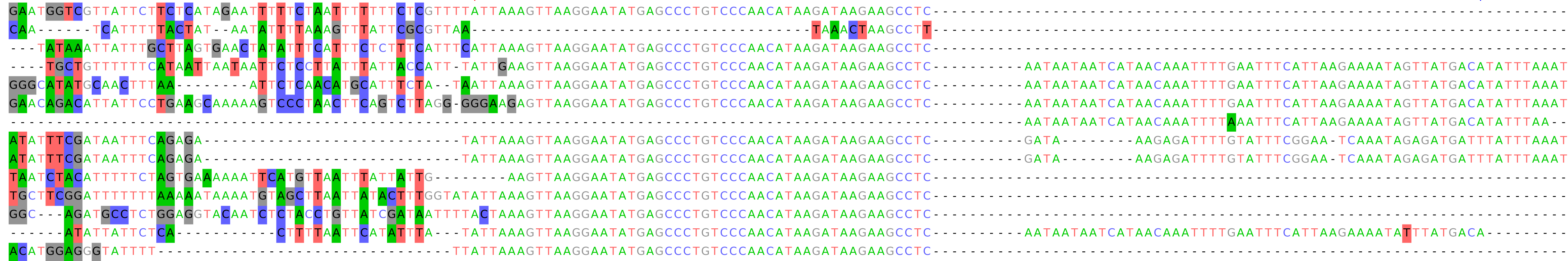


Start crop Point

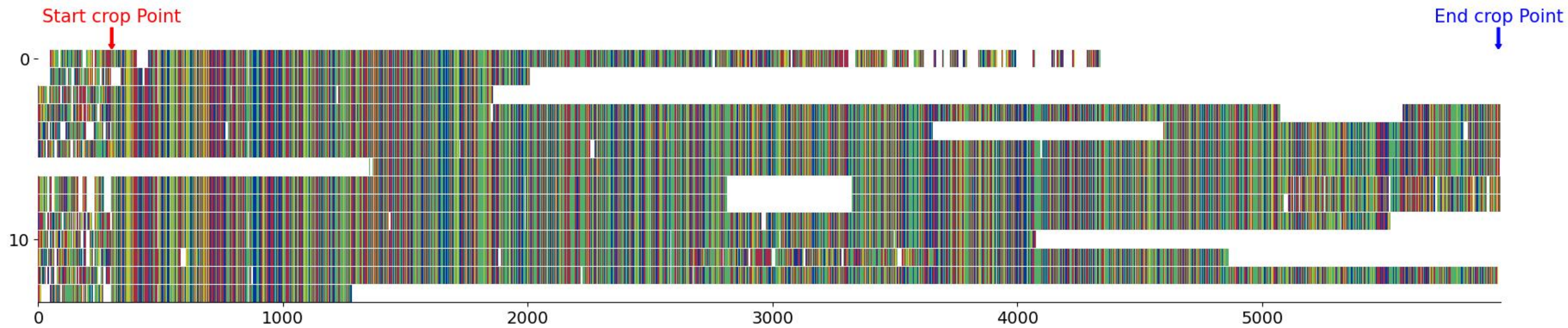
End crop Point

MSA length = 5667



