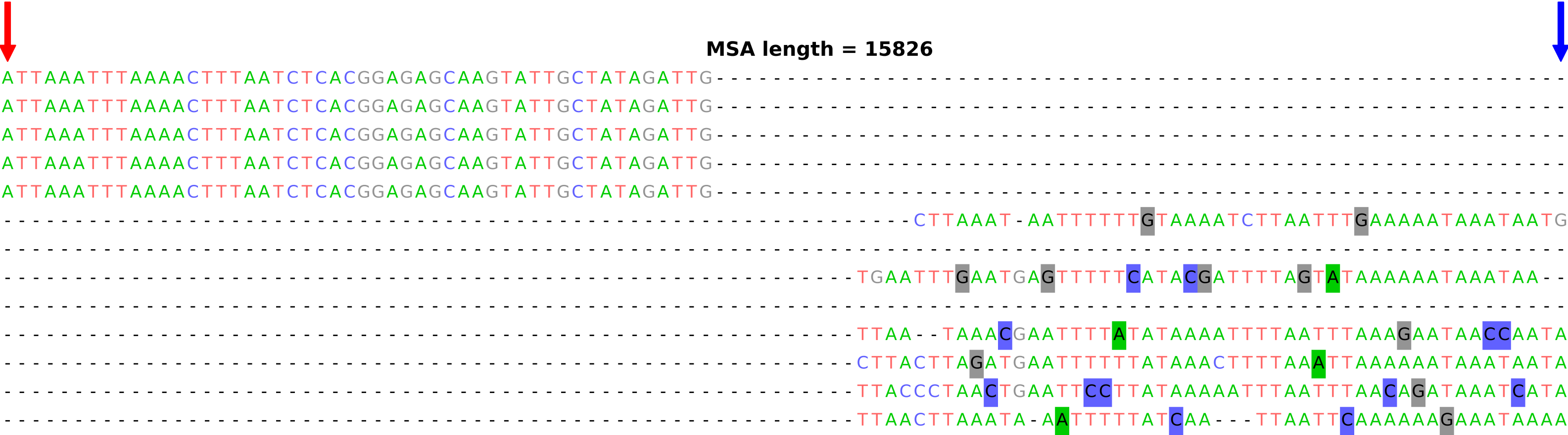
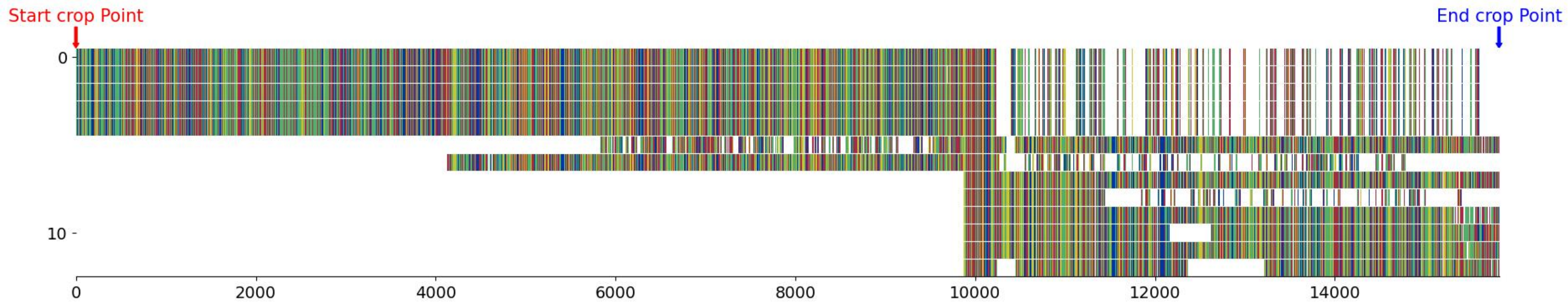


Start crop Point

End crop Point

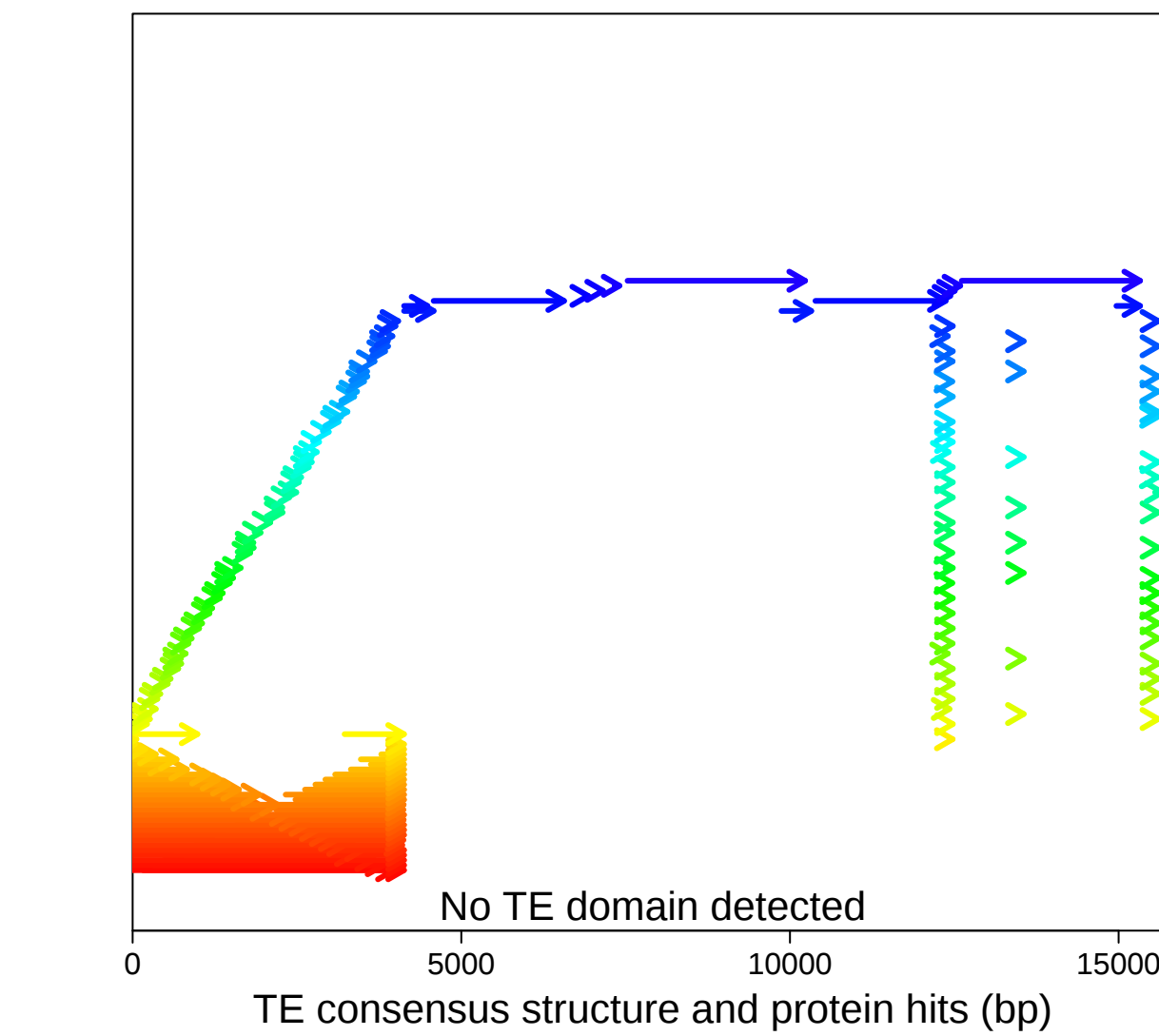
