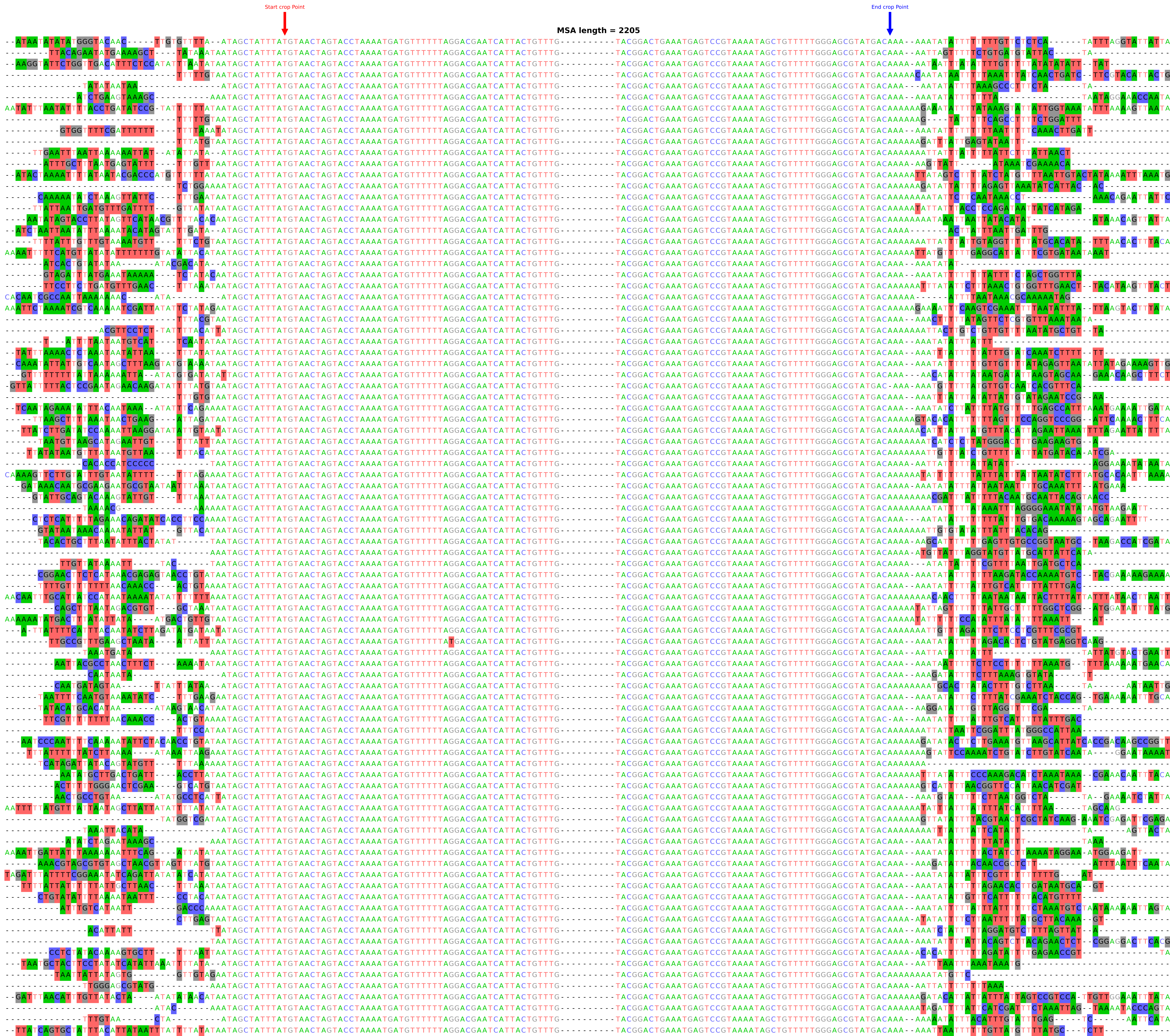
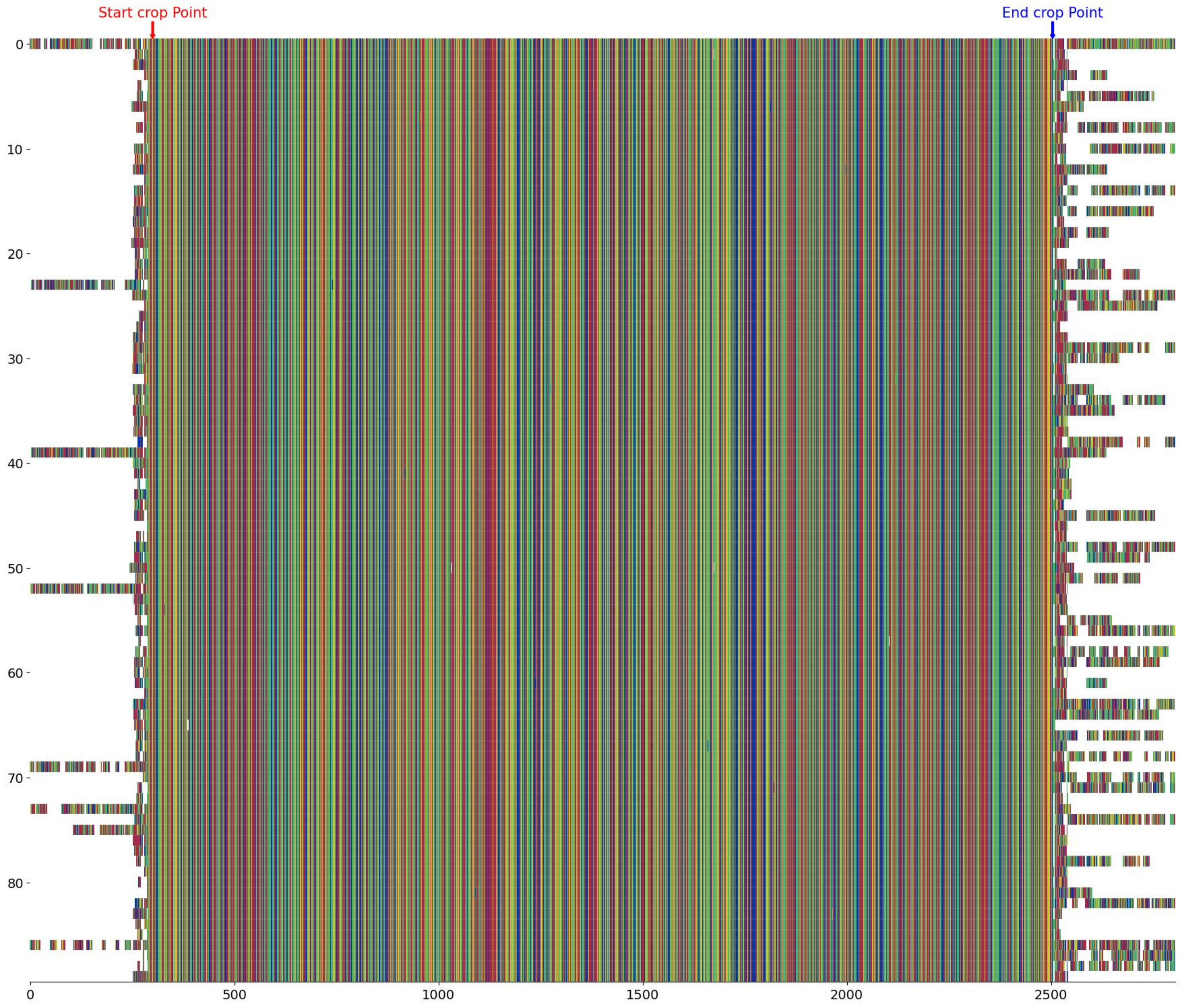
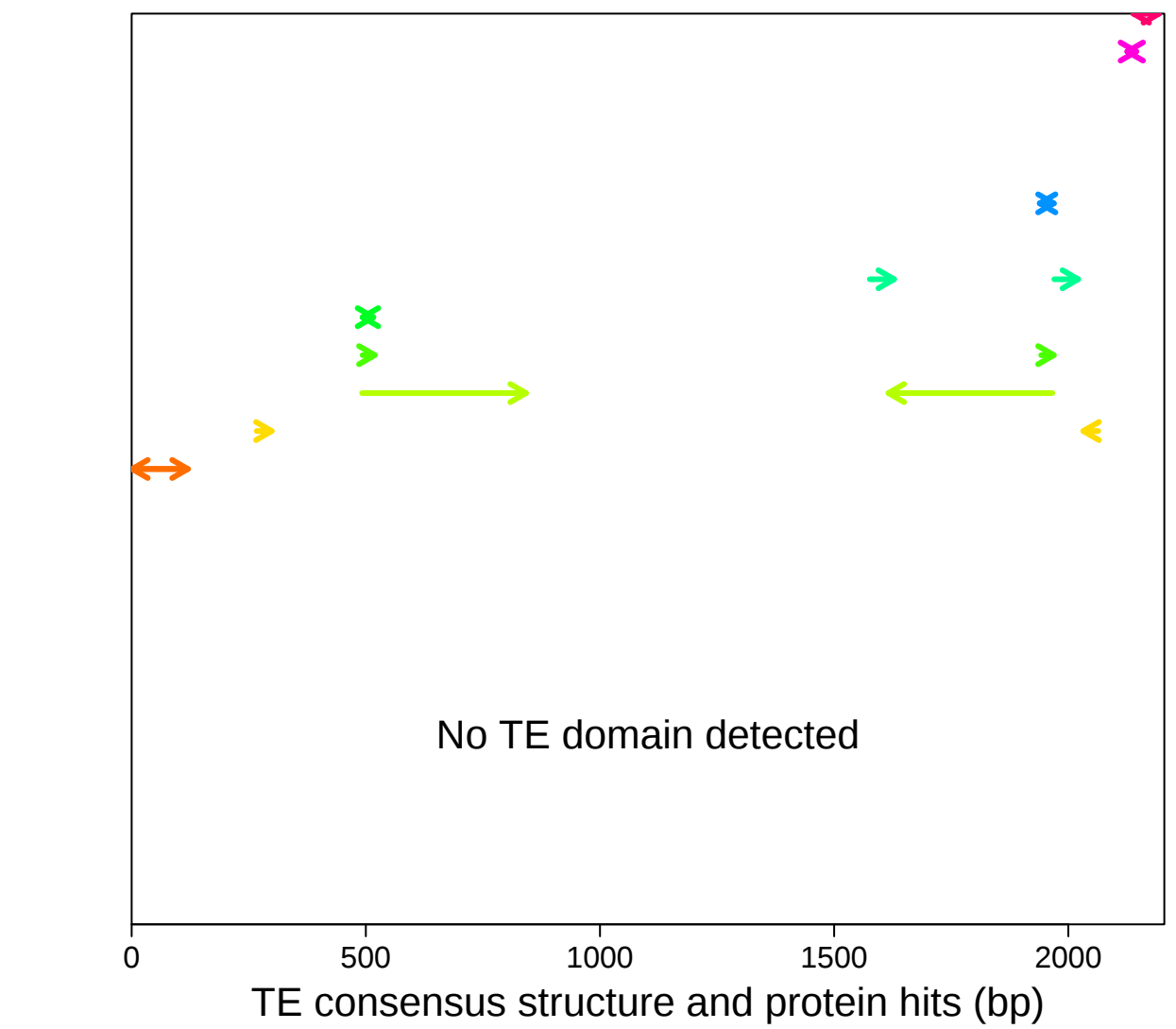
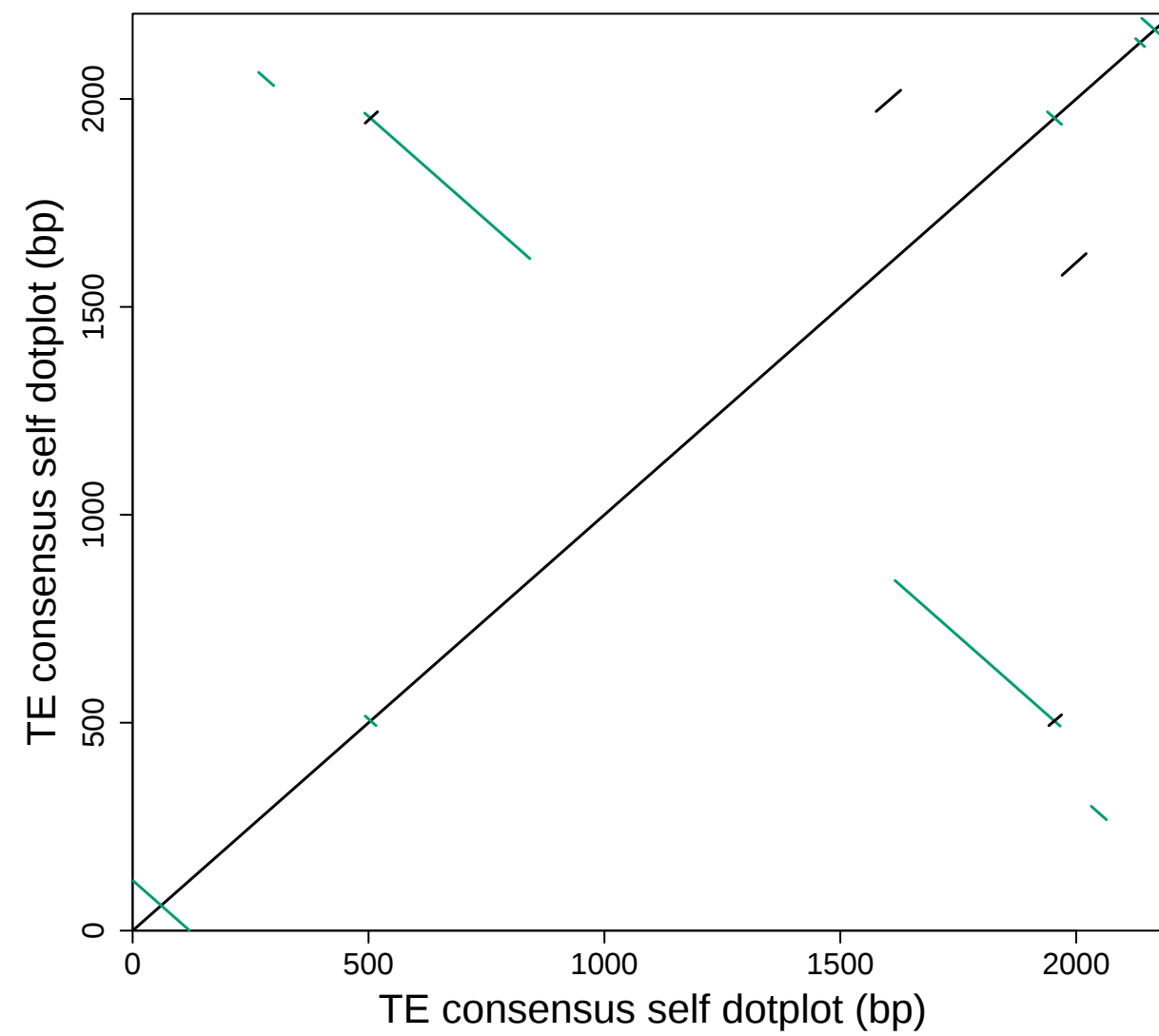
