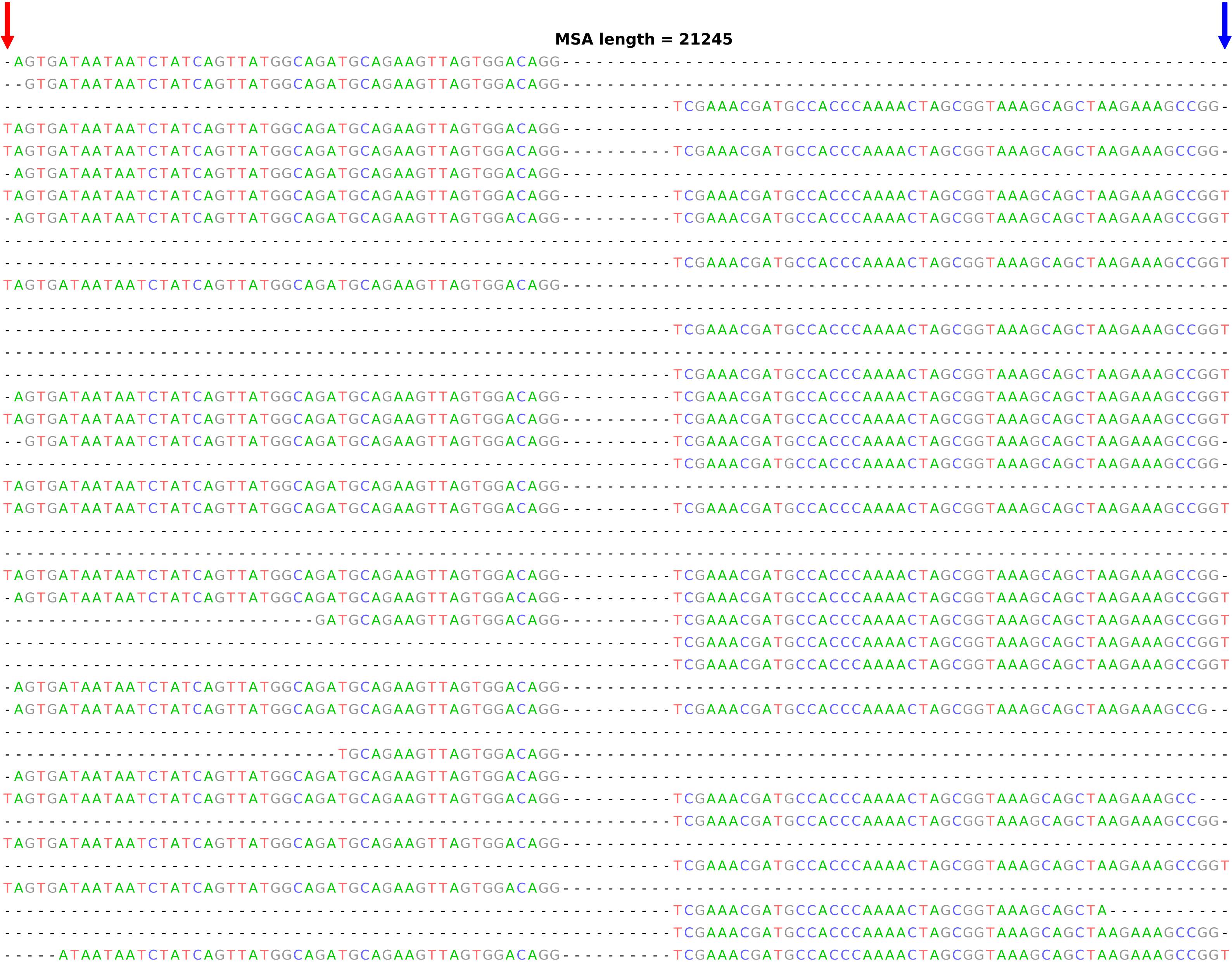
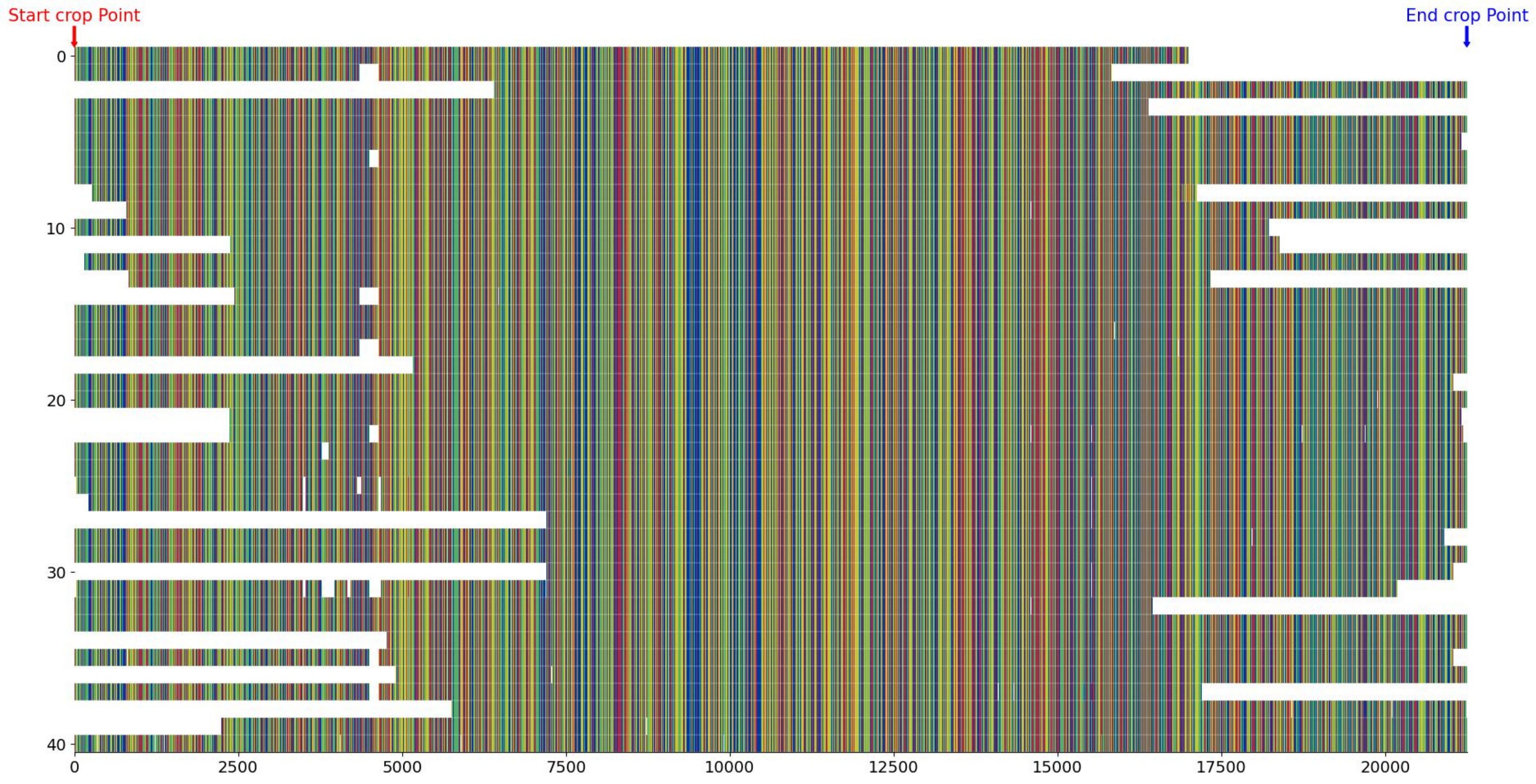


Start crop Point

End crop Point

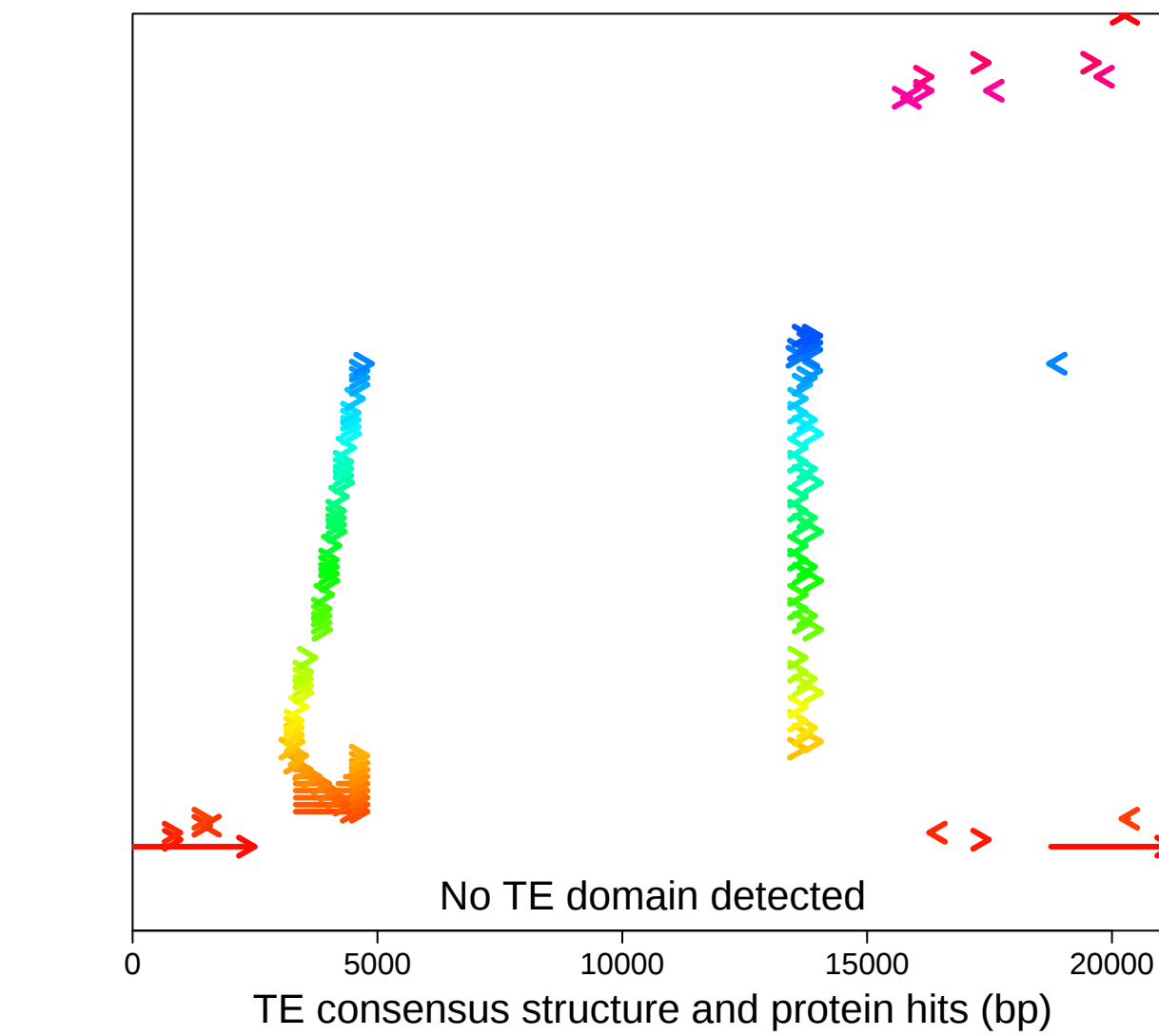
