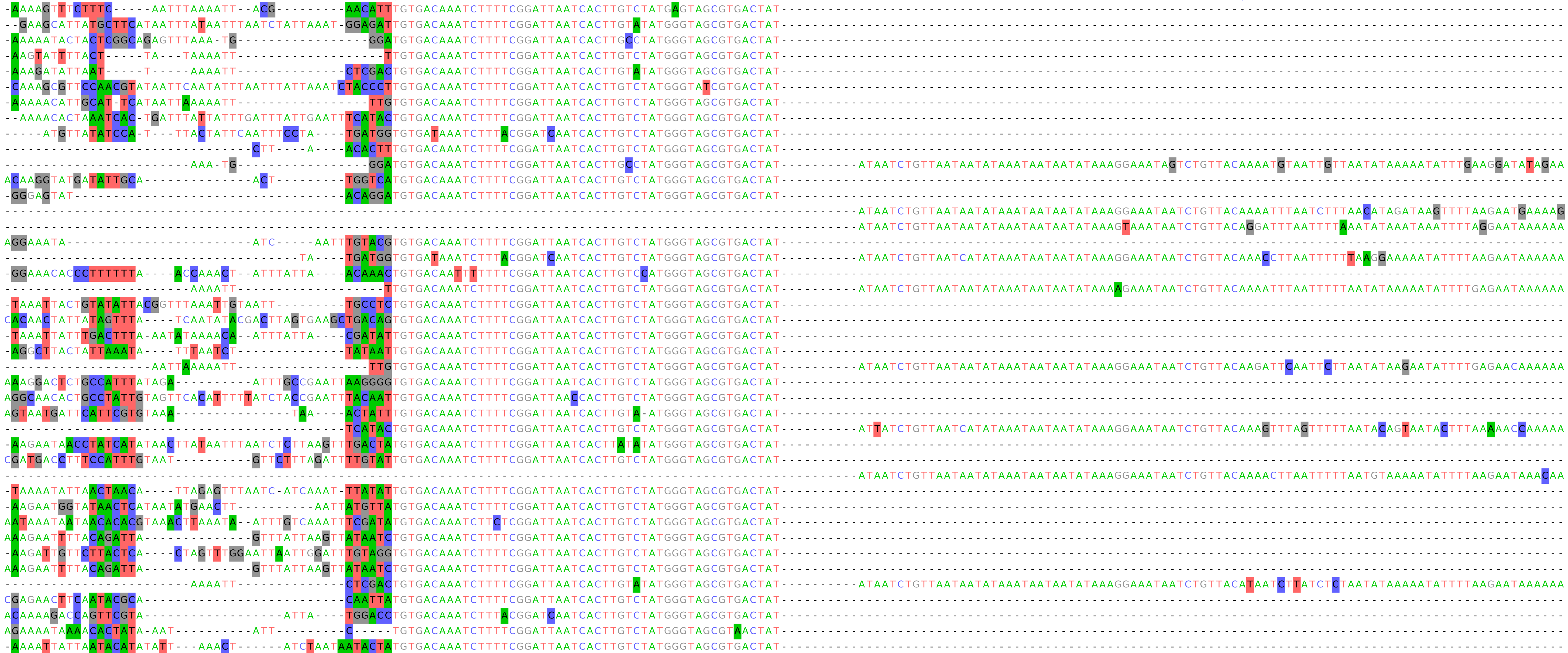
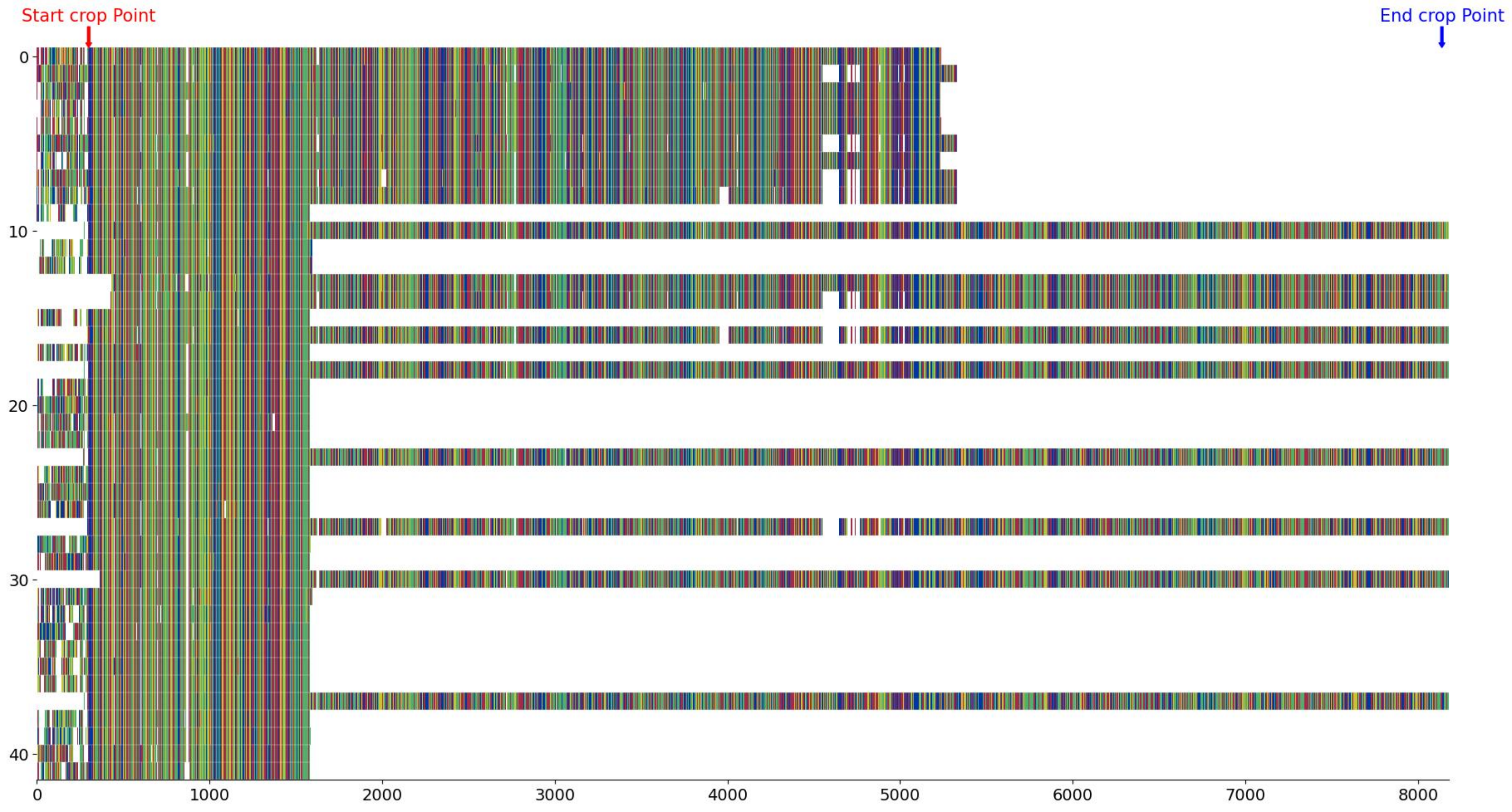


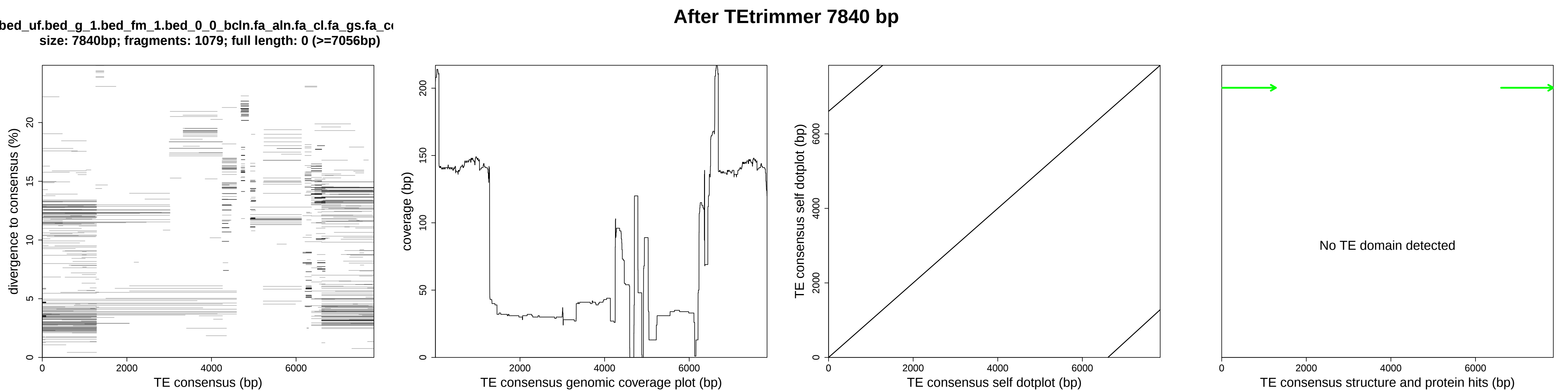