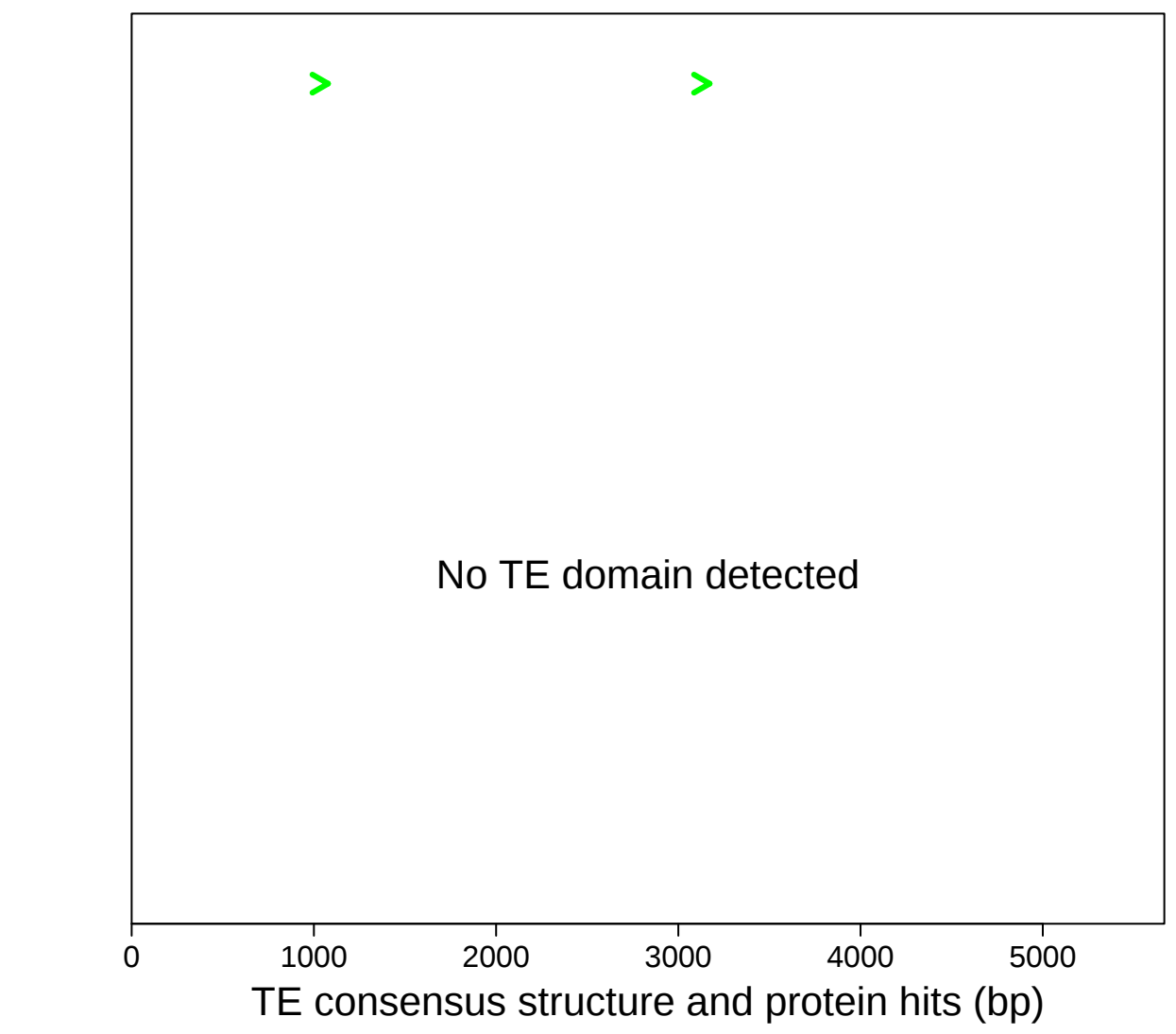
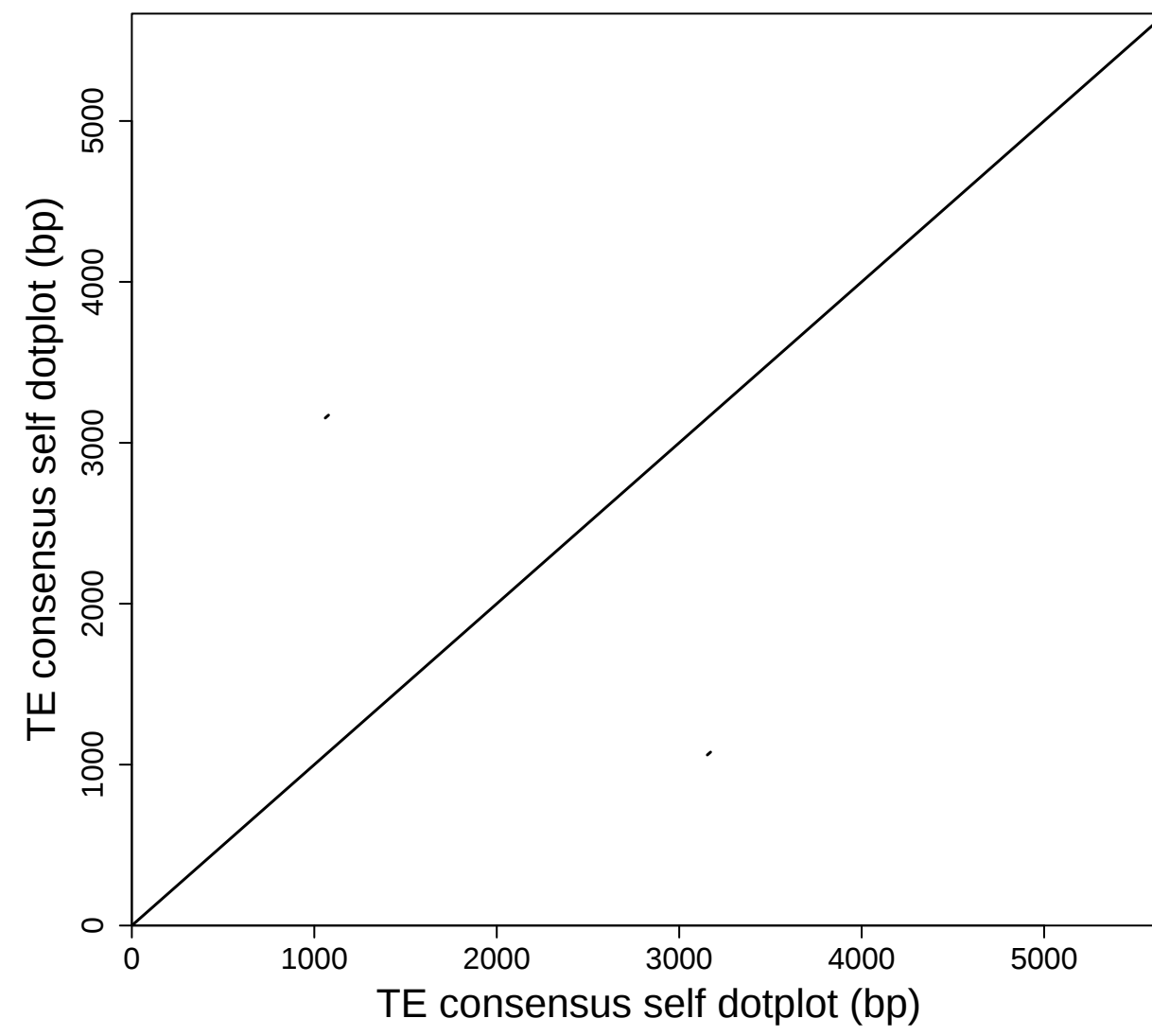
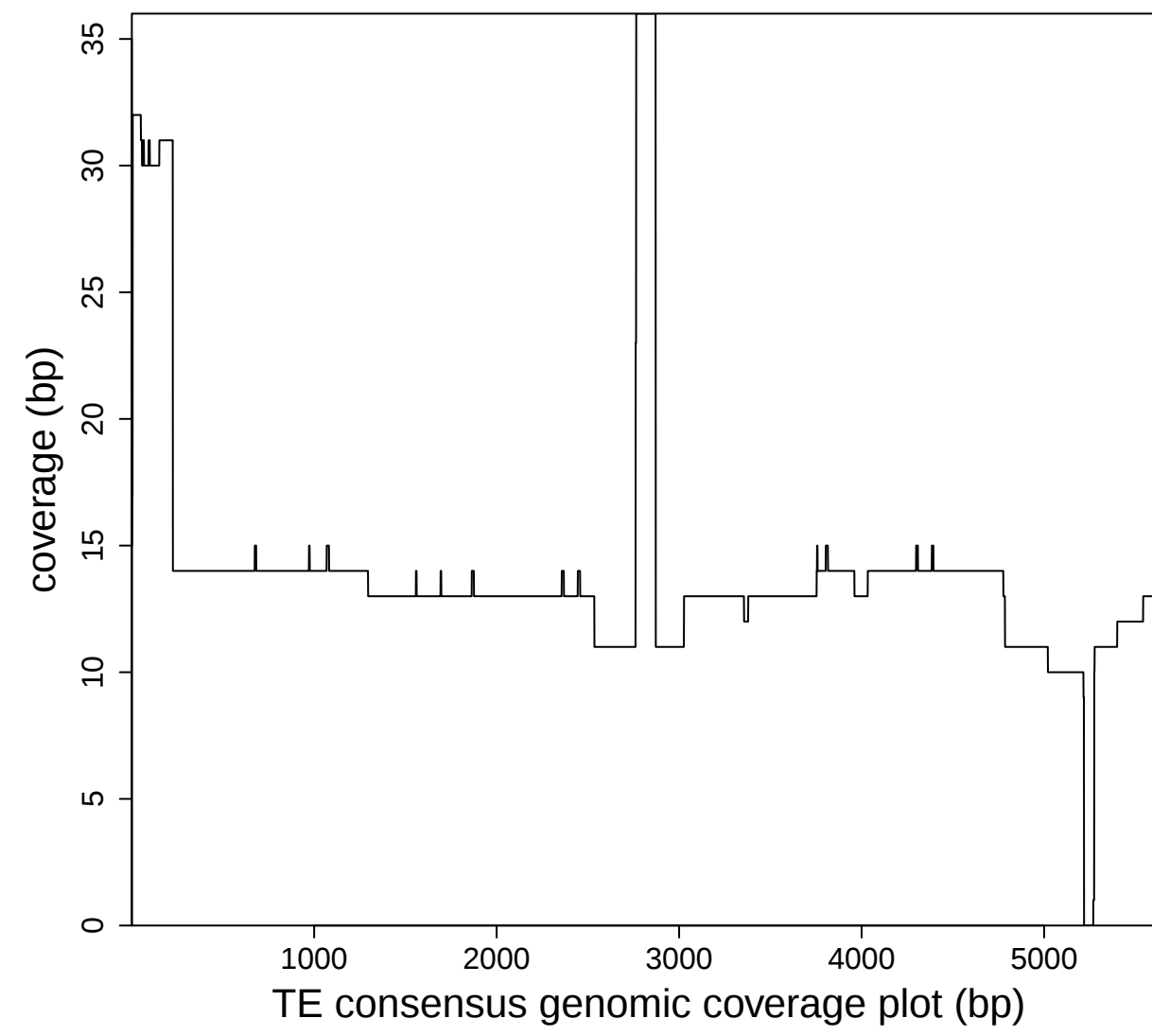
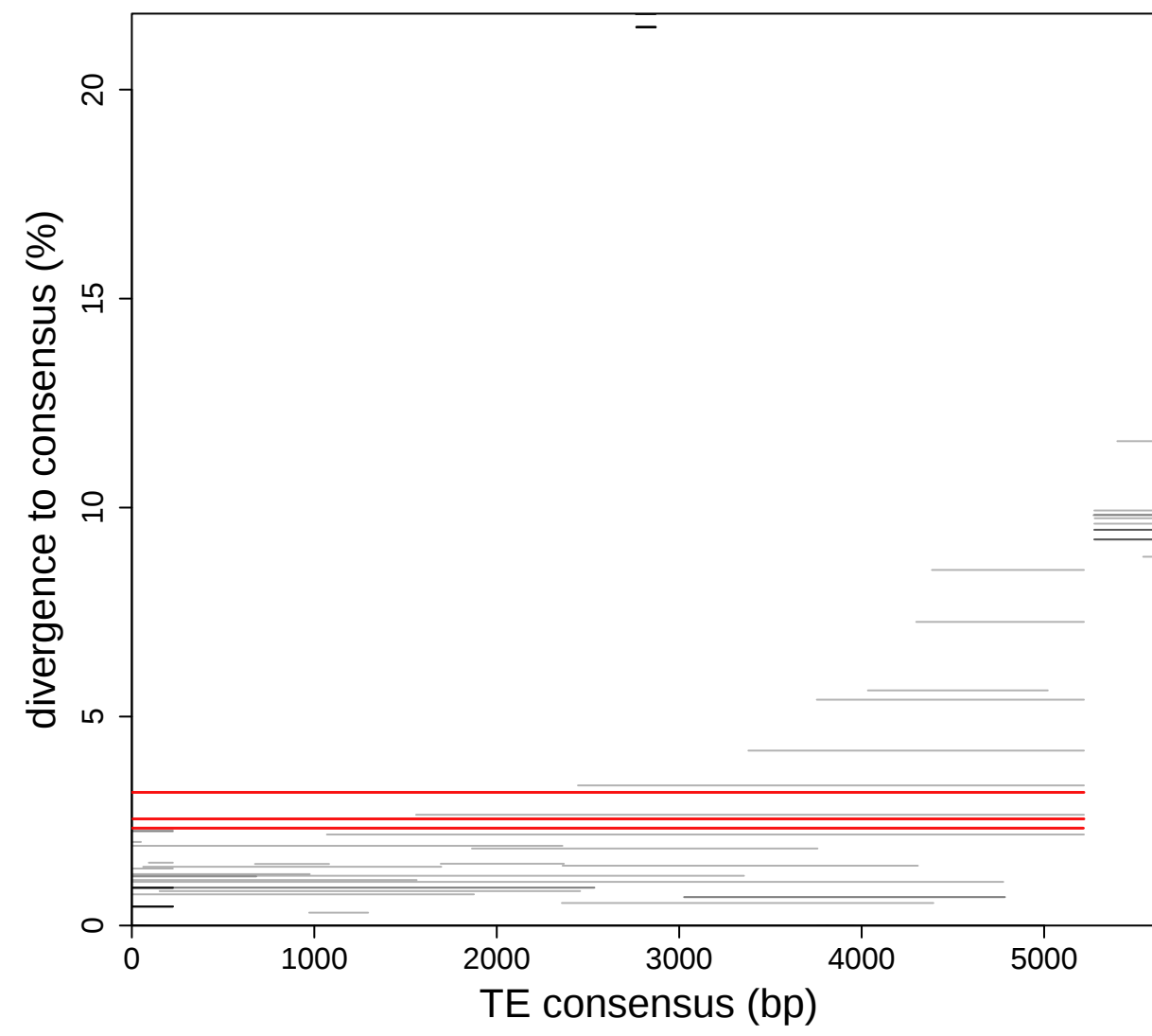