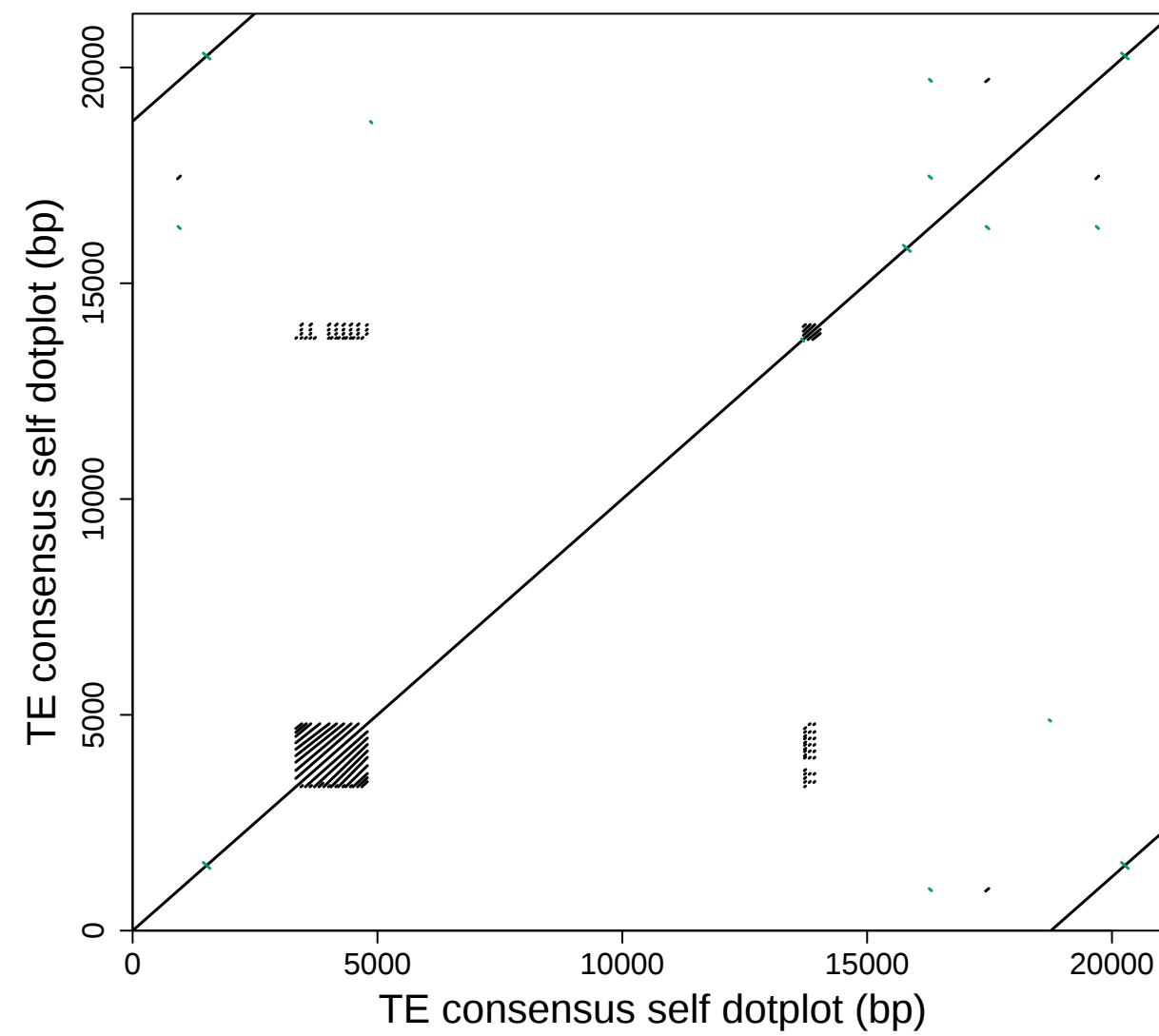
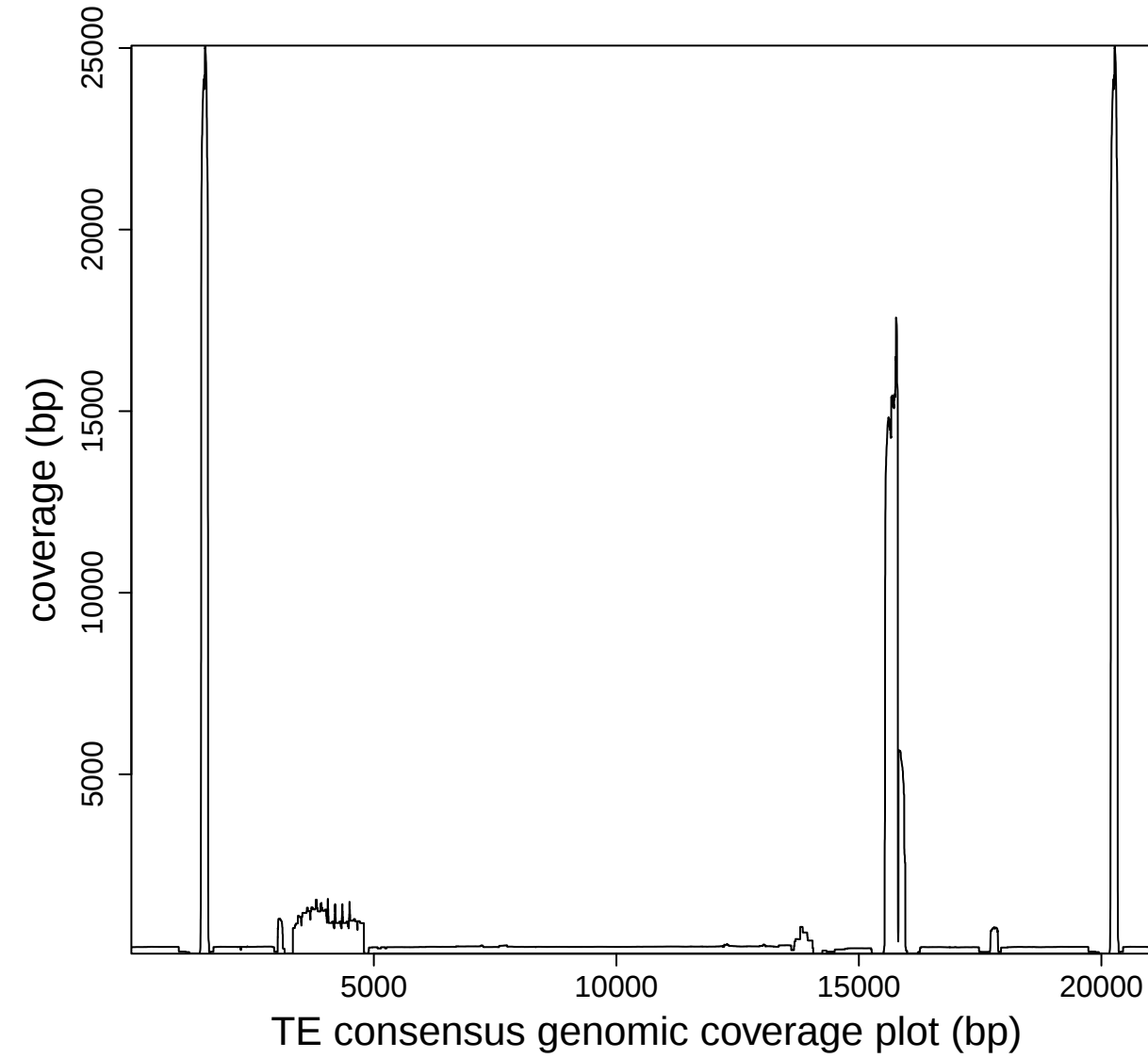
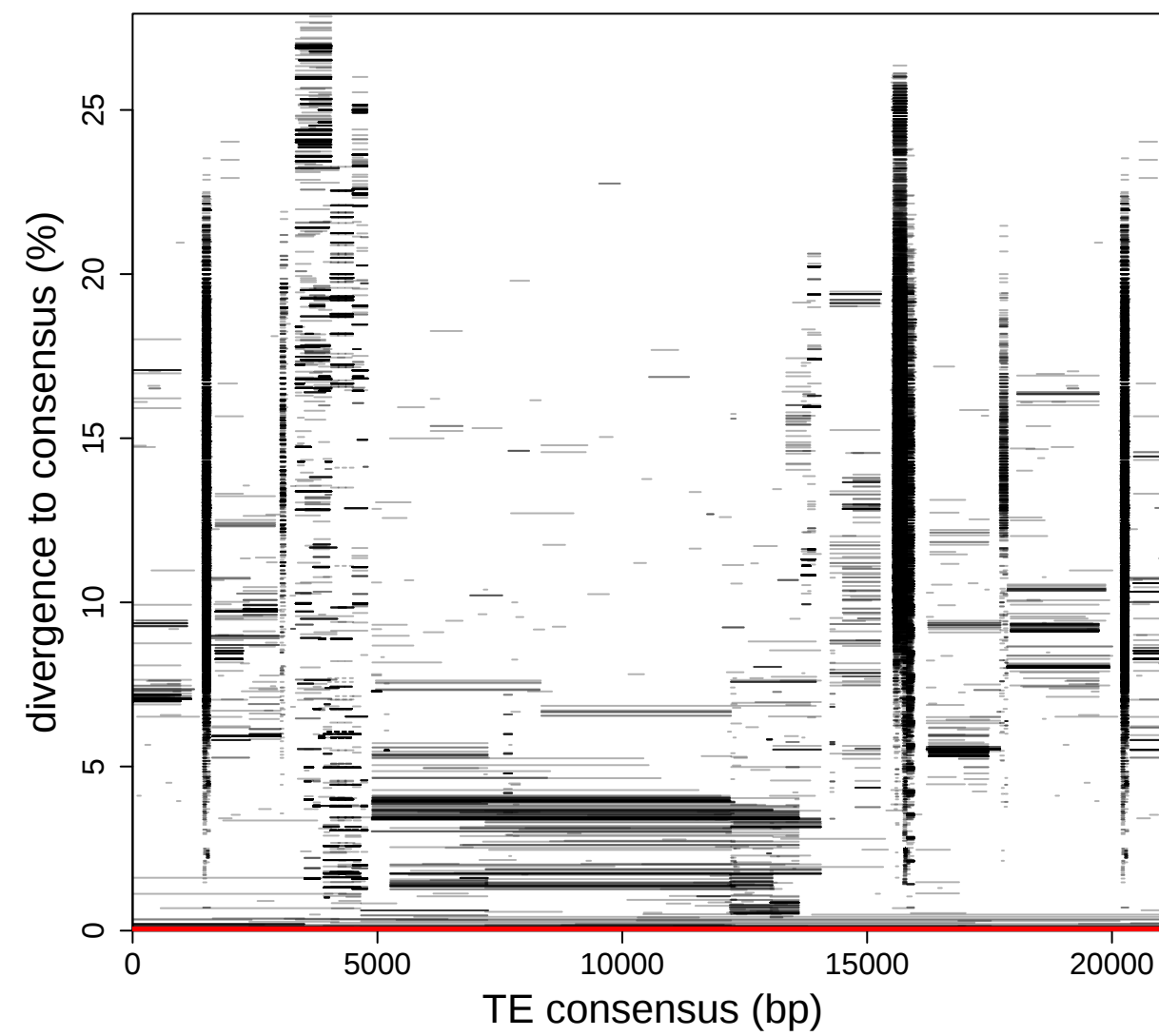
MSA length = 21245





