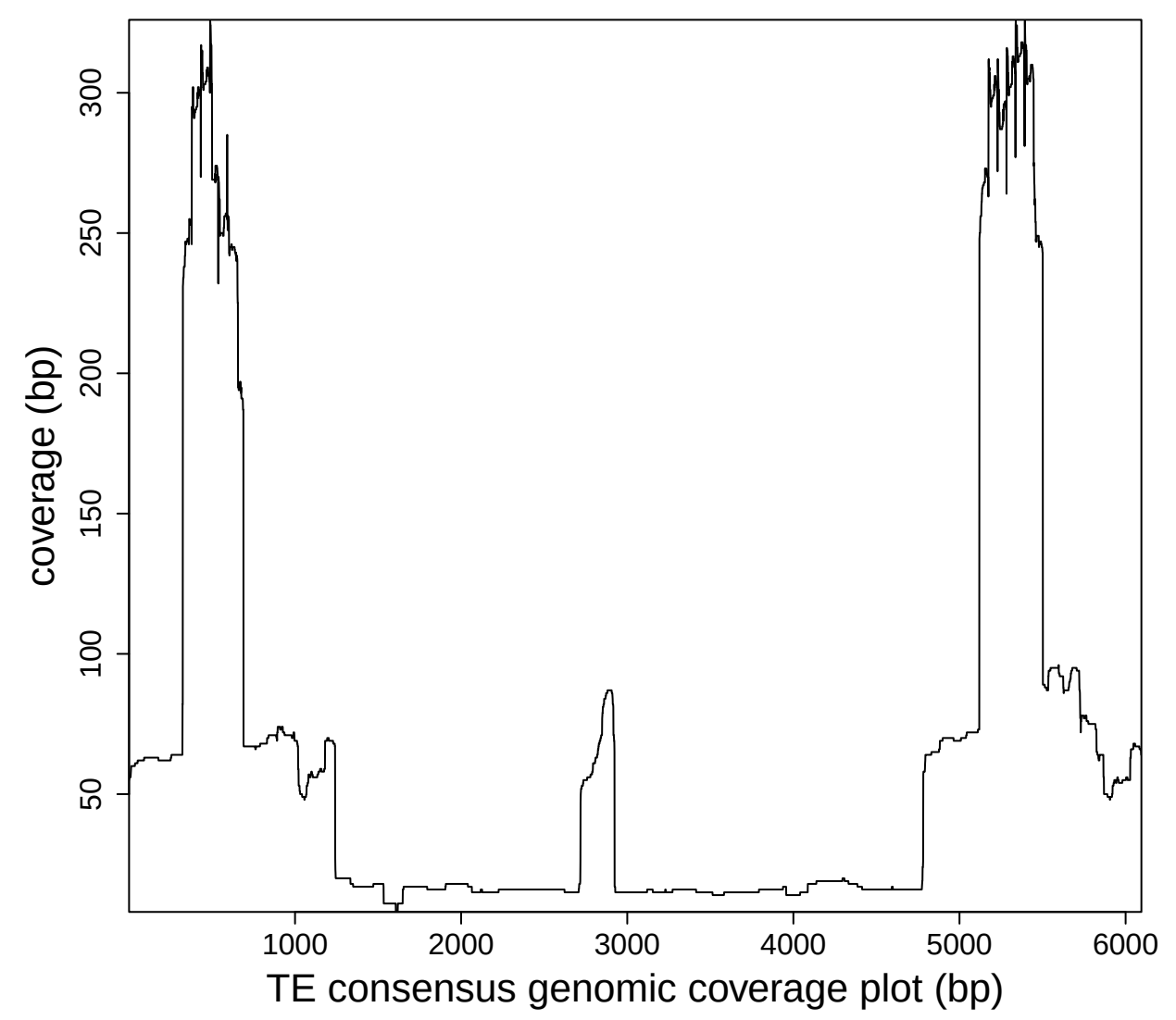
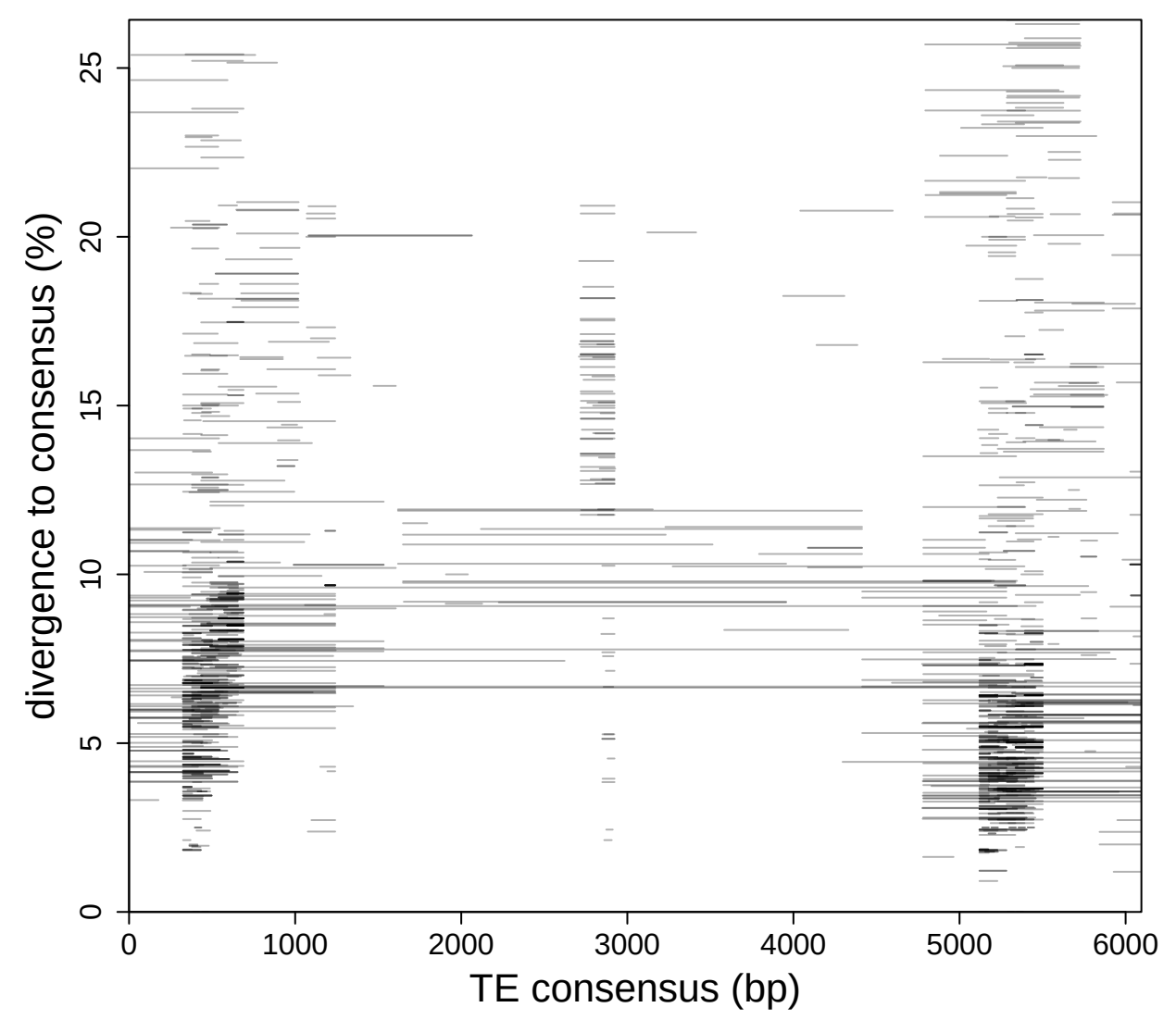