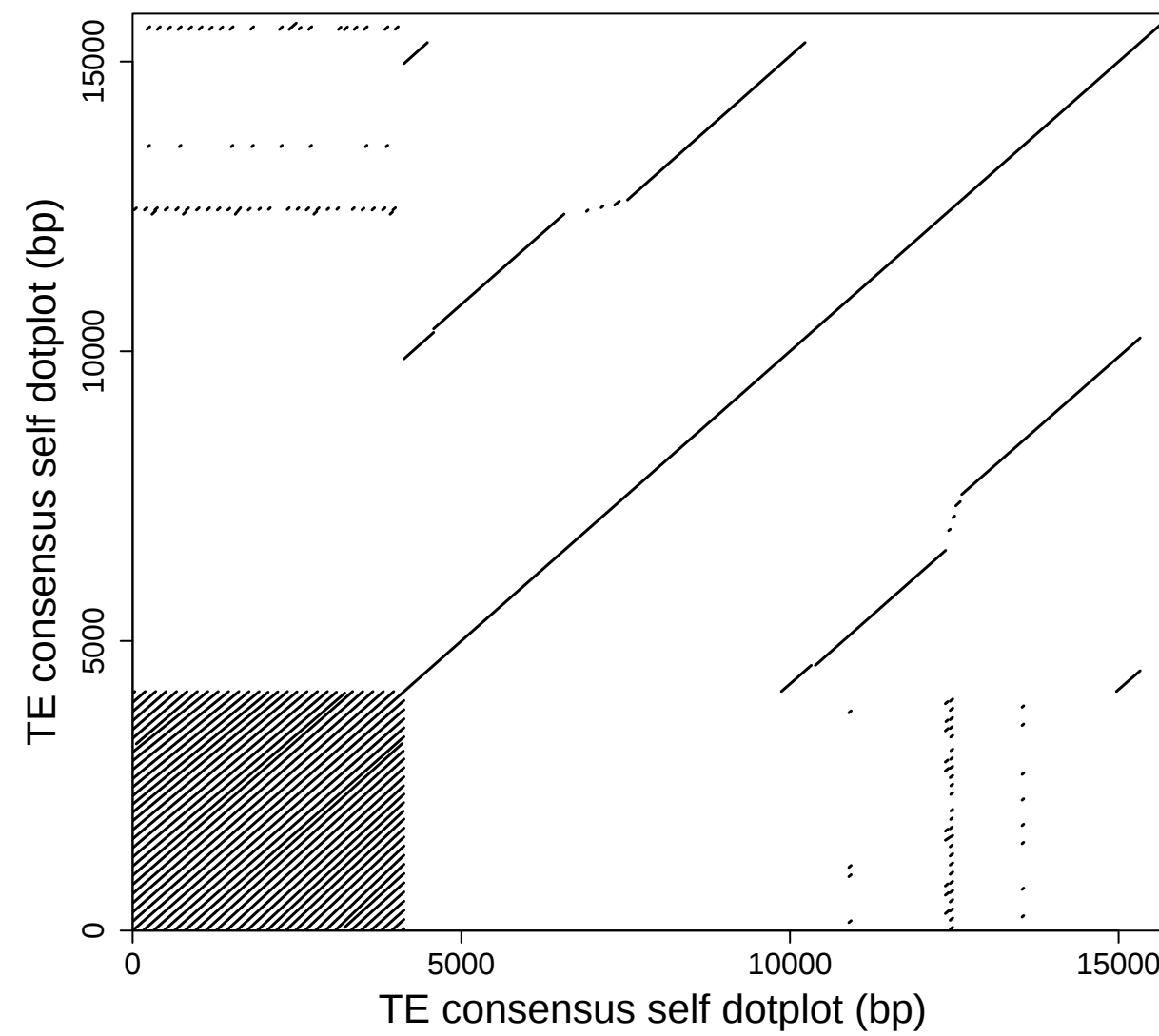
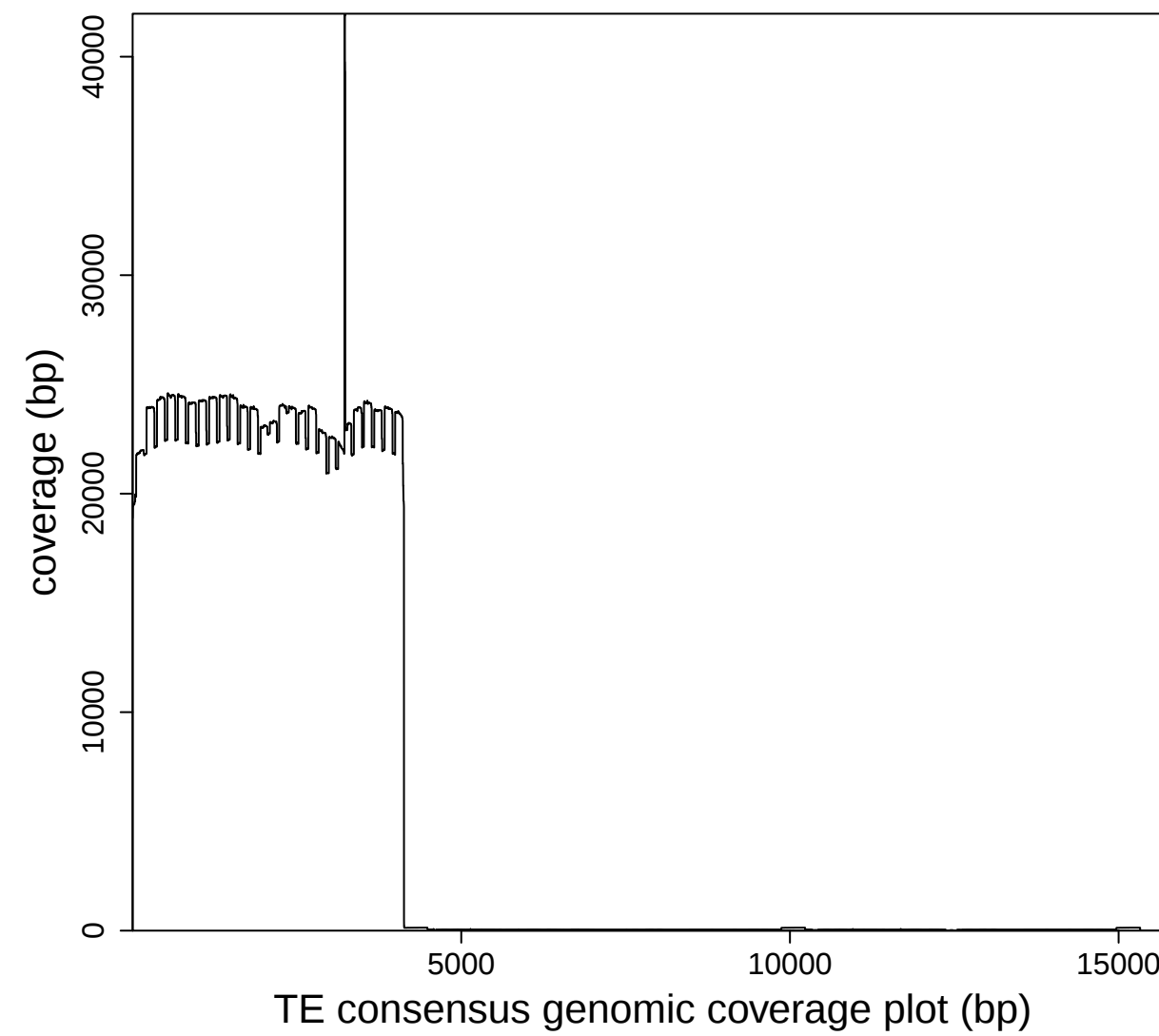
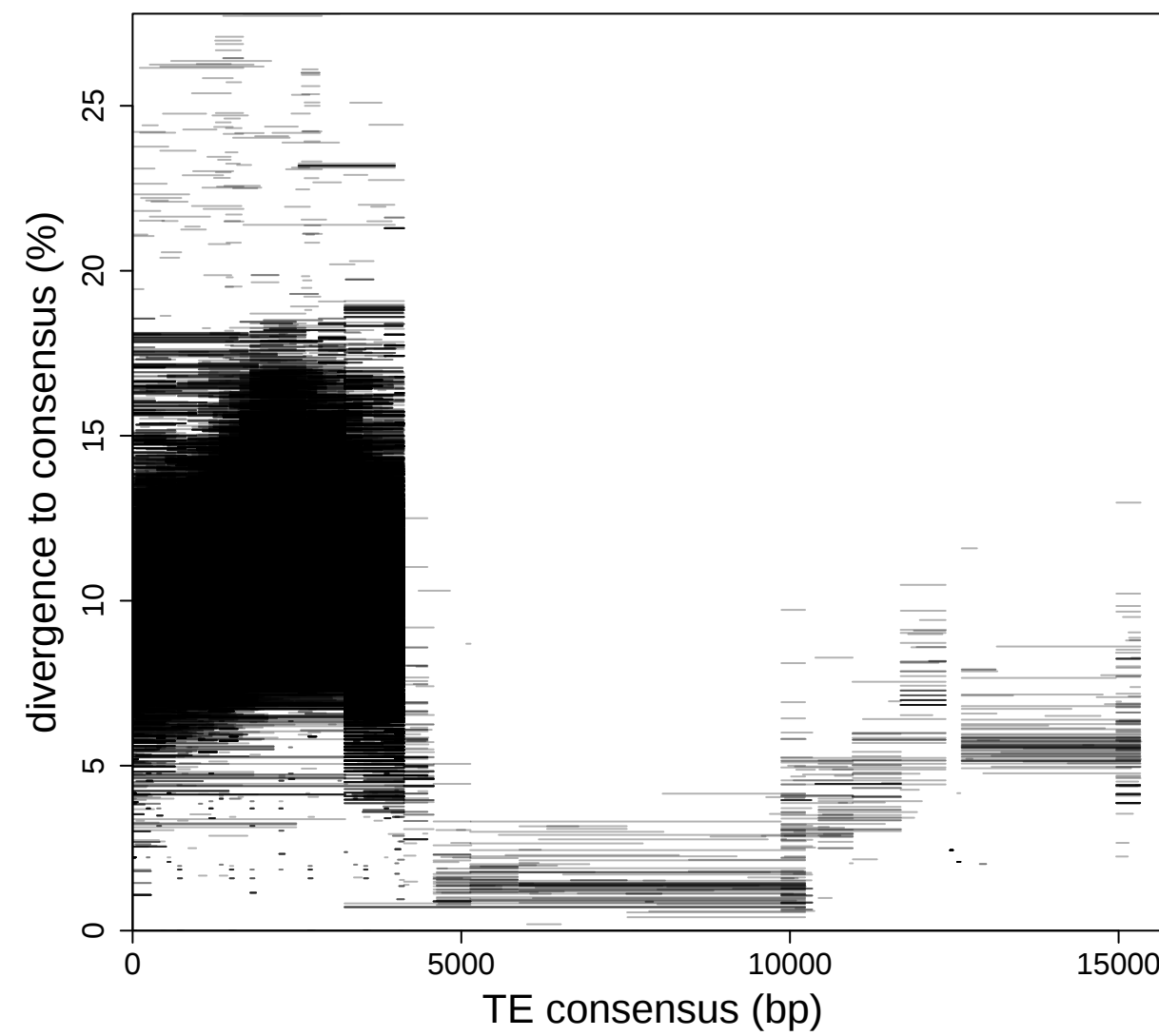
MSA length = 15826





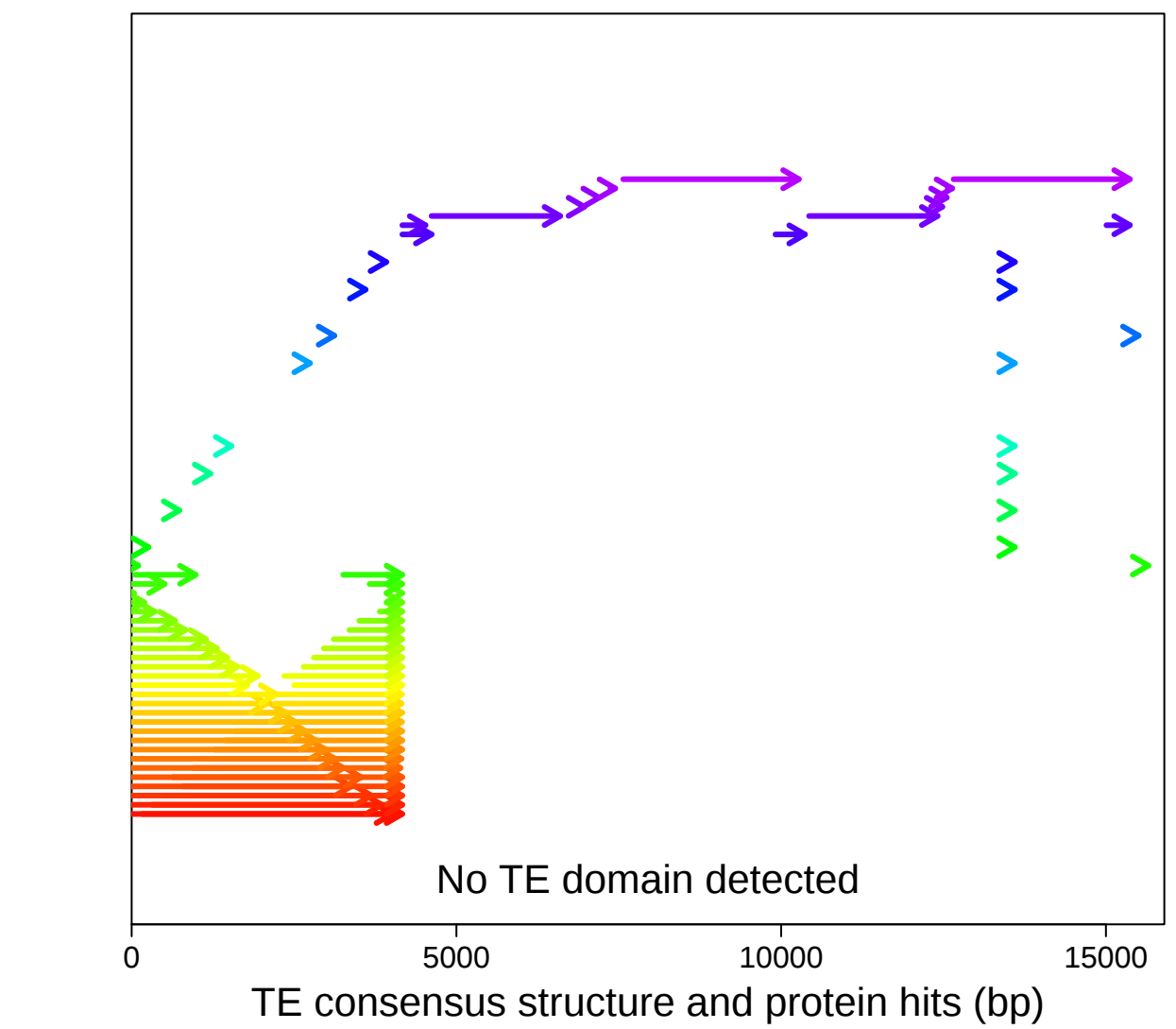