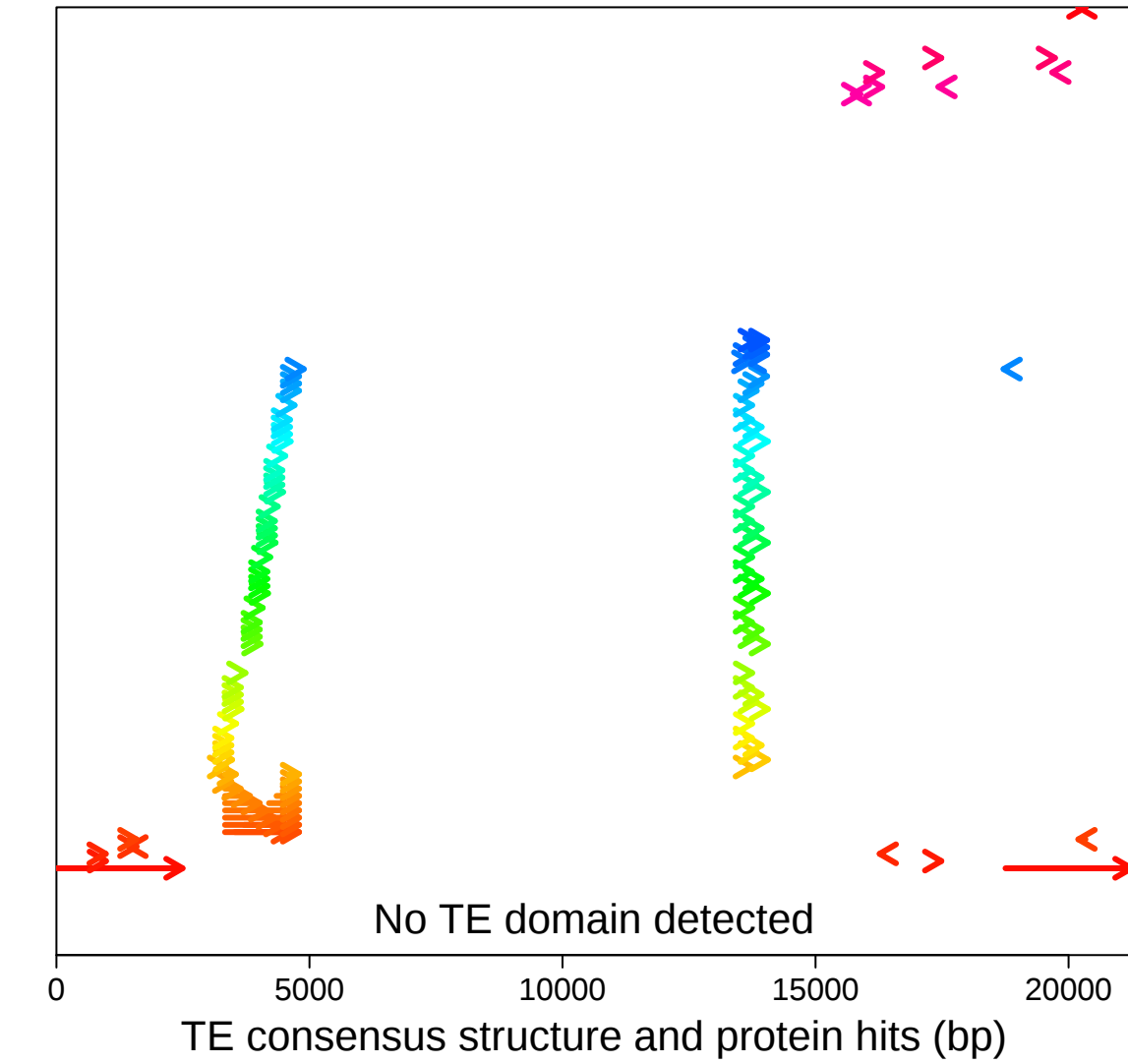
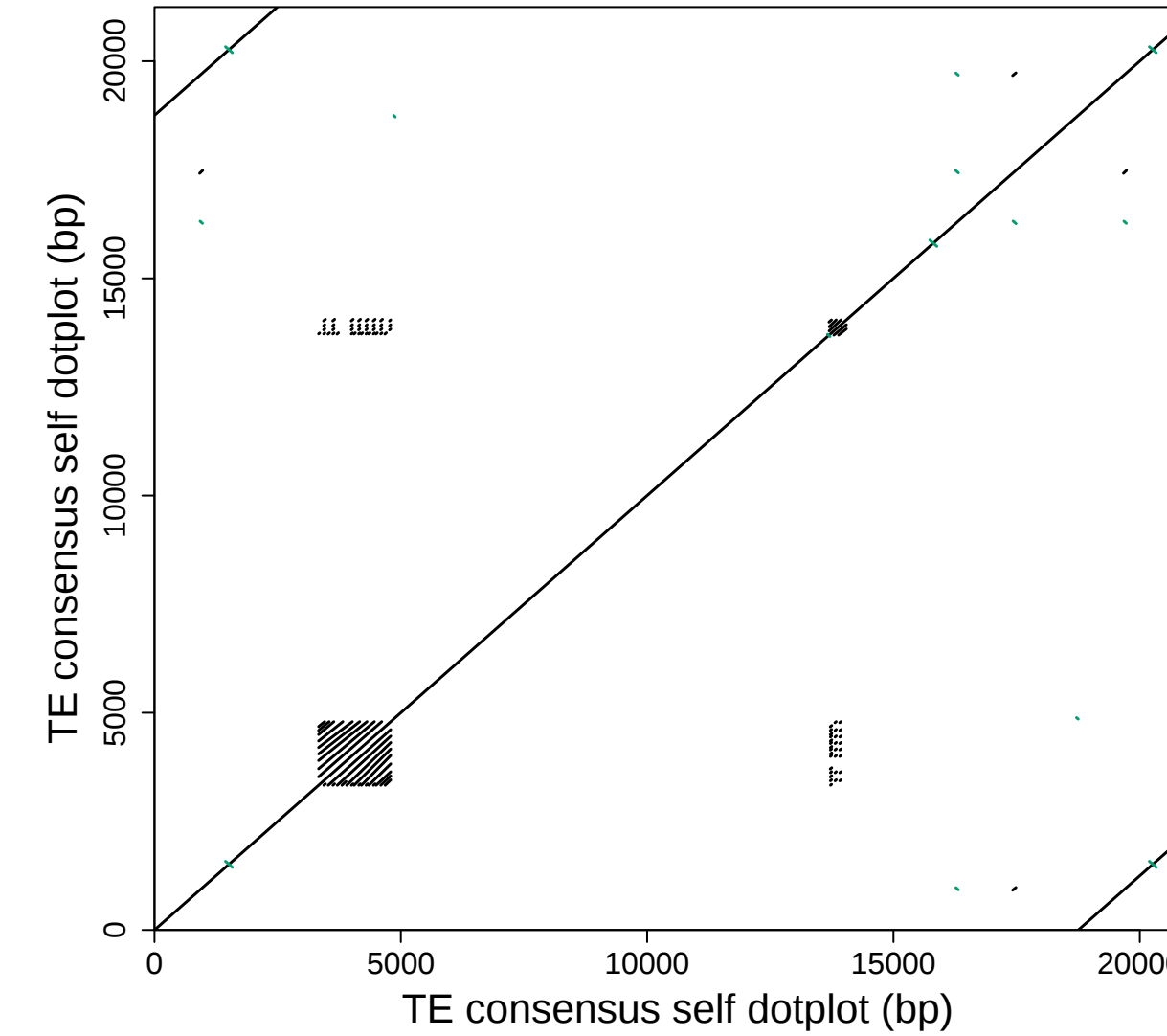
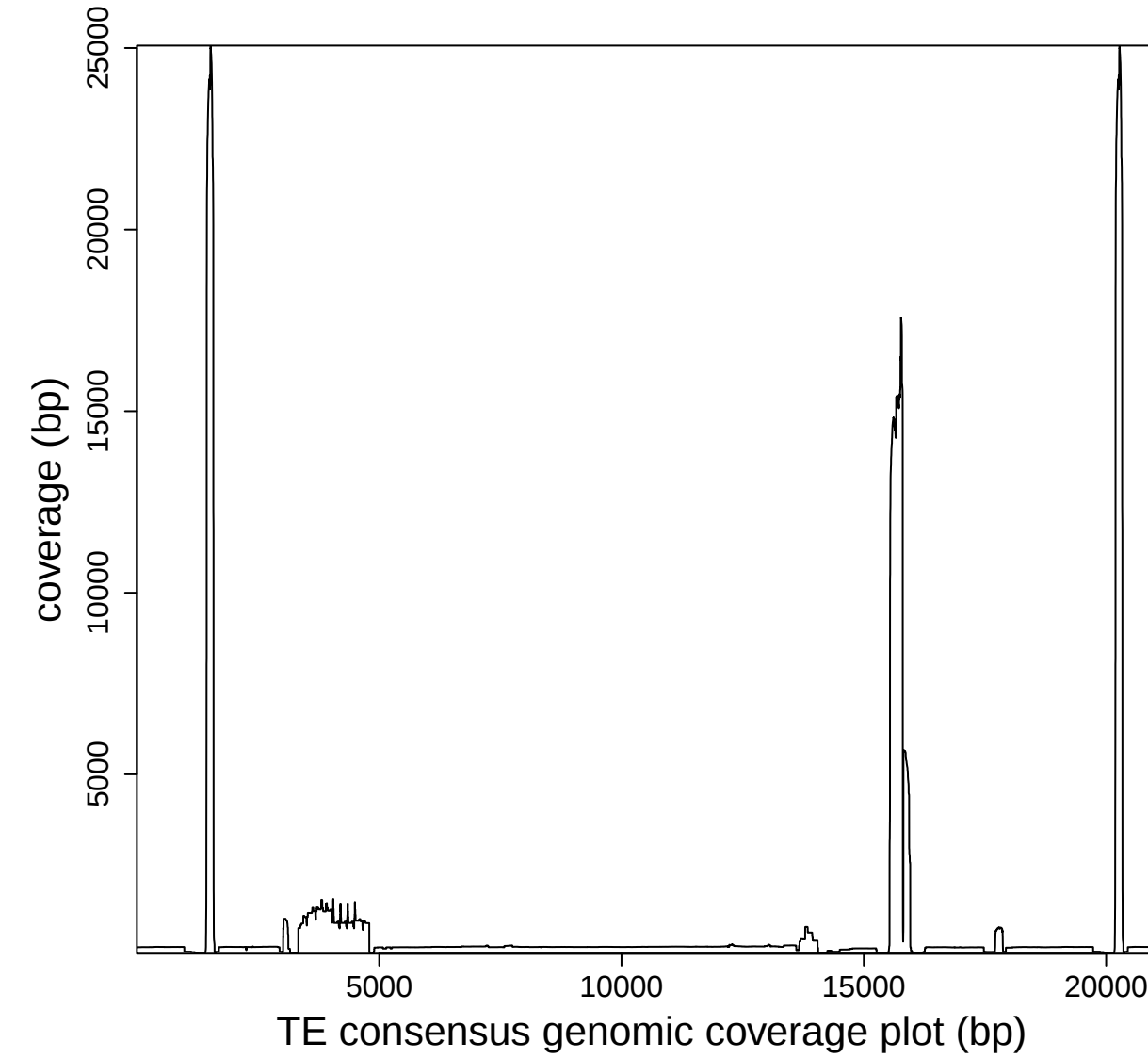
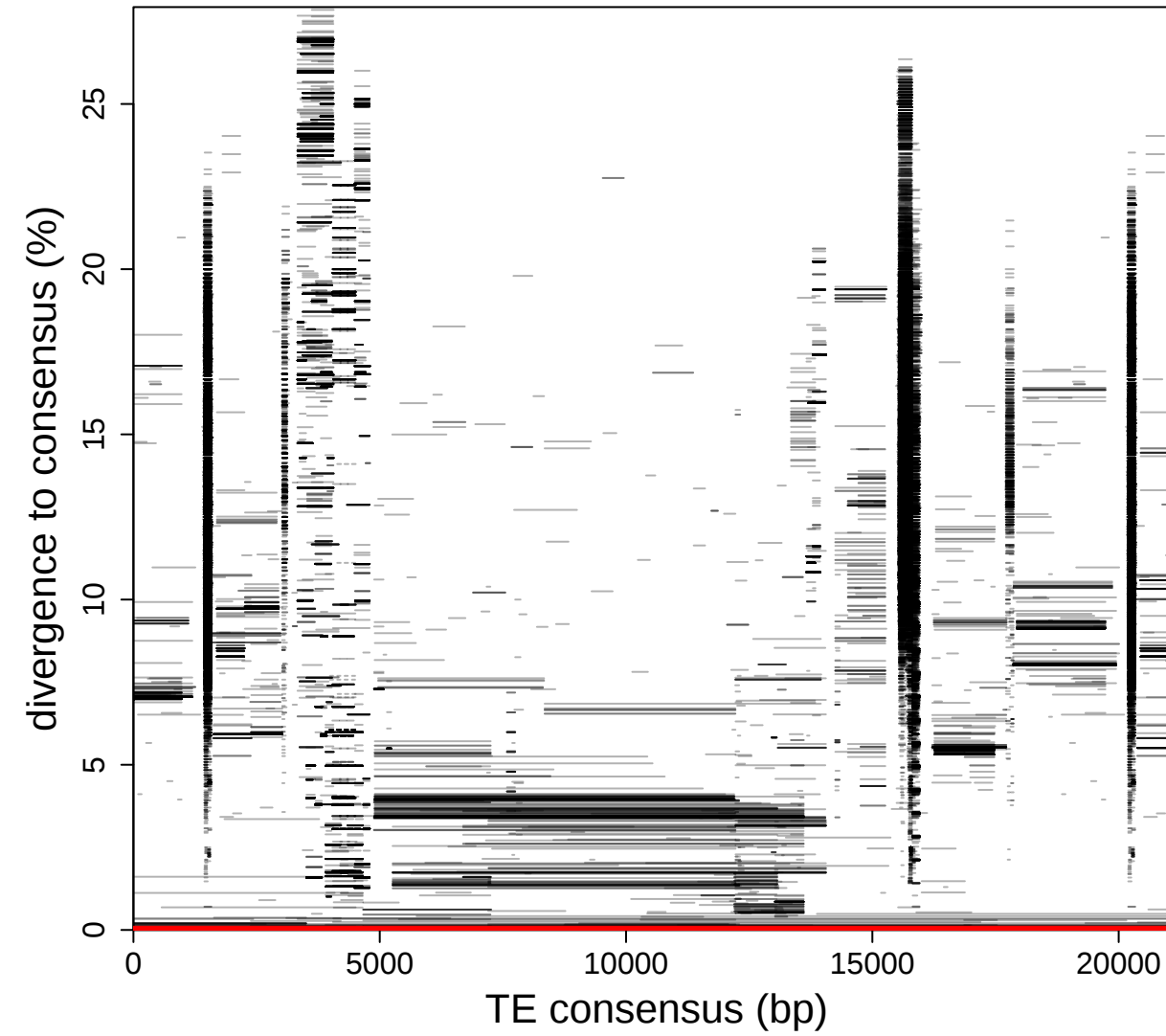