Start crop Point

End crop Point



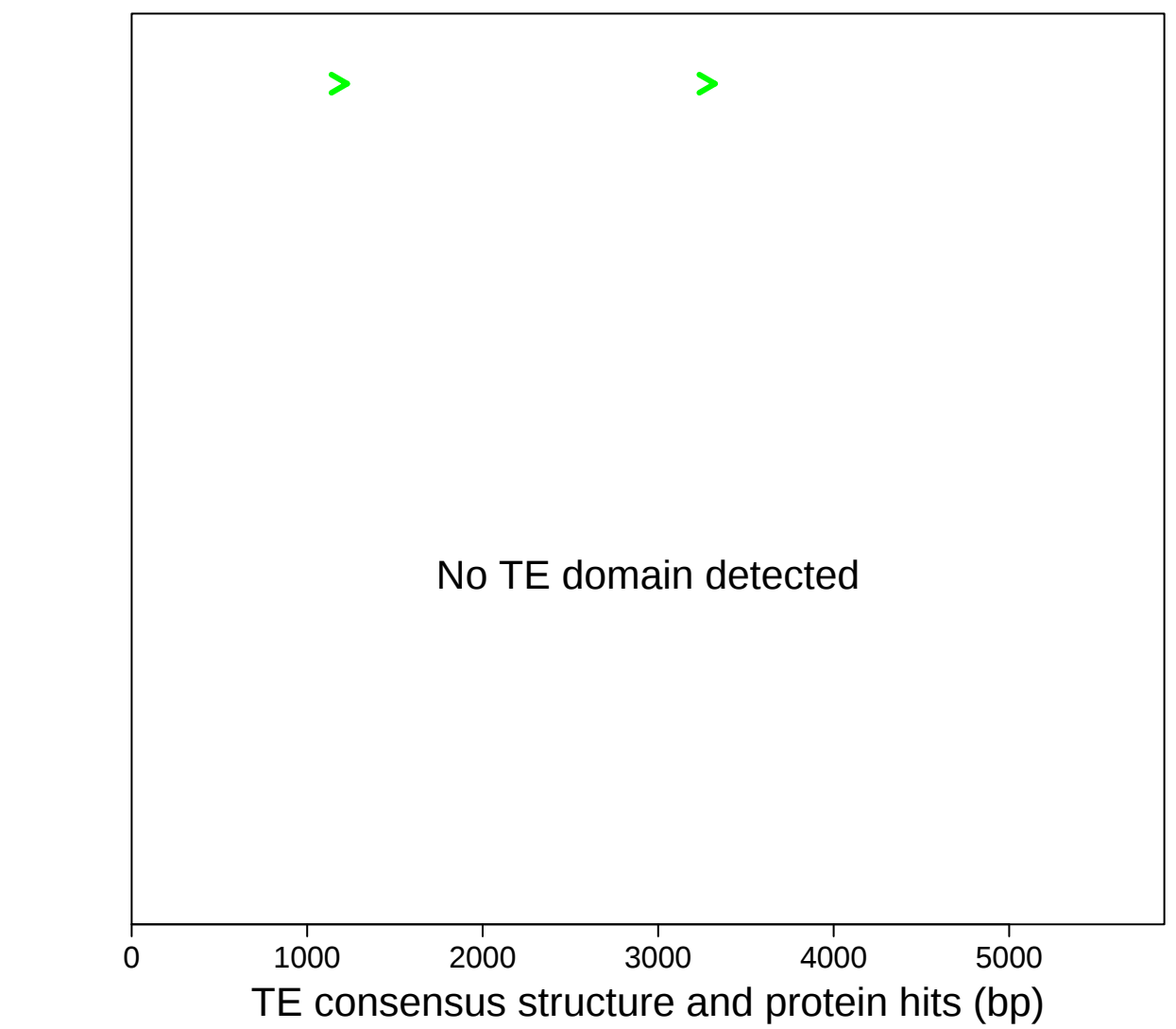
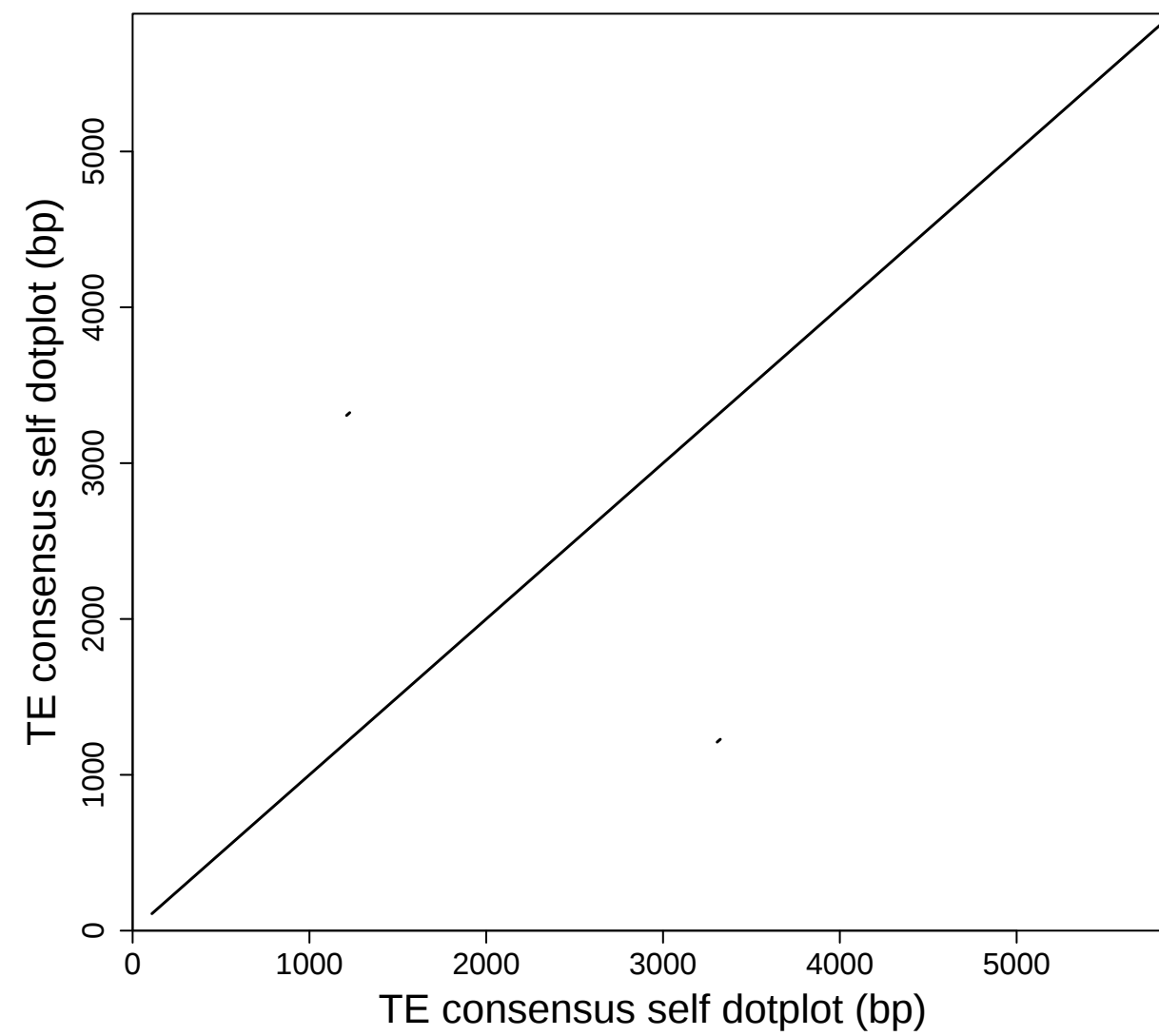
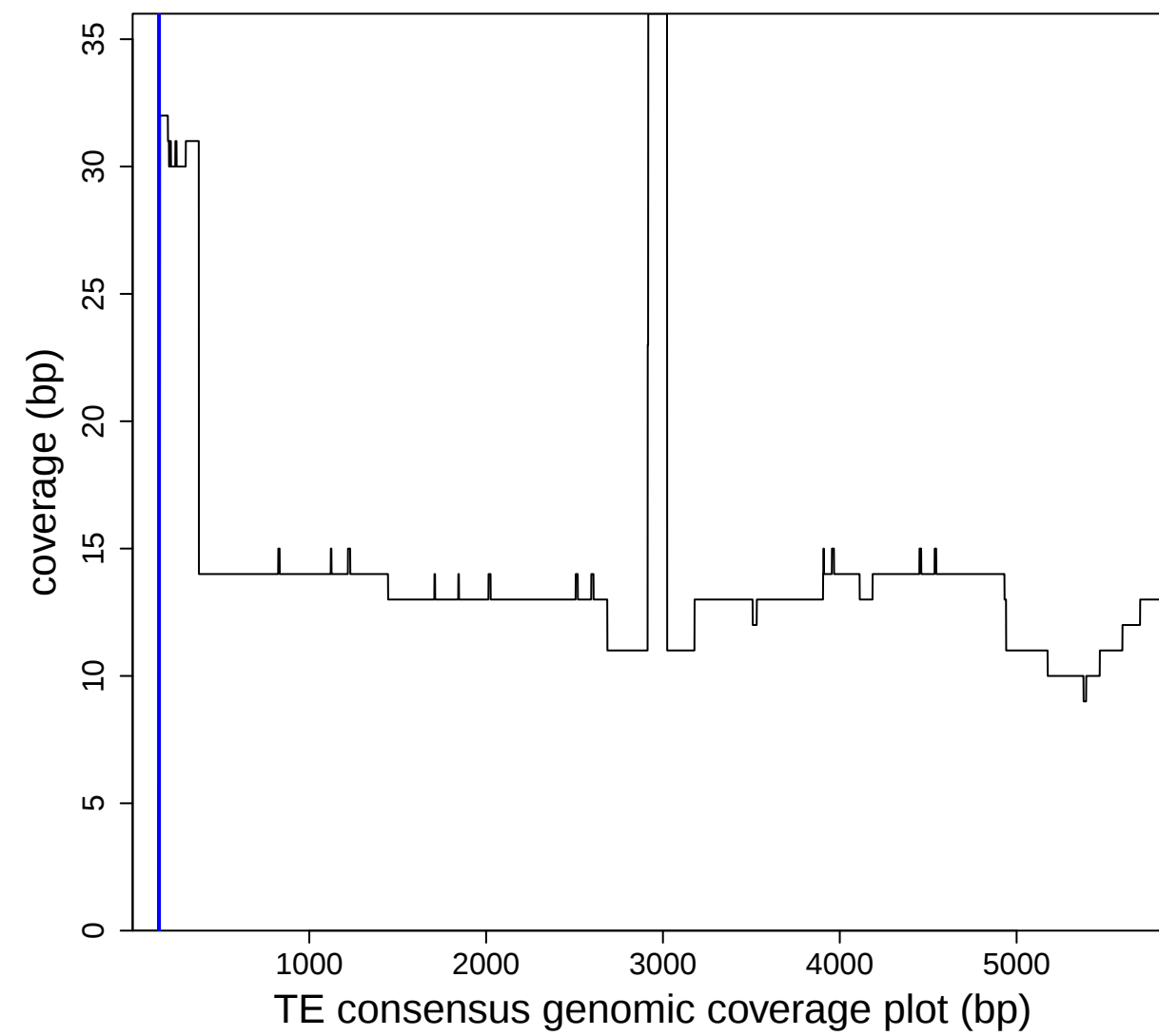
