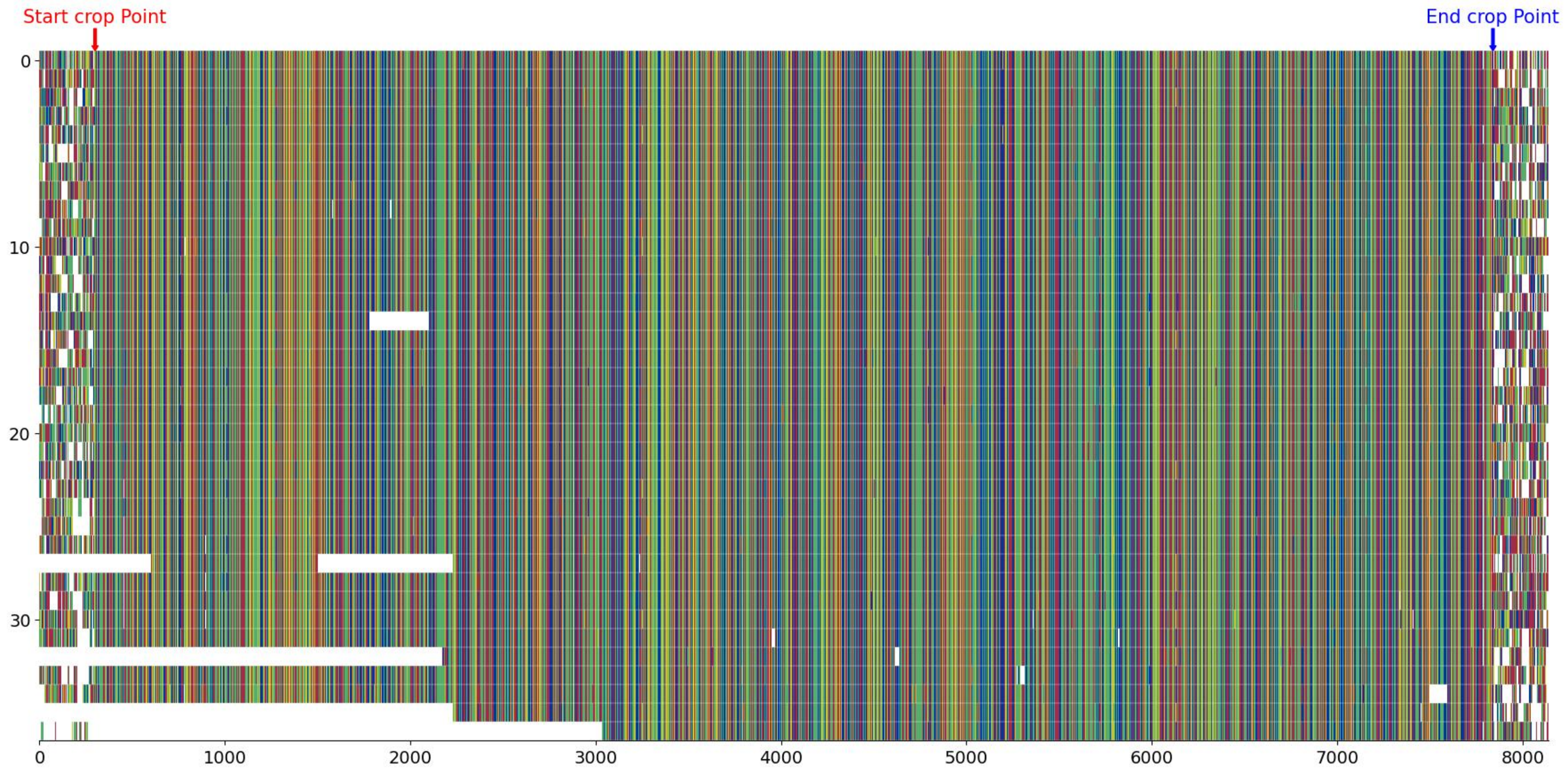


MSA length = 7541

Start crop Point

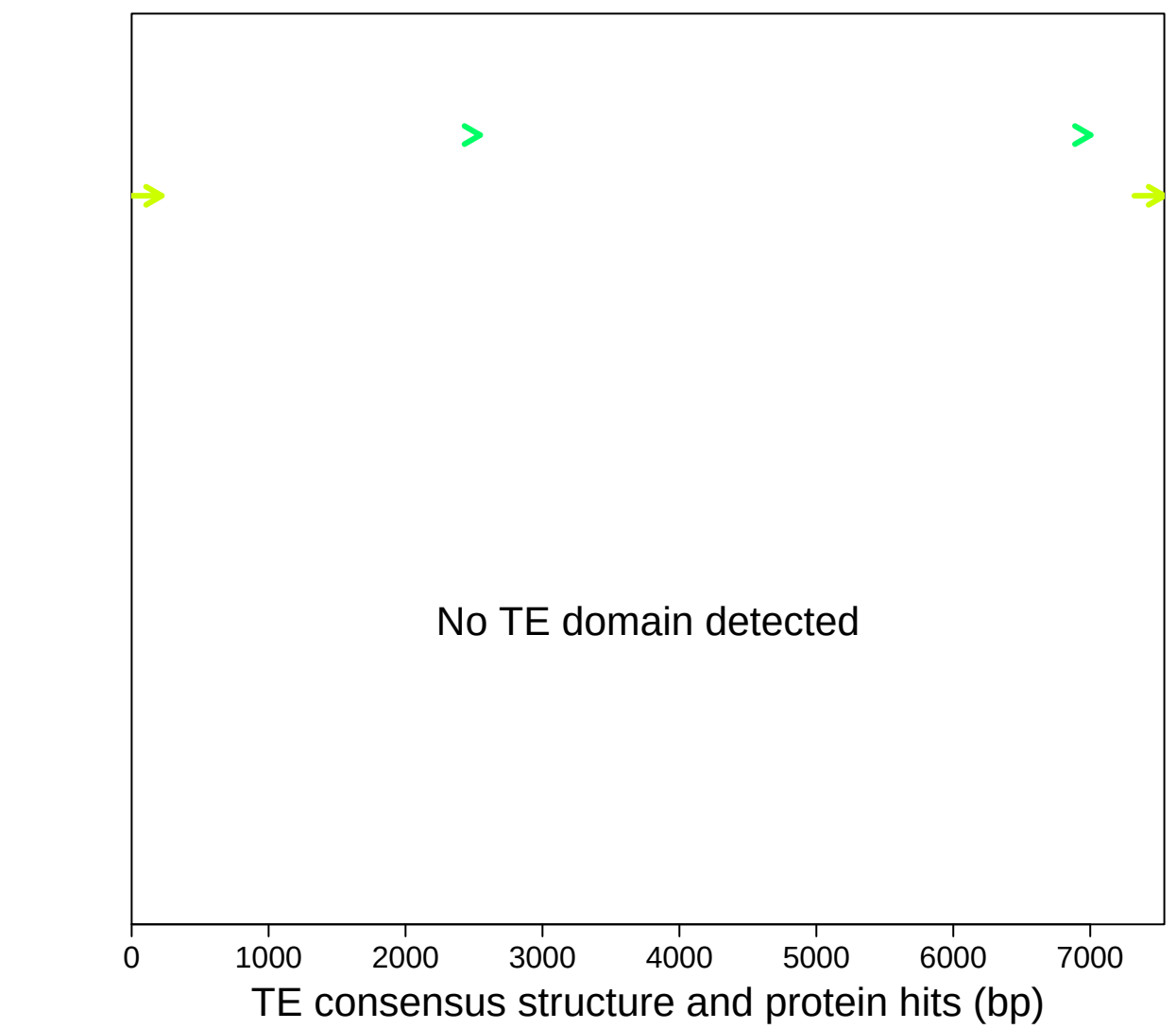
