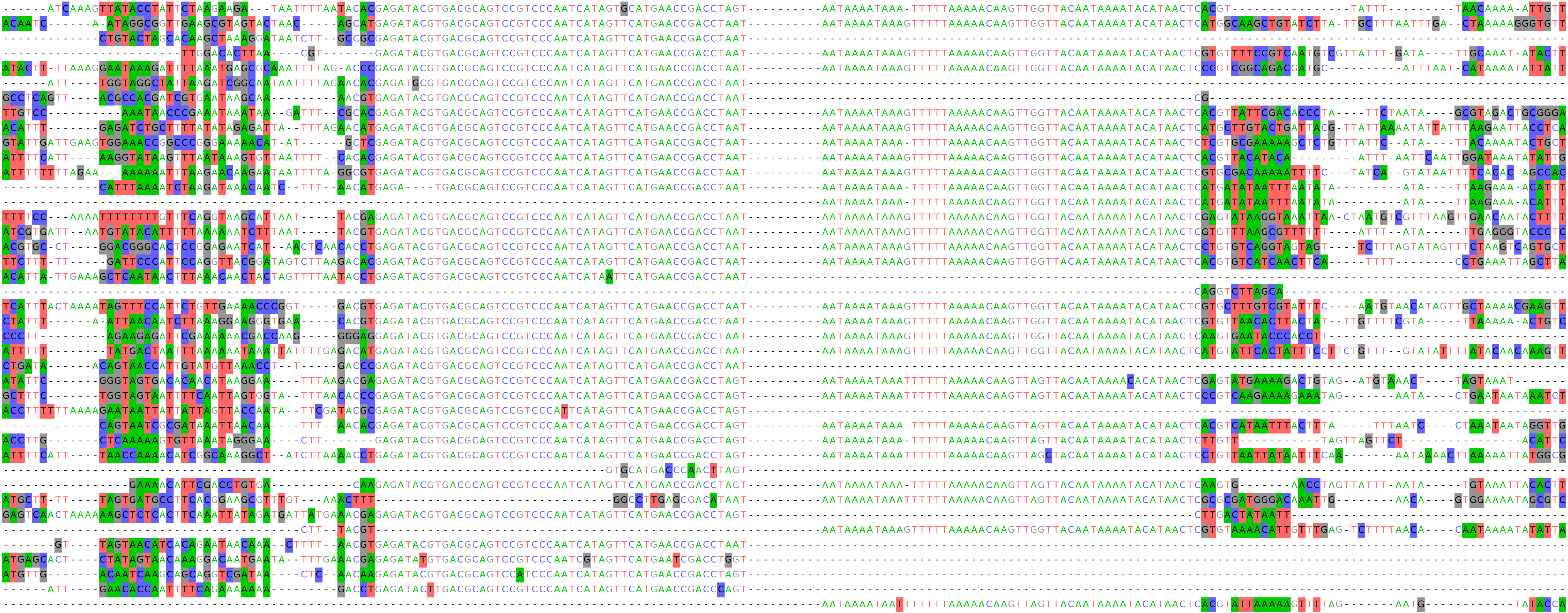
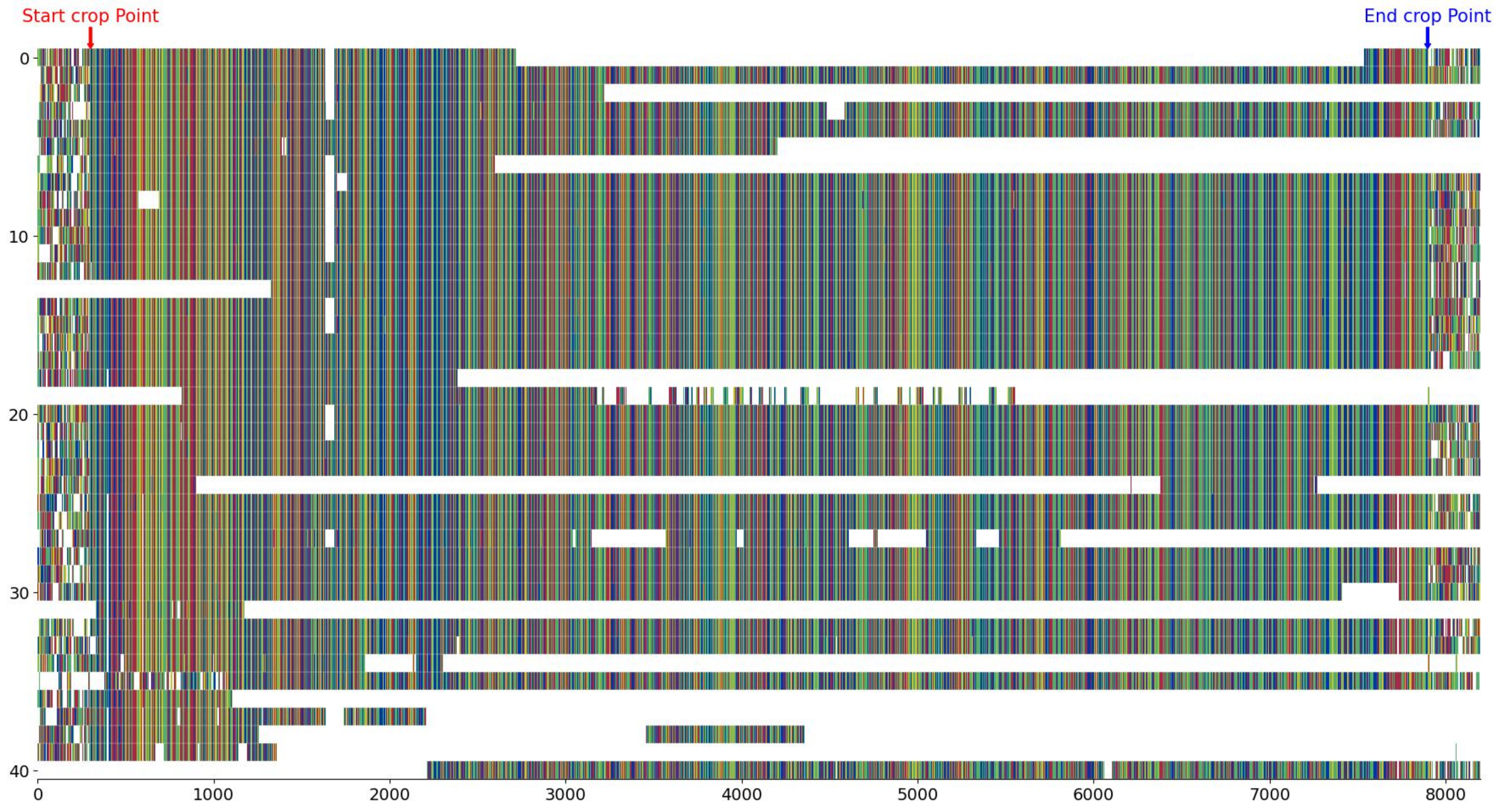


MSA length = 7599

Start crop Point

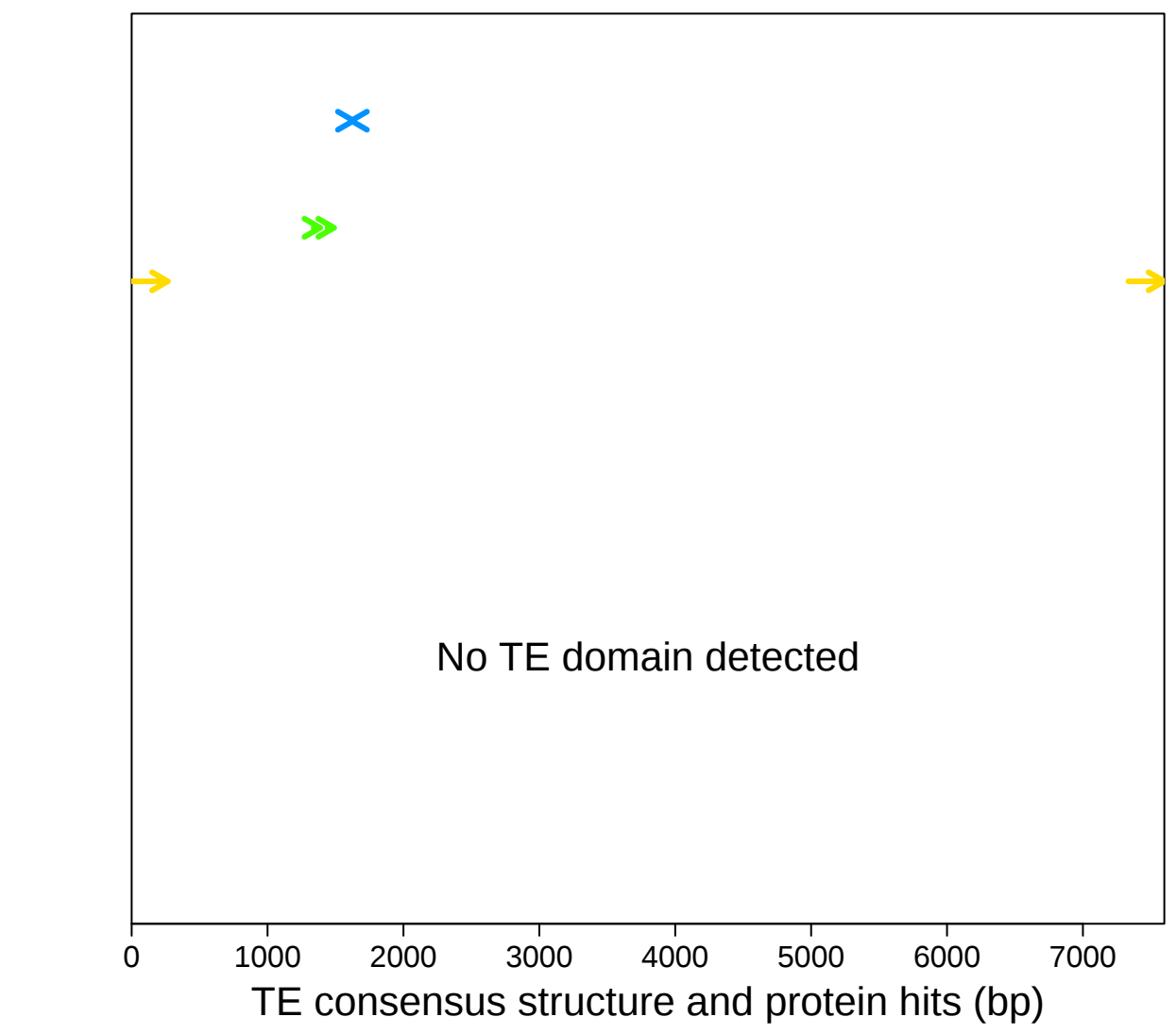