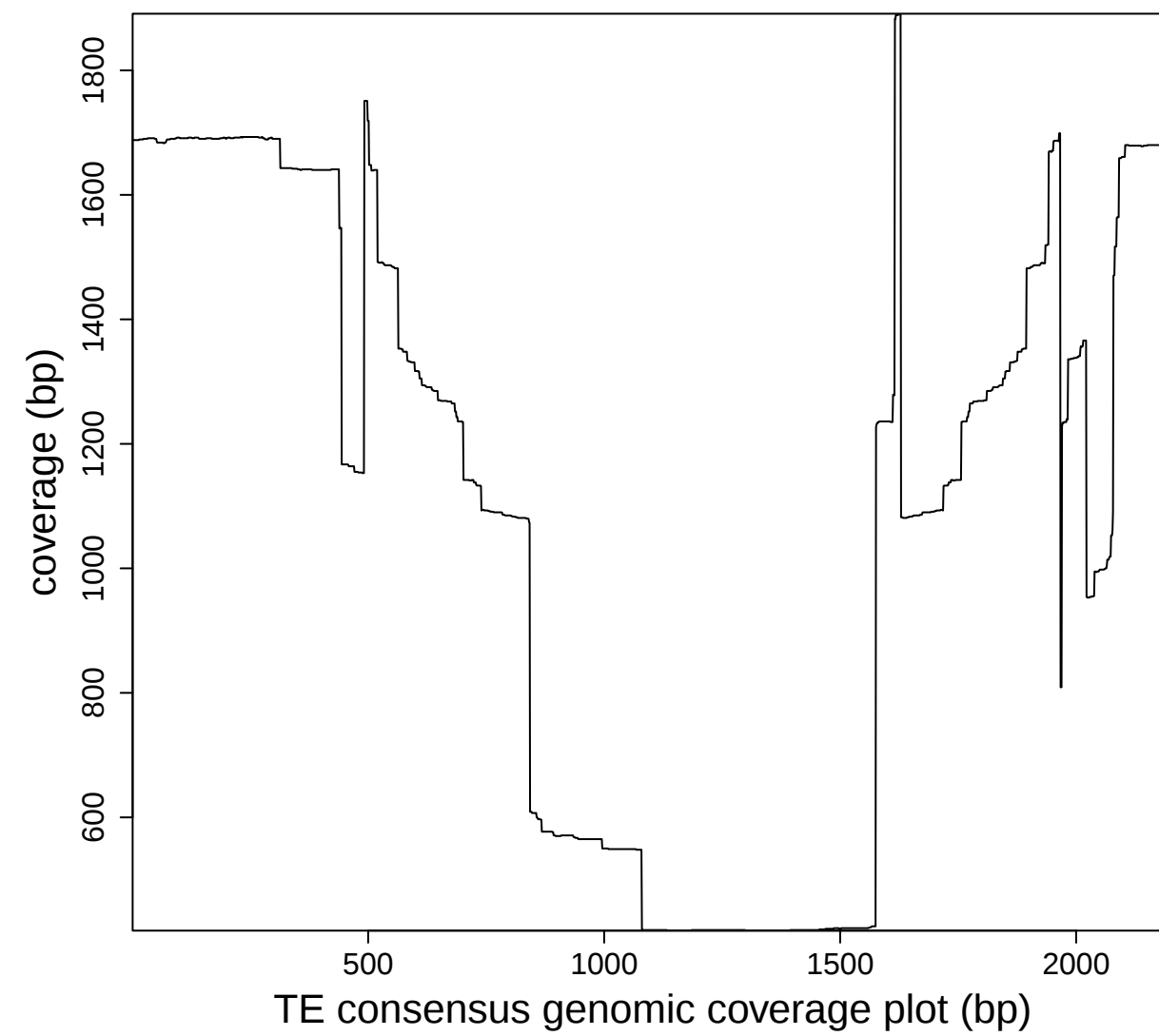
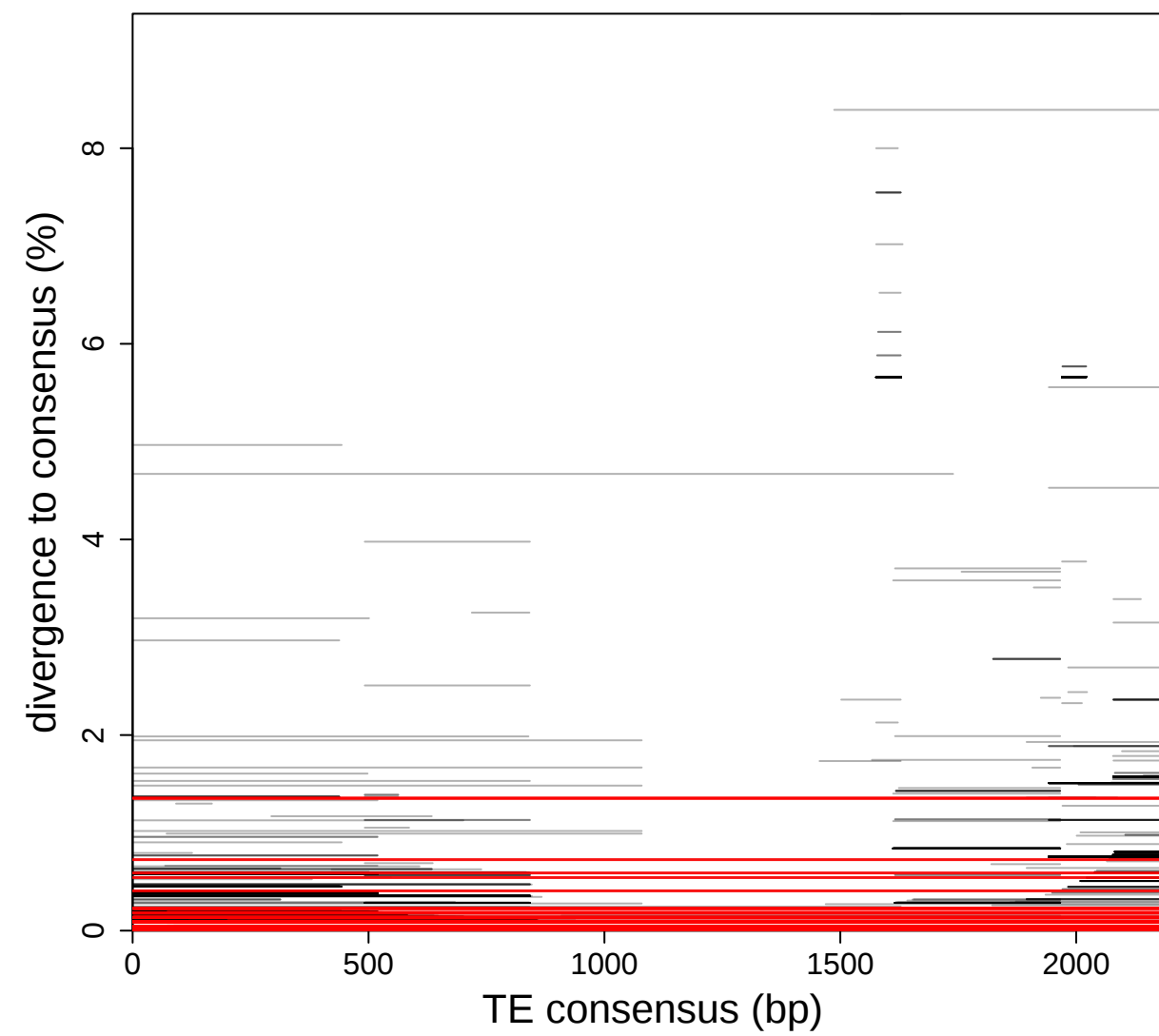






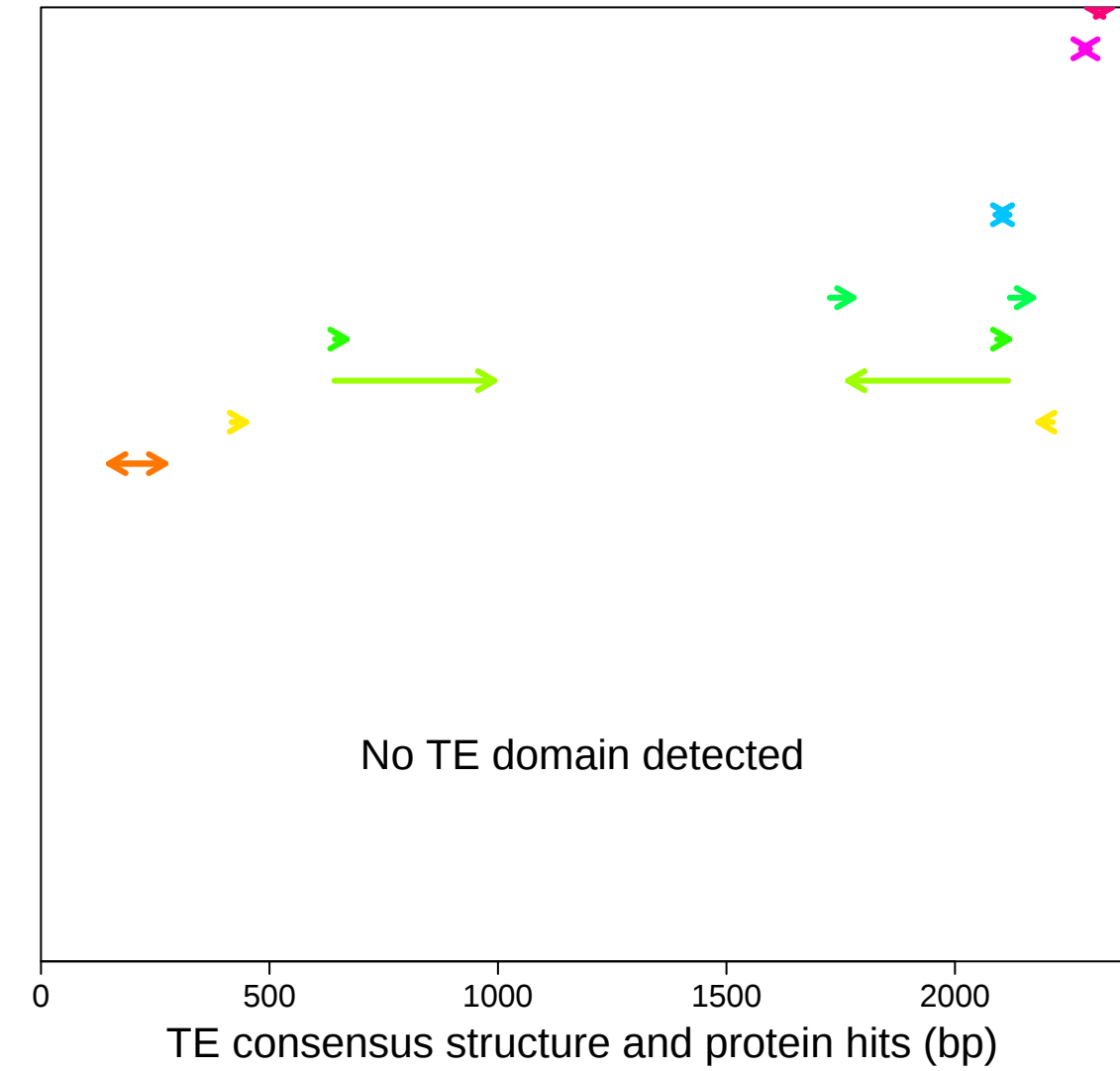
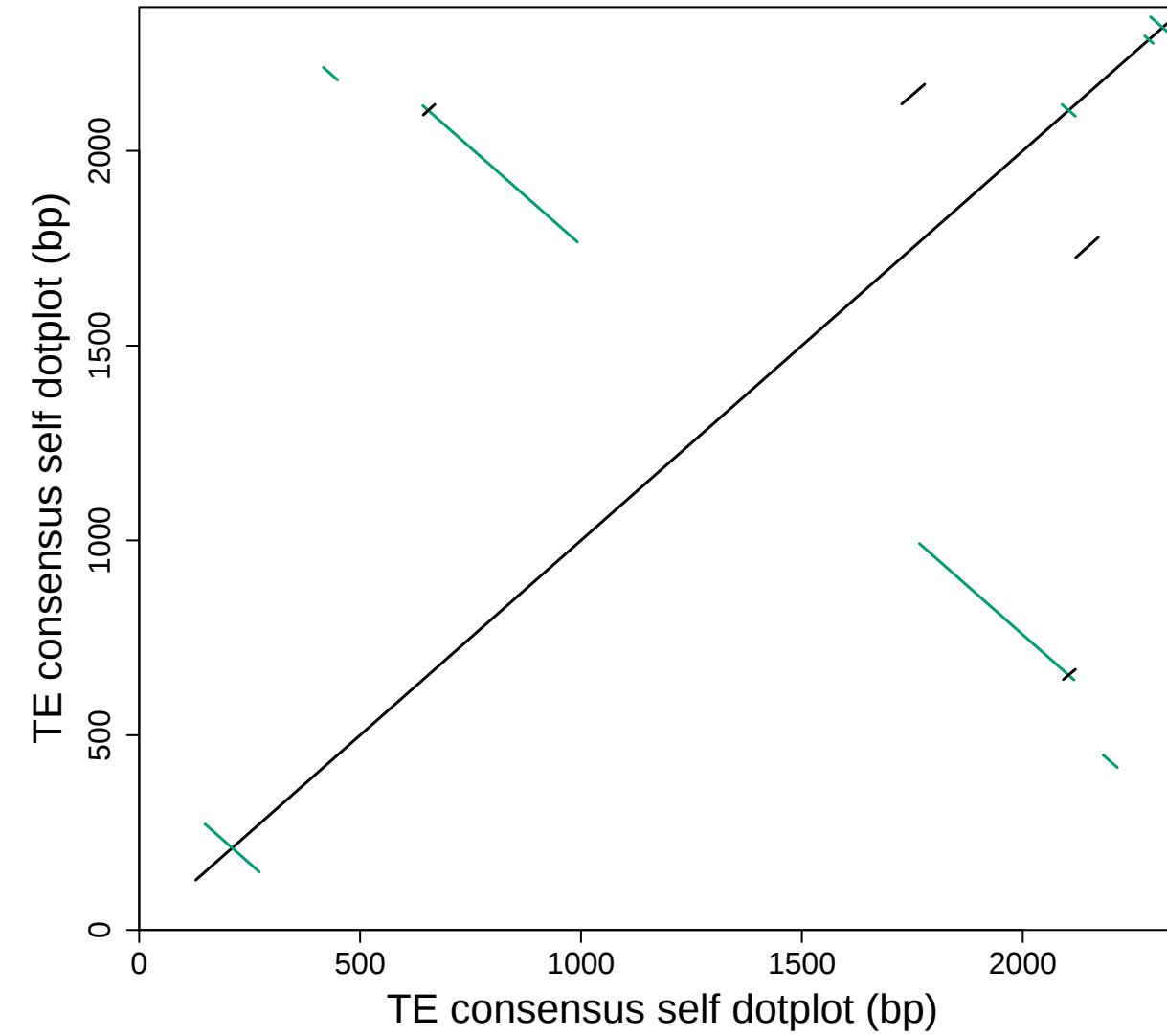