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce
size: 21245bp; fragments: 87983; full length: 16 (>=19120.5bp)

After TEtrimmer 21245 bp



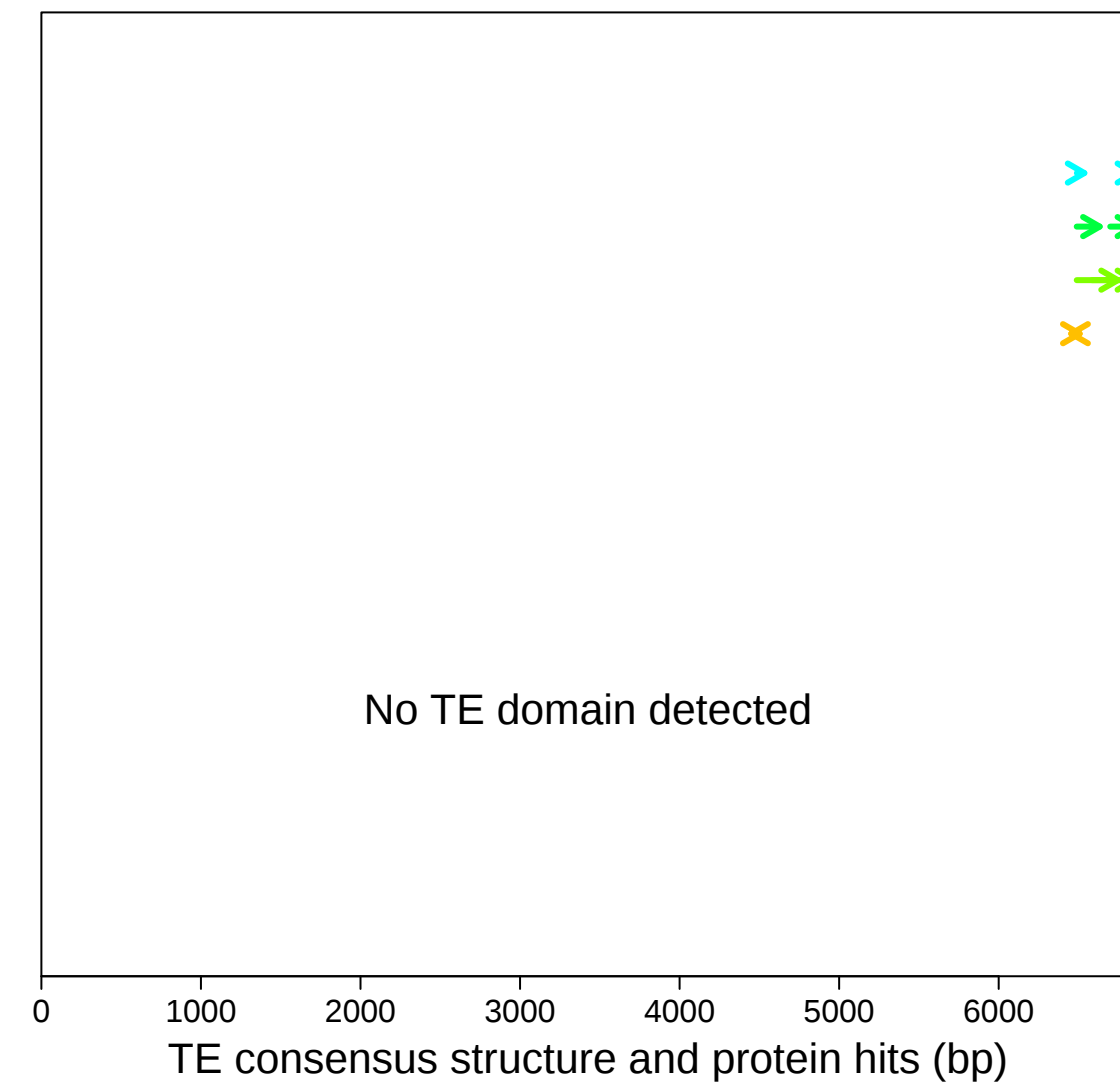
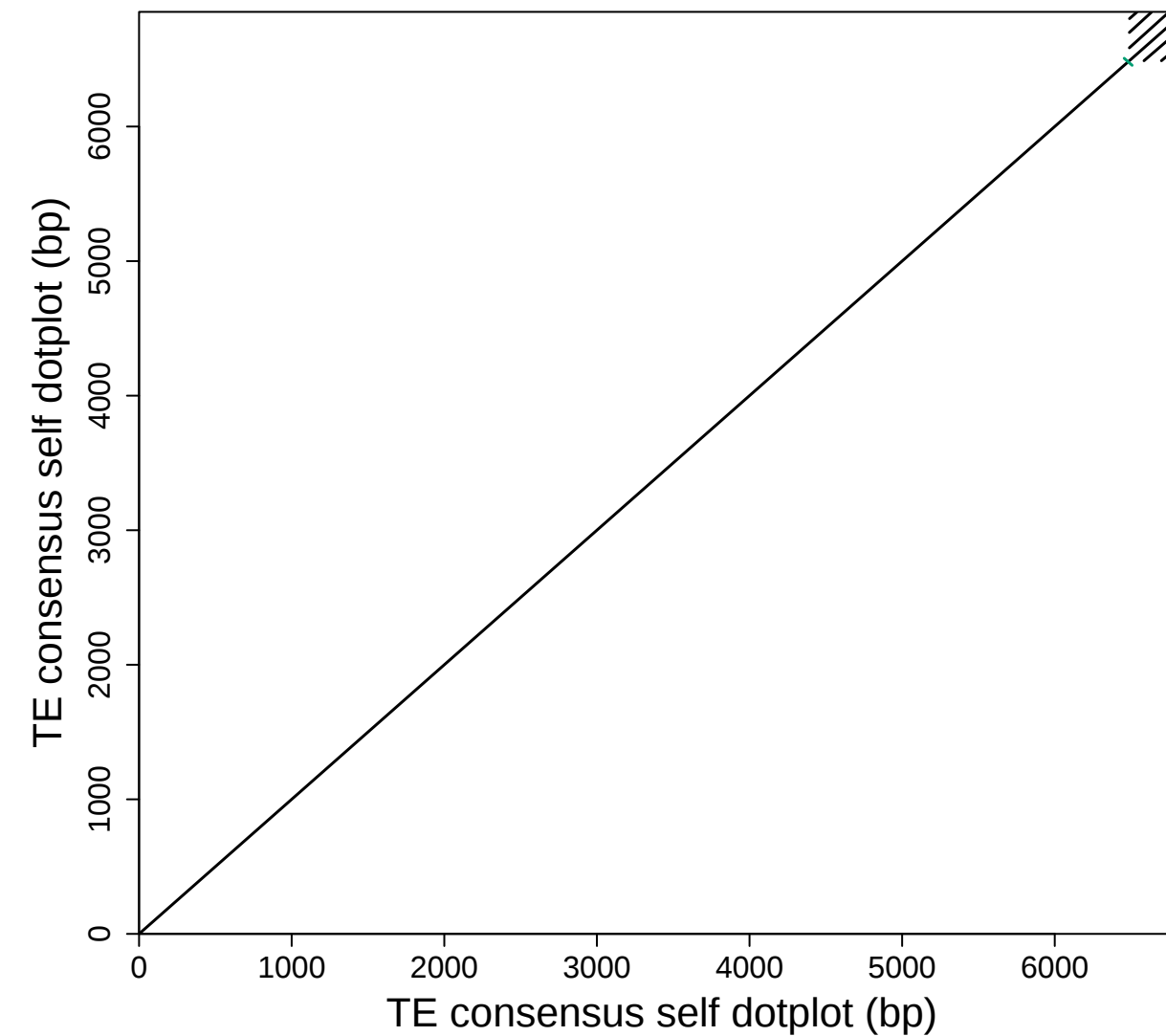
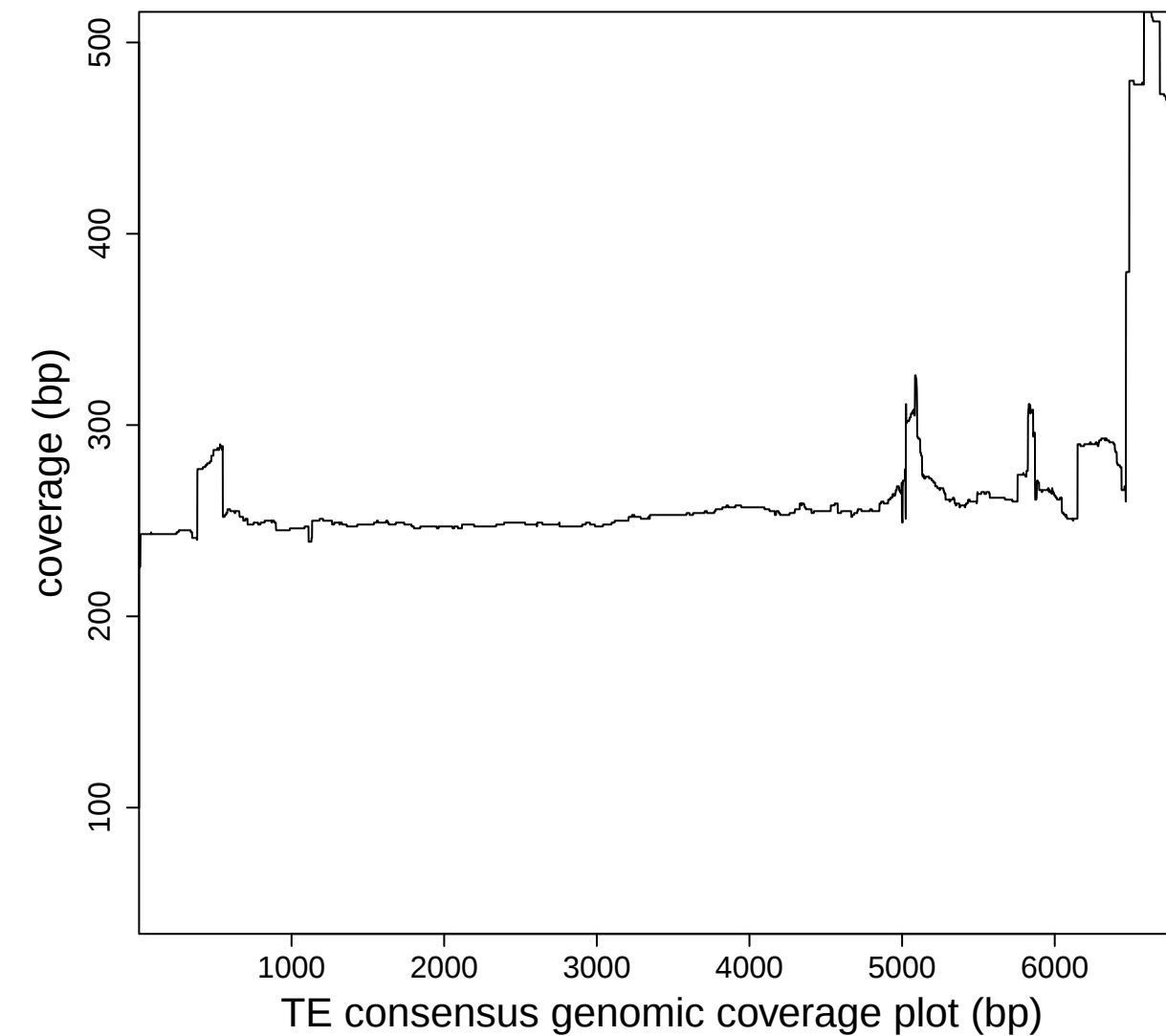