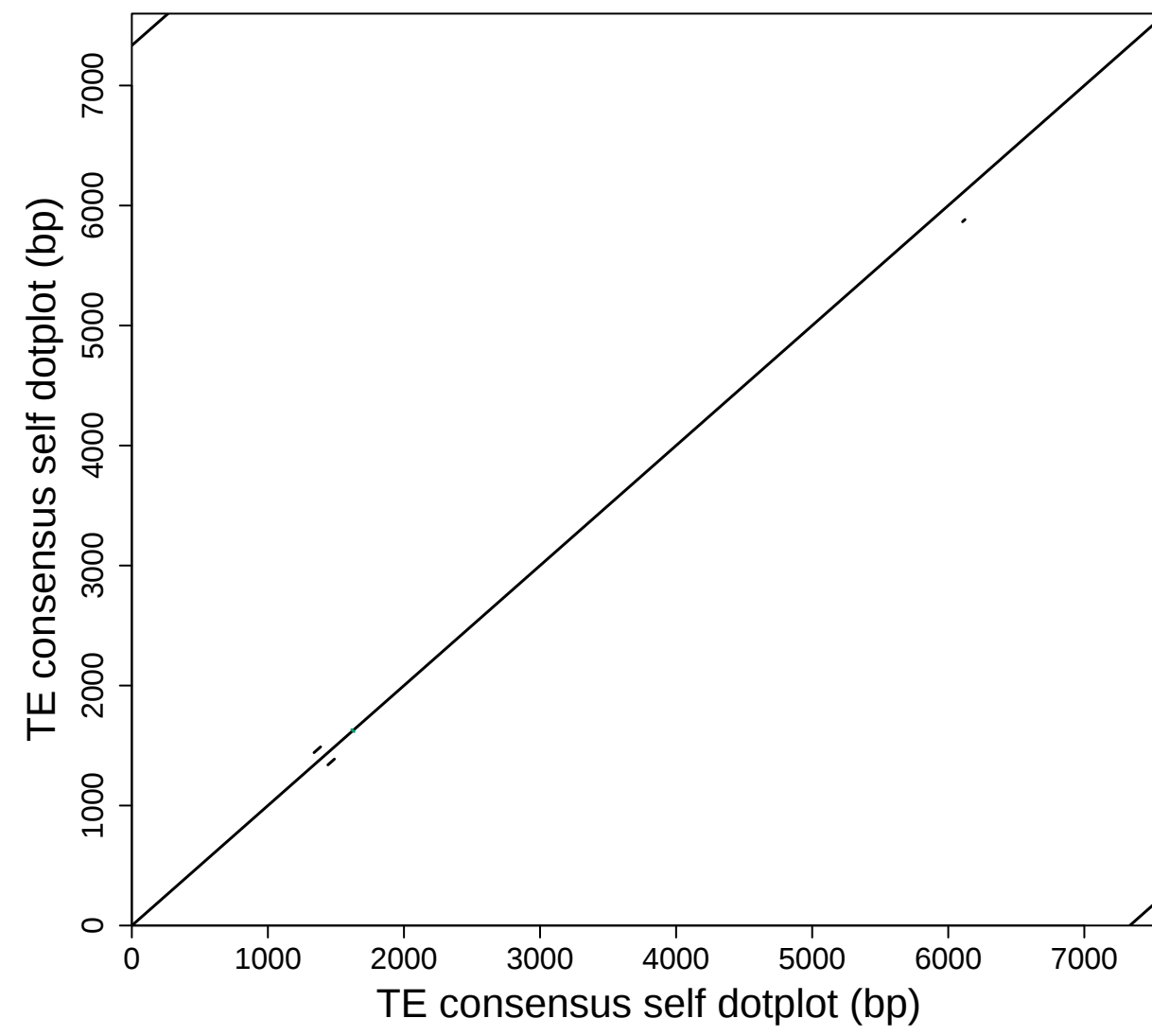
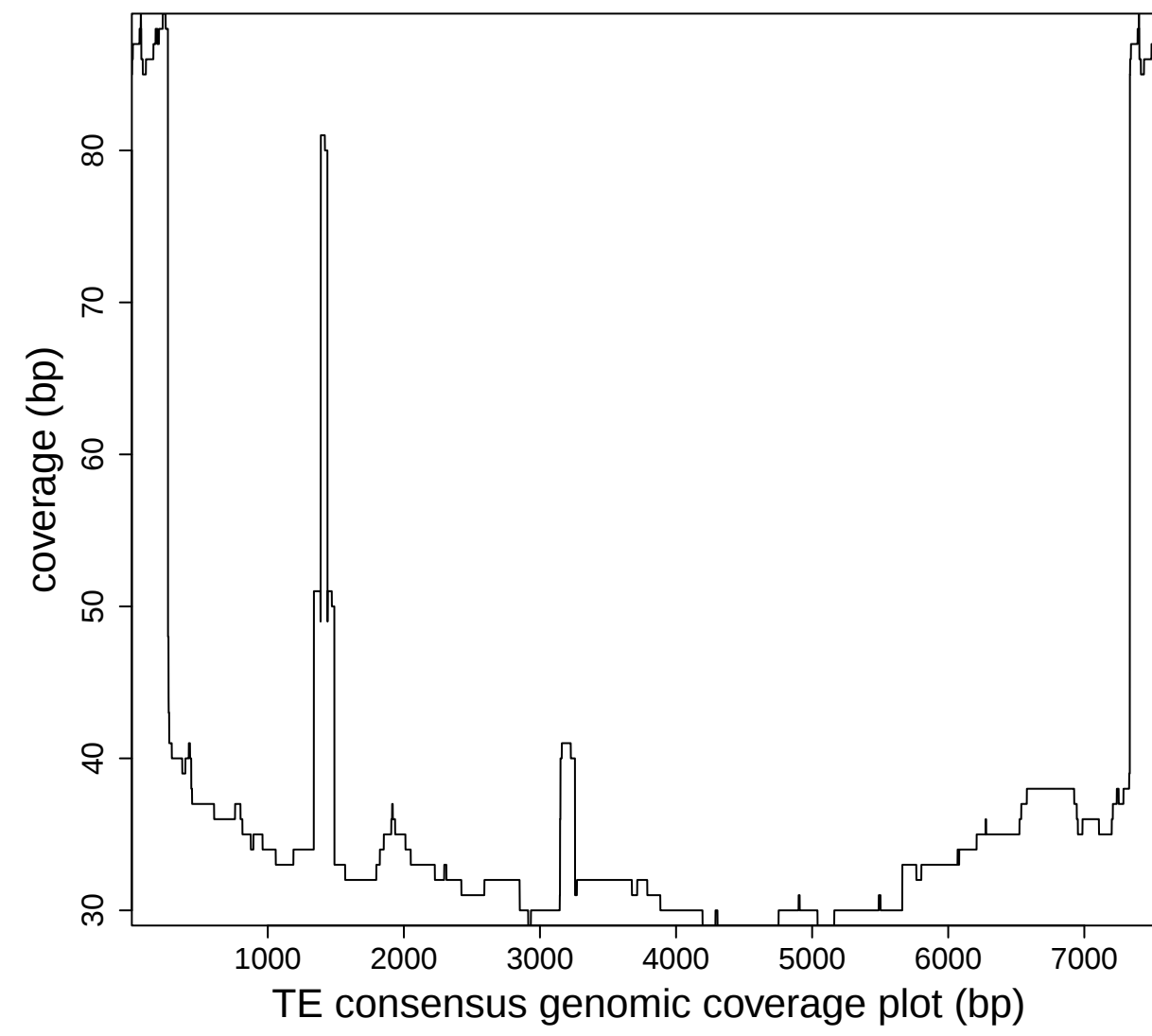
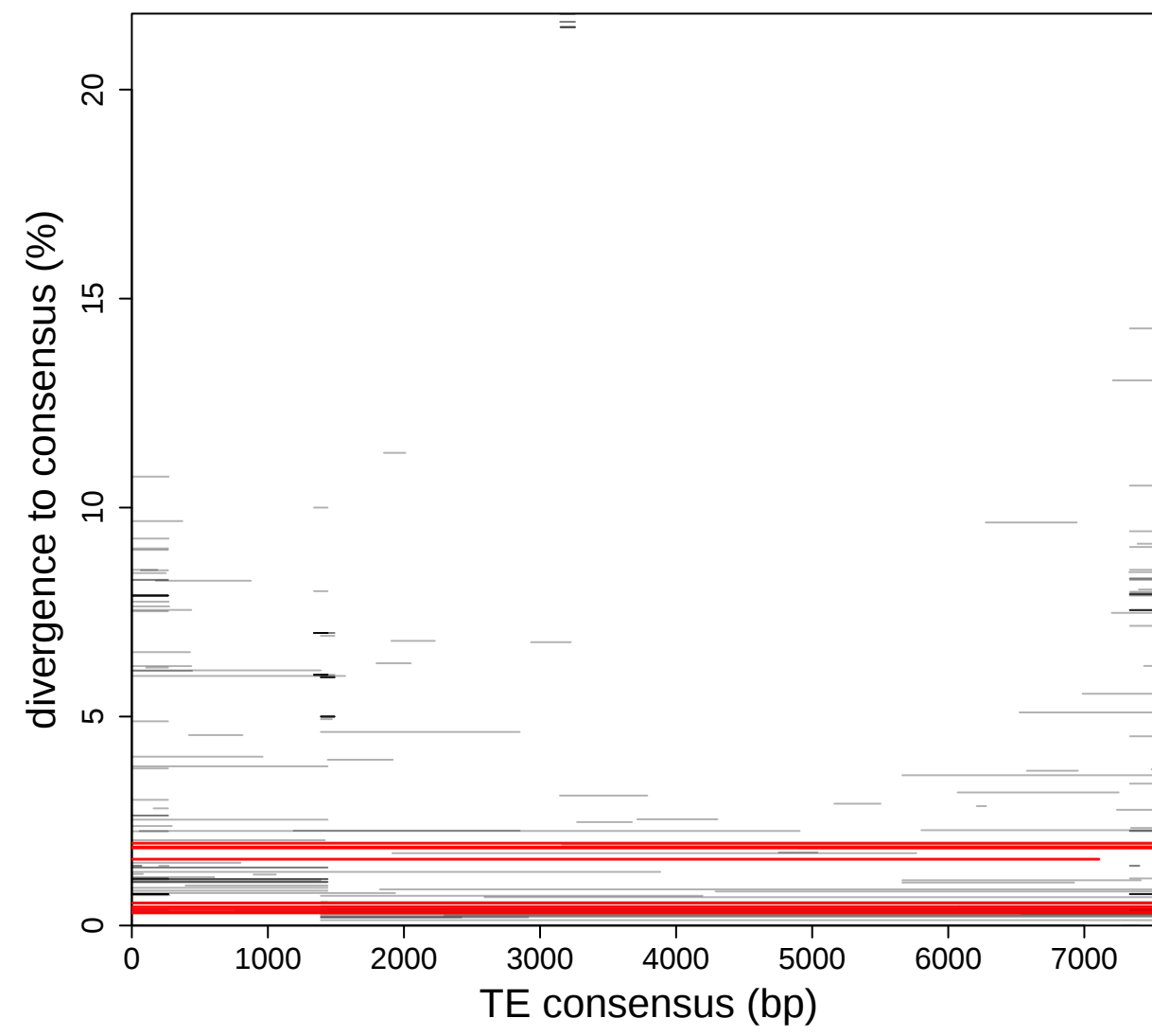
End crop Point