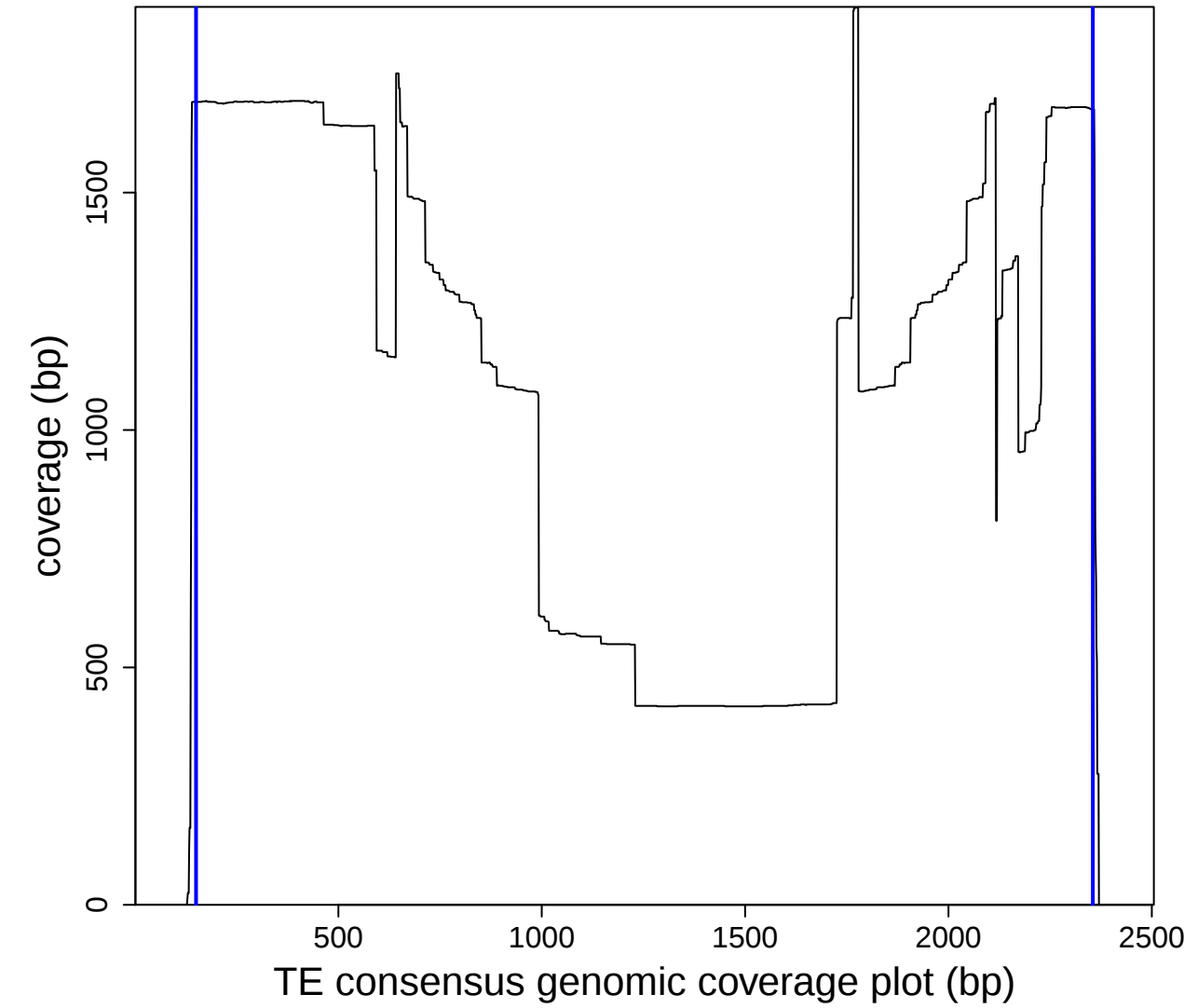
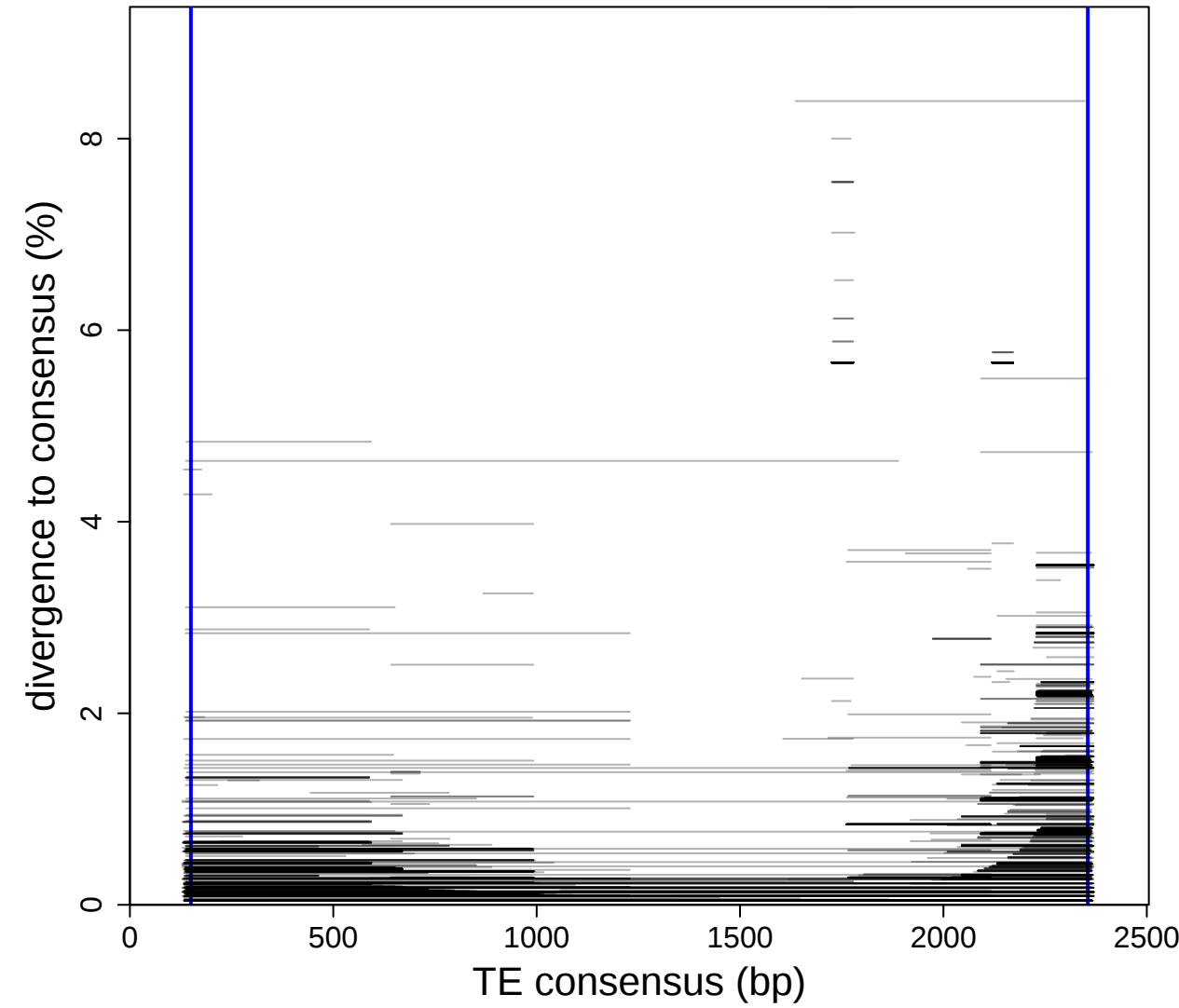
l.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 2205bp; fragments: 