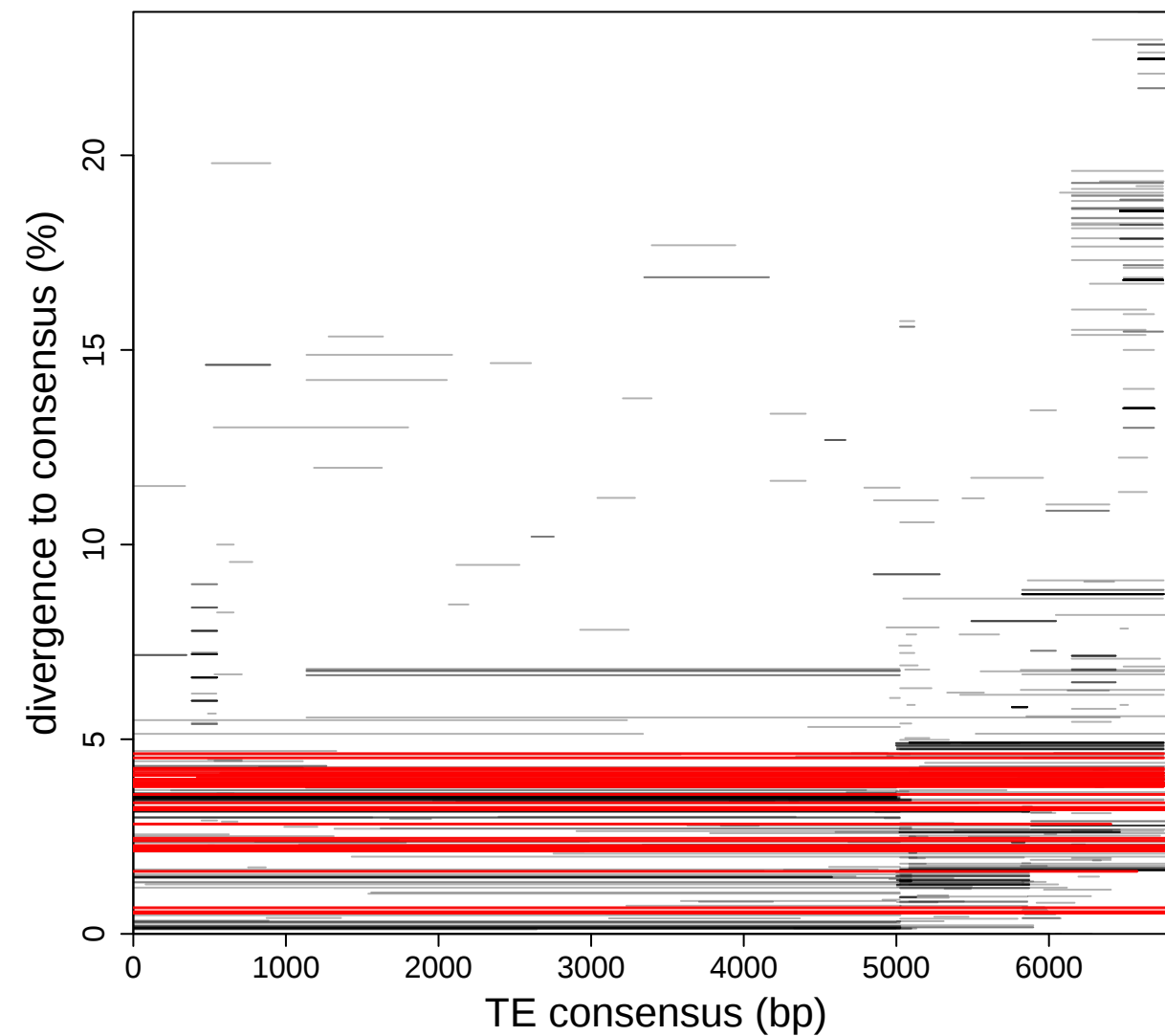
ily_522.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.f
size: 21245bp; fragments: 87983; full length: 16 (>=19120.