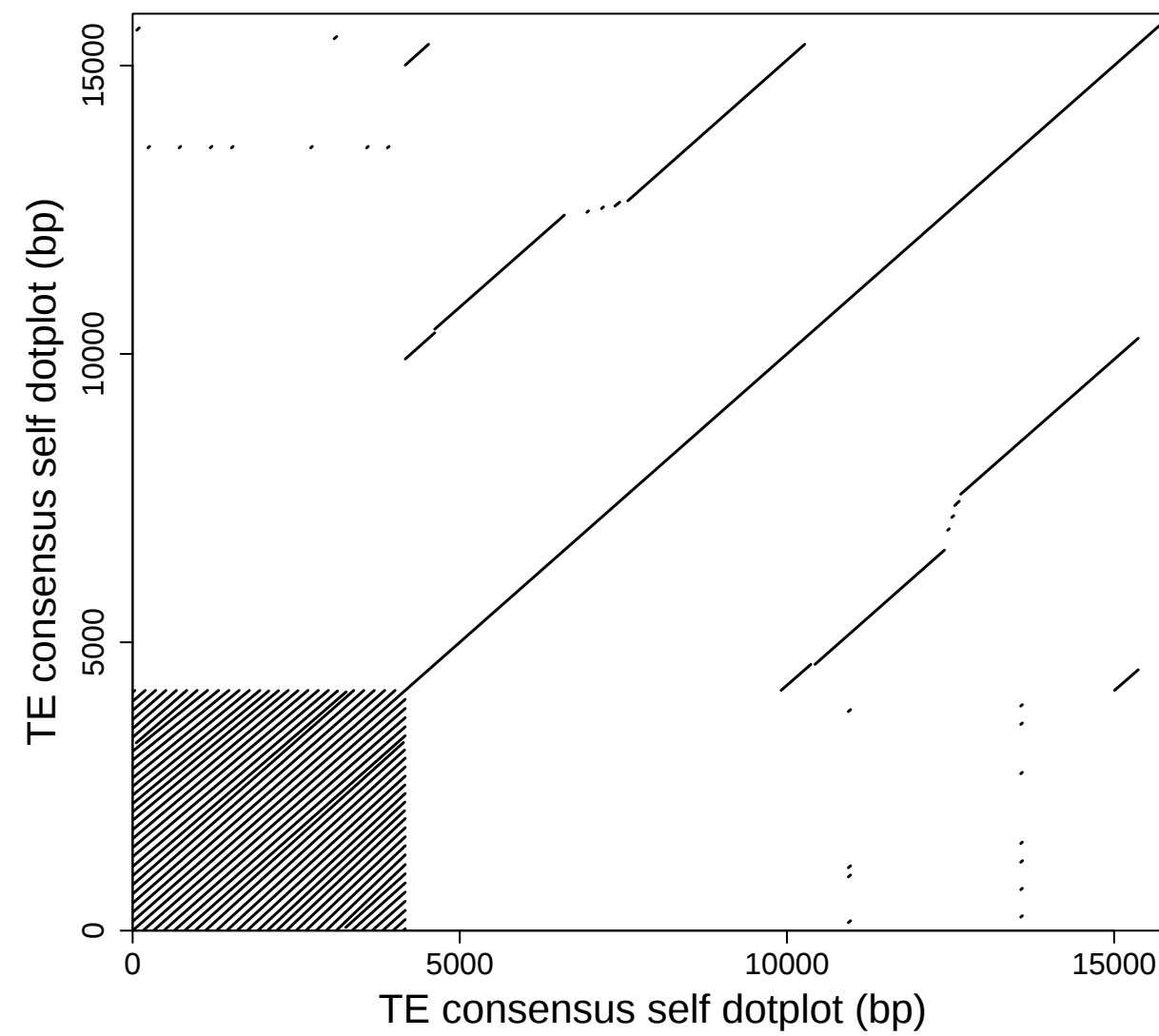
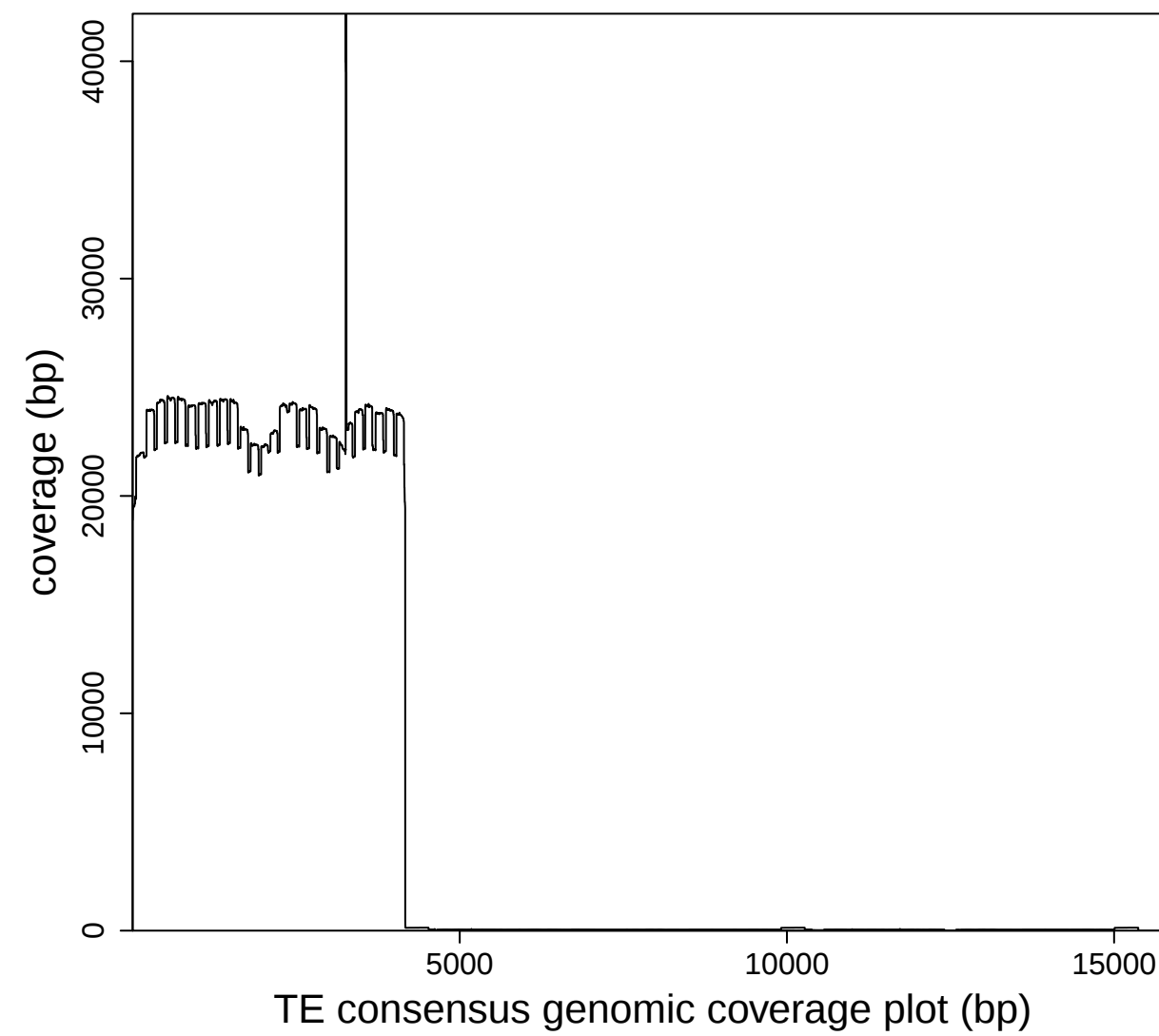
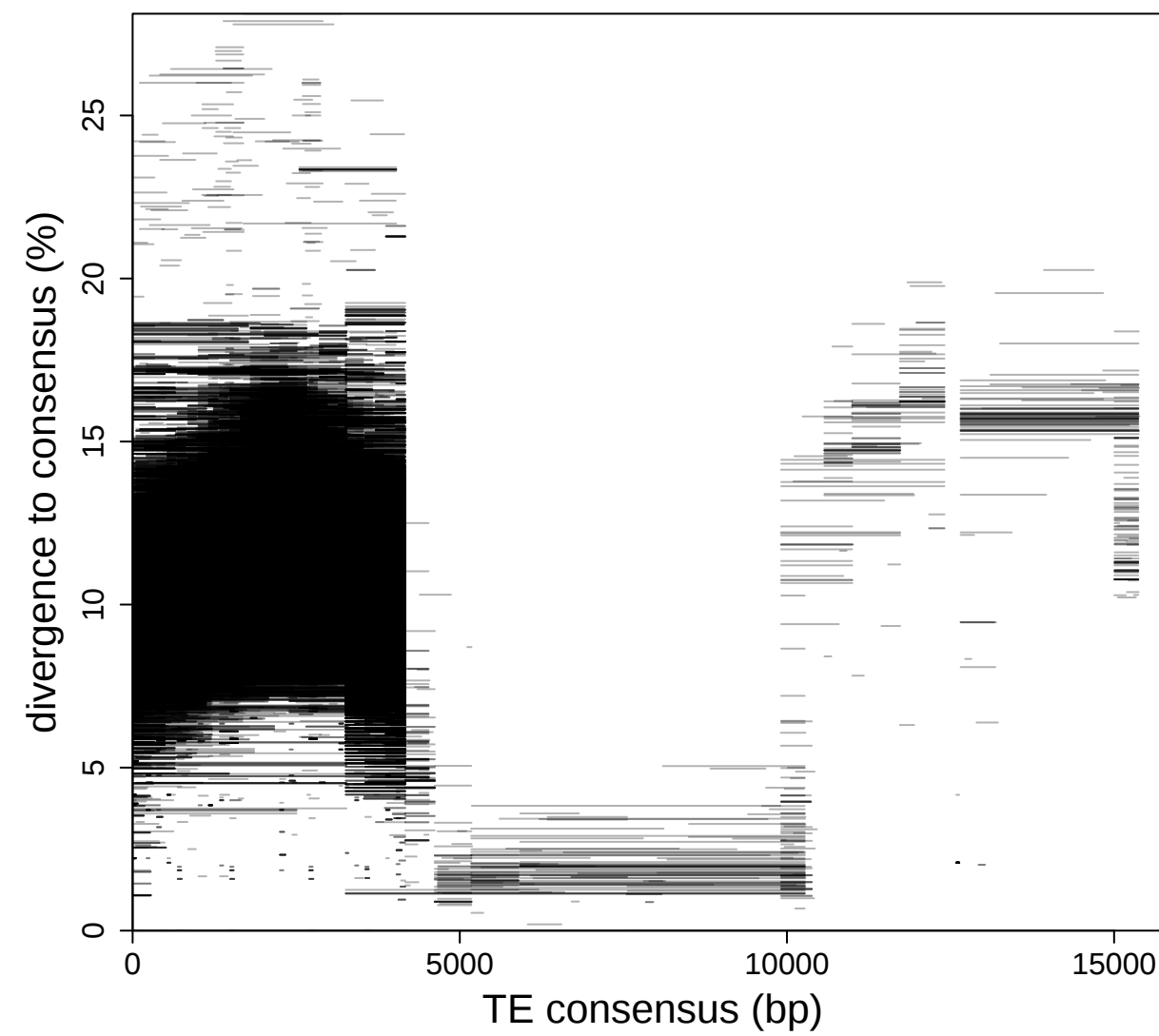
b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 