5727; full length: 399 (>=1984.5bp)

After TEtrimmer 2205 bp



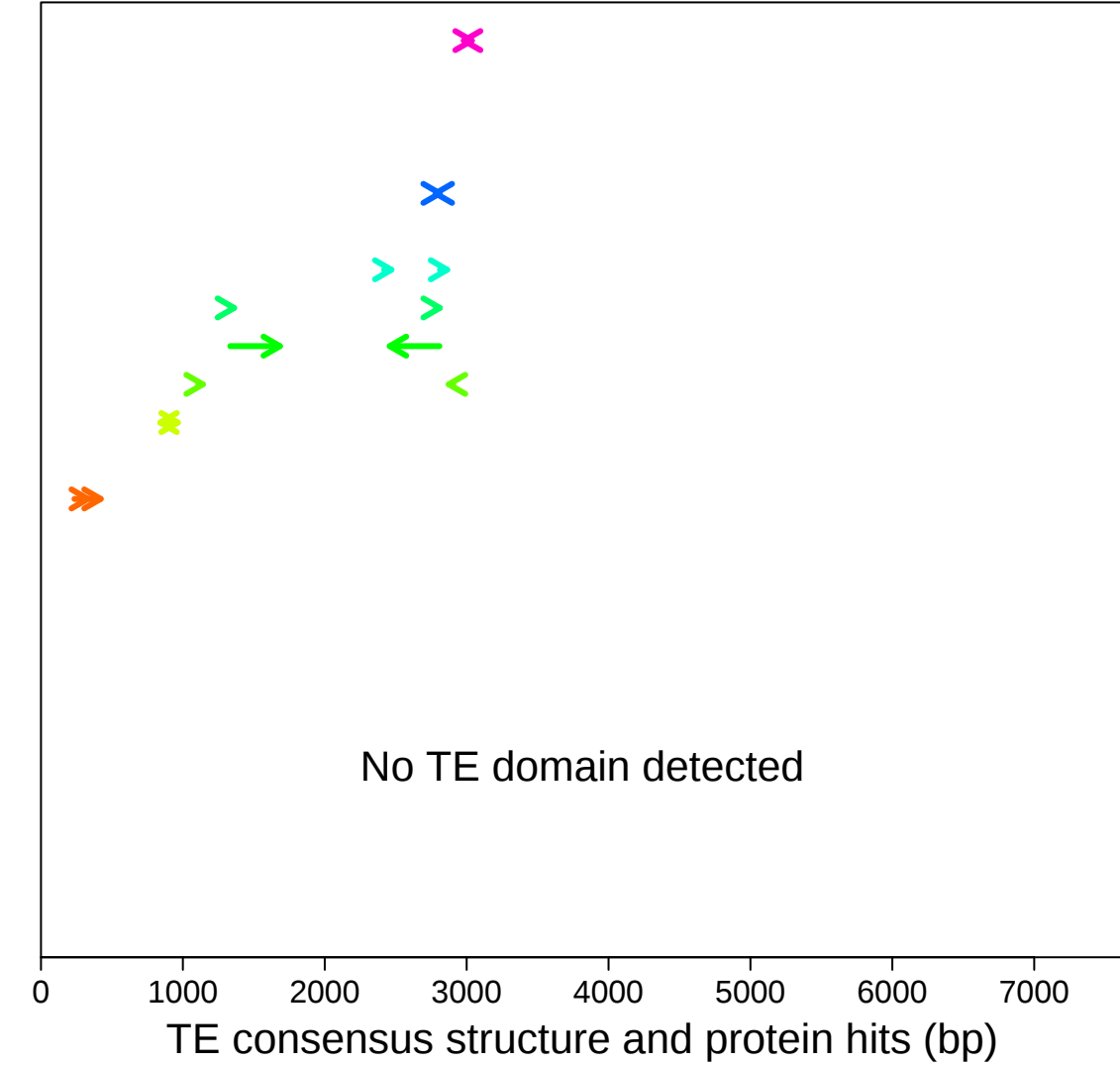
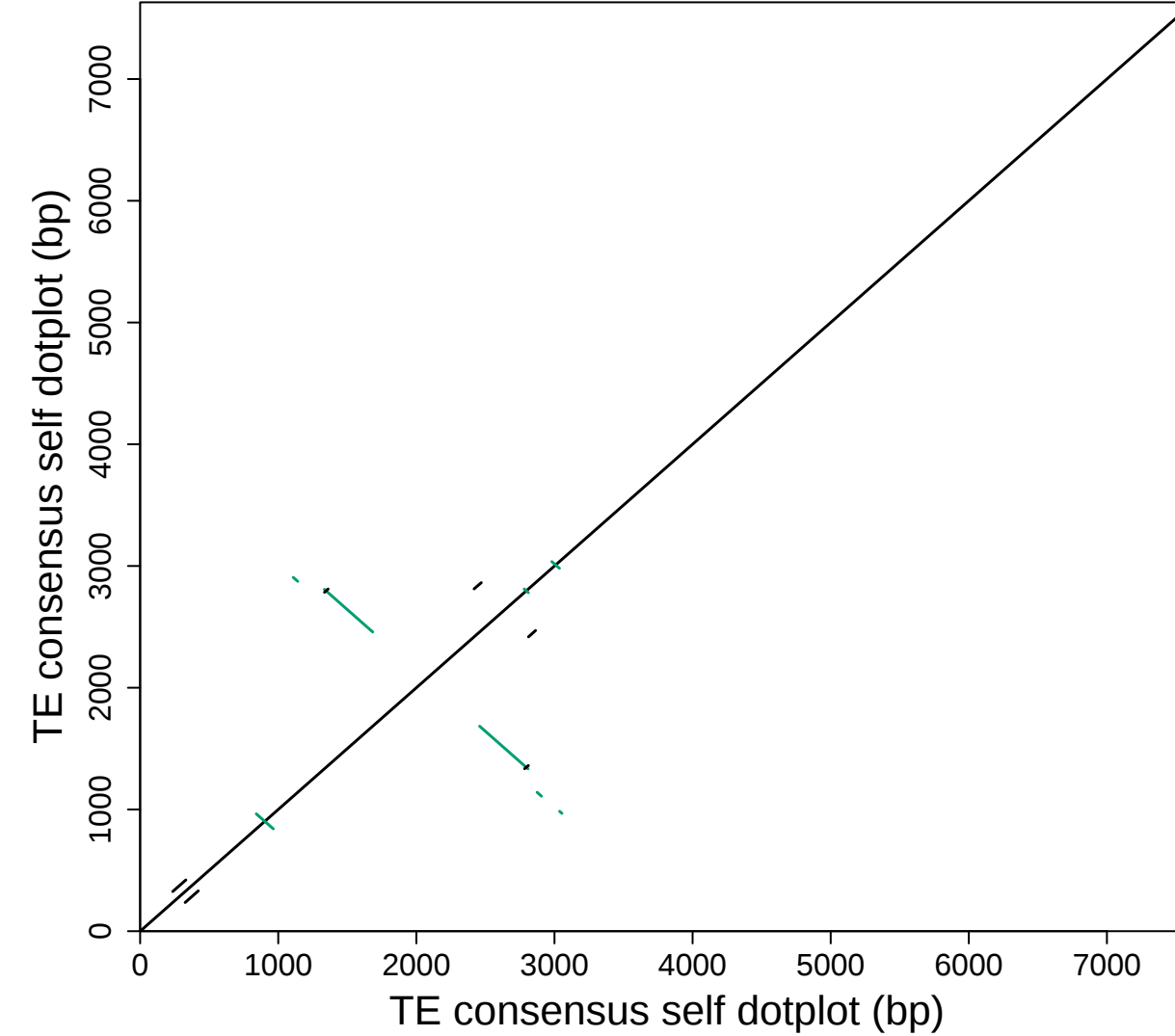