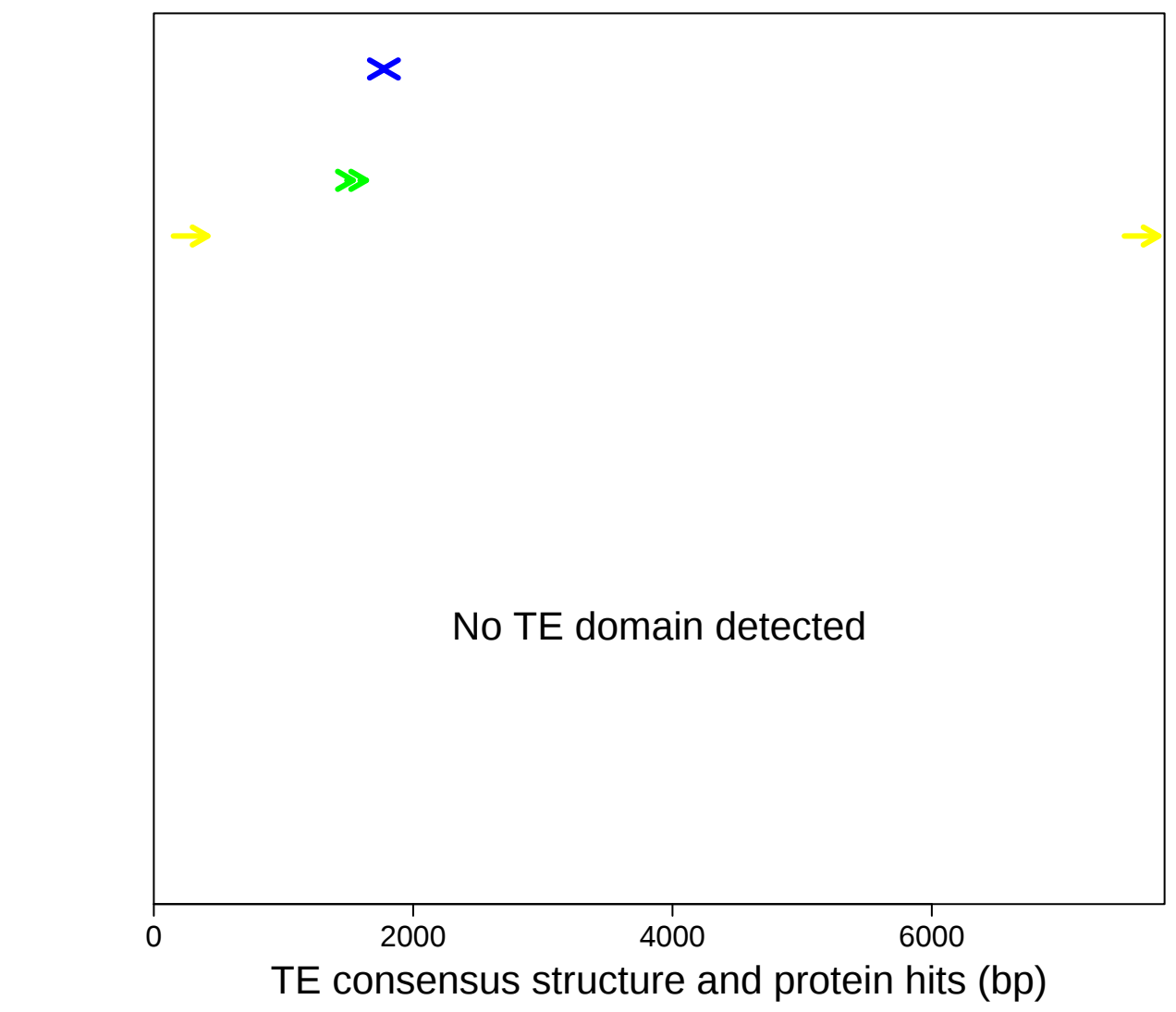
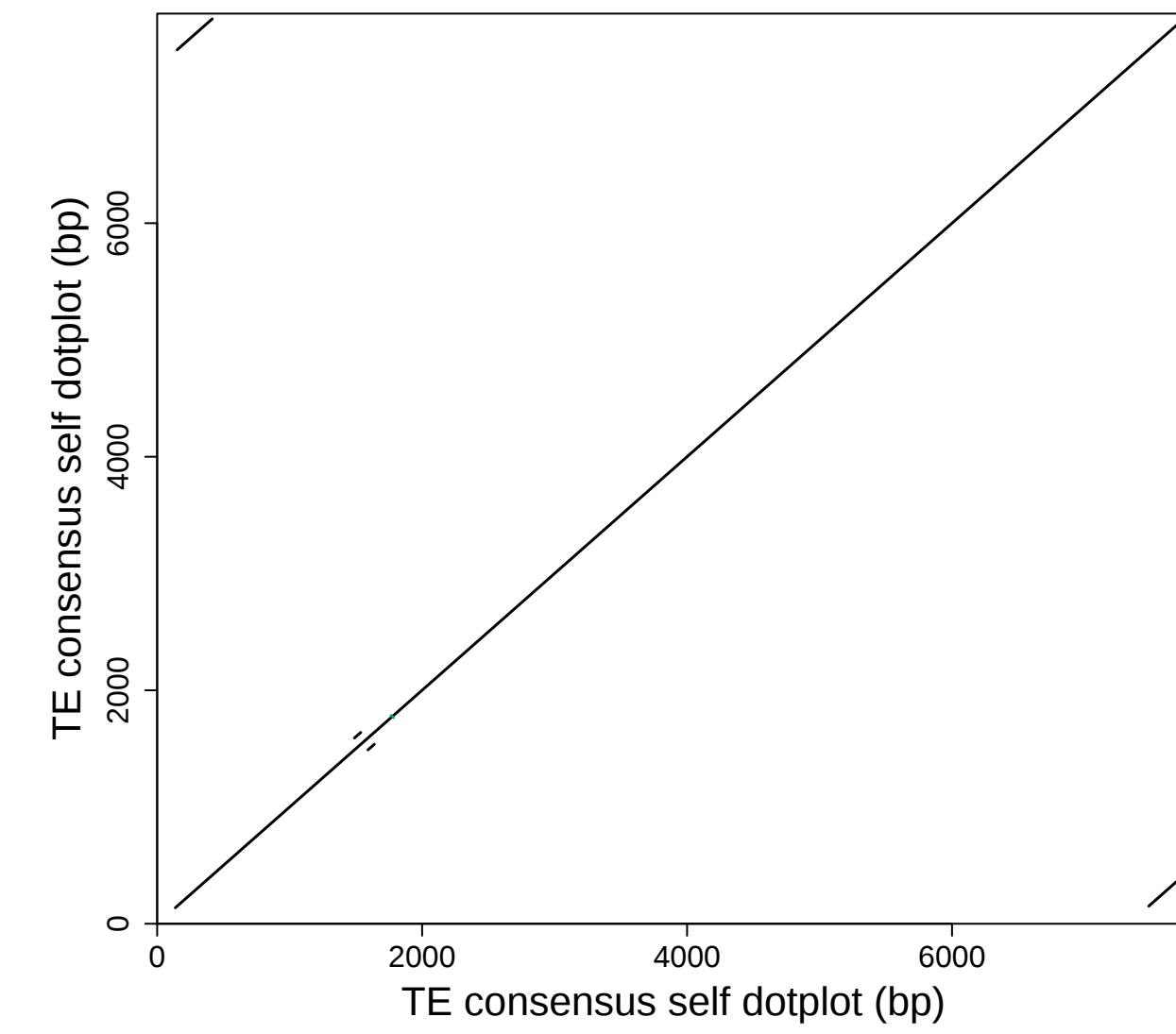
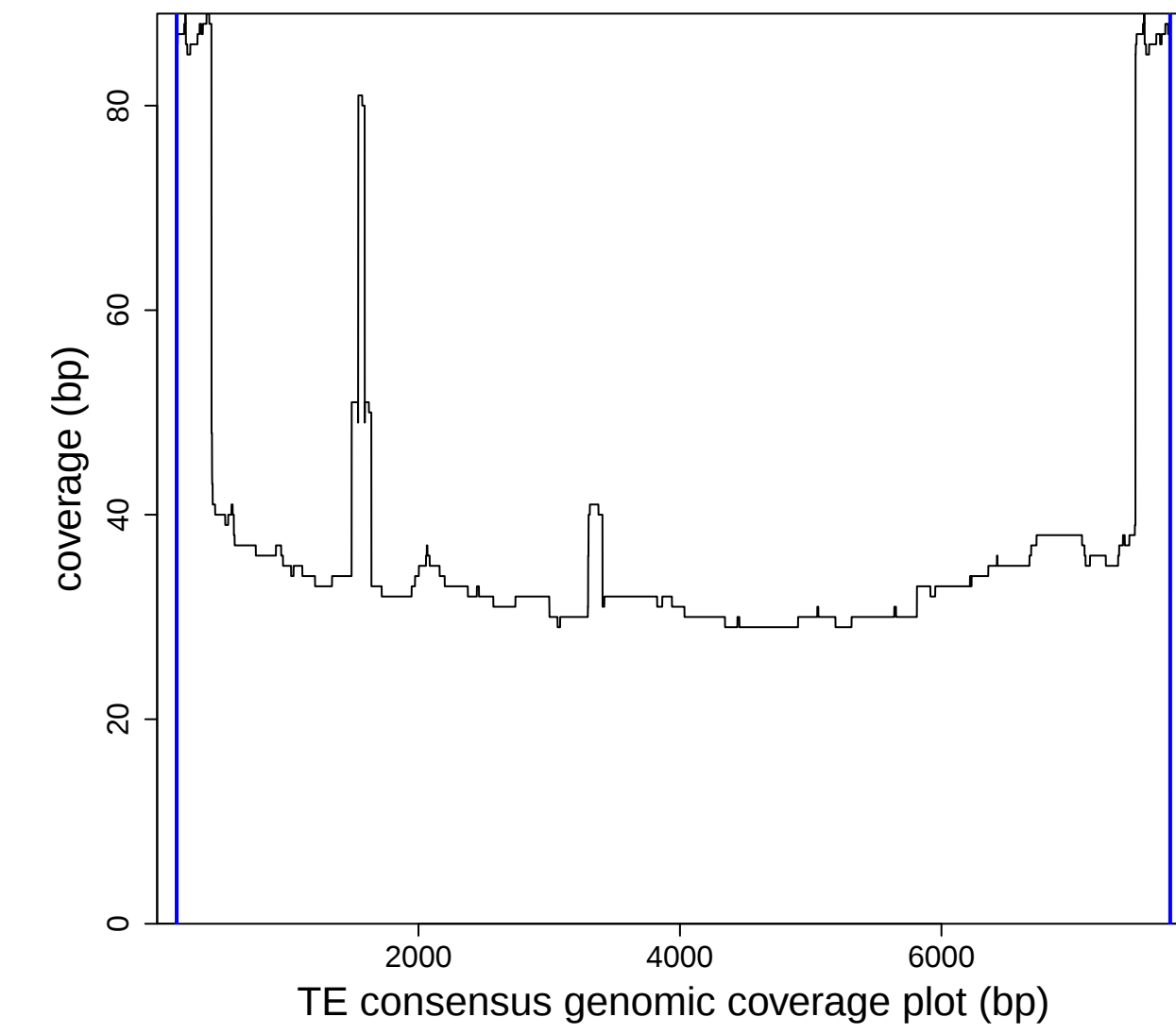
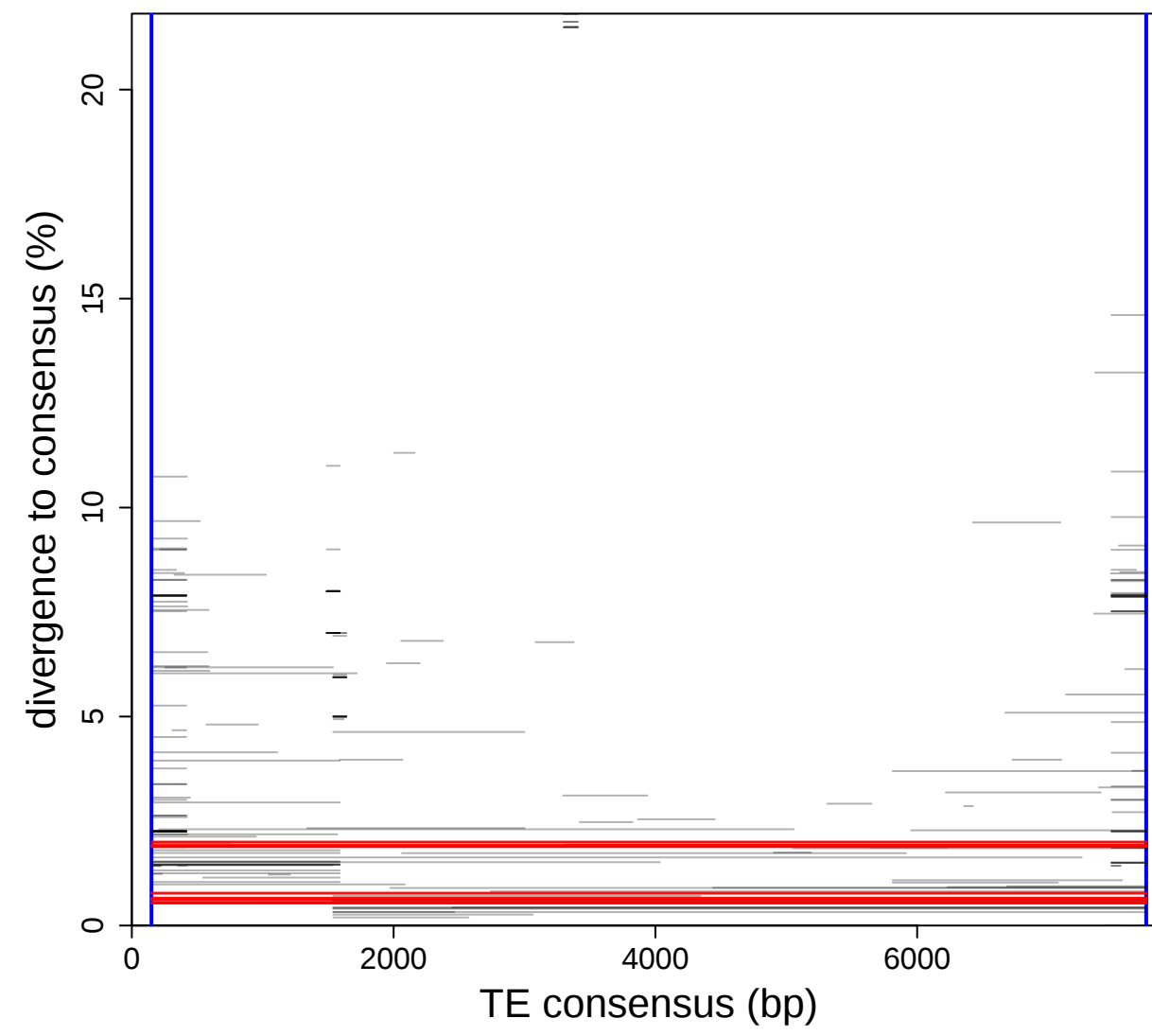