15826bp; fragments: 96074; full length: 0 (>=14243.4bp)

After TEtrimmer 15826 bp



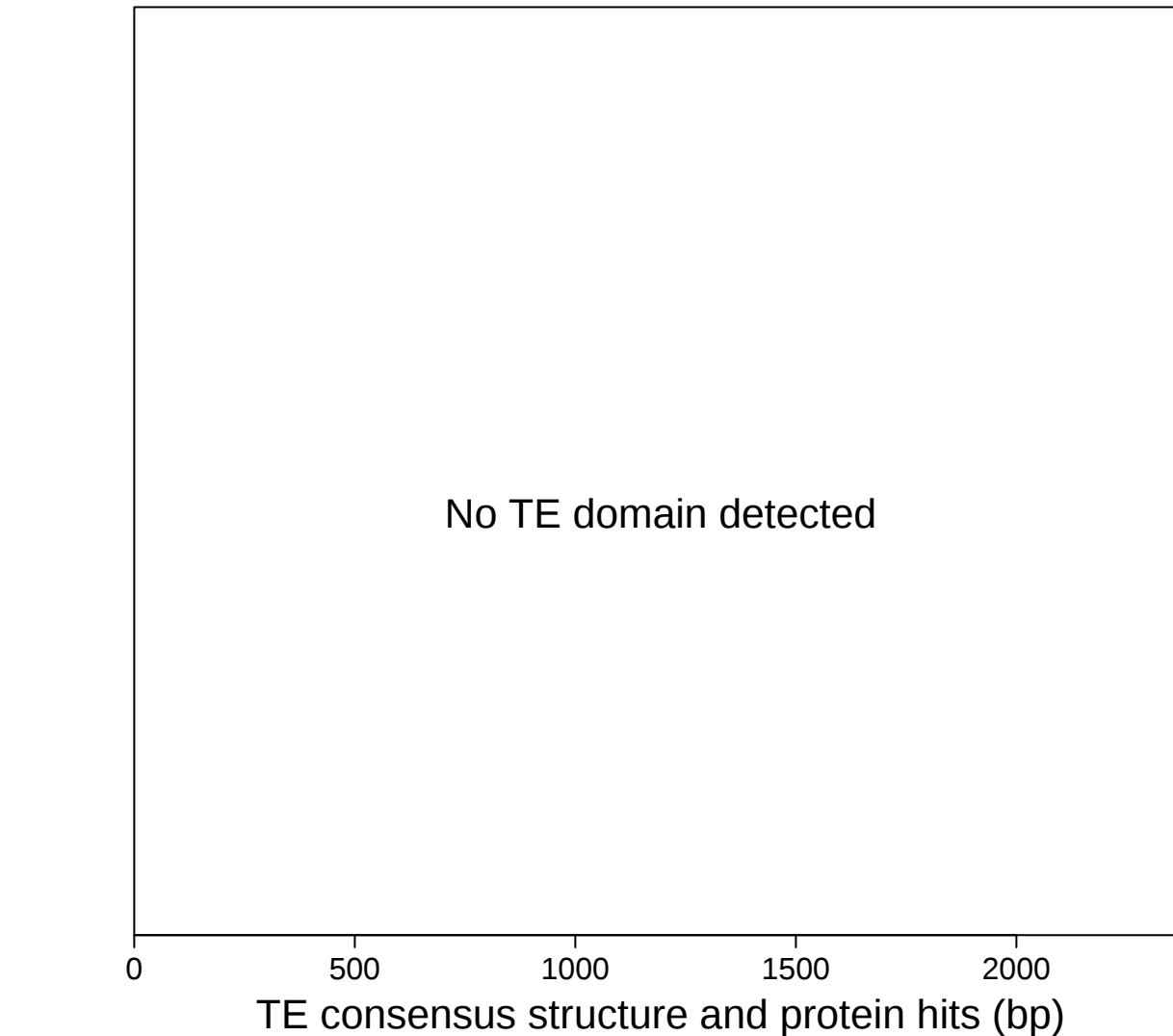
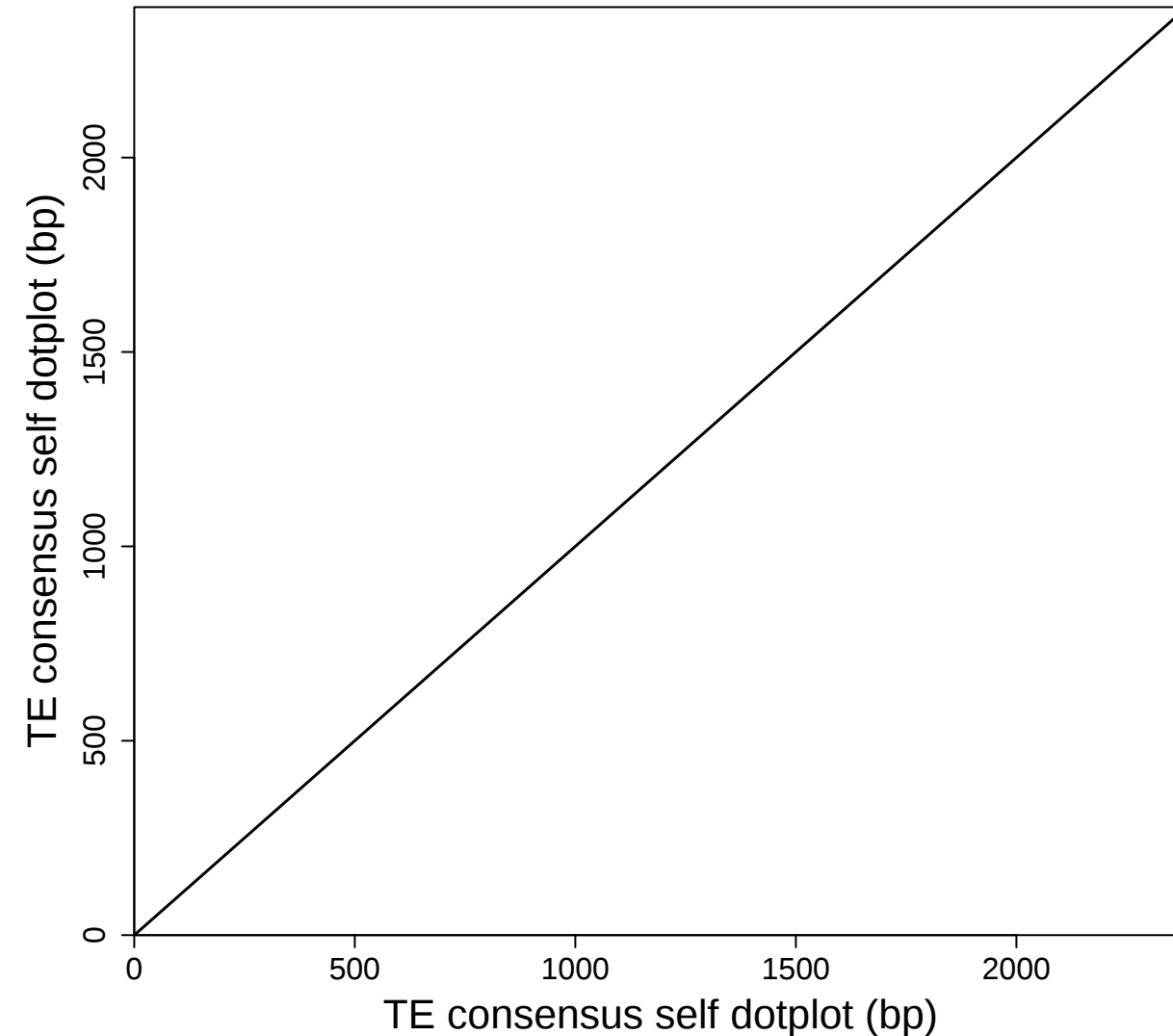