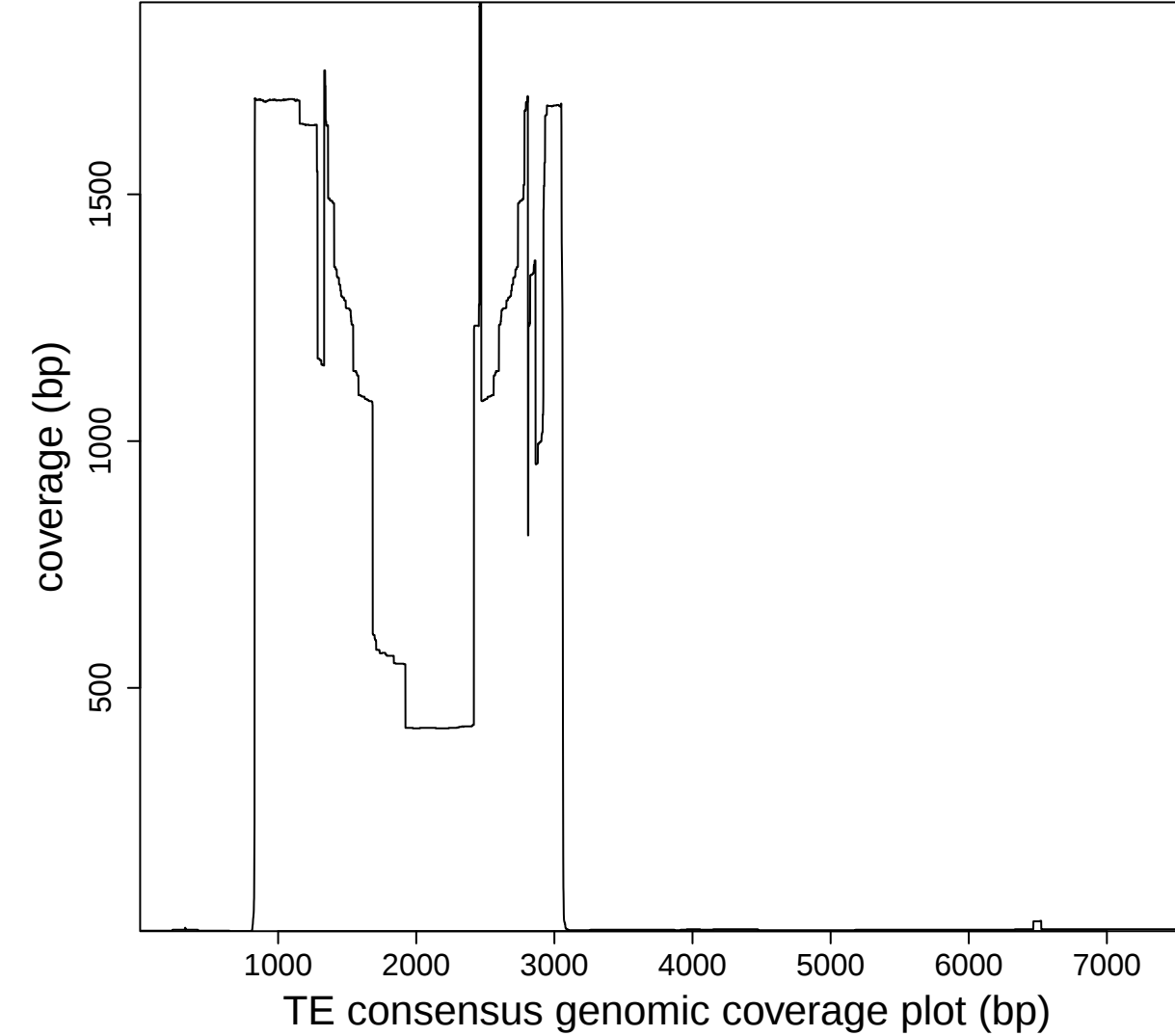
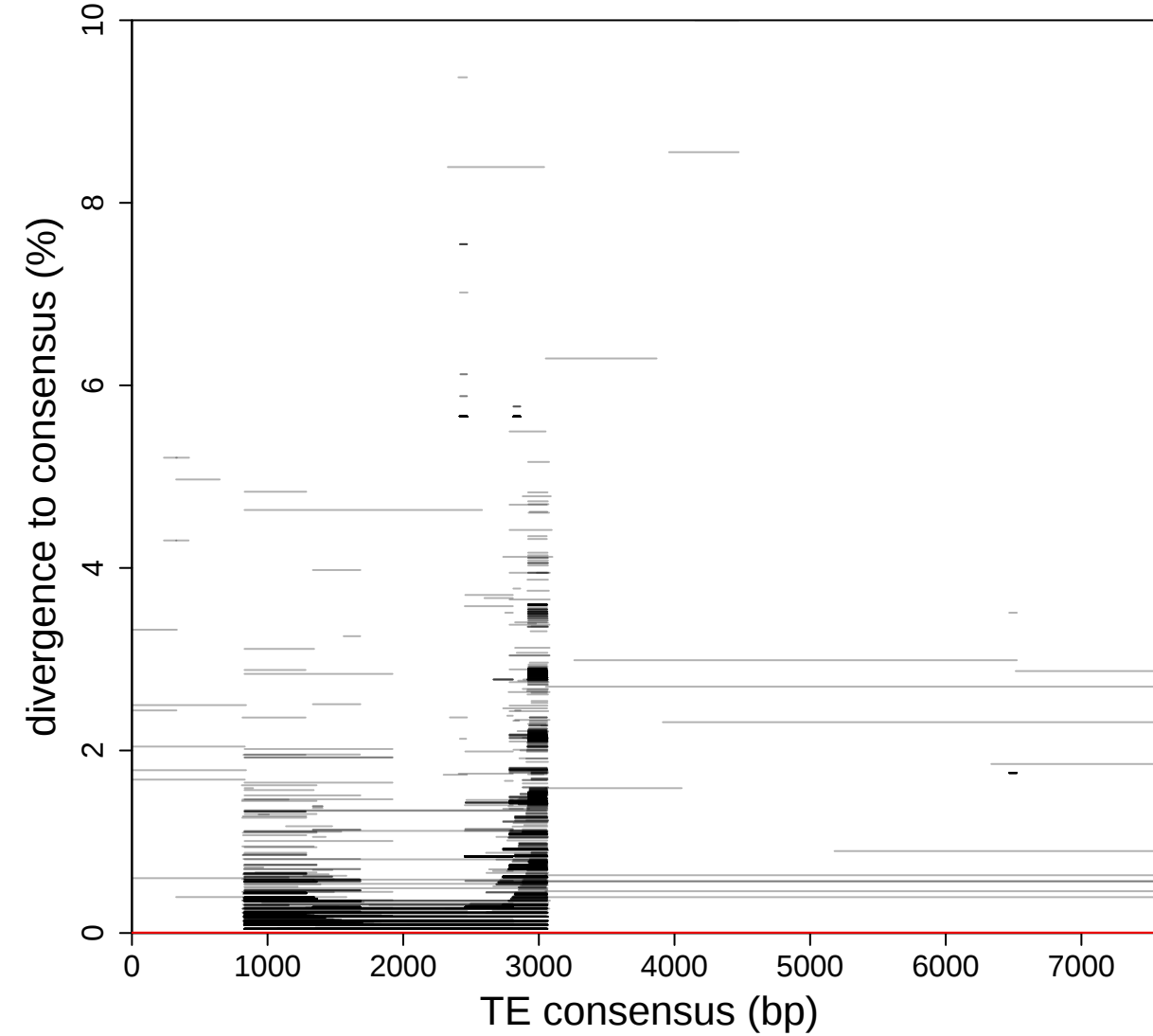
3.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 2505bp; fragments: 5726; full length: 0 (>=2254.5bp)