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

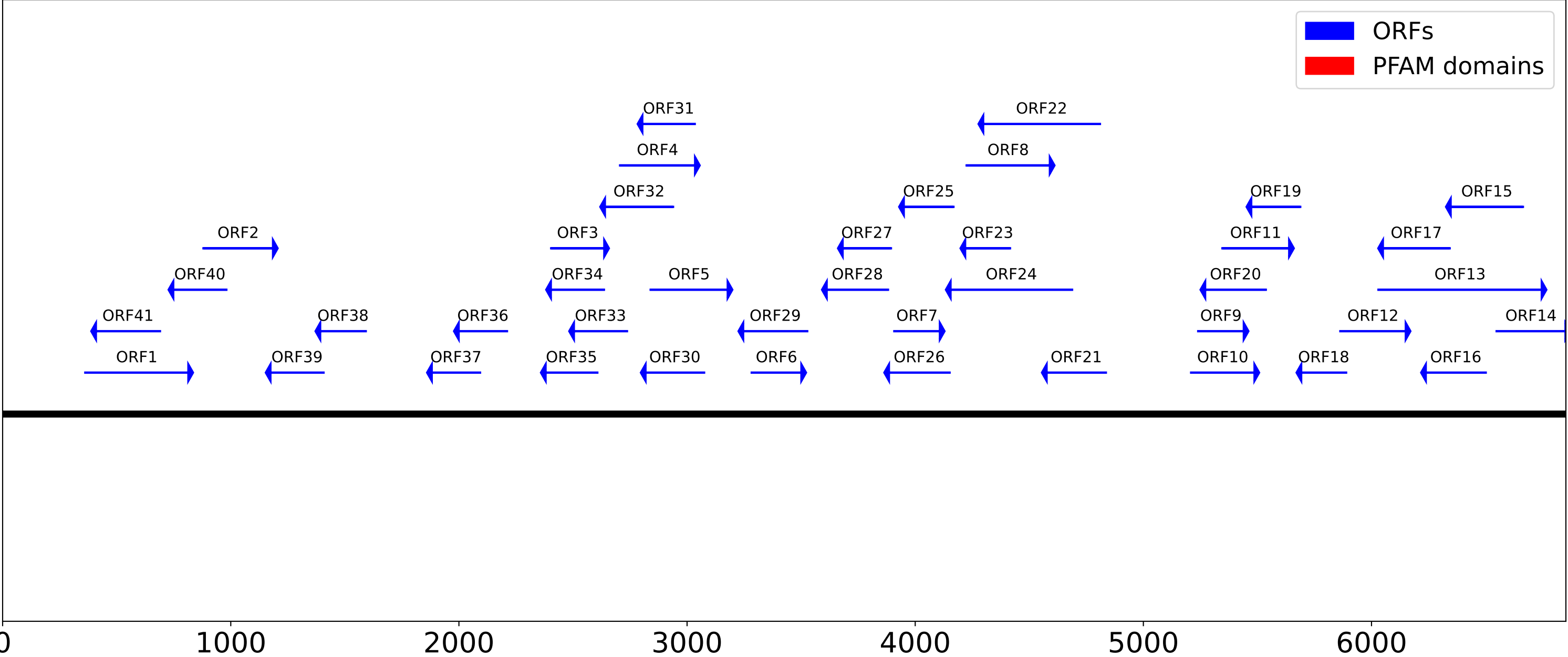


TE: rnd_1_family_522
size: 6852bp; fragments: 956; full