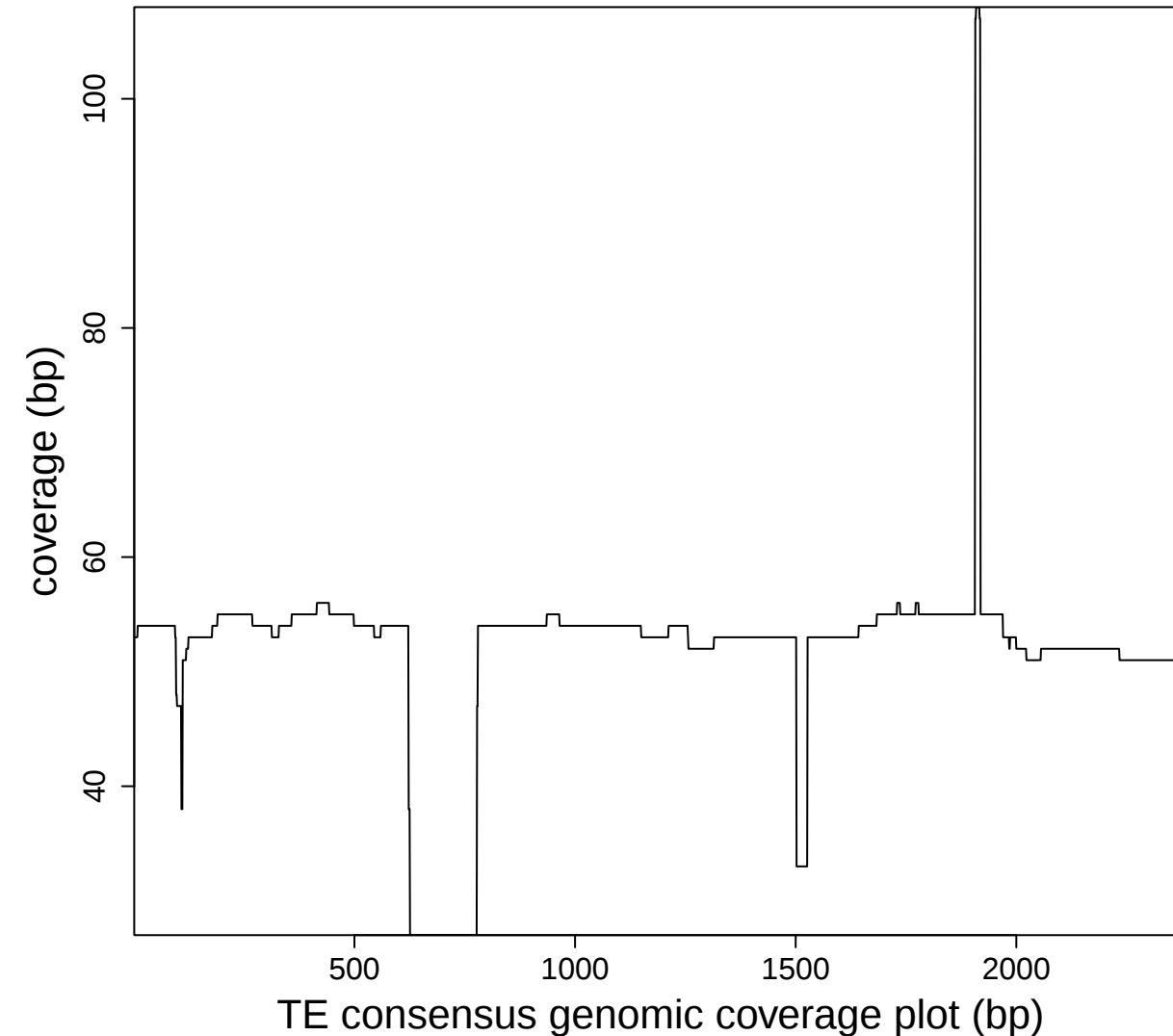
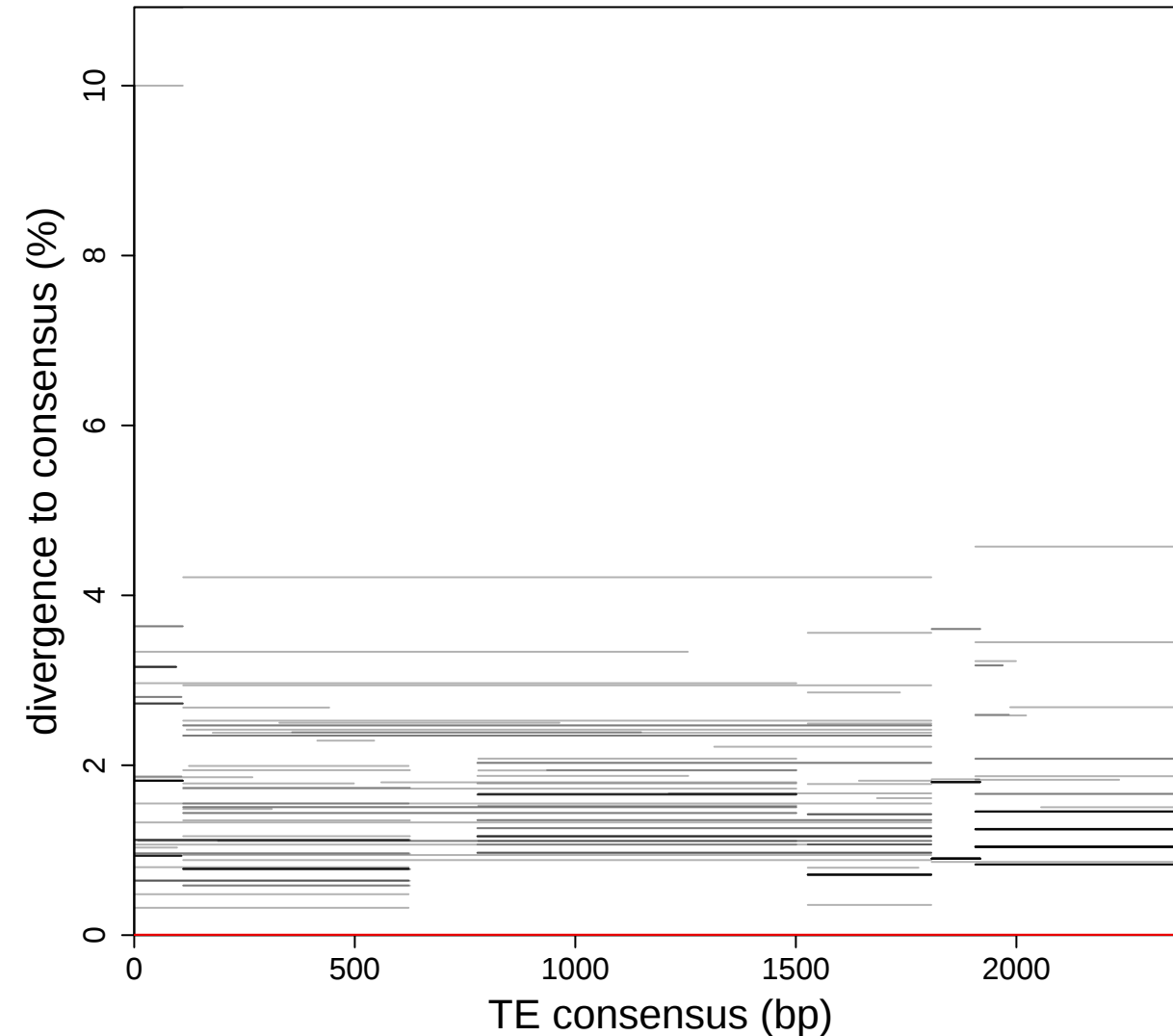
.fasta.b.bed_uf.bed_g_3.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 15899bp; fragments: 95981; full length: 0 (>=14309.1bp)

After TEtrimmer Extended plot Blue lines are boundaries

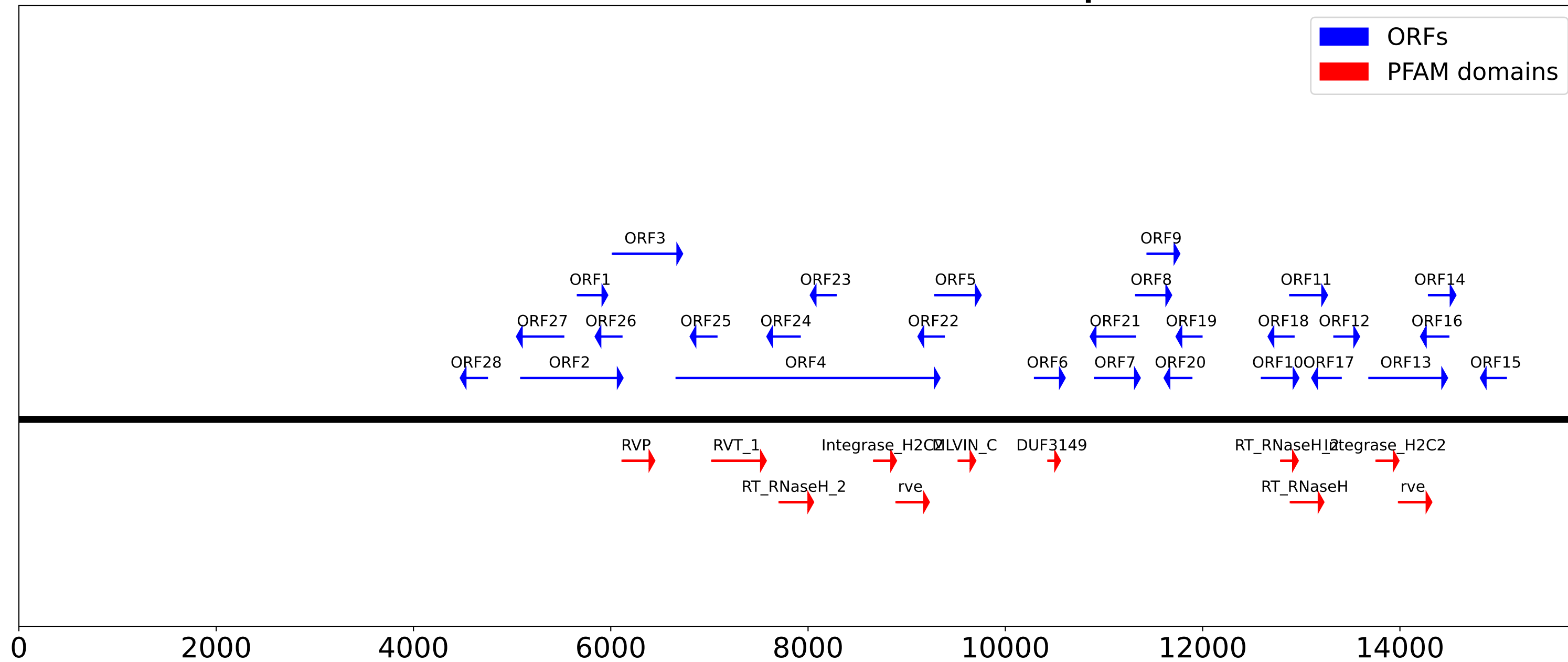


TE: ltr_1_family_426
size: 2387bp; fragments: 256; full length: 1 (≥ 2148.3 bp)

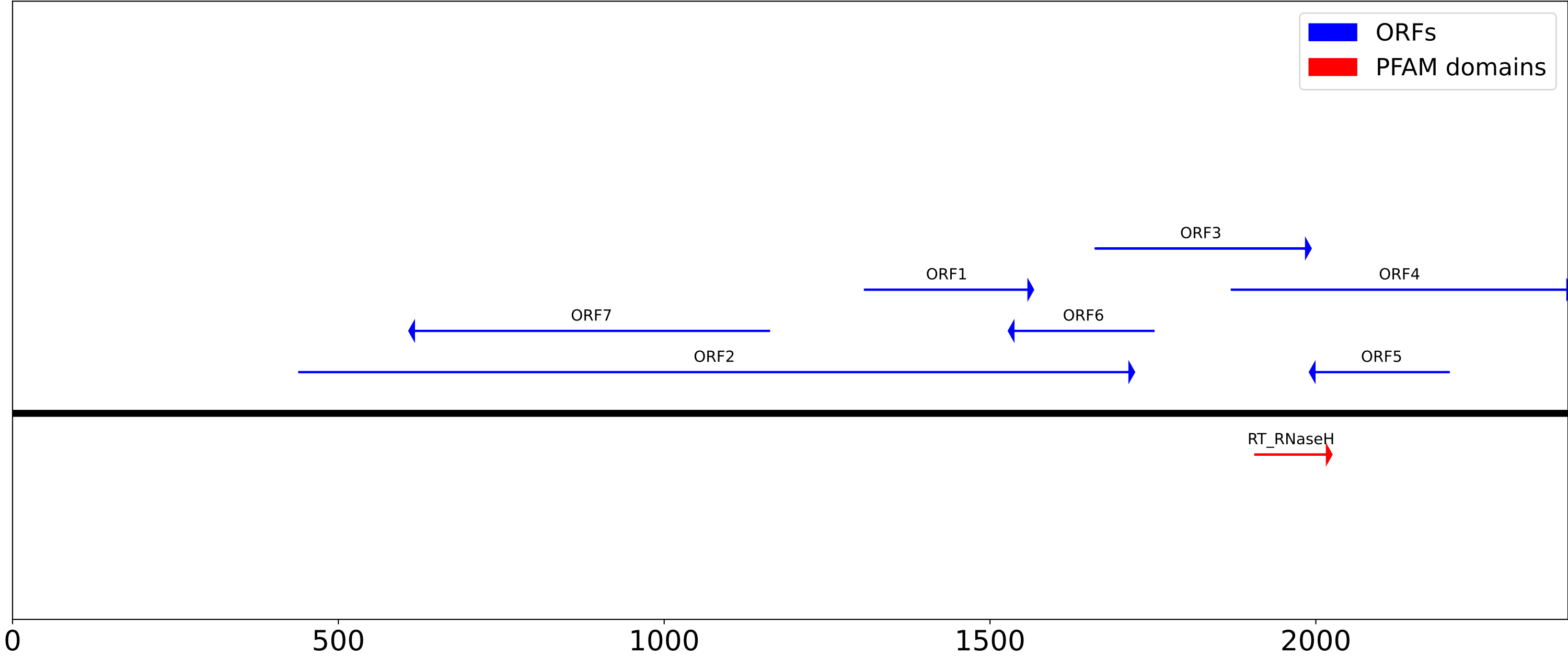
Before TEtrimmer 2387 bp



After TETrimmer ORF and PFAM domain plot