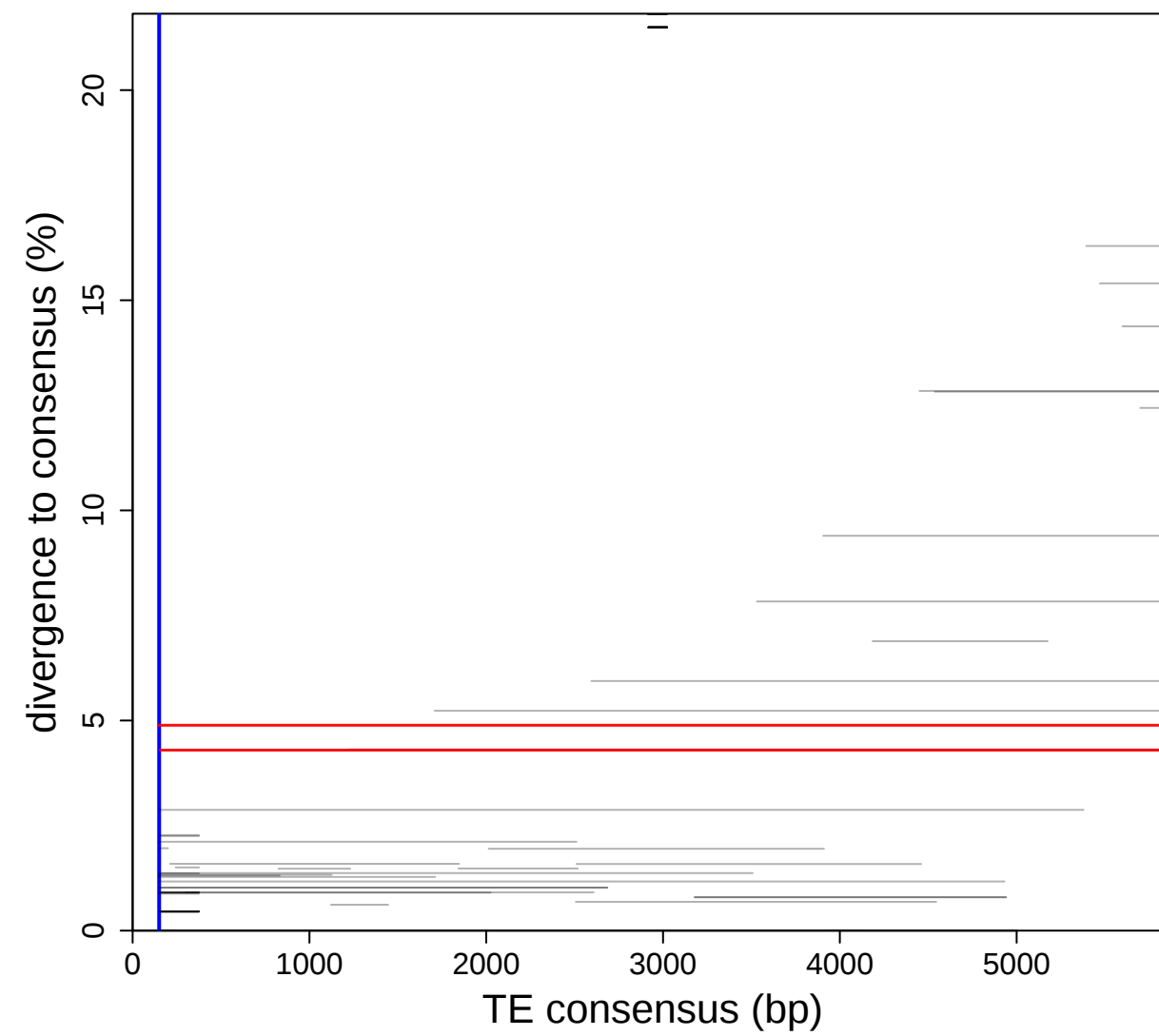
b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa
size: 5667bp; fragments: 91; full length: 3 (≥ 5100.3 bp)