length: 131 (≥ 6166.8 bp)

Before TEtrimmer 6852 bp



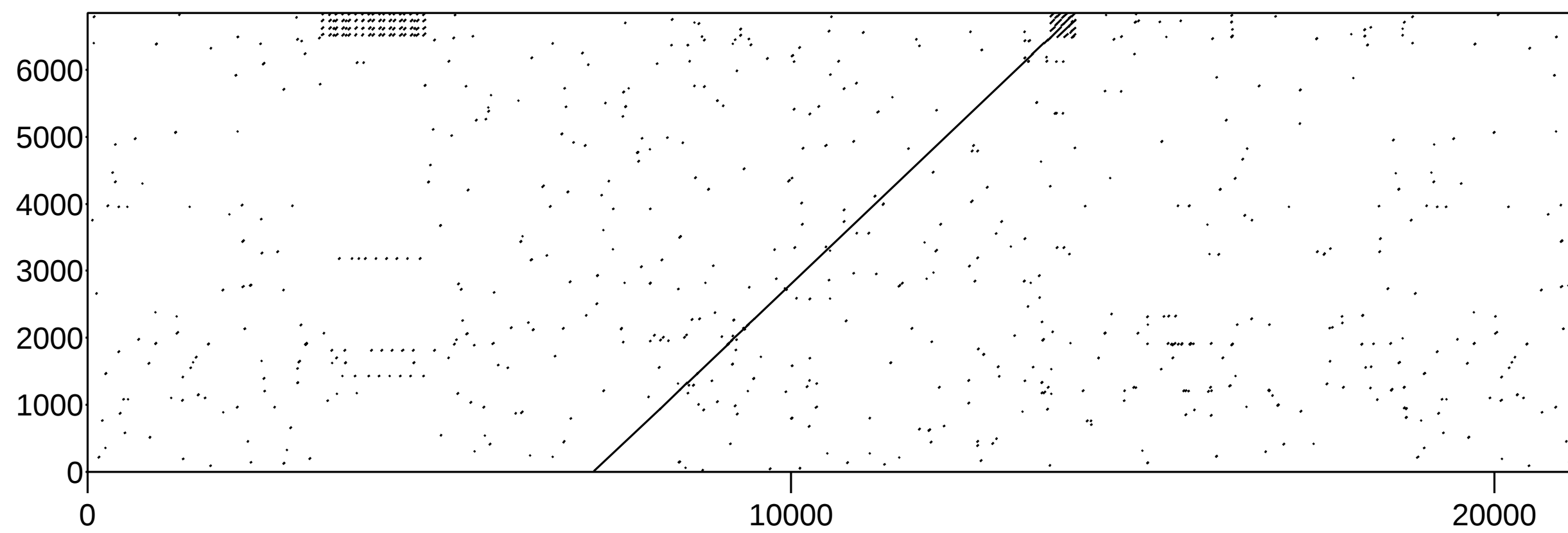
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)