ed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 7599bp; fragments: 247; full length: 11 (>=6839.1bp)

After TETrimmer 7599 bp



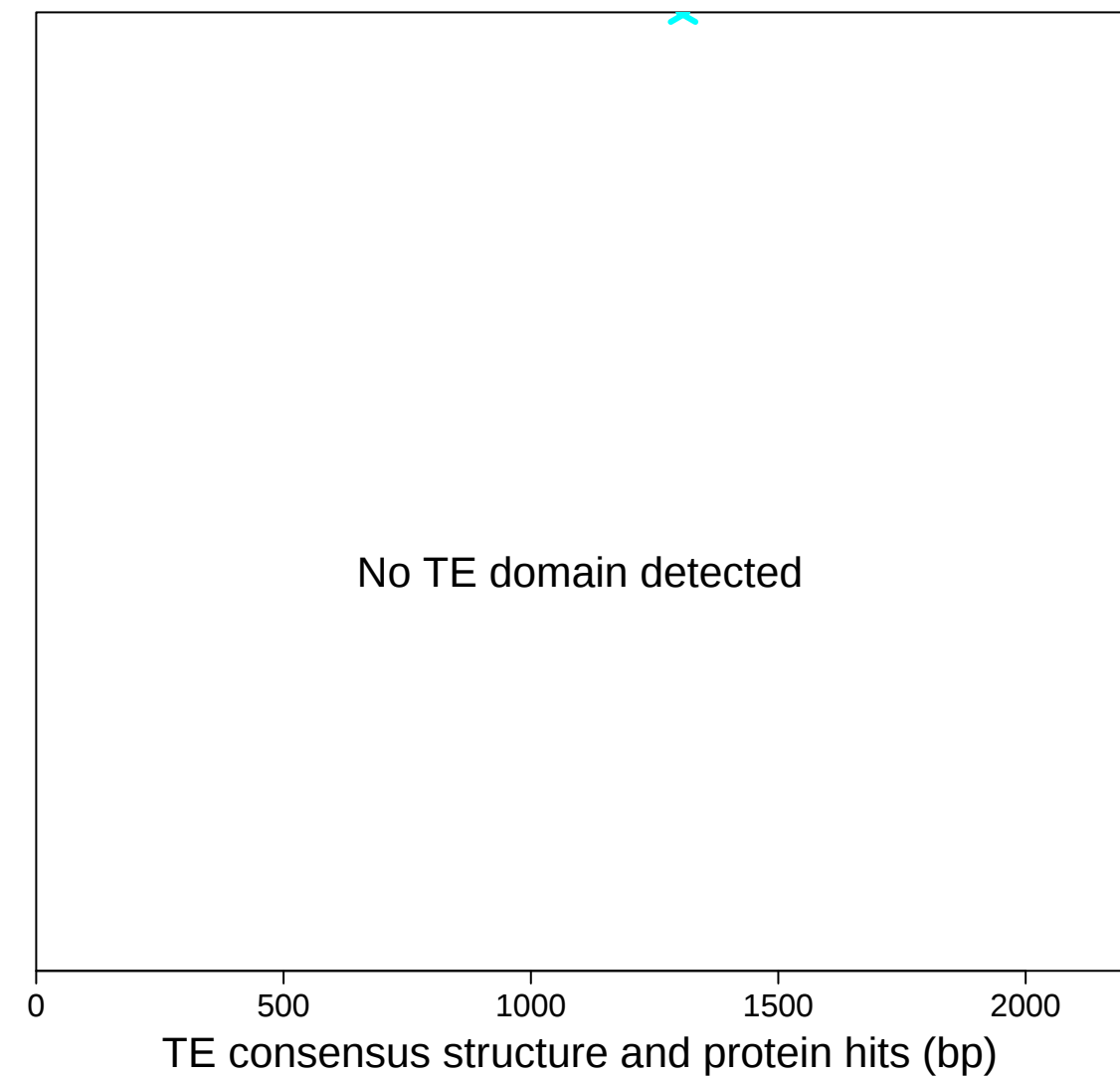