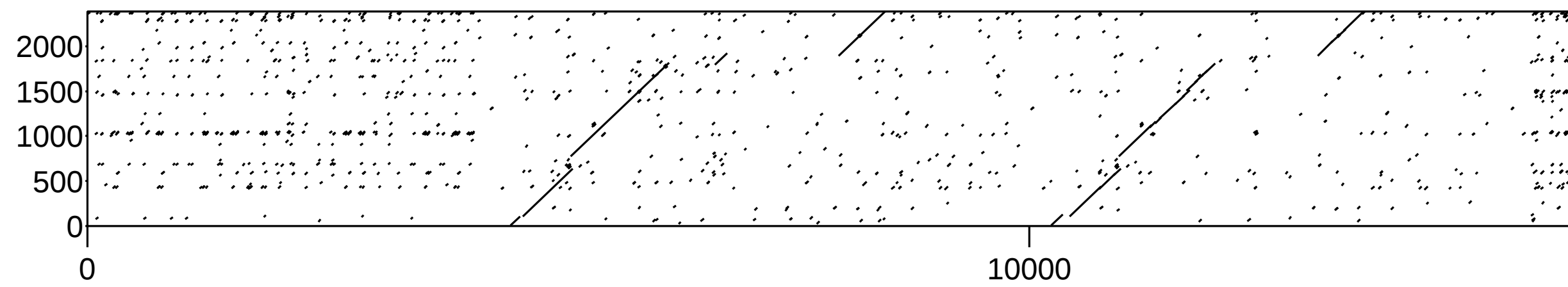
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

Isensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)