MSA length = 7840

Start crop Point

End crop Point

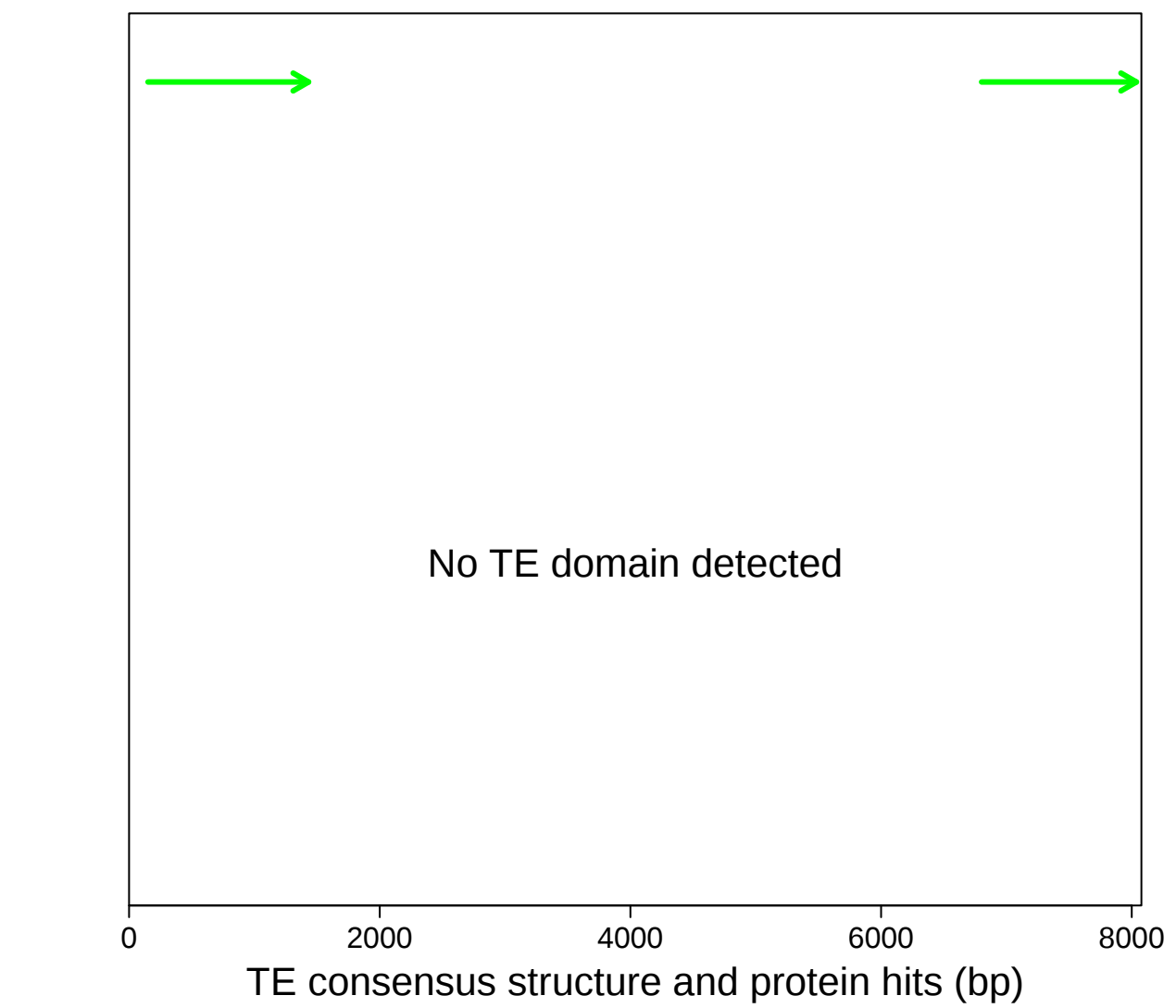
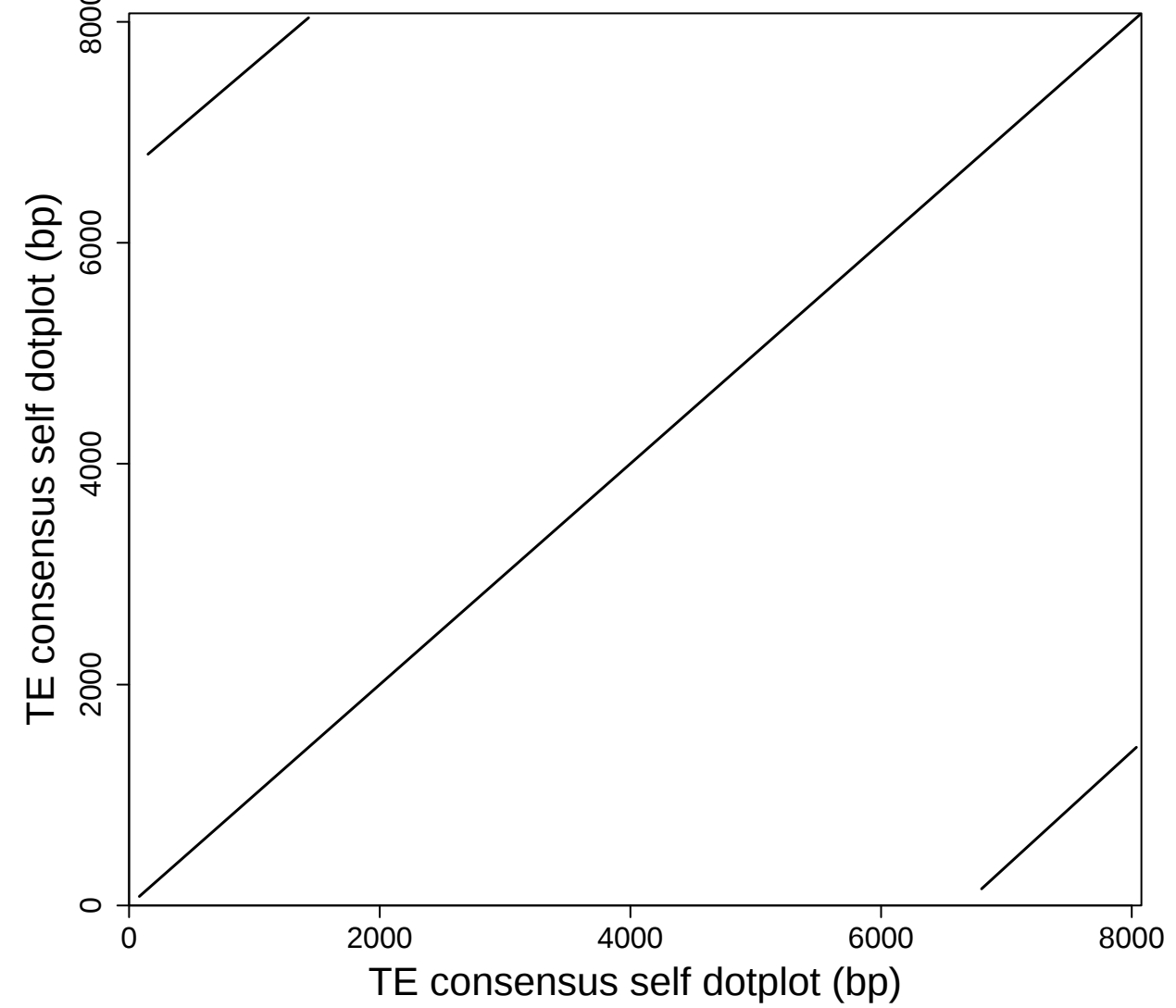
