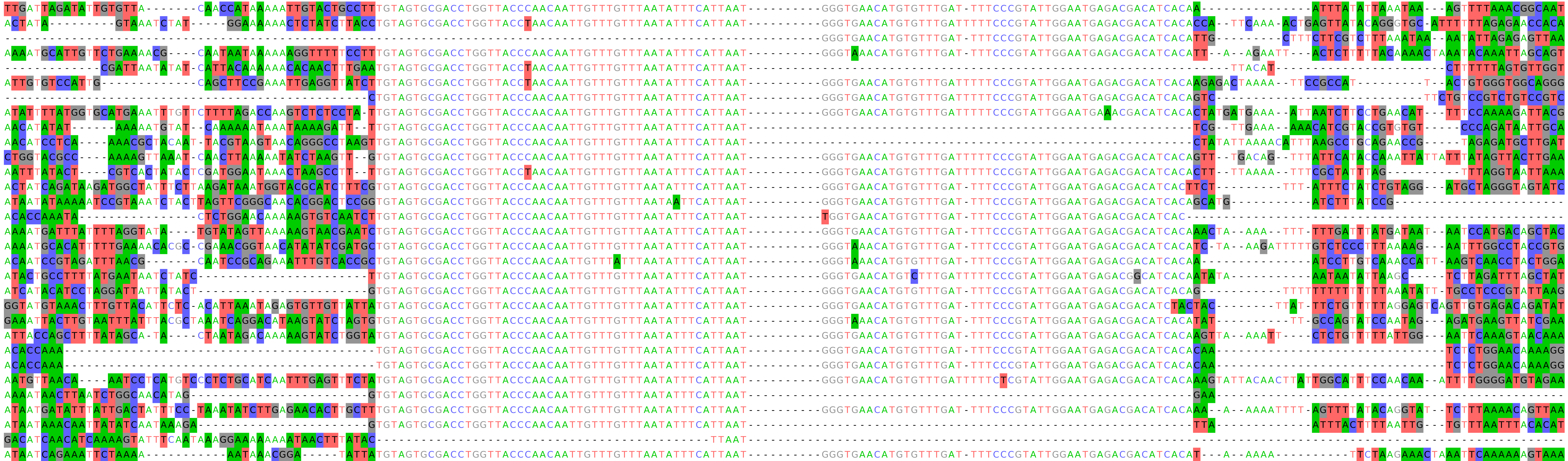


MSA length = 6545

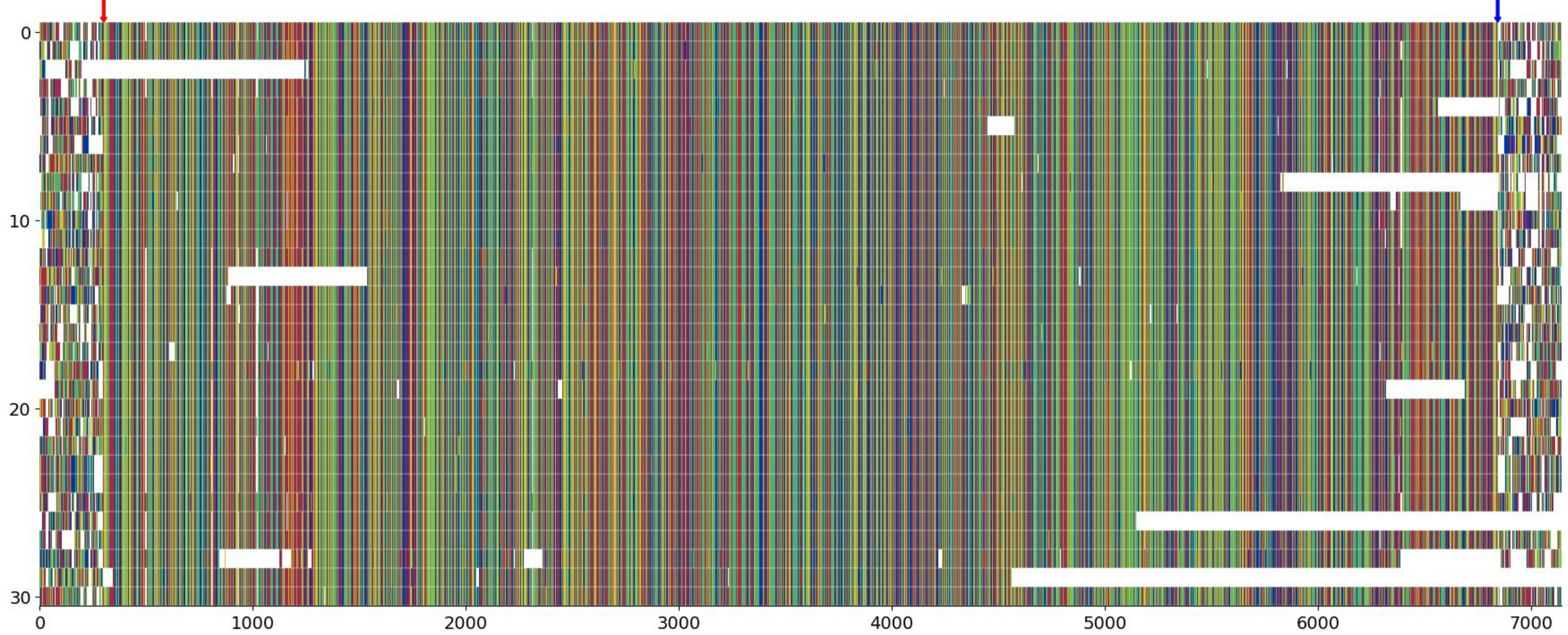
Start crop Point

End crop Point



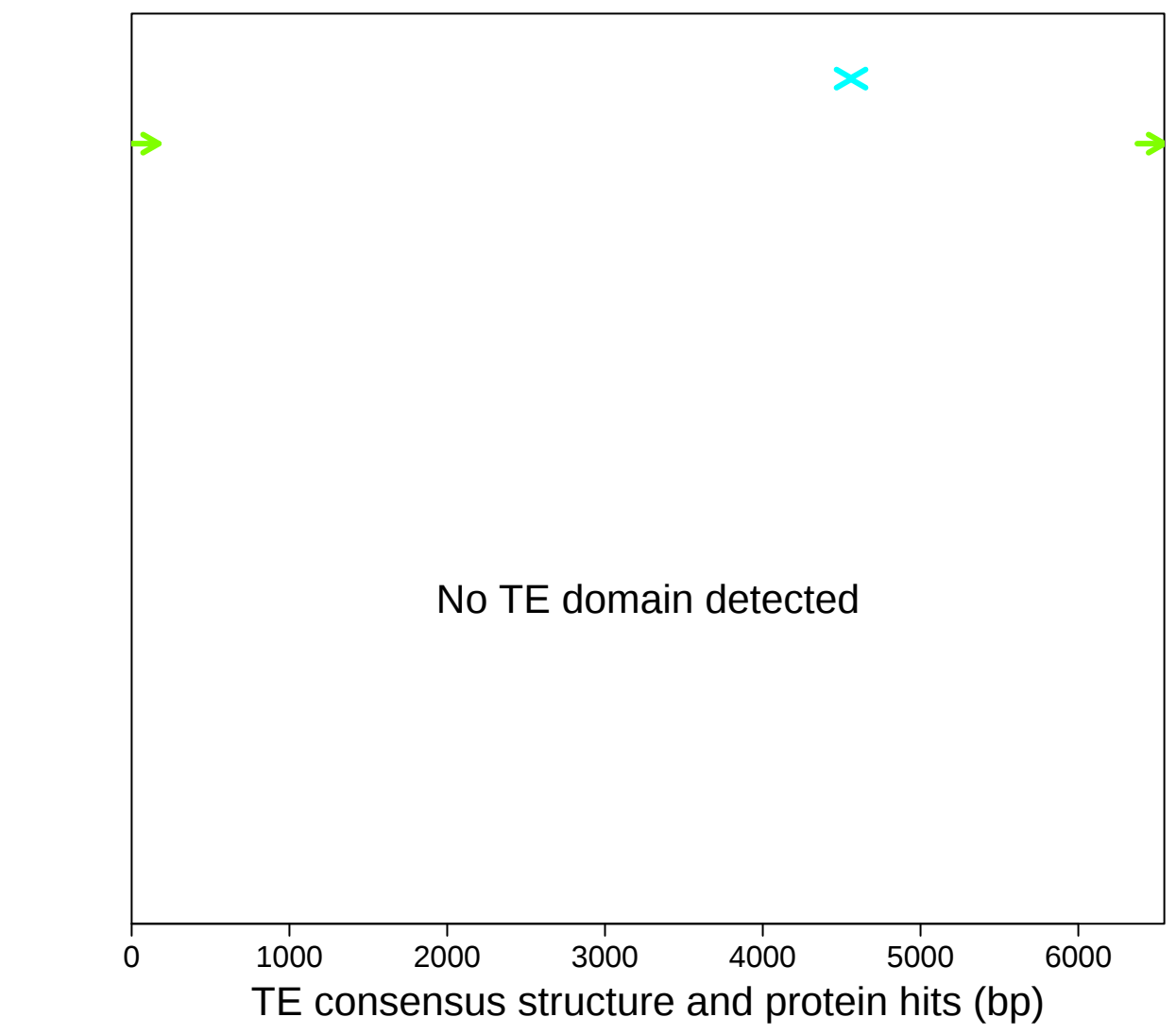
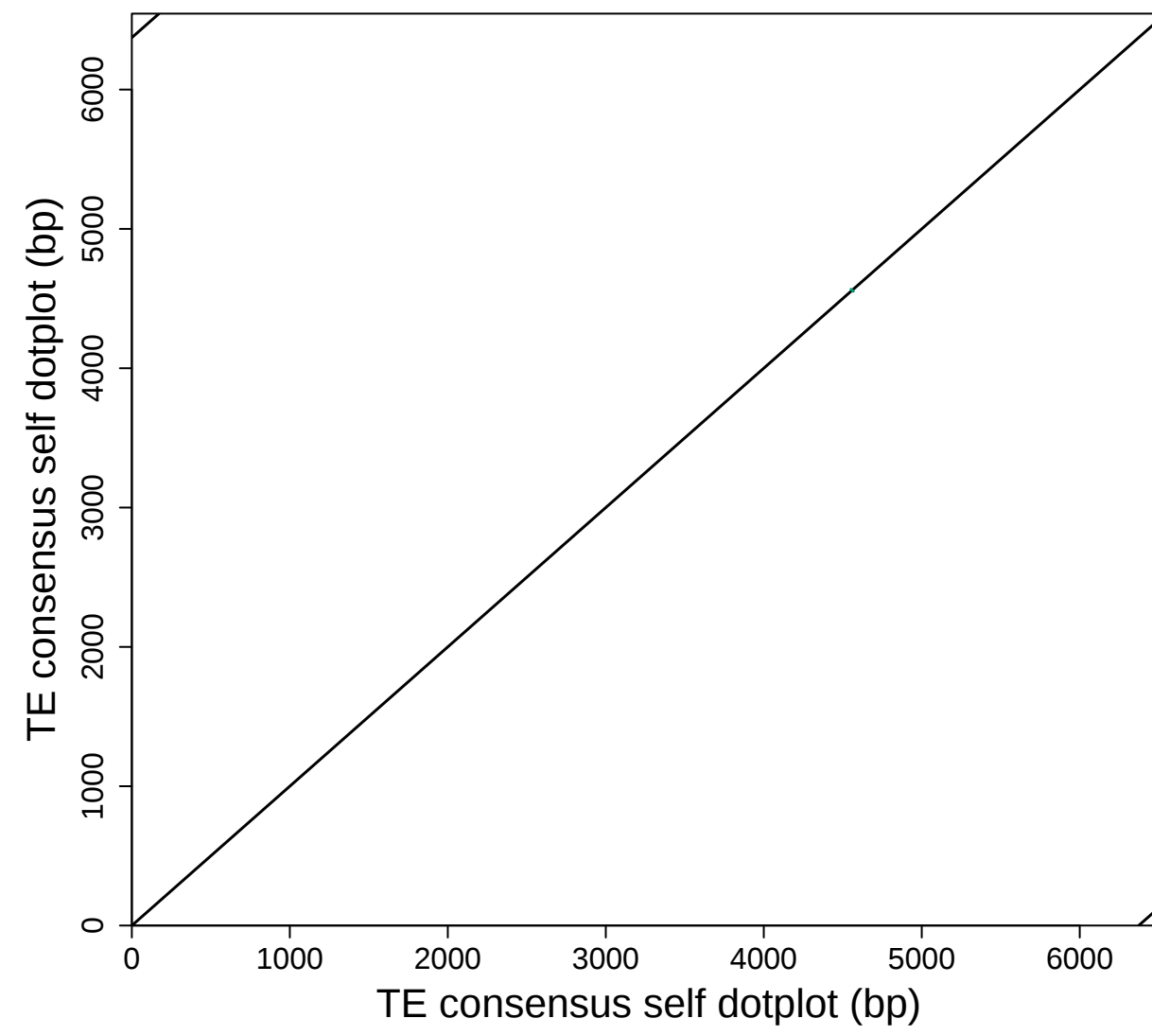
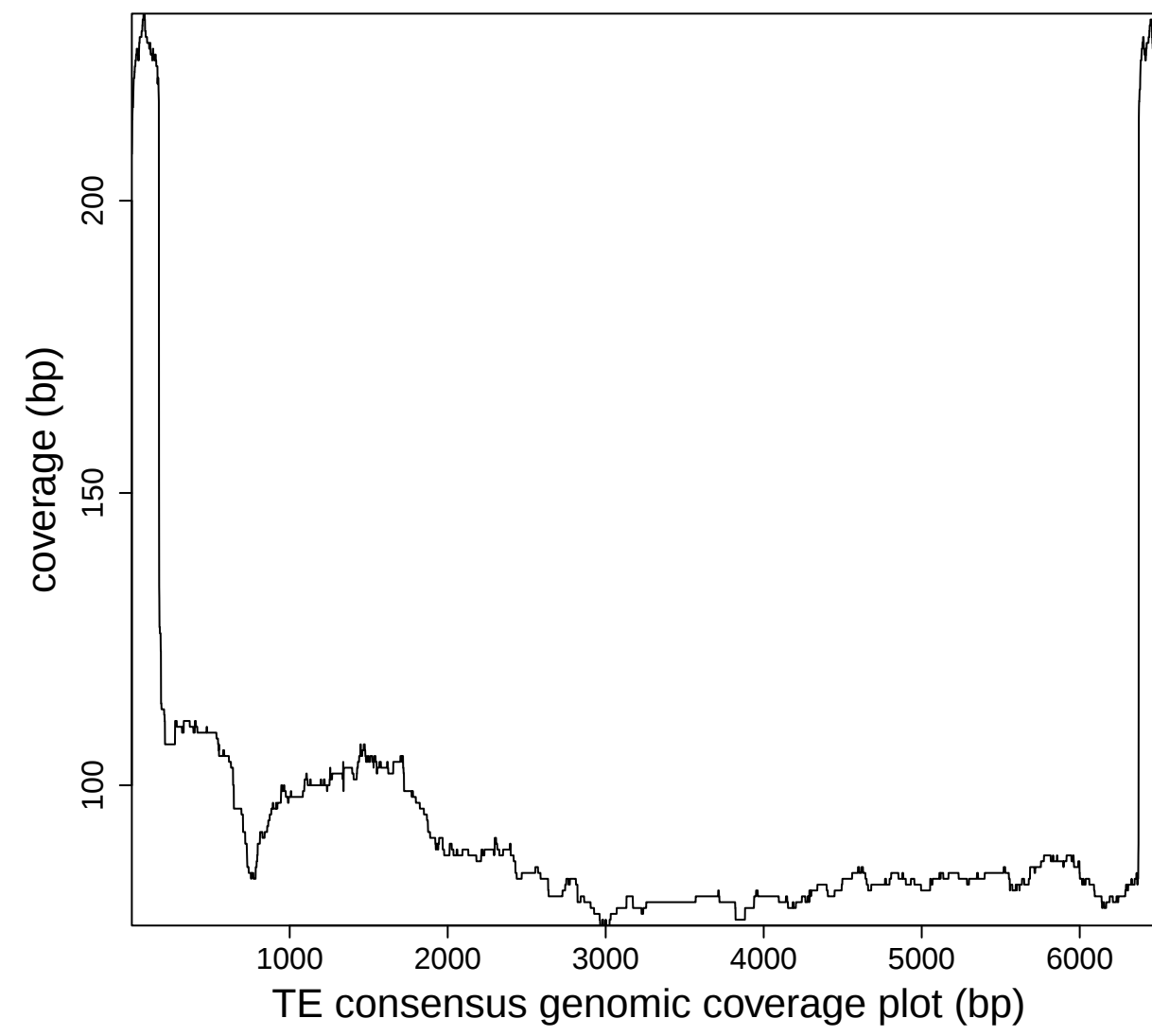
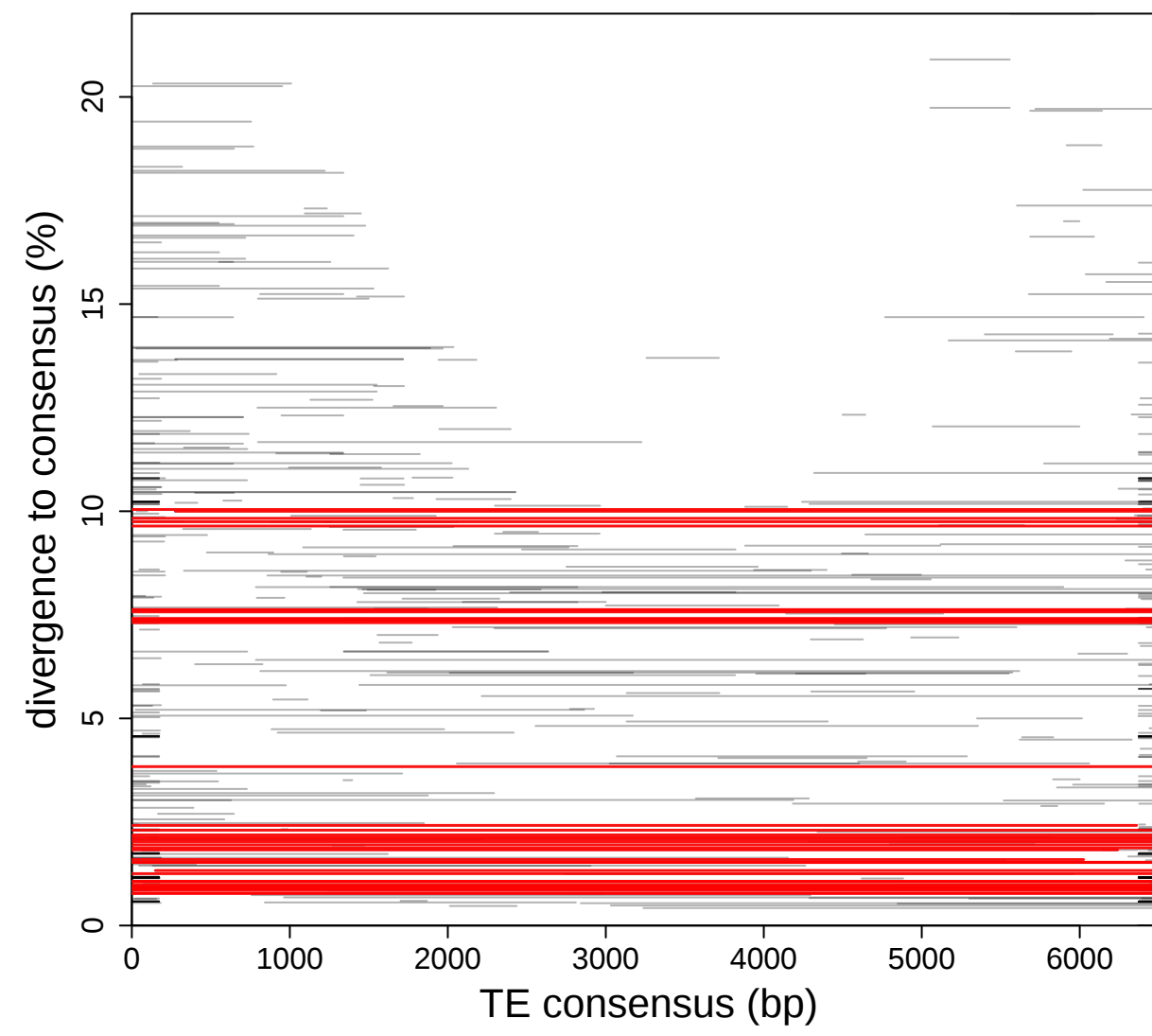
Start crop Point

End crop Point



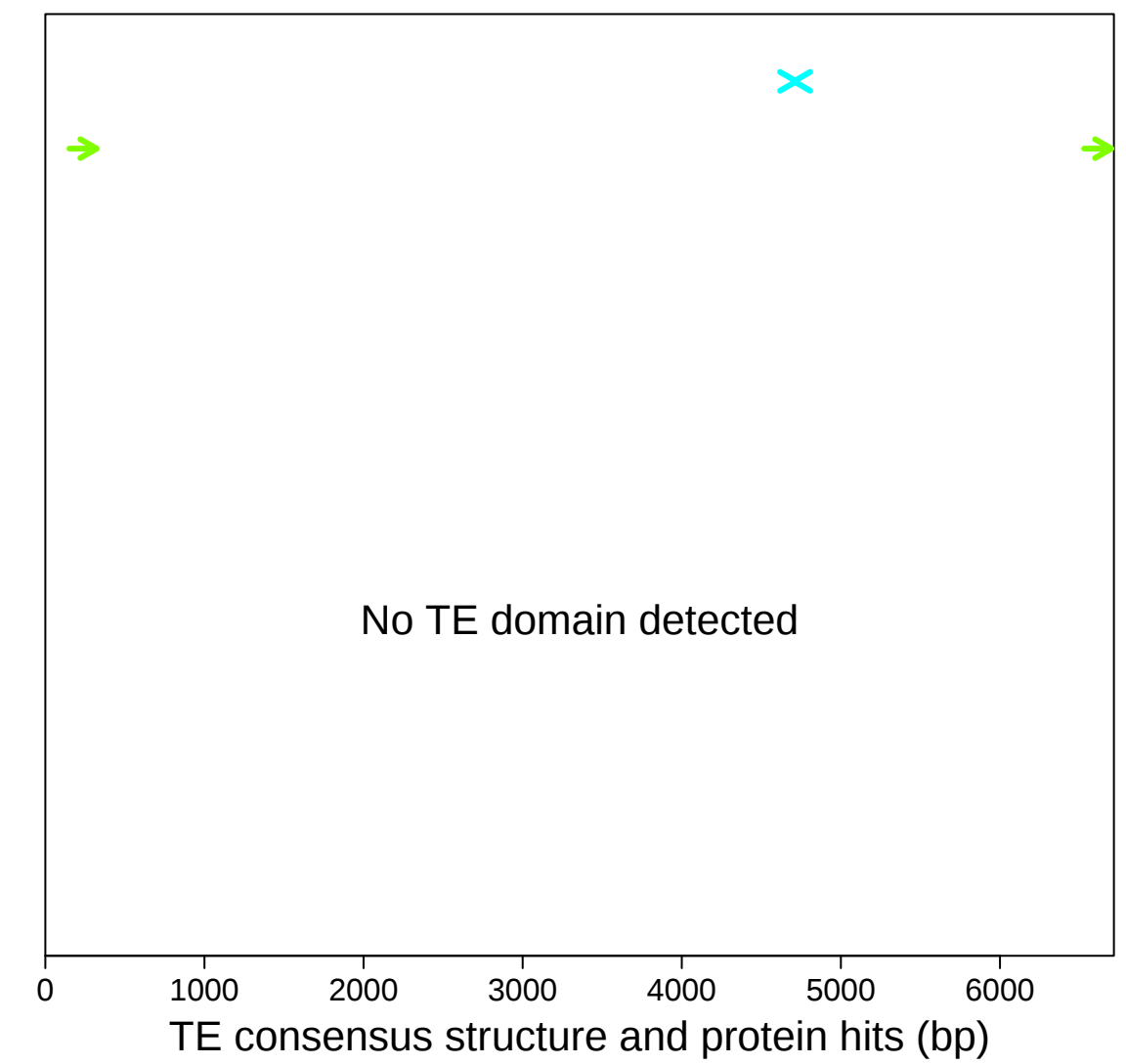
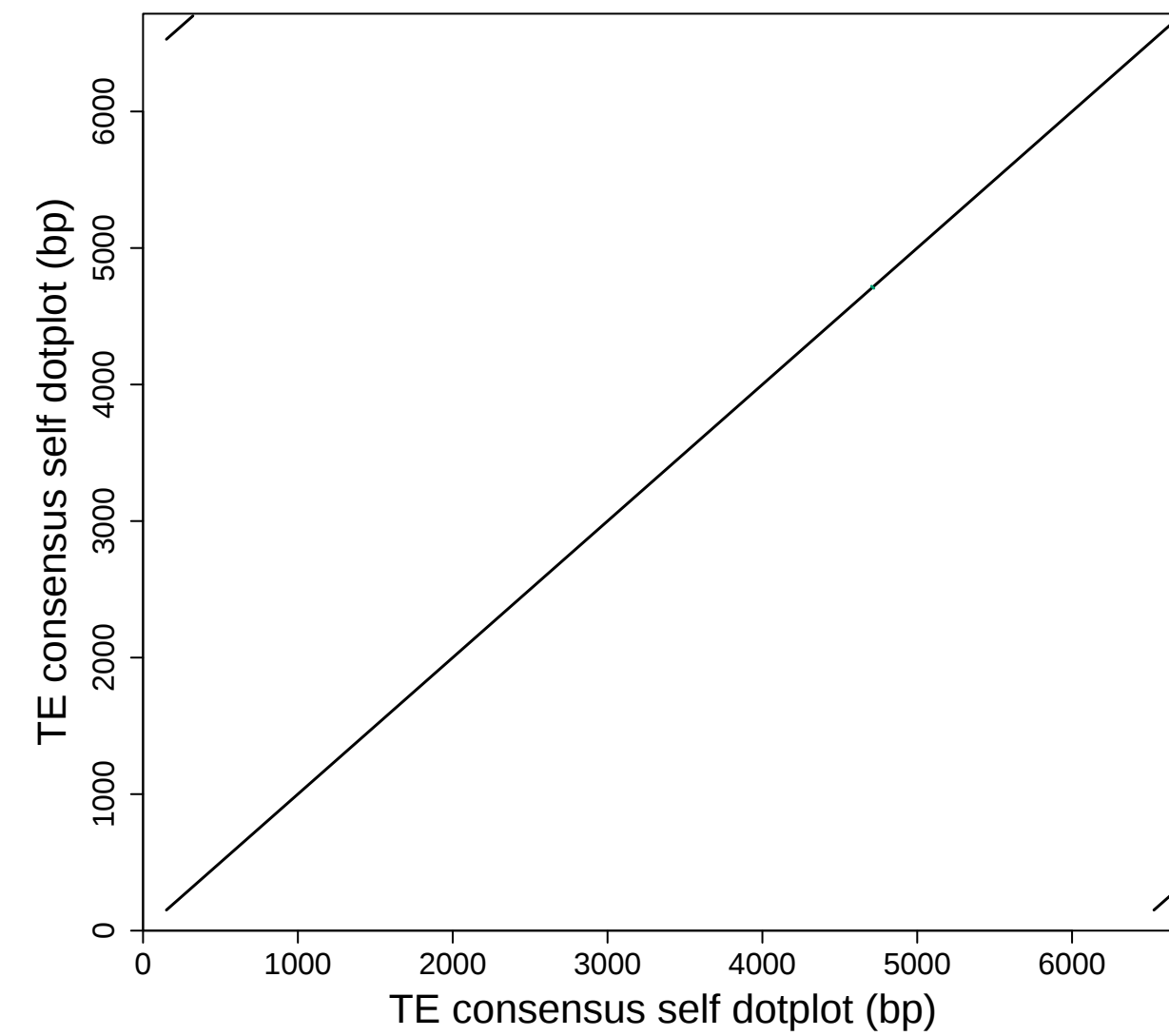
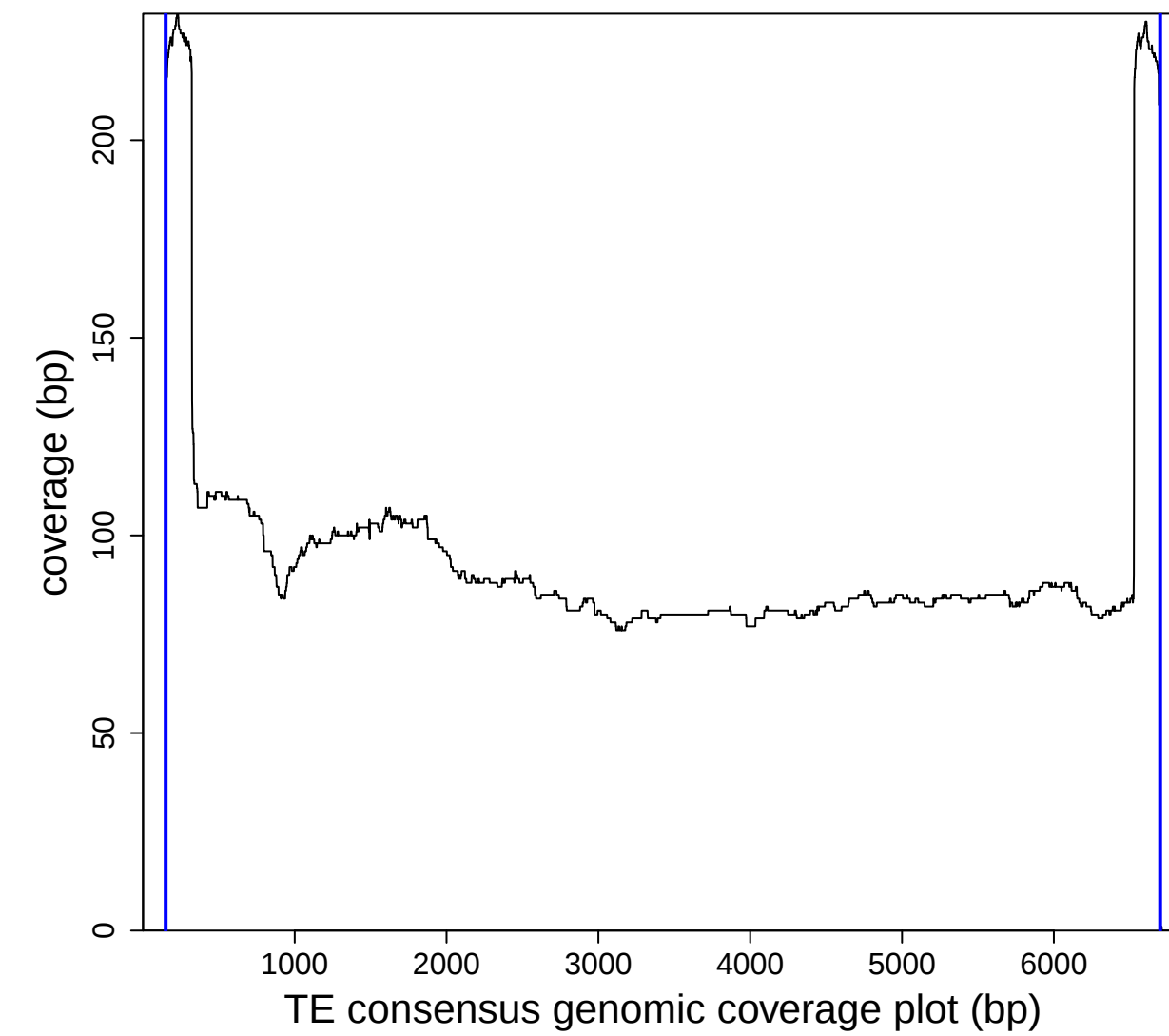
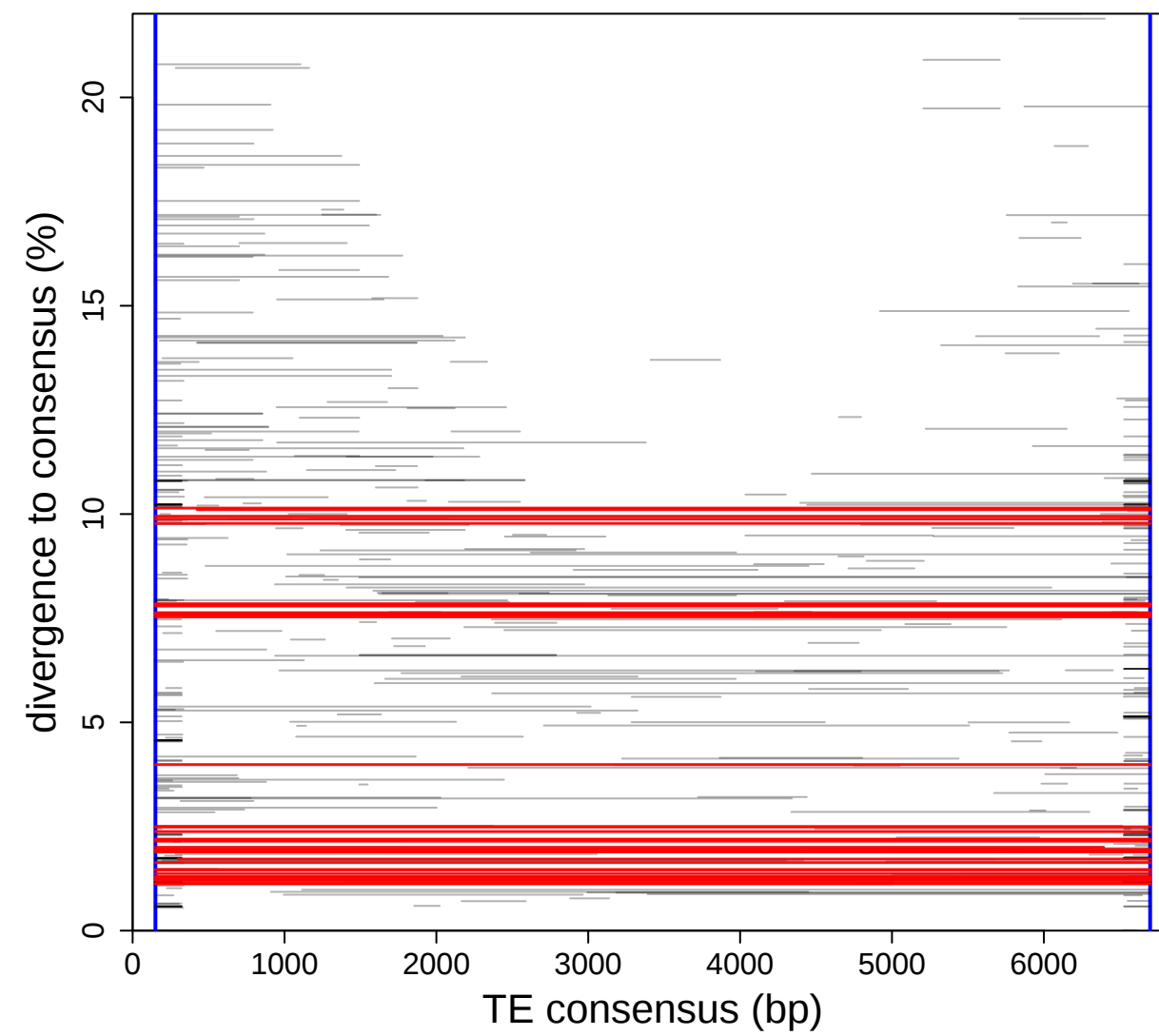
ed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_ce.
size: 6545bp; fragments: 572; full length: 39 (≥ 5890.5 bp)

After TETrimmer 6545 bp



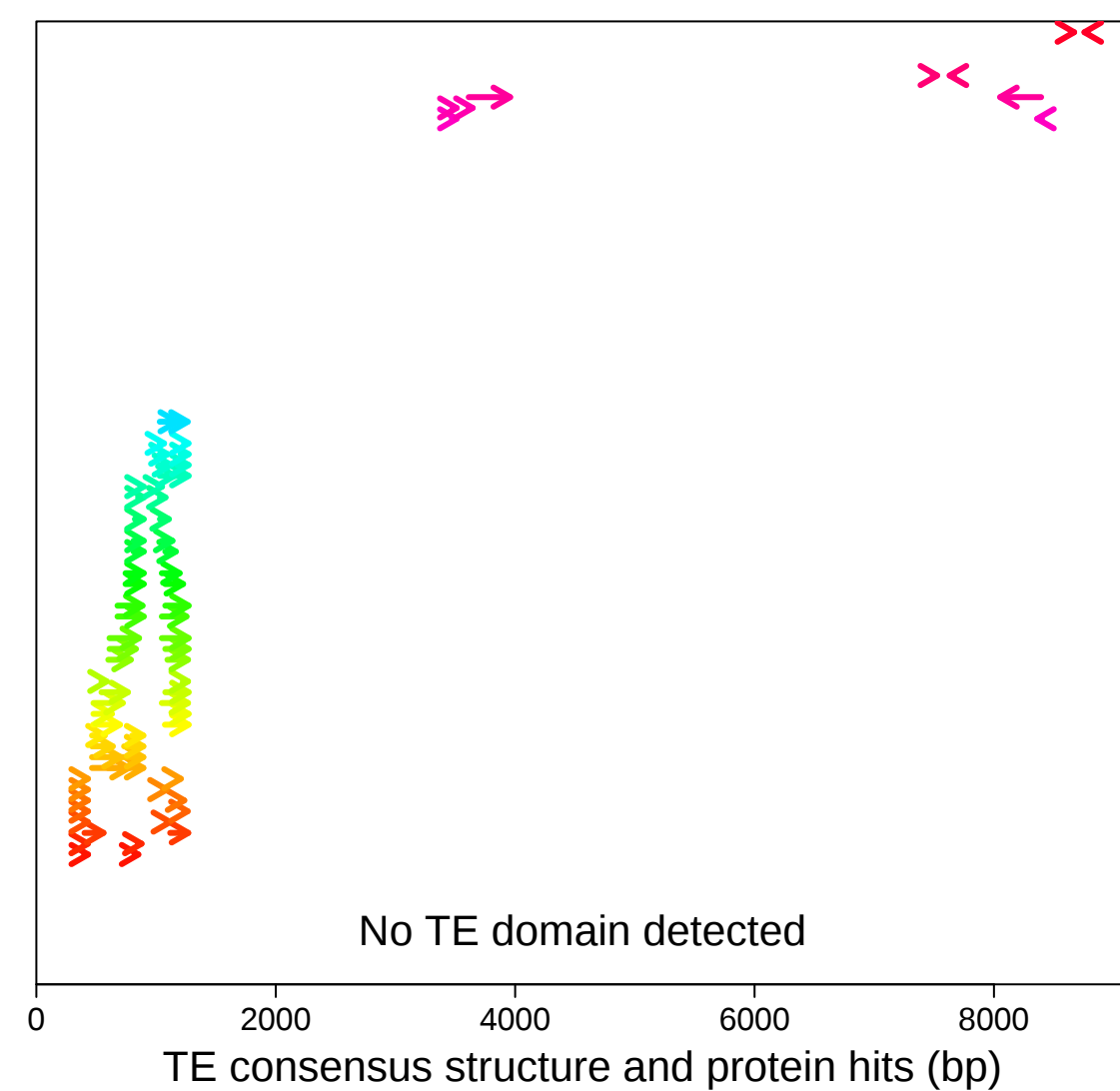
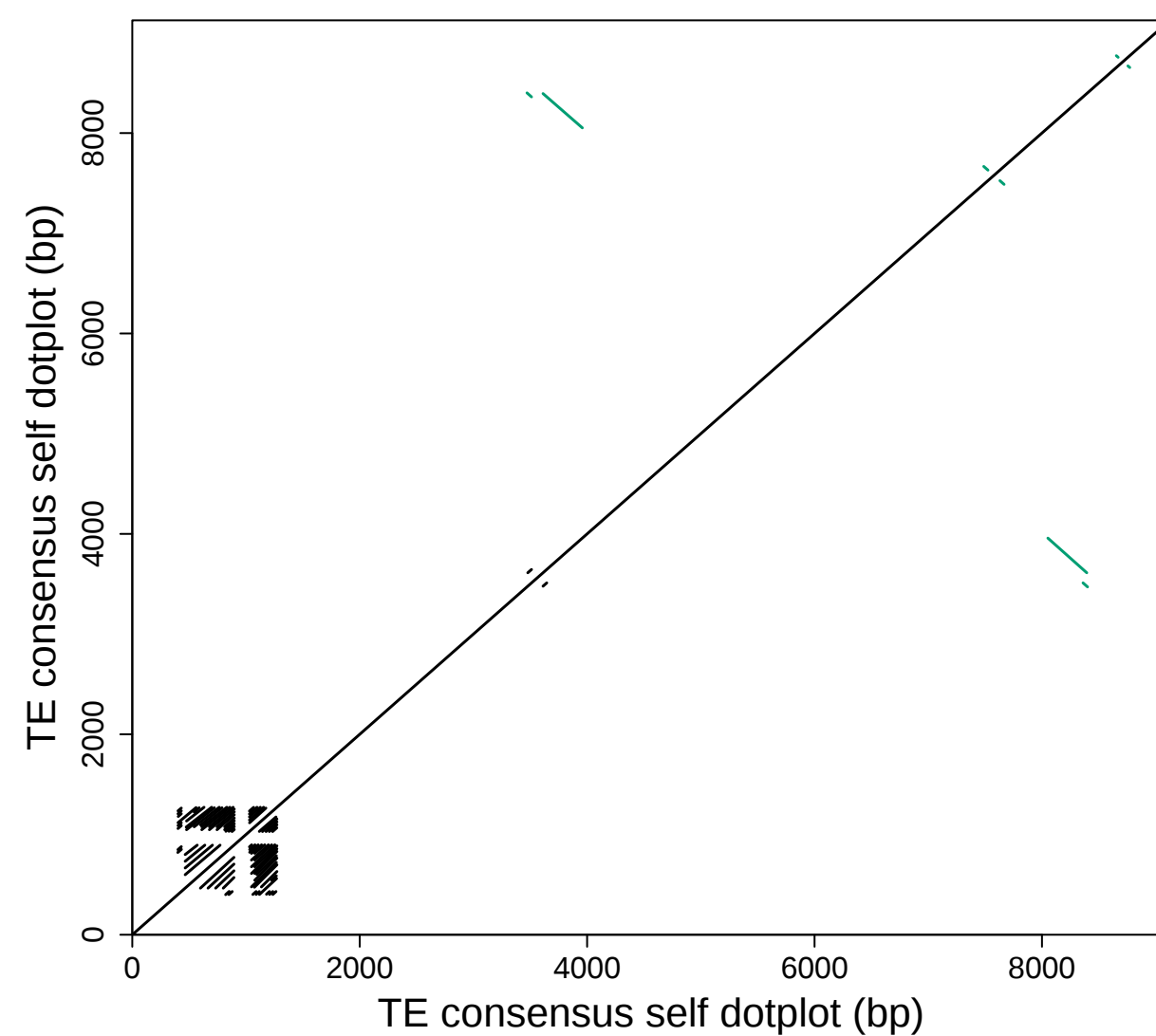
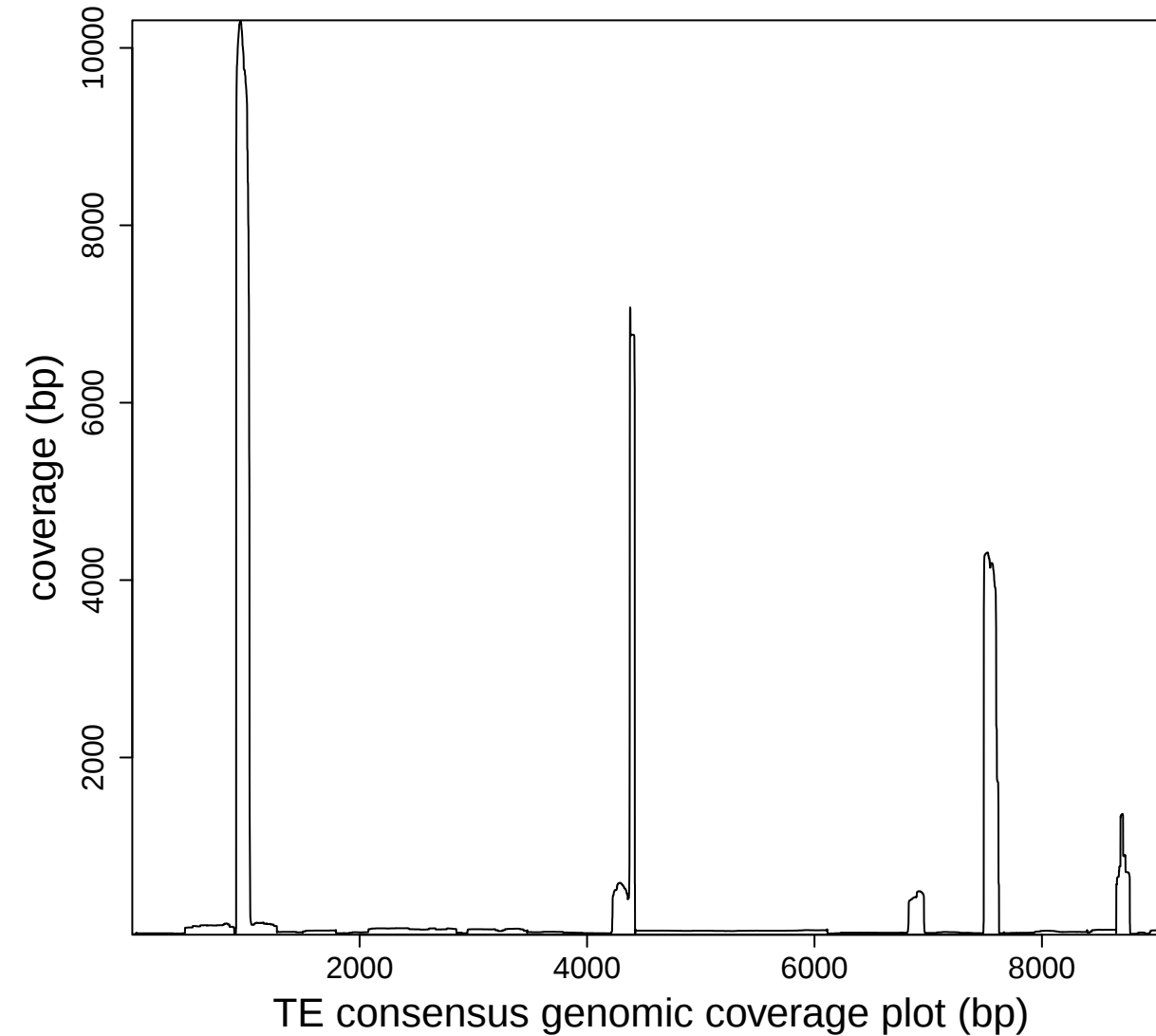
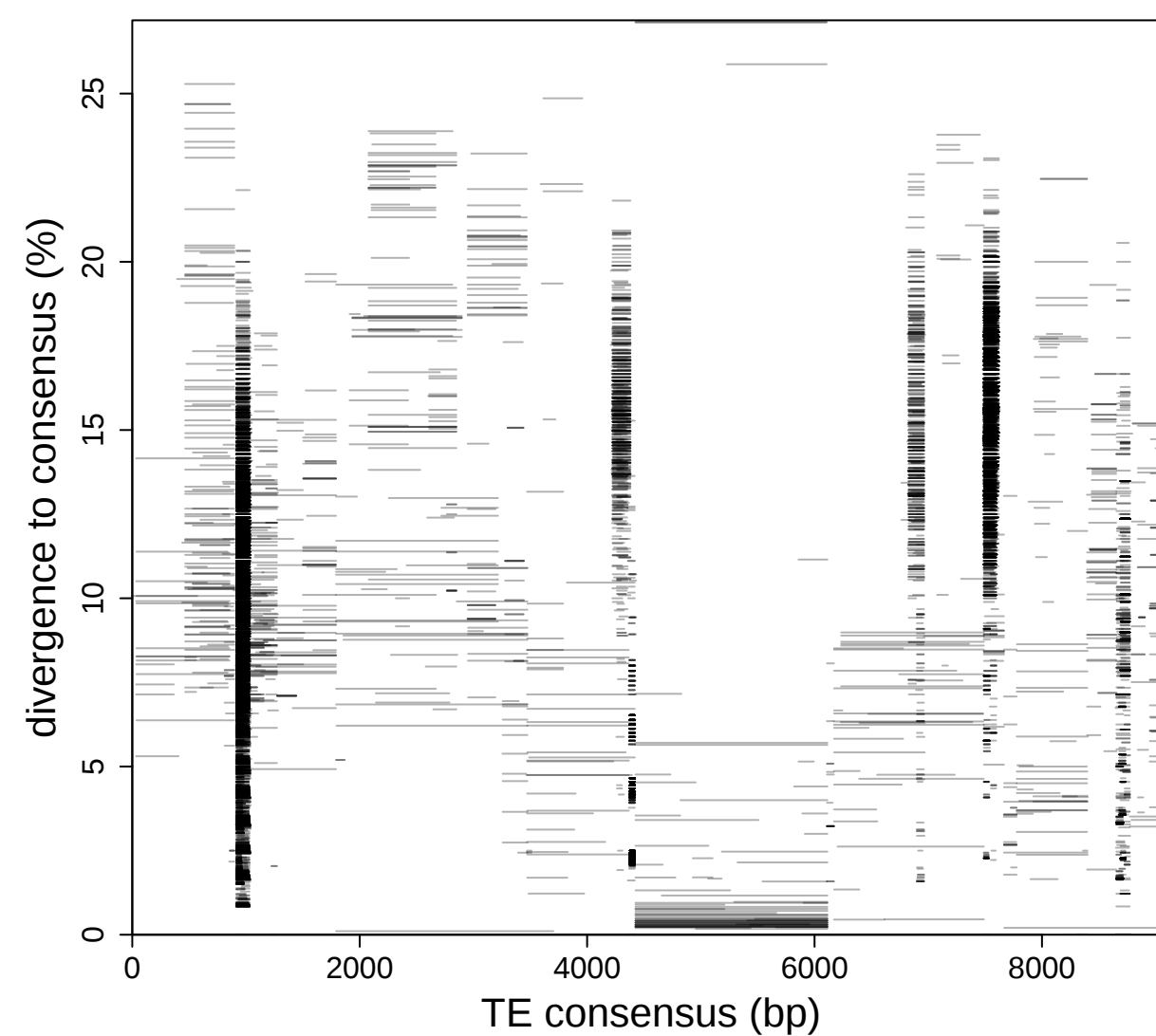
ly_576.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 6850bp; fragments: 571; full length: 38 (>=6165bp)

After TEtrimmer Extended plot Blue lines are boundaries

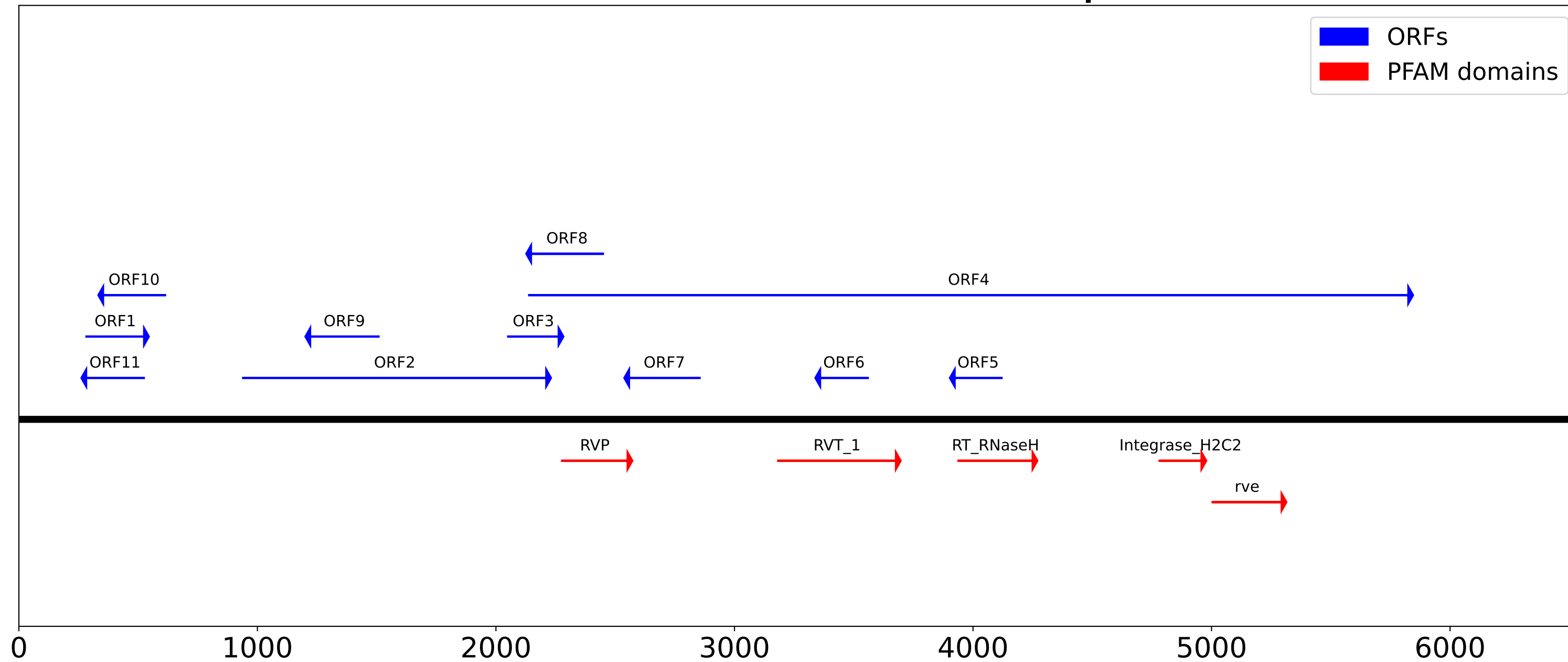


TE: ltr_1_family_576
size: 9125bp; fragments: 24910; full length: 0 (>=8212.5bp)

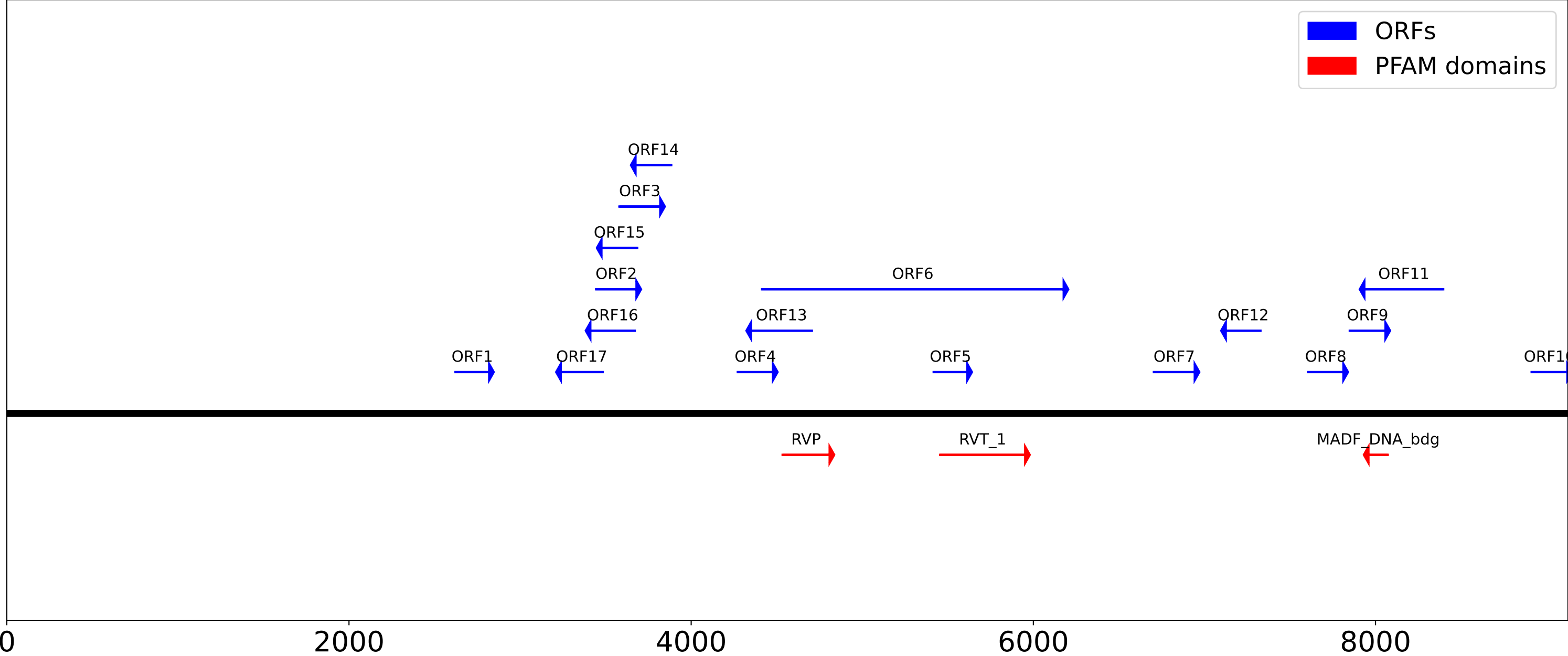
Before TEtrimmer 9125 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