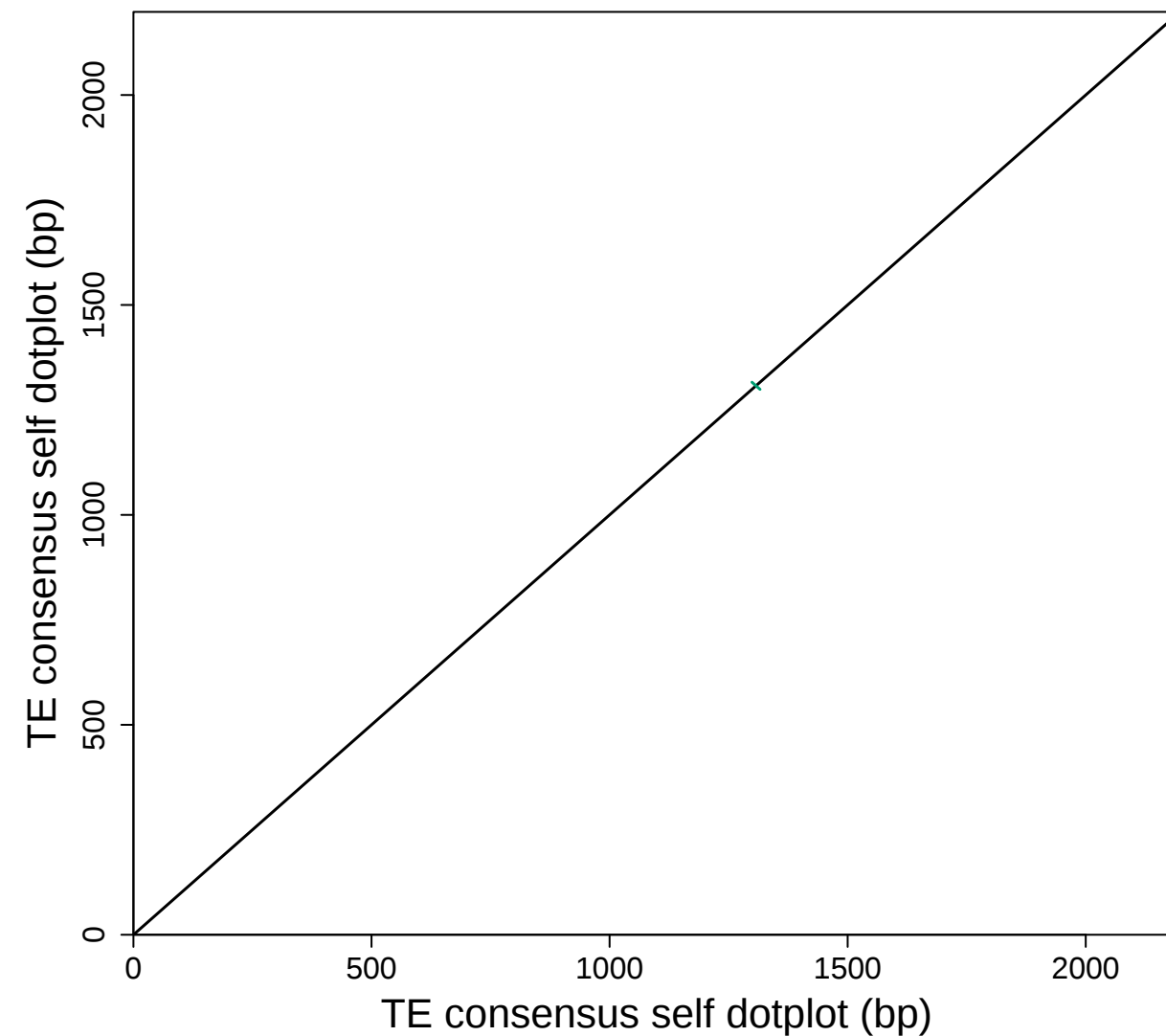
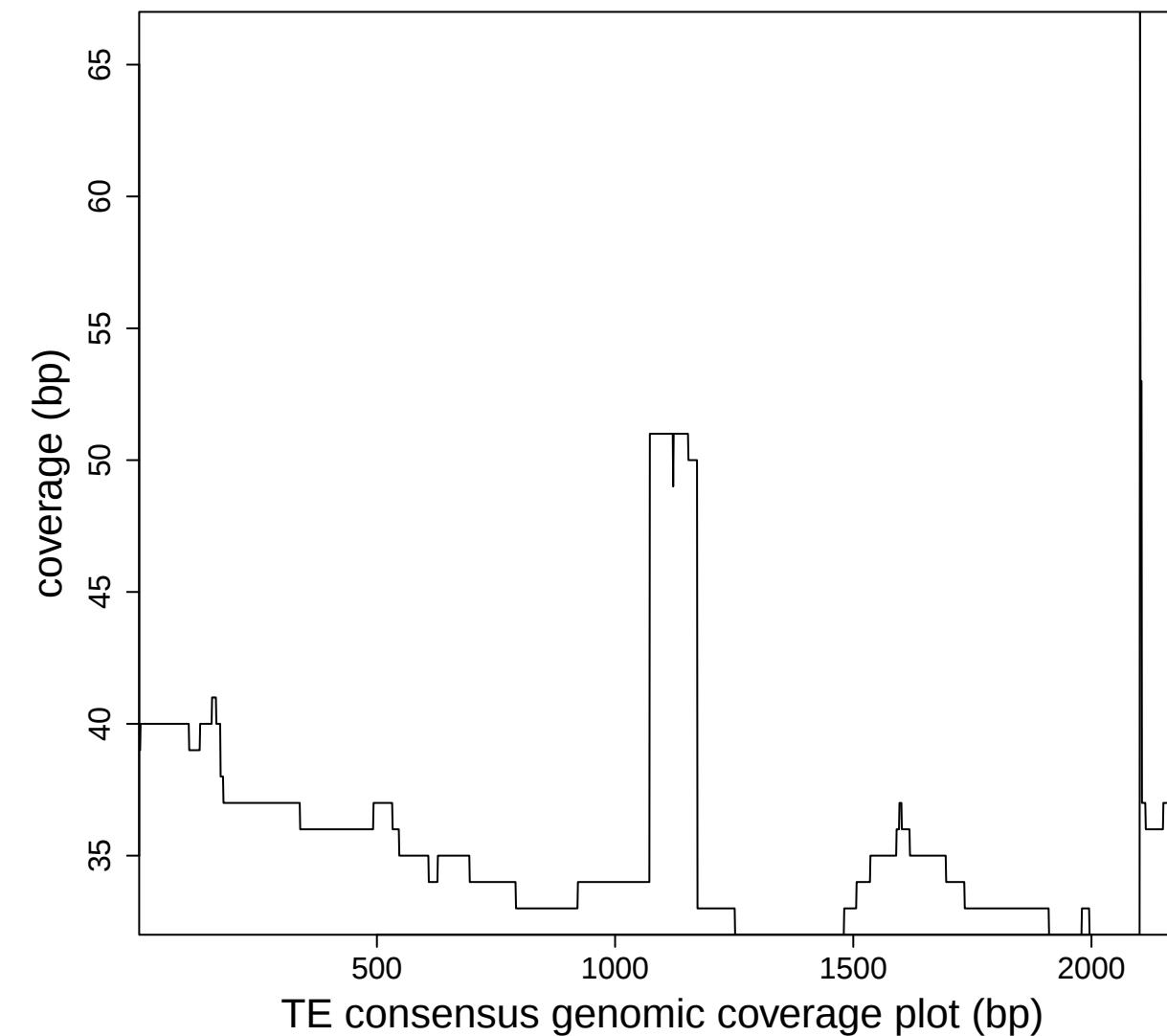
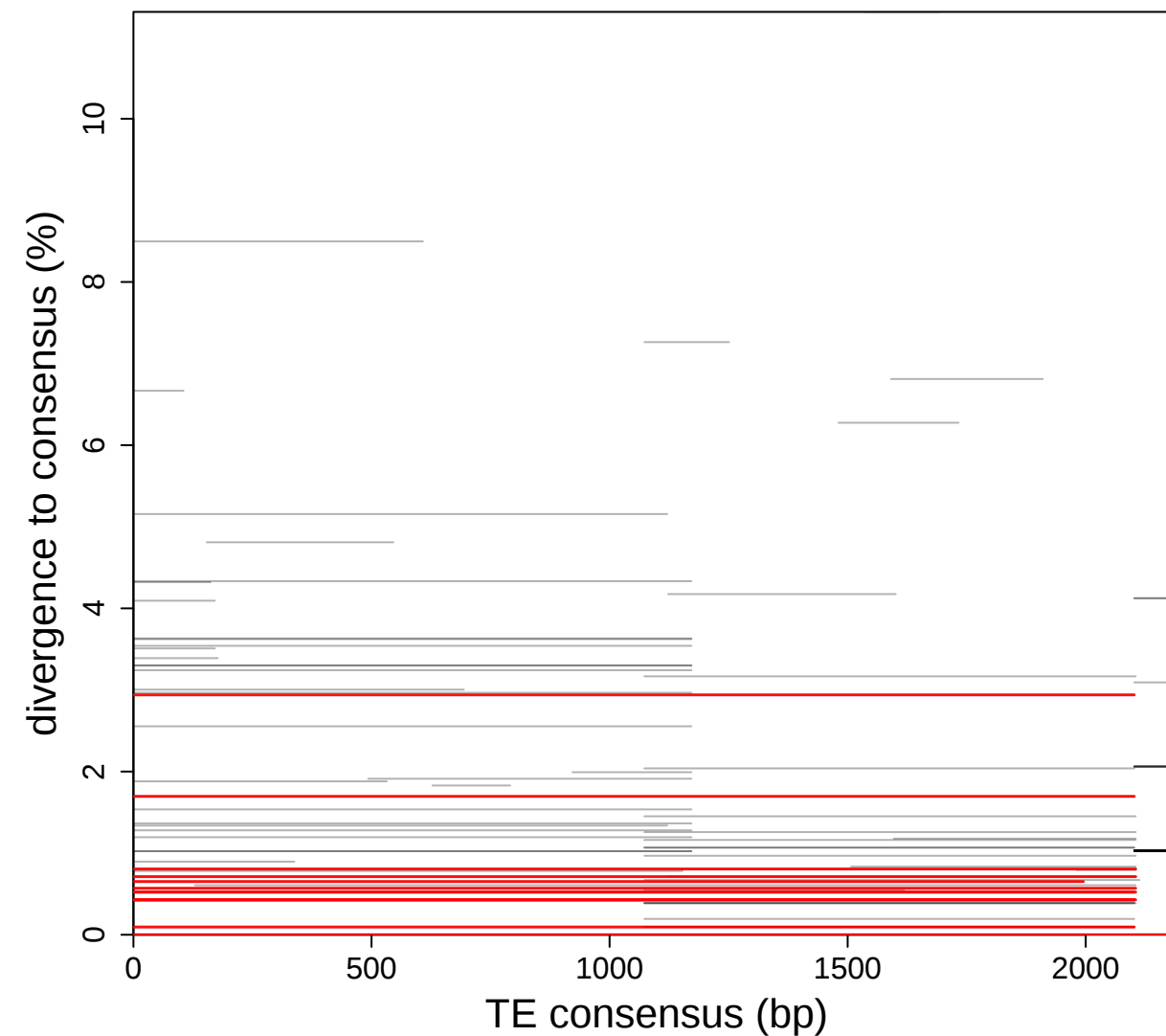
ly_623.