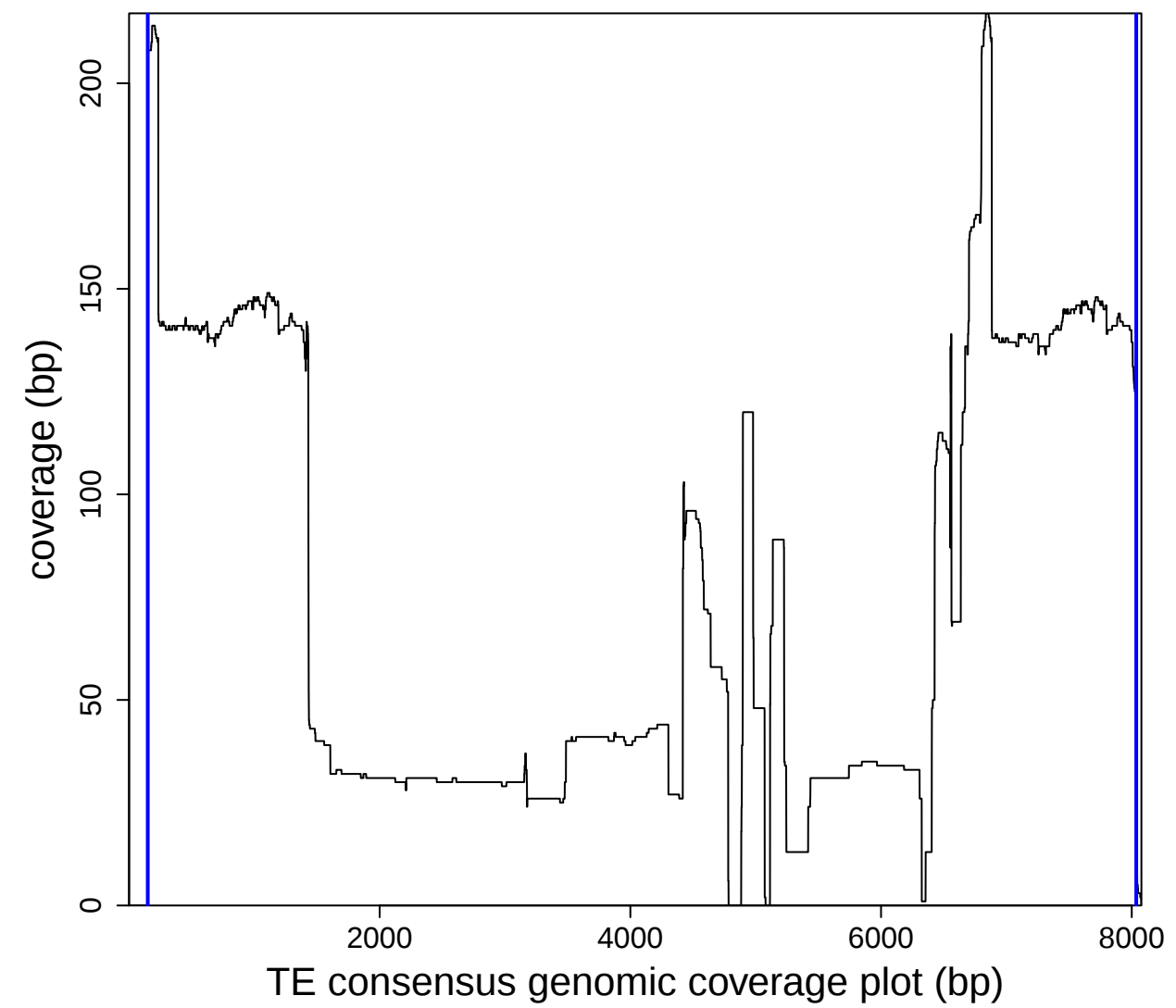
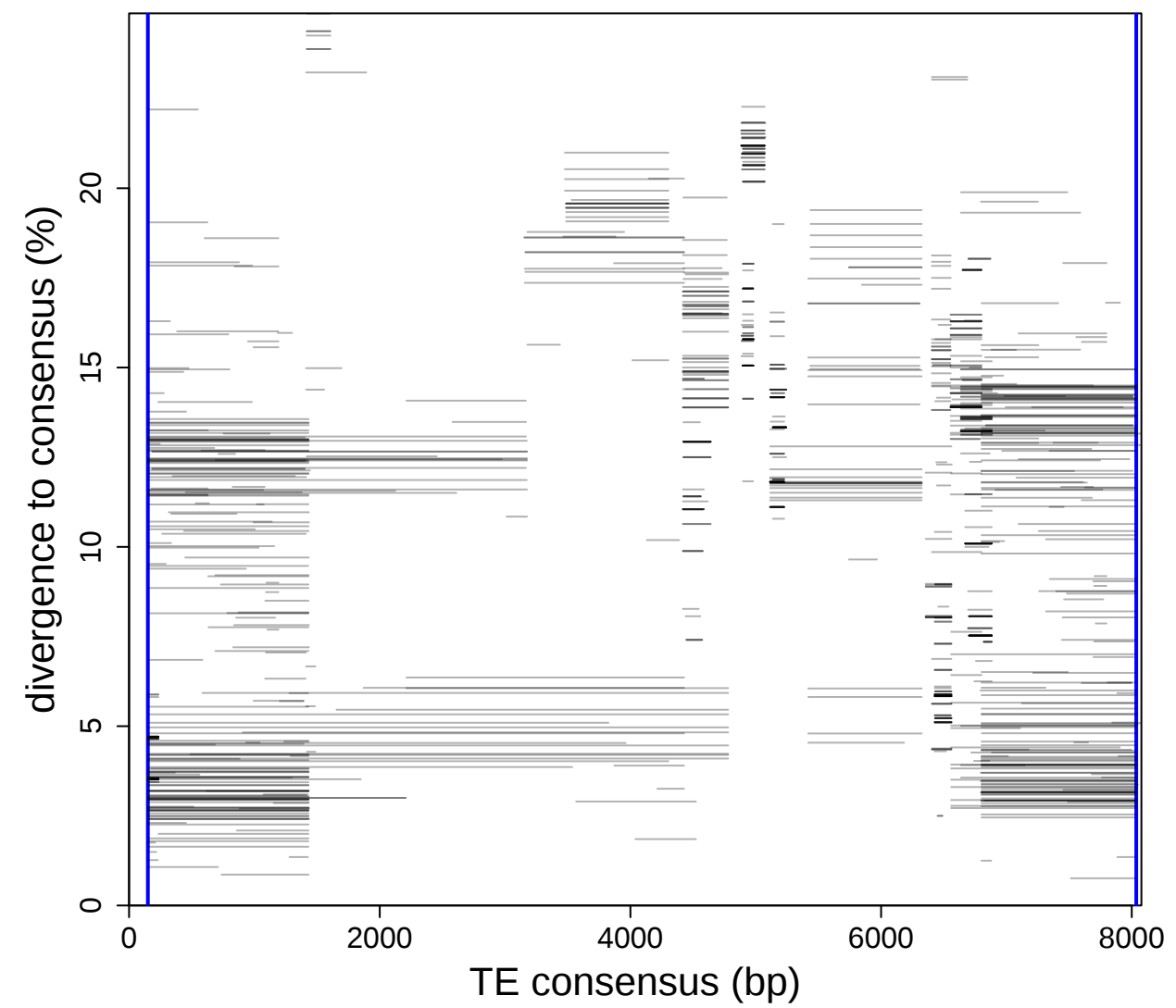






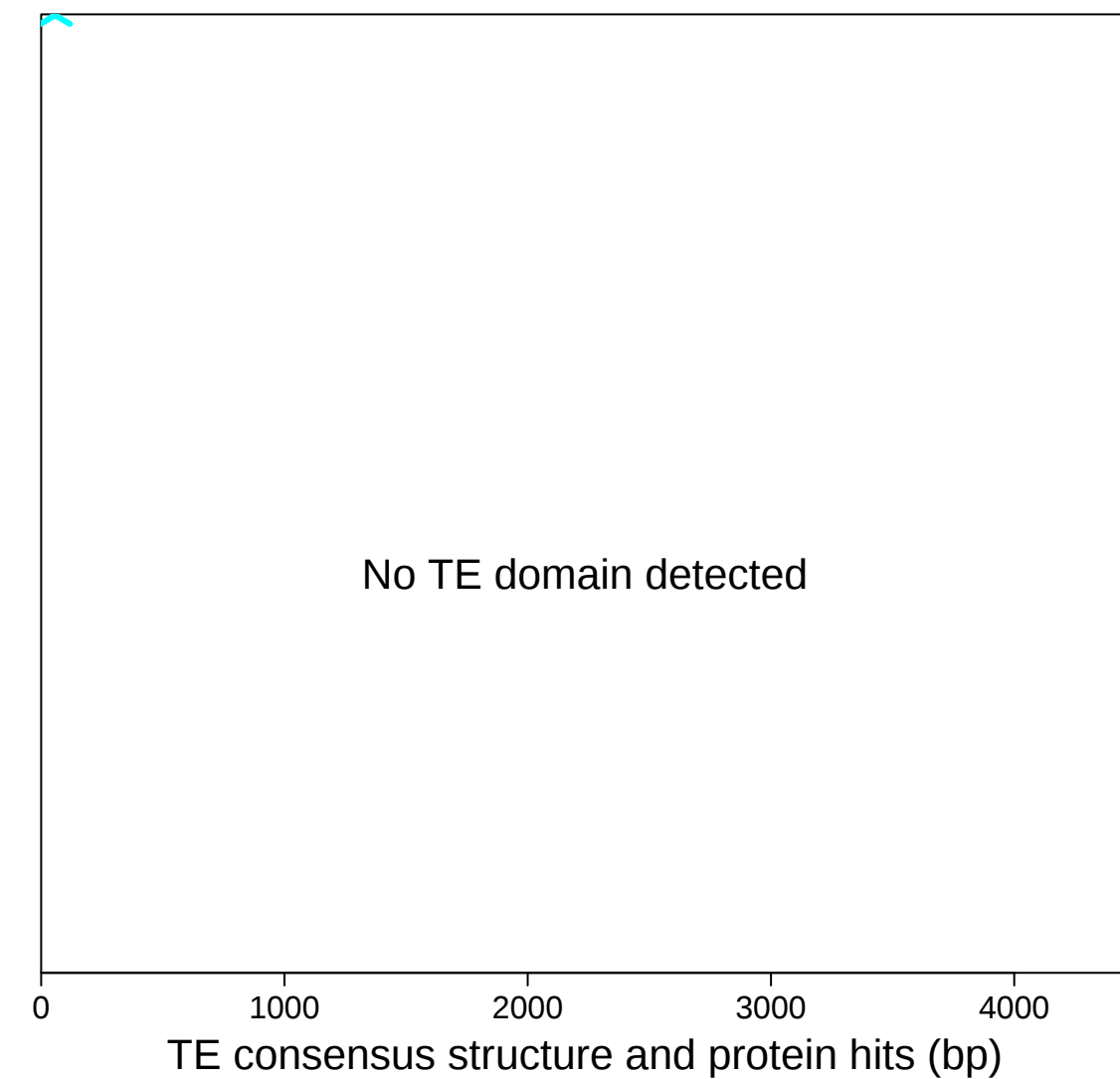