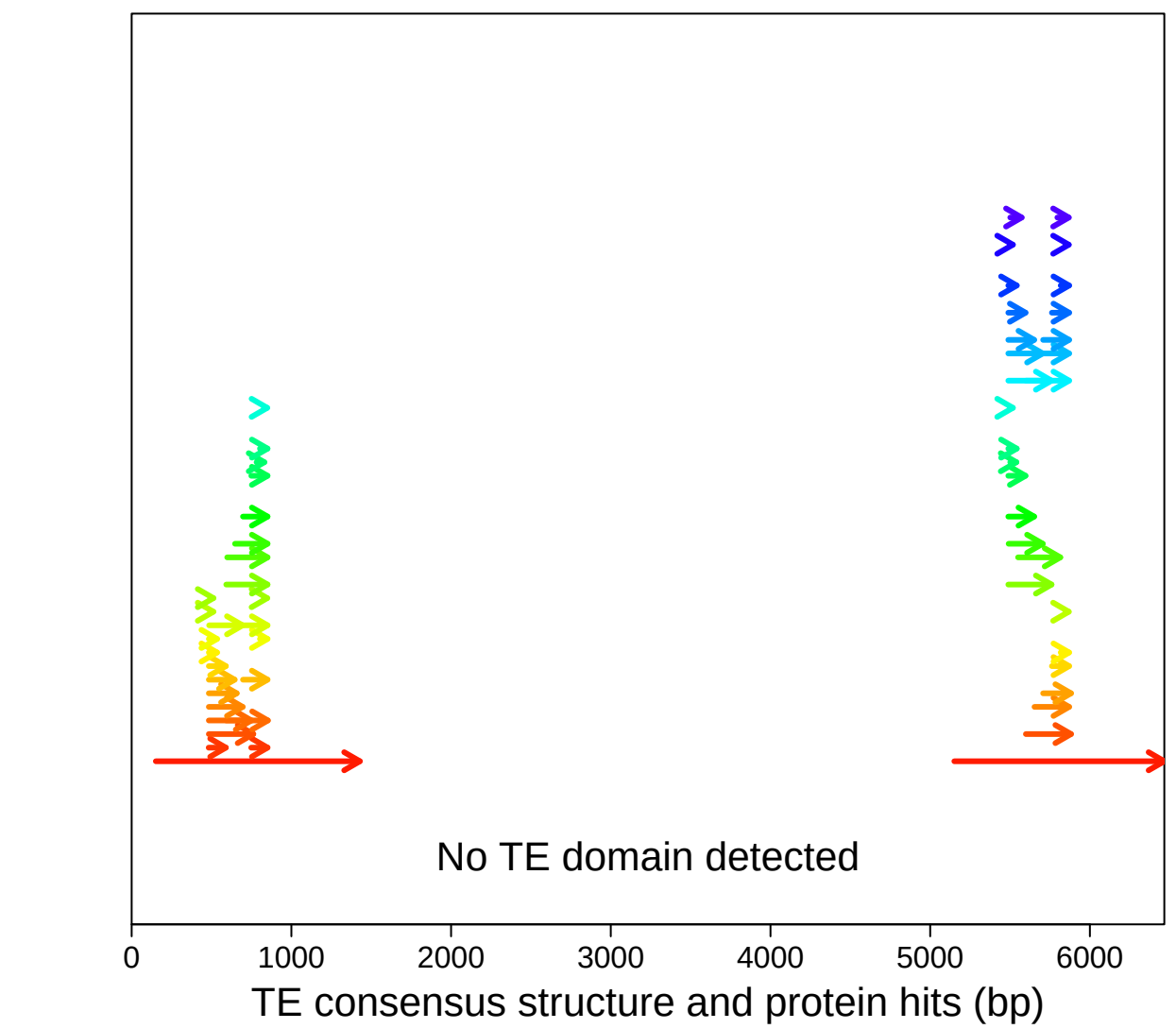
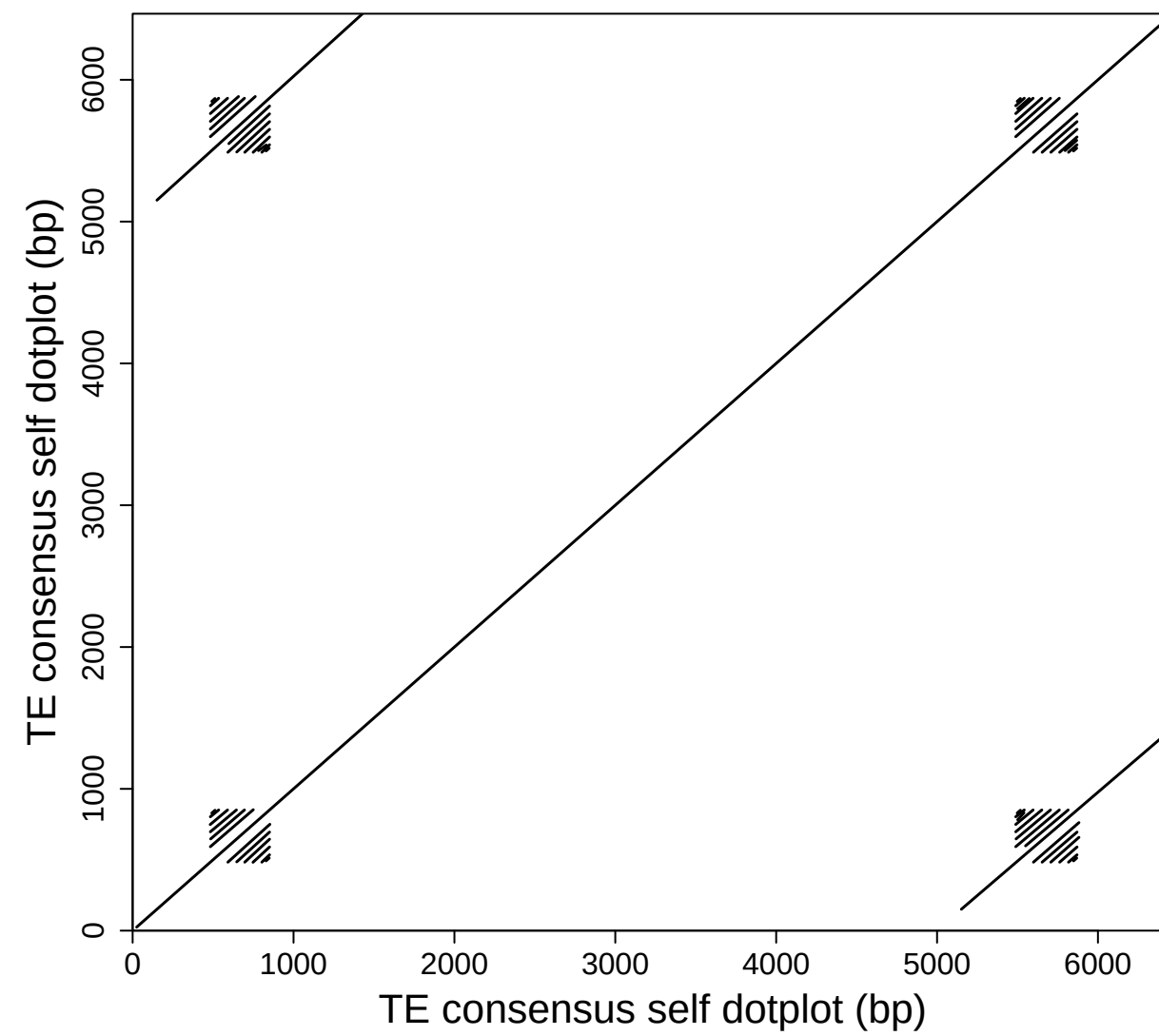