After TEtrimmer Extended plot Blue lines are boundaries

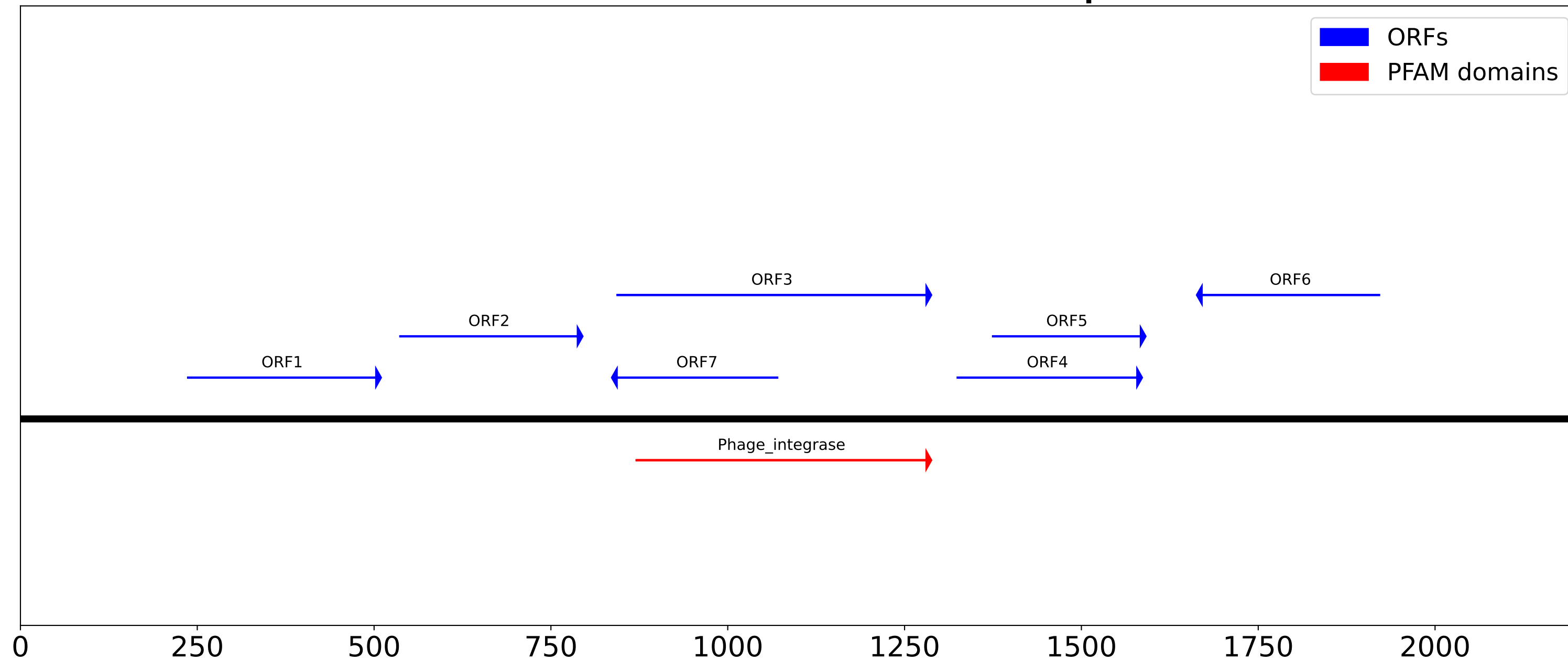


TE: ltr_1_family_1008
size: 7630bp; fragments: 