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 7900bp; fragments: 247; full length: 10 (>=7110bp)

After TEtrimmer Extended plot Blue lines are boundaries

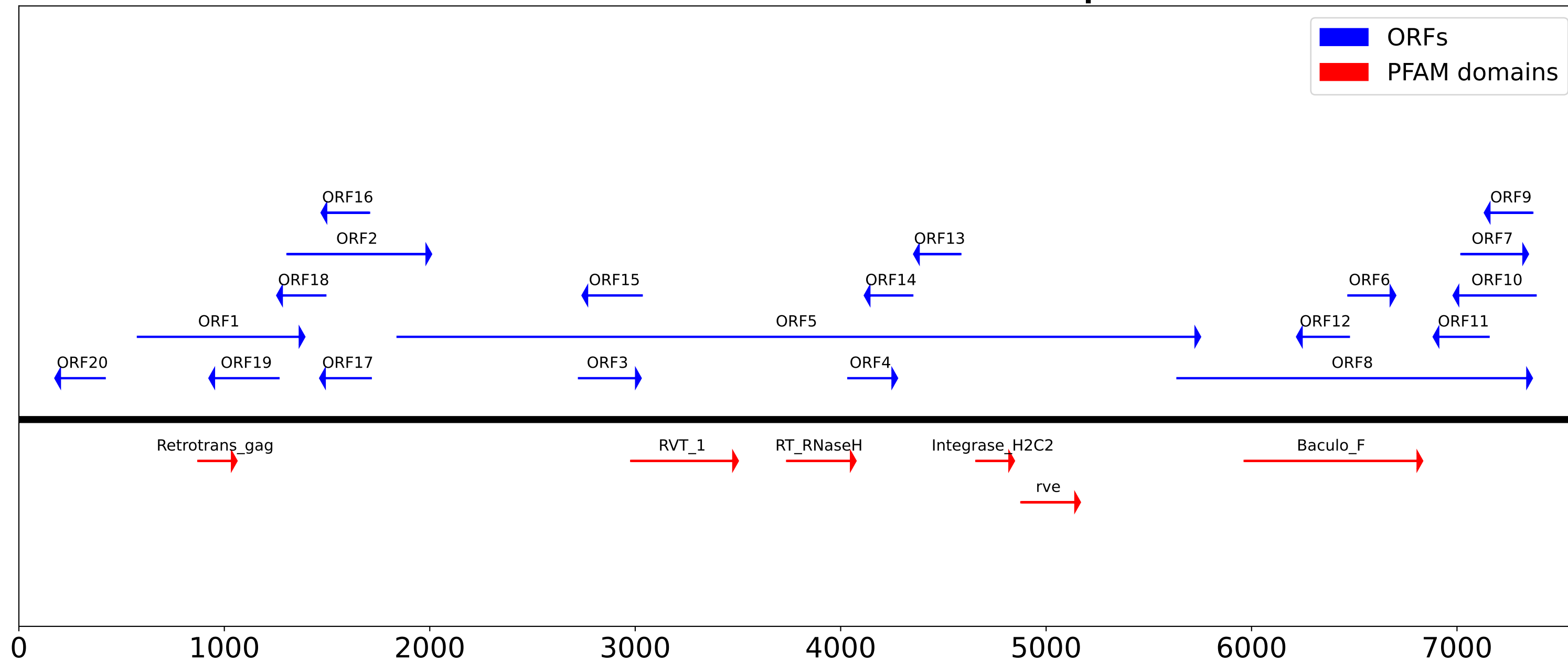


TE: ltr_1_family_623
size: 2198bp; fragments: 106; full length: 13 (≥ 1978.2 bp)

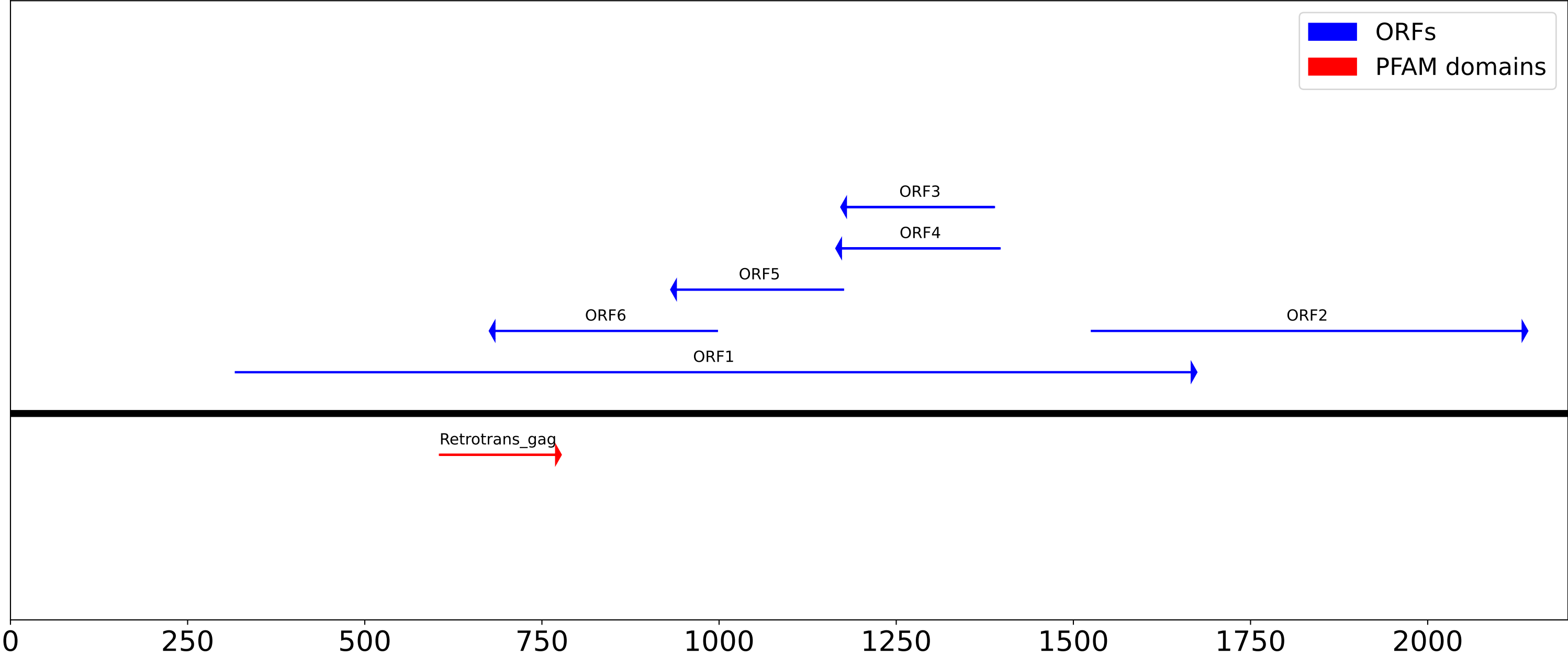