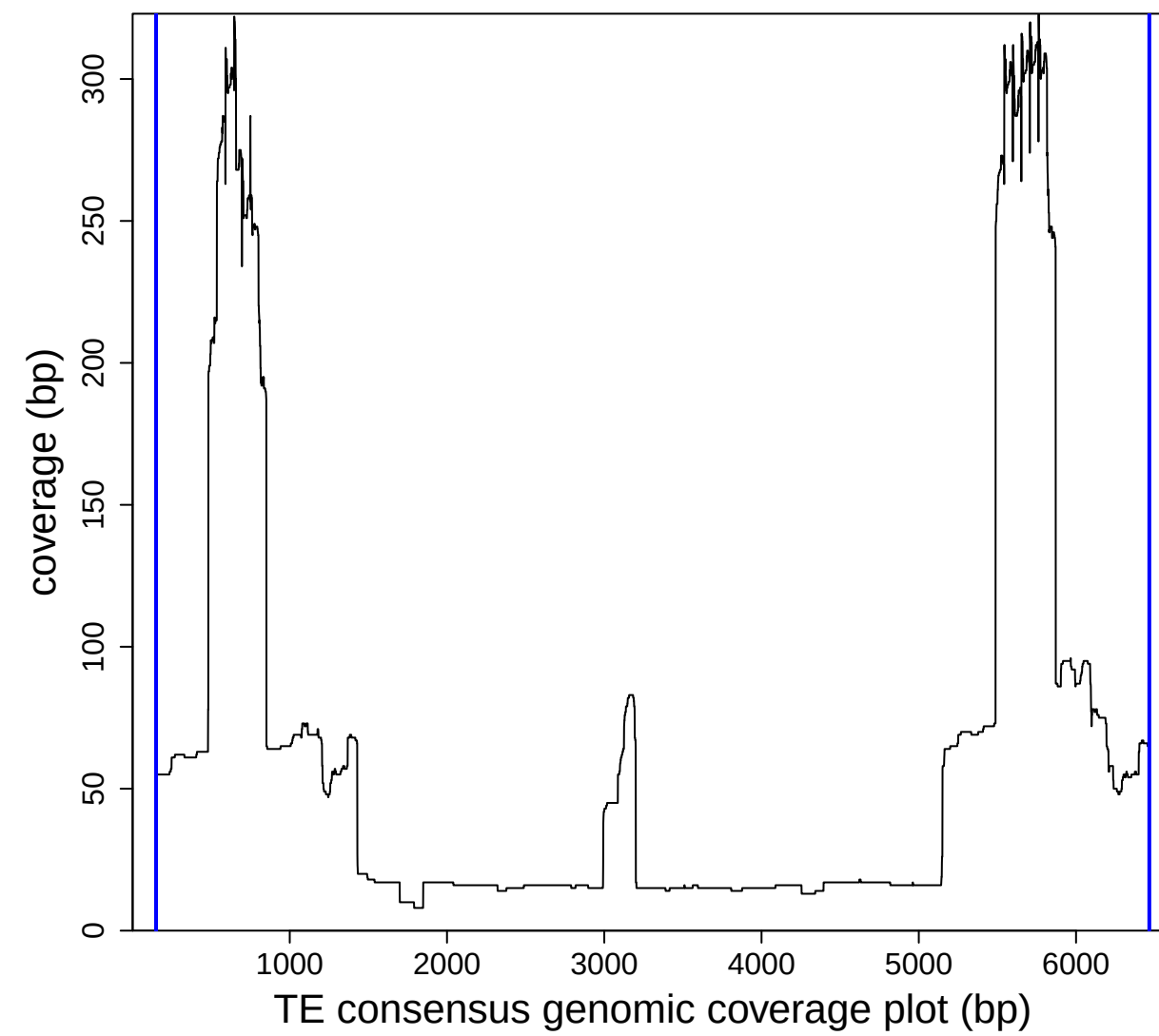
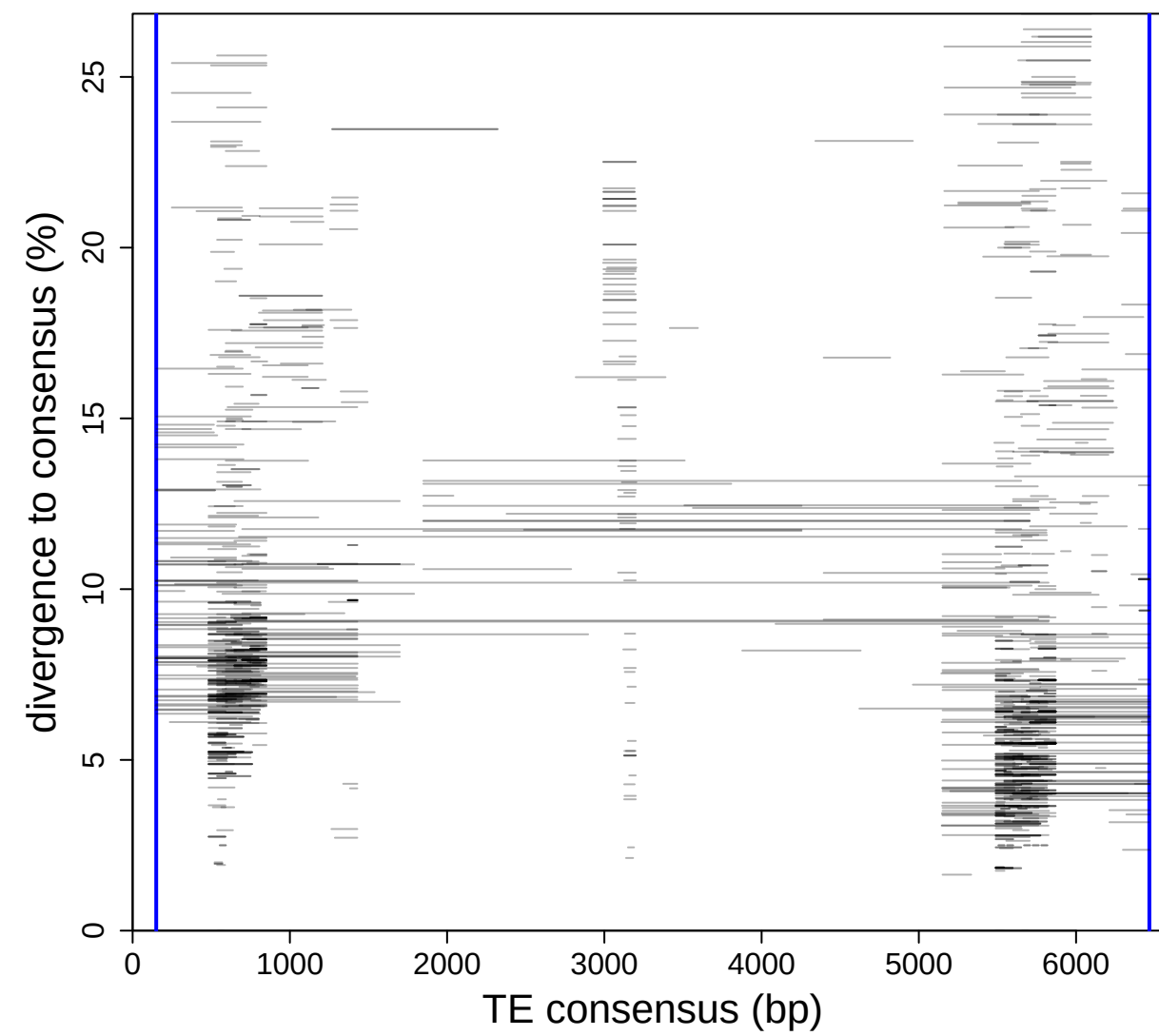
d_u.f.