5769; full length: 1 (>=6867bp)

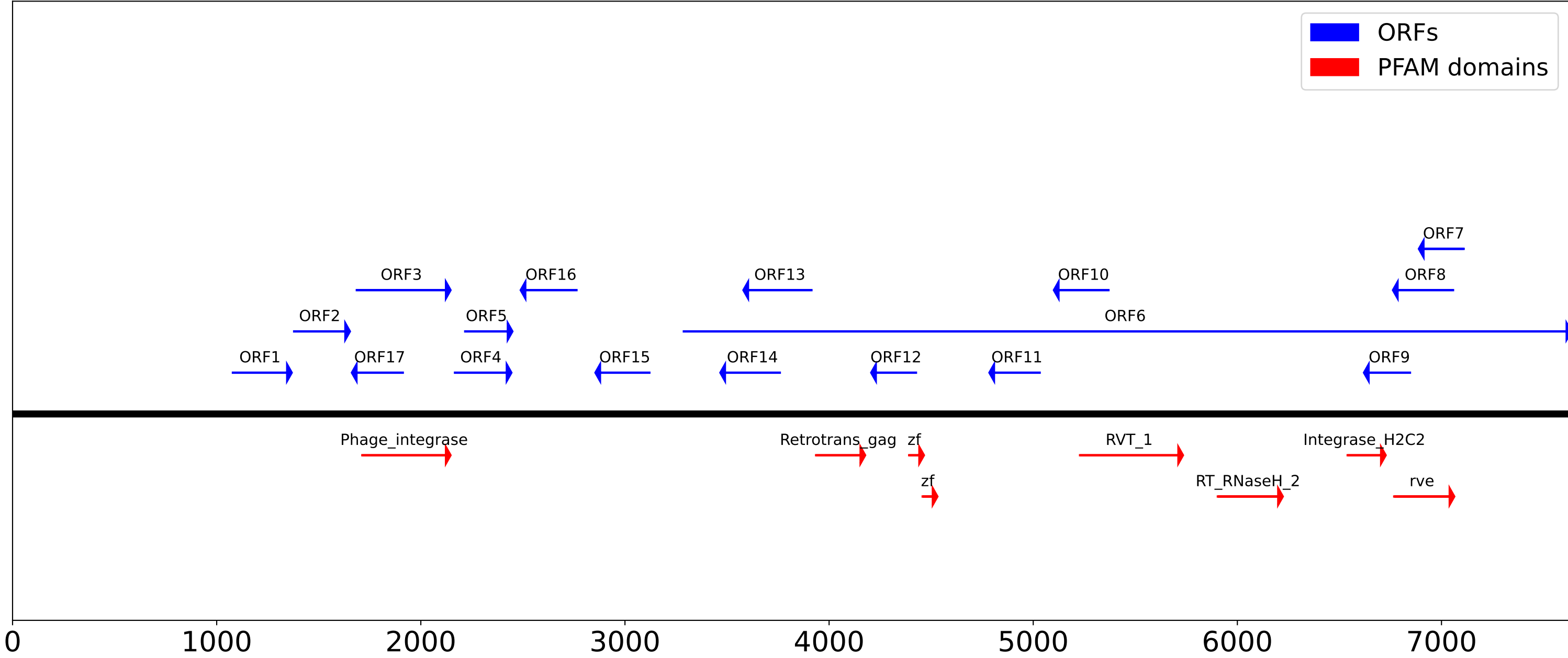
Before TEtrimmer 7630 bp



After TEtrimmer ORF and PFAM domain plot

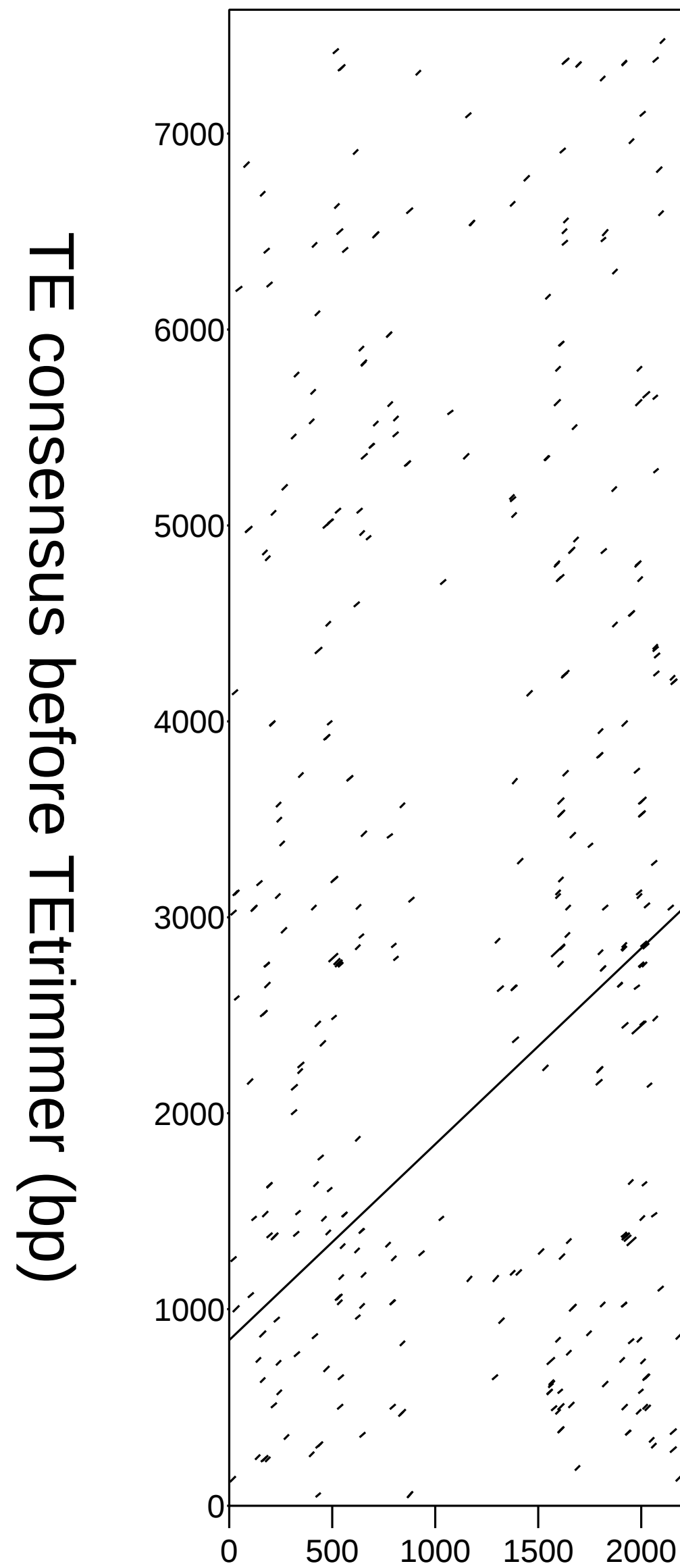


Before TEtrimmer ORF and PFAM domain plot



Dotplot

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



E consensus after TETrimmer (bp)