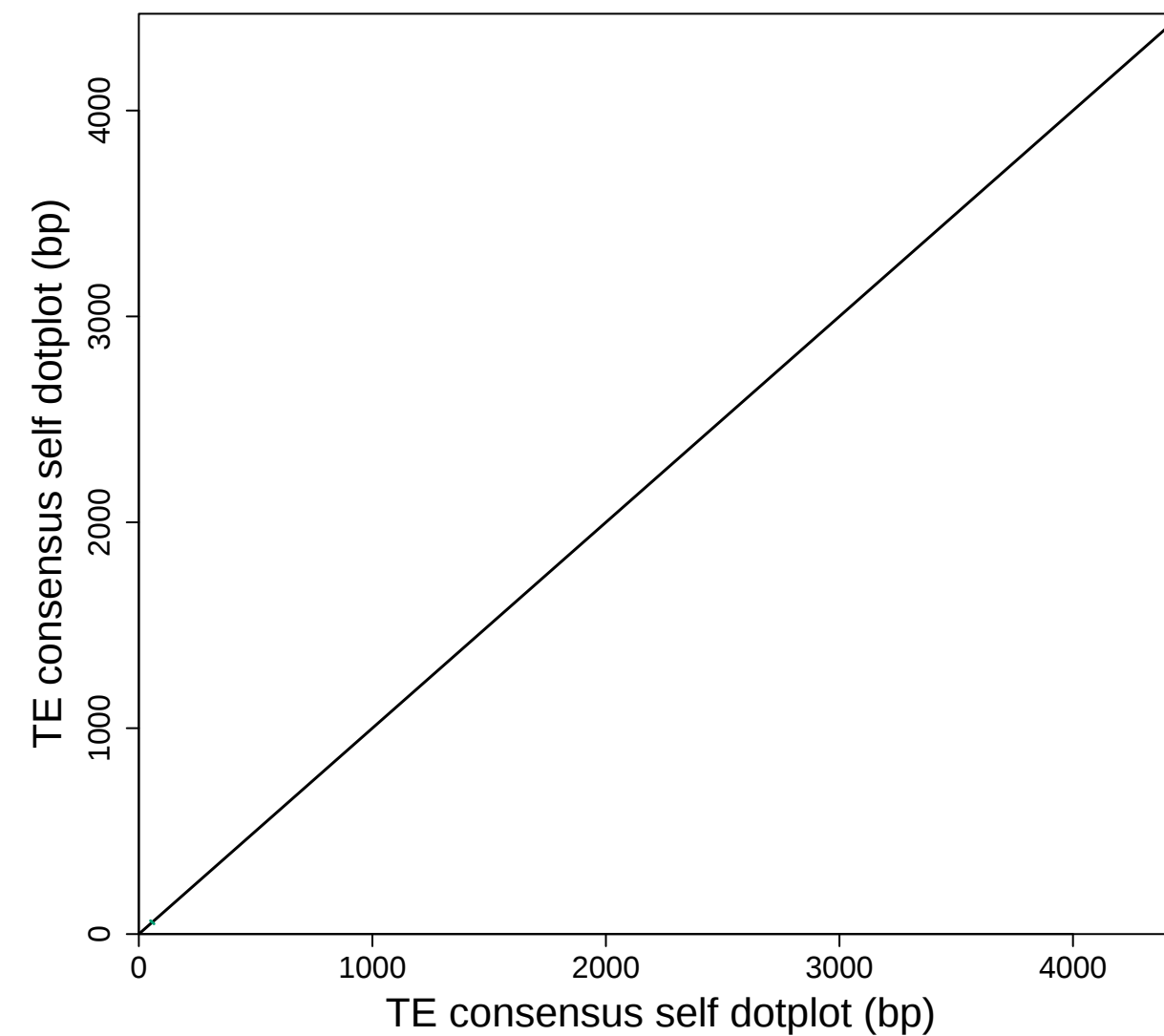
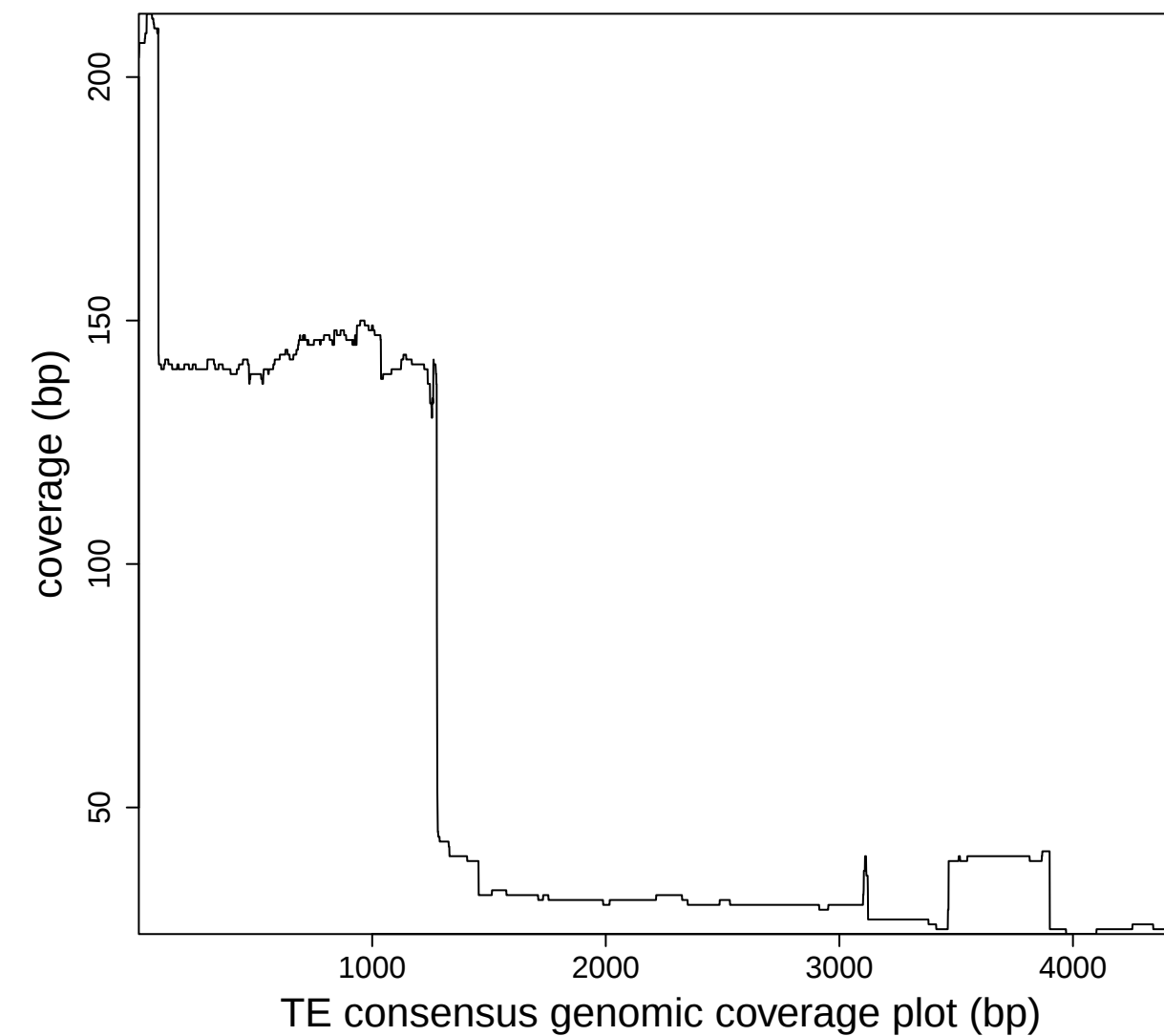
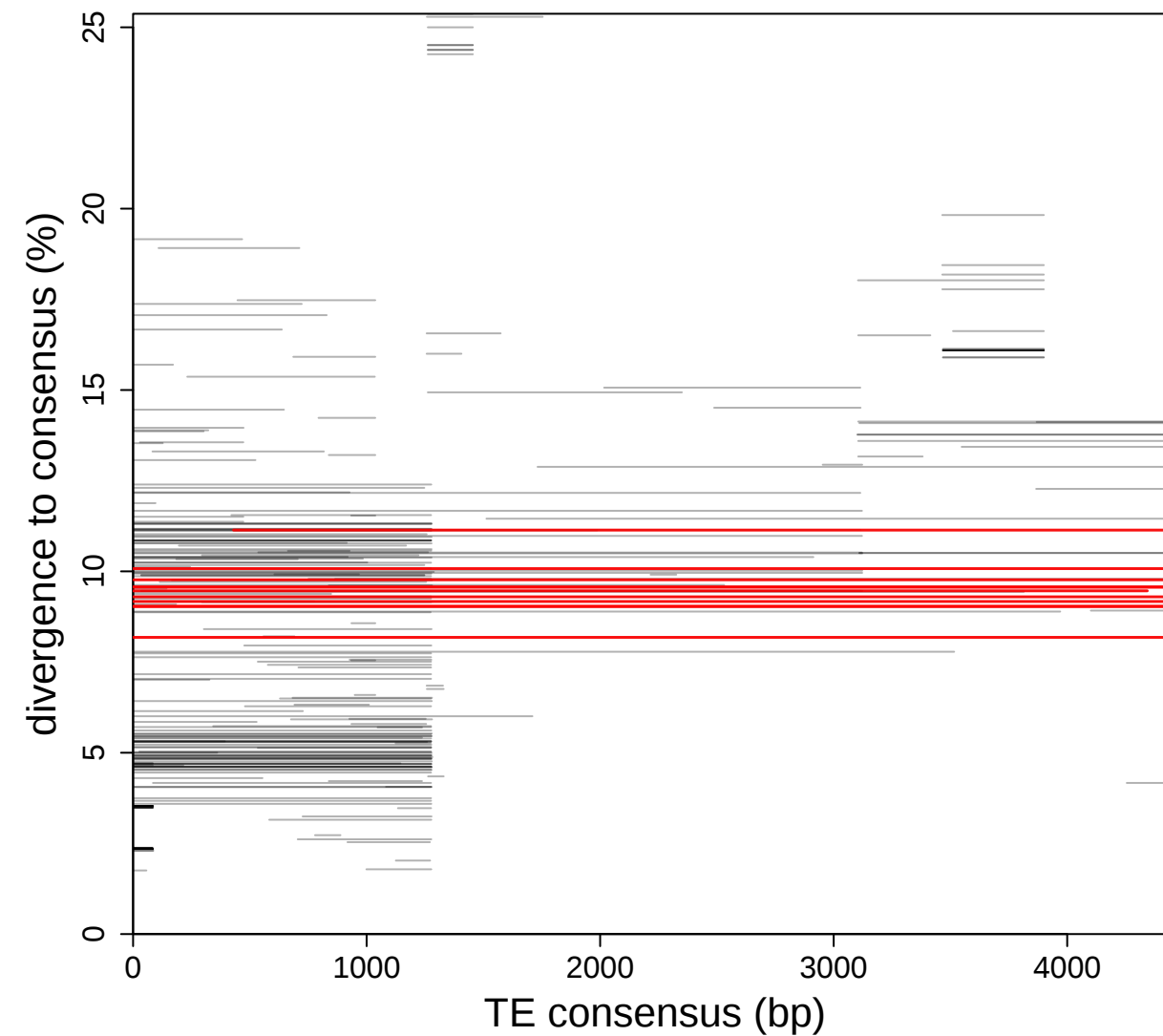
ly_2686.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.f
size: 8077bp; fragments: 1078; full length: 0 (>=7269.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

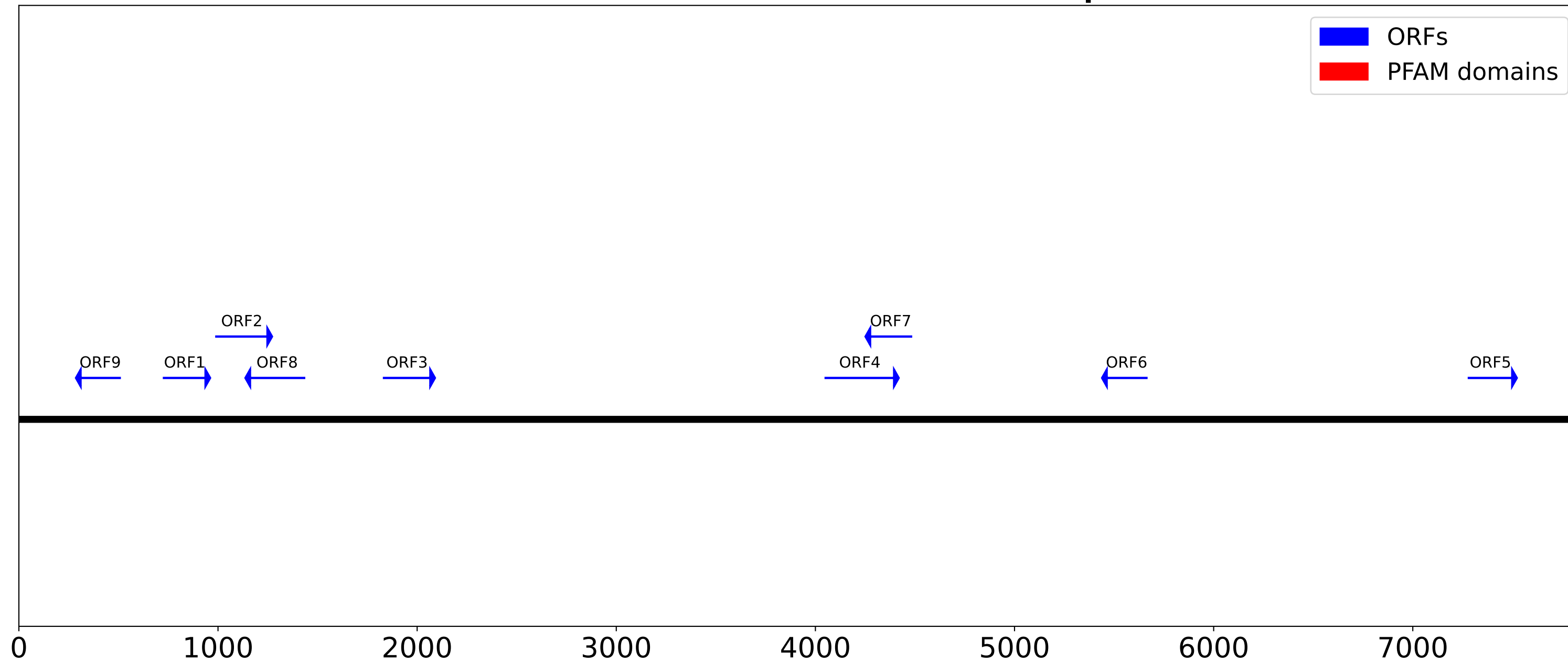


TE: rnd_5_family_2686
size: 4470bp; fragments: 317; full length: 11 (≥ 4023 bp)

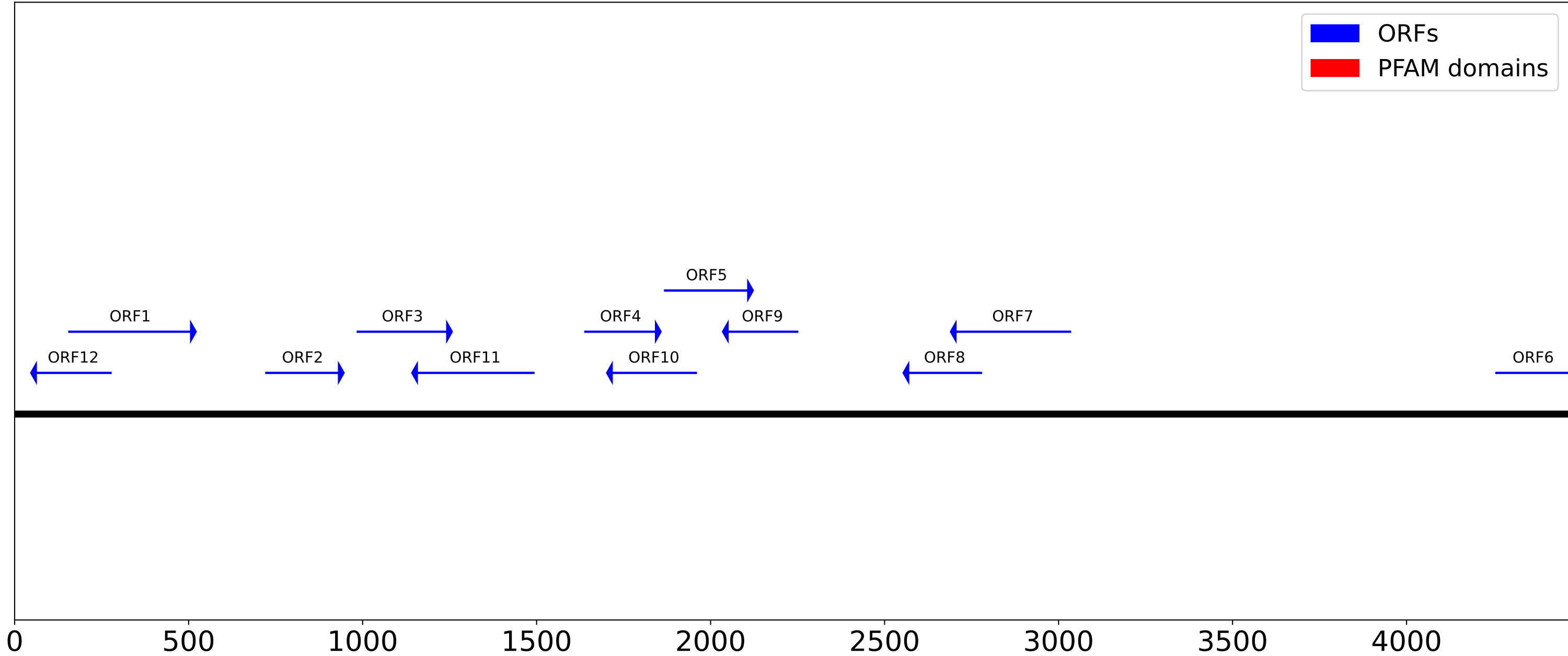
Before TEtrimmer 4470 bp



After TEtrimmer ORF and PFAM domain plot



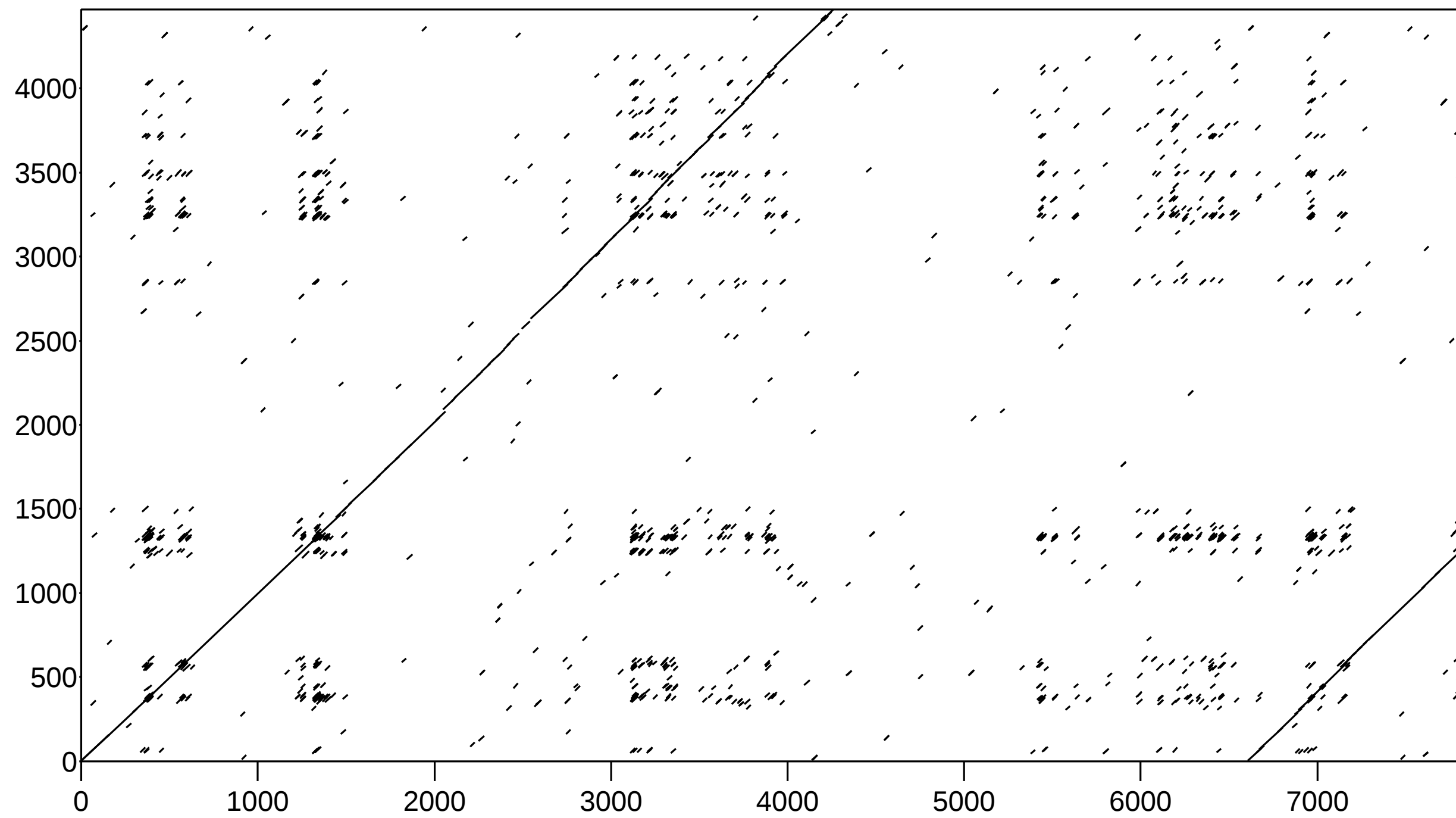
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)