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.f
size: 6095bp; fragments: 1373; full length: 0 (>=5485.5bp)

After TEtrimmer 6095 bp

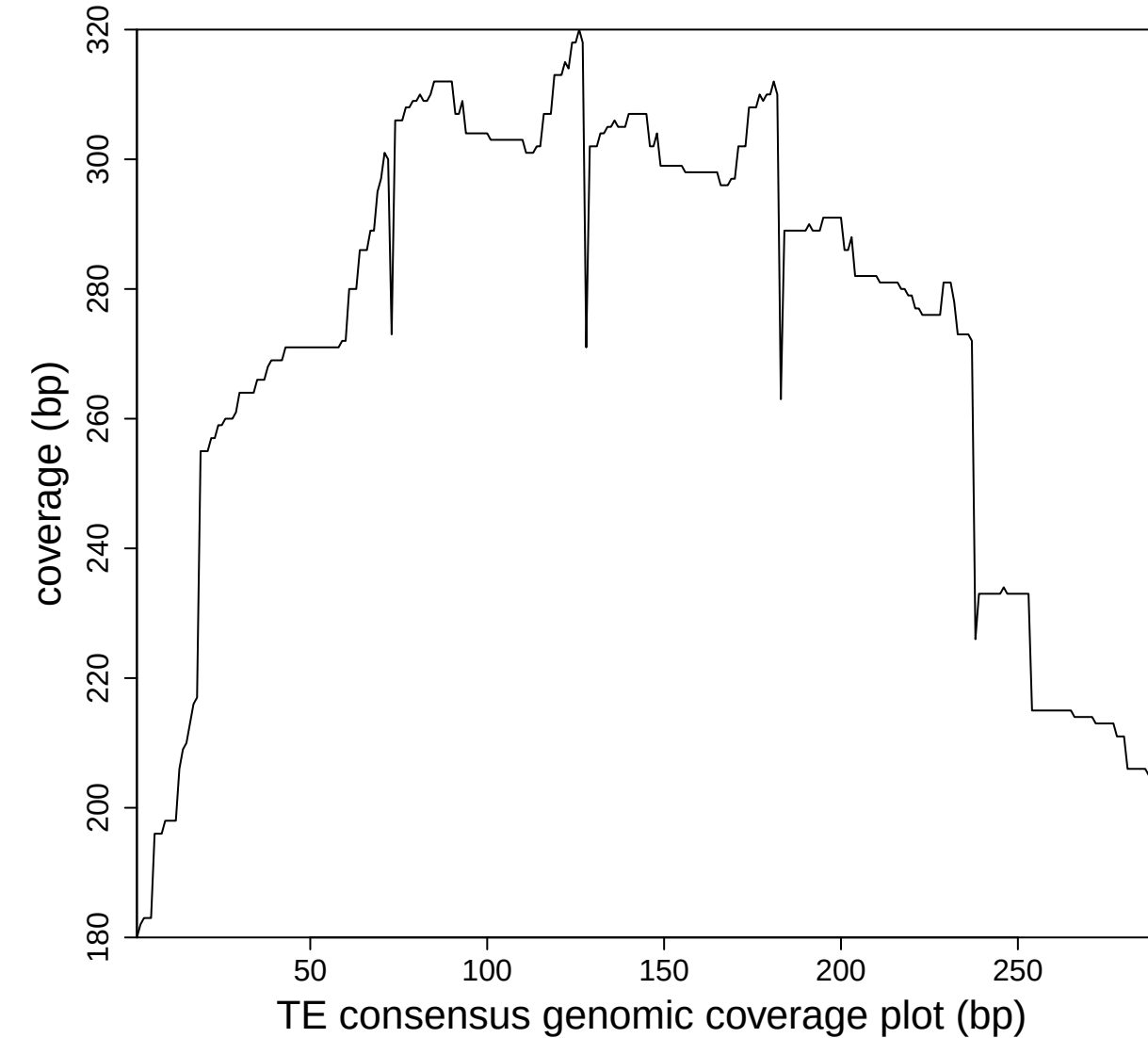