After TEtrimmer 5667 bp



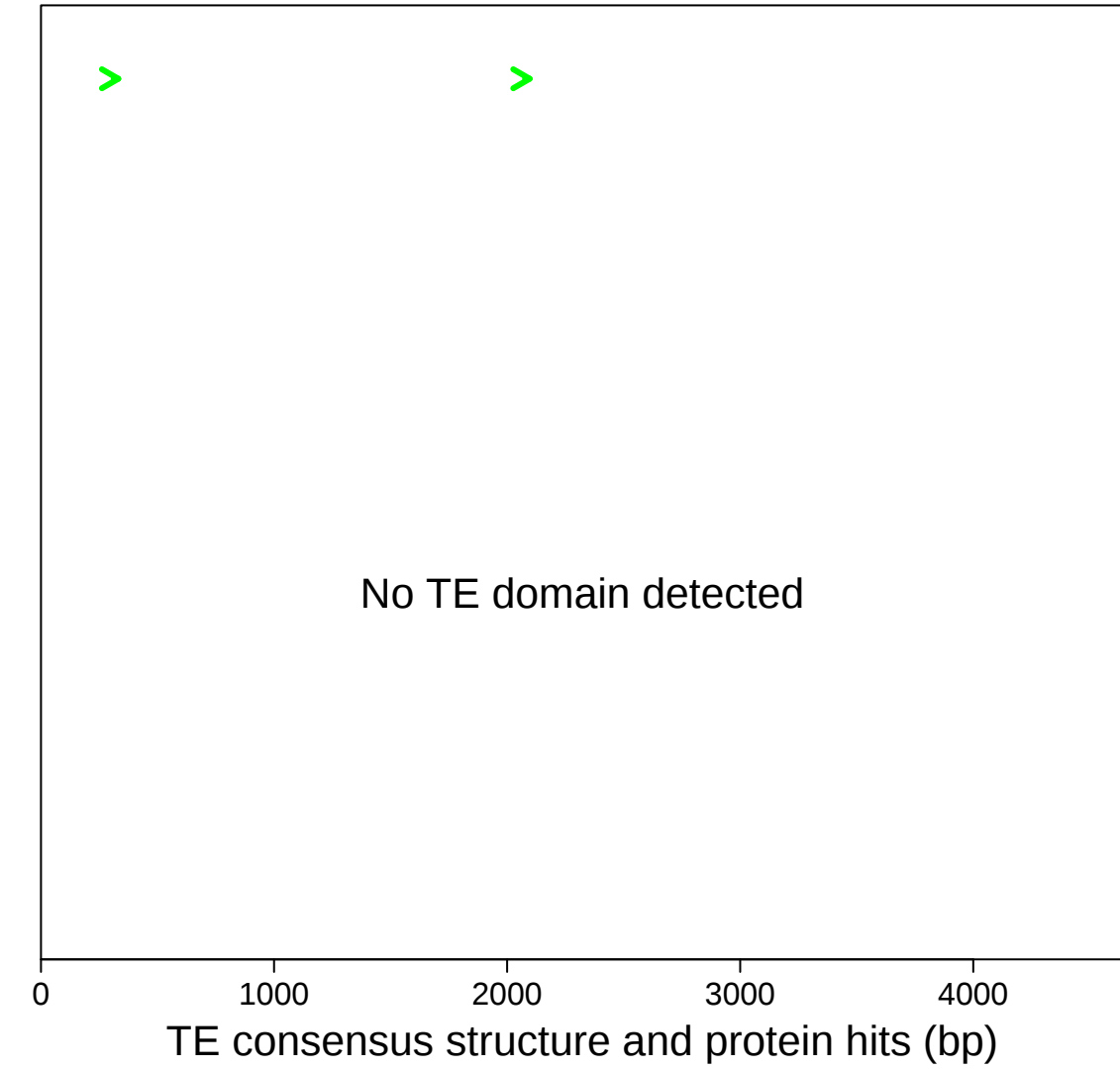
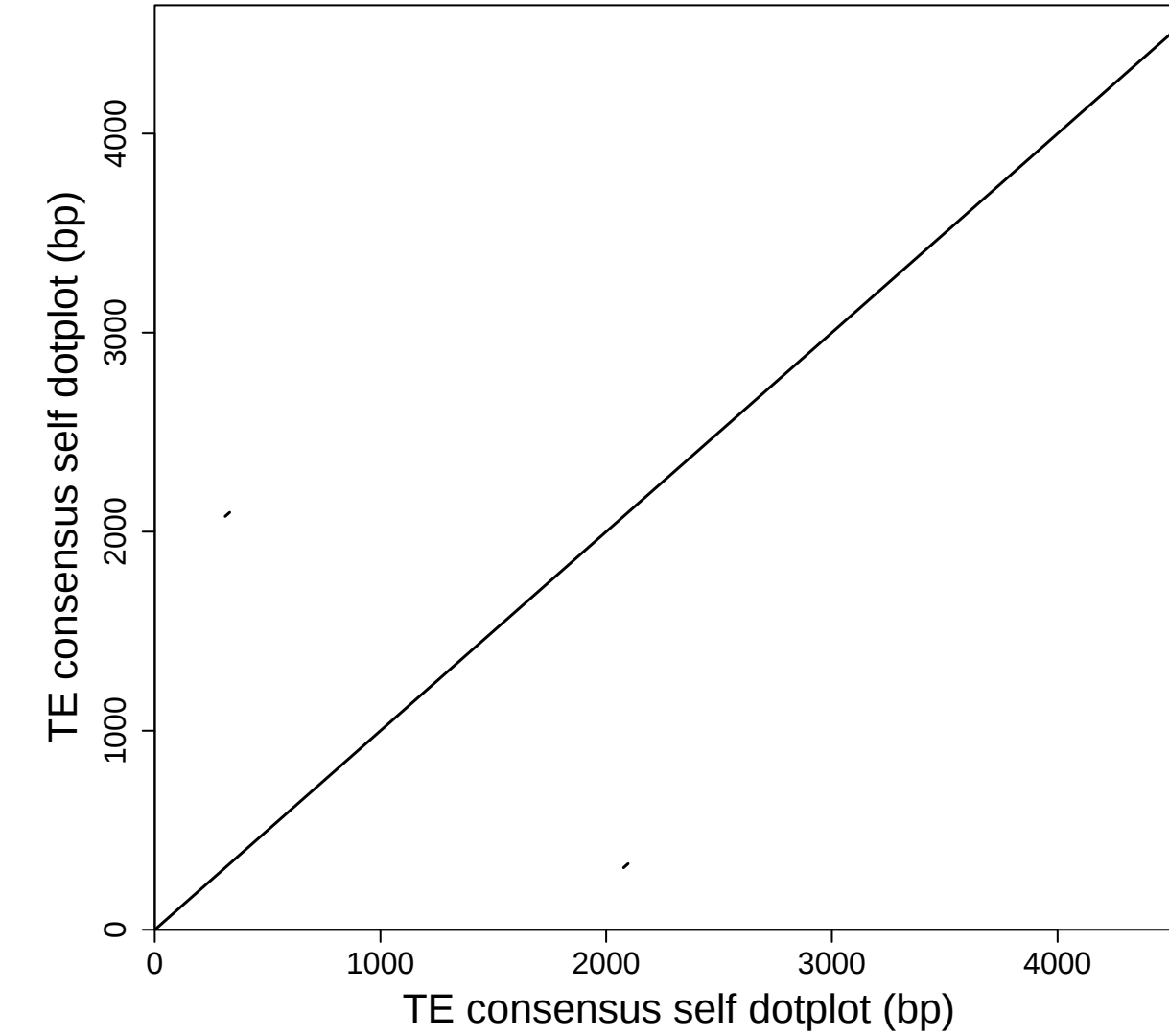
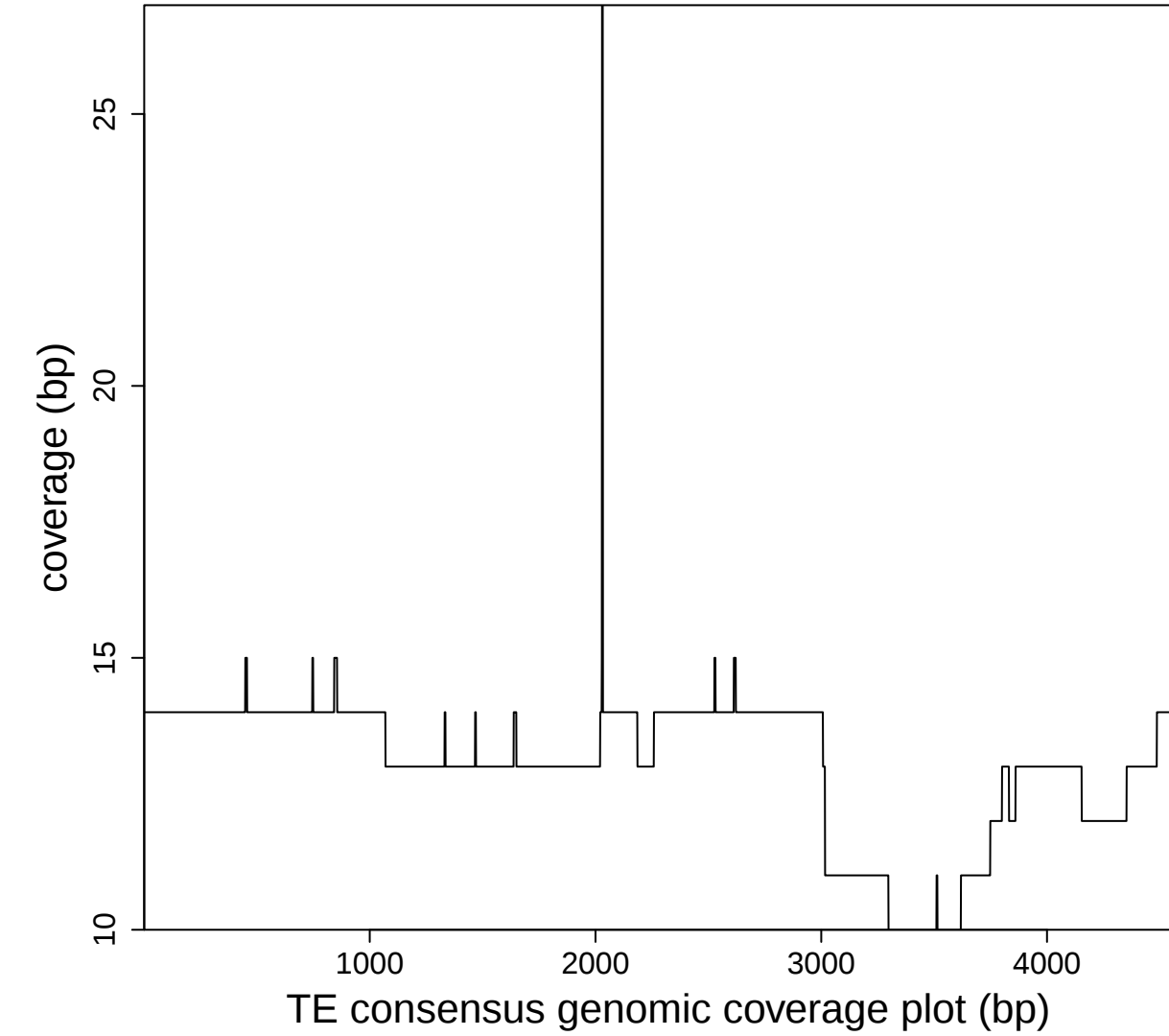