Before TEtrimmer 2198 bp



After TETrimmer ORF and PFAM domain plot



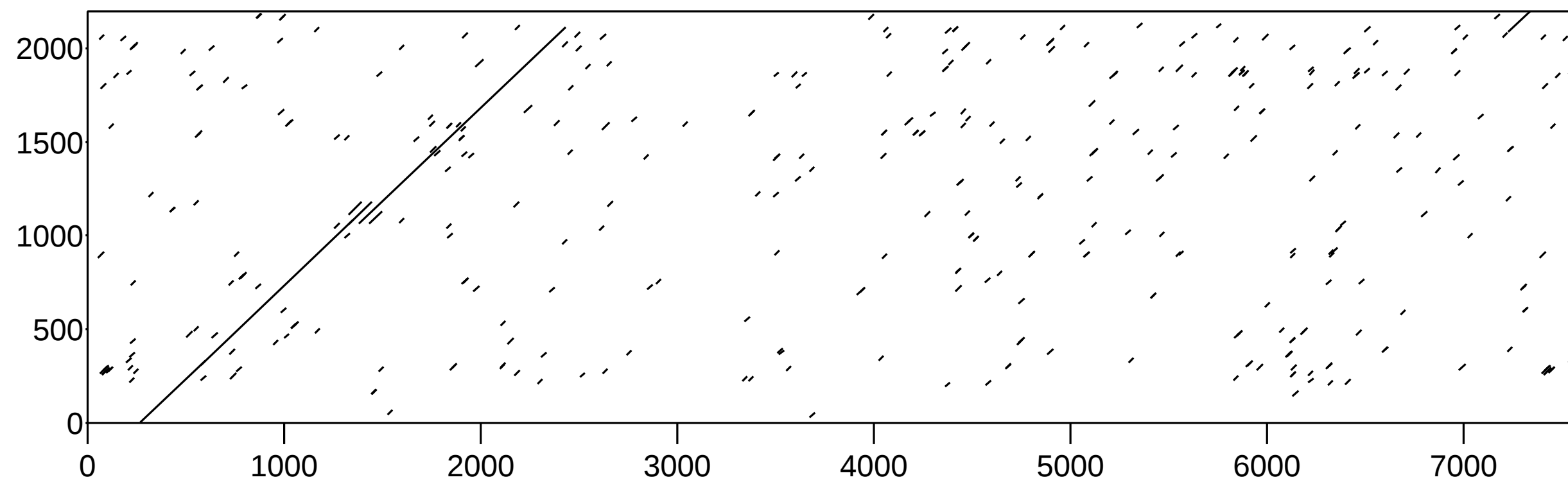
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)