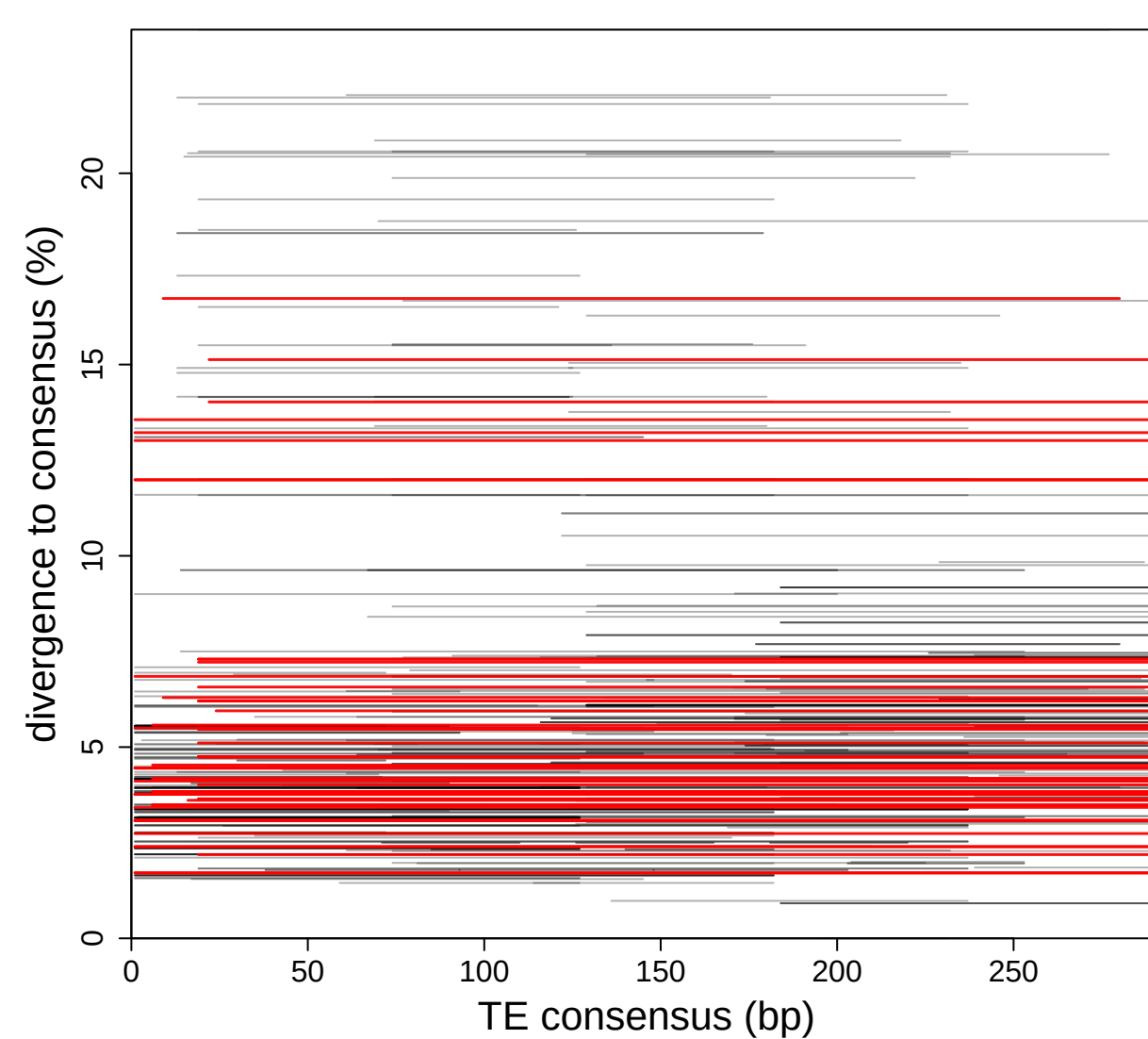


1.fasta.b.bed uf.bed g 1.bed fm 1.bed 0 0 bcIn.fa aln.fa cl.
size: 6616bp; fragments: 1314; full length: 0 (>=5954.4bp)

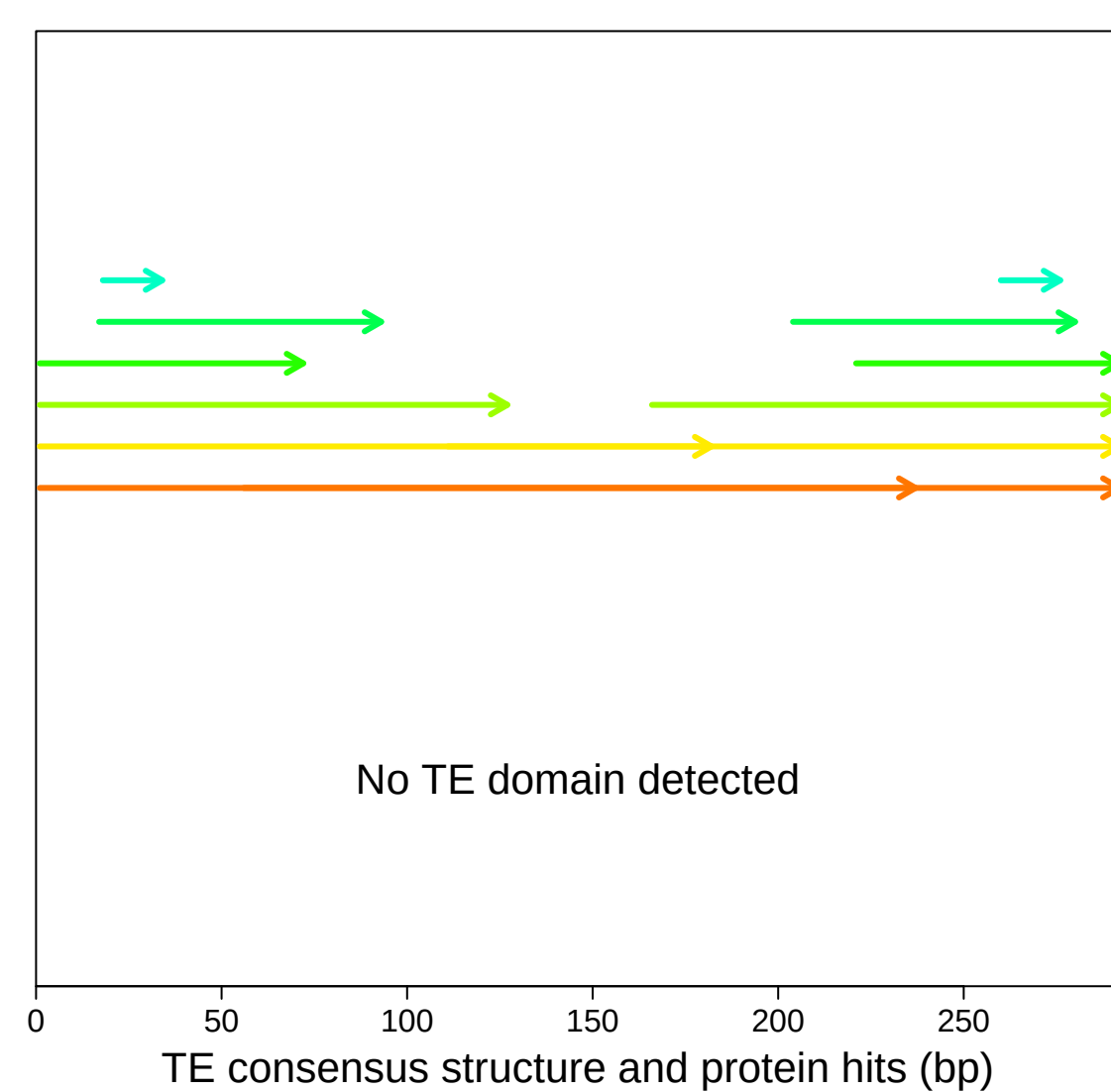
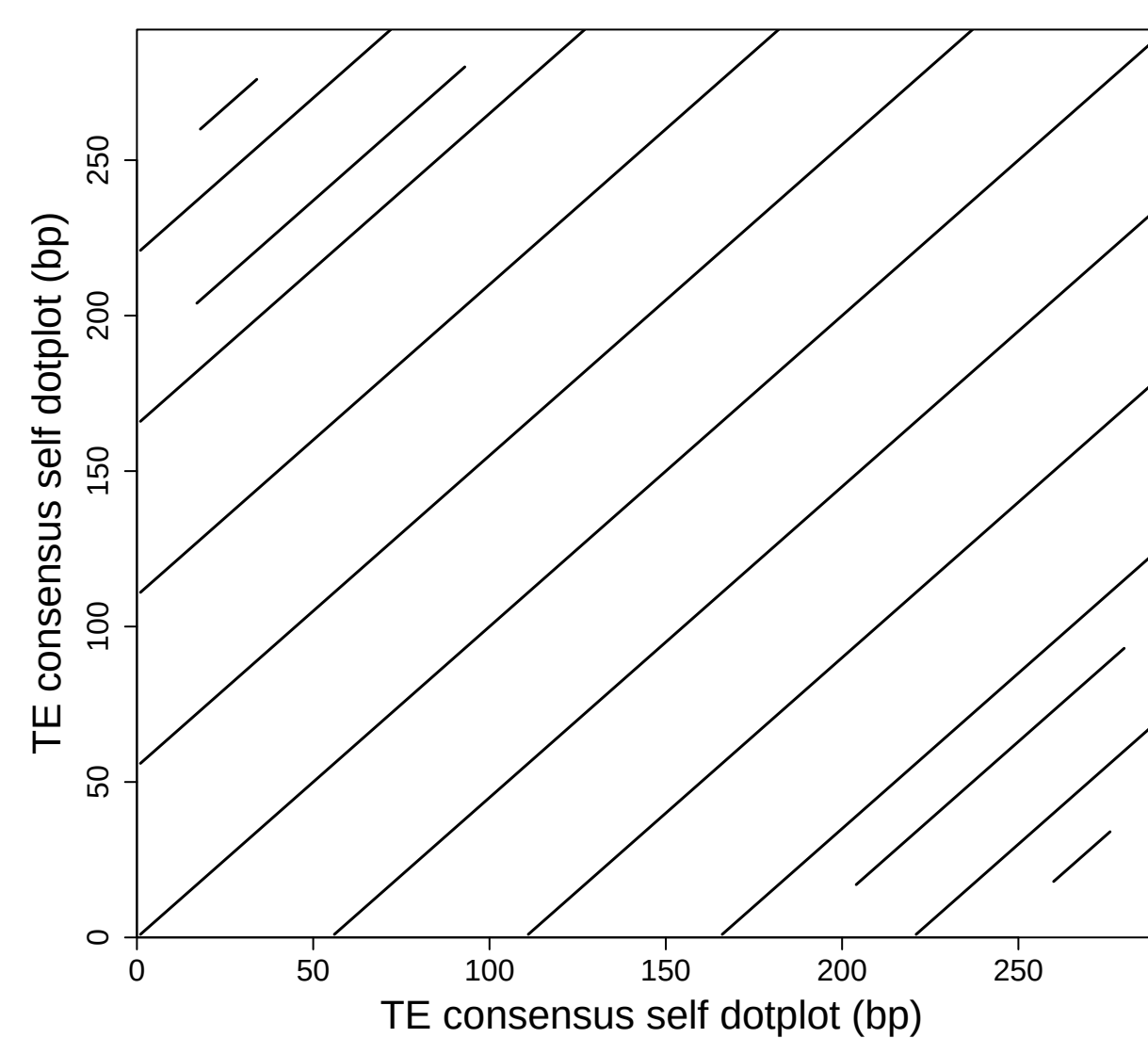
After TEtrimmer Extended plot Blue lines are boundaries



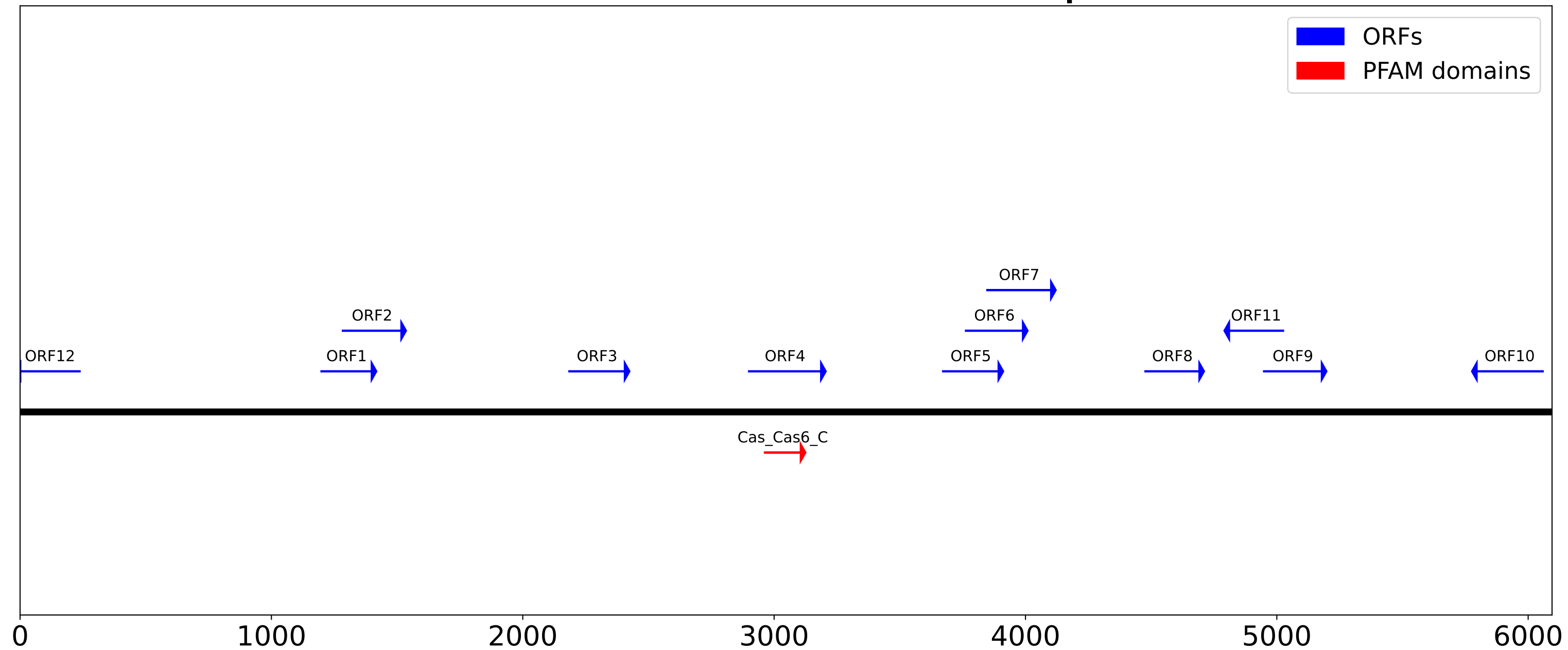
TE: ltr_1_family_1
size: 292bp; fragments: 494; full length: 57 (≥ 262.8 bp)



Before TEtrimmer 292 bp



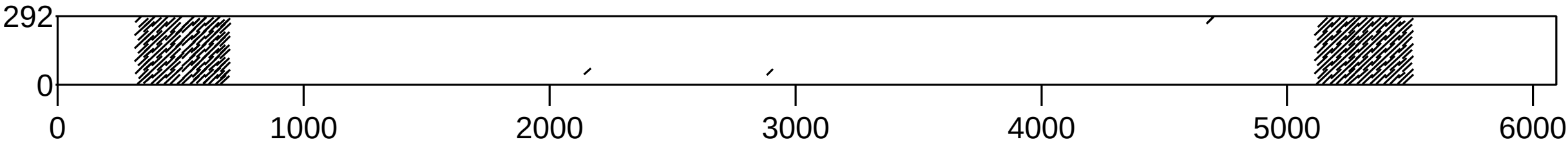
After TEtrimmer ORF and PFAM domain plot



Dotplot

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

nsus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)