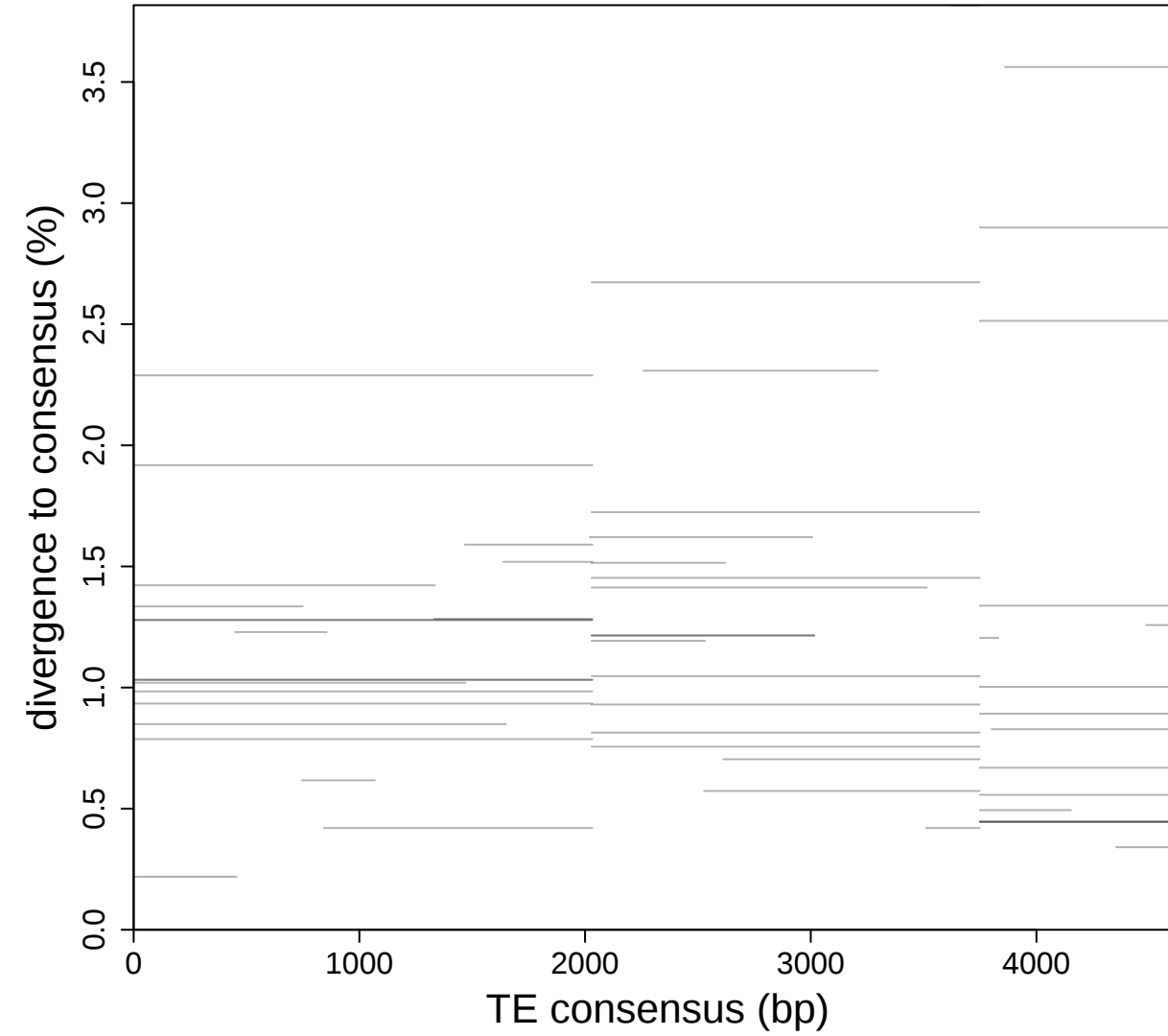
3.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa
size: 5884bp; fragments: 82; full length: 2 (>=5295.6bp)

After TEtrimmer Extended plot Blue lines are boundaries

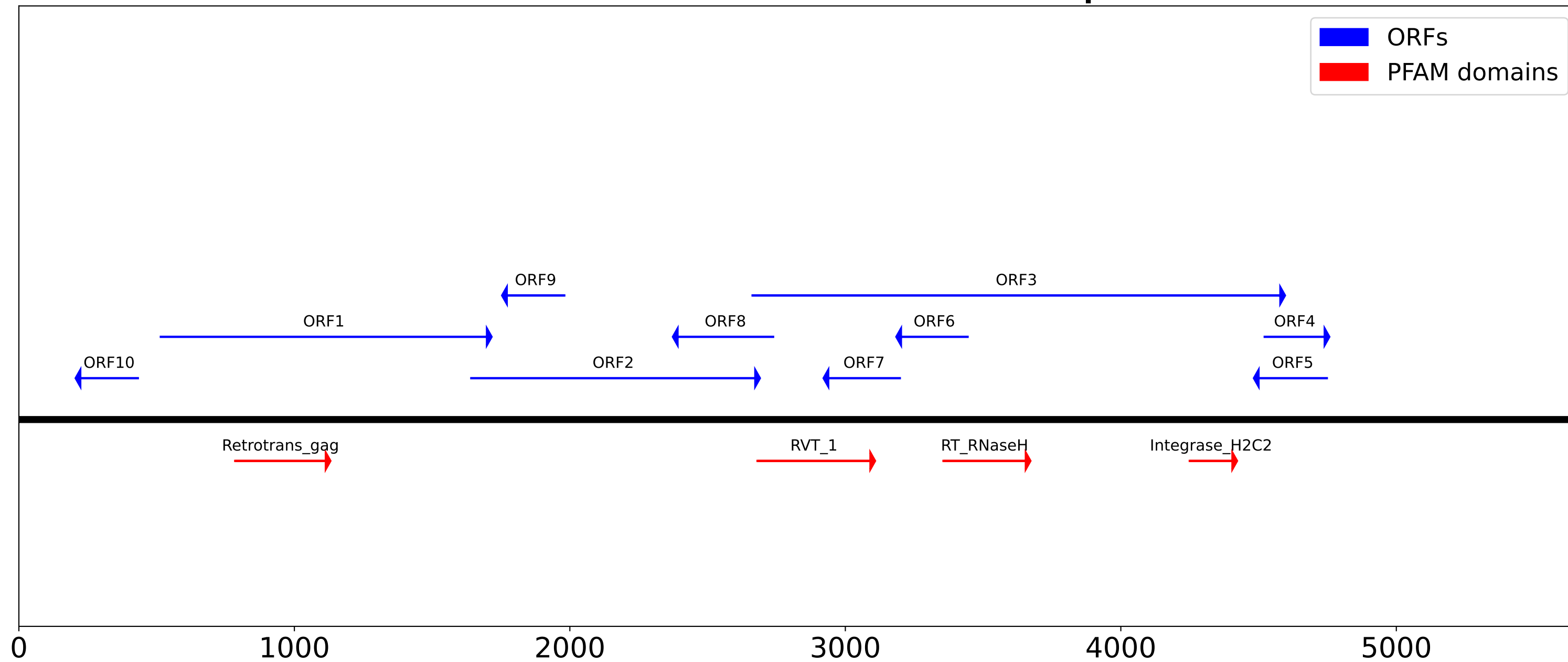


TE: ltr_1_family_613
size: 4645bp; fragments: 55; full length: 0 (≥ 4180.5 bp)

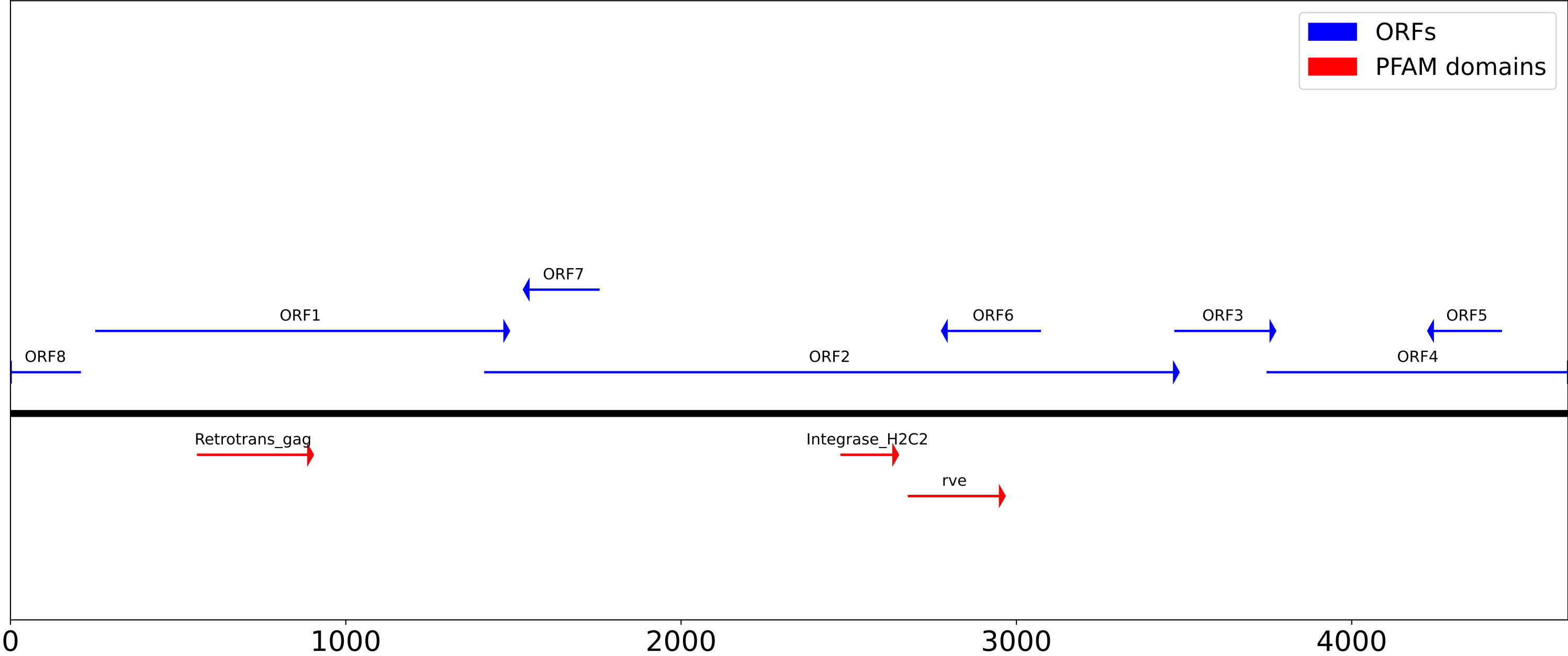
Before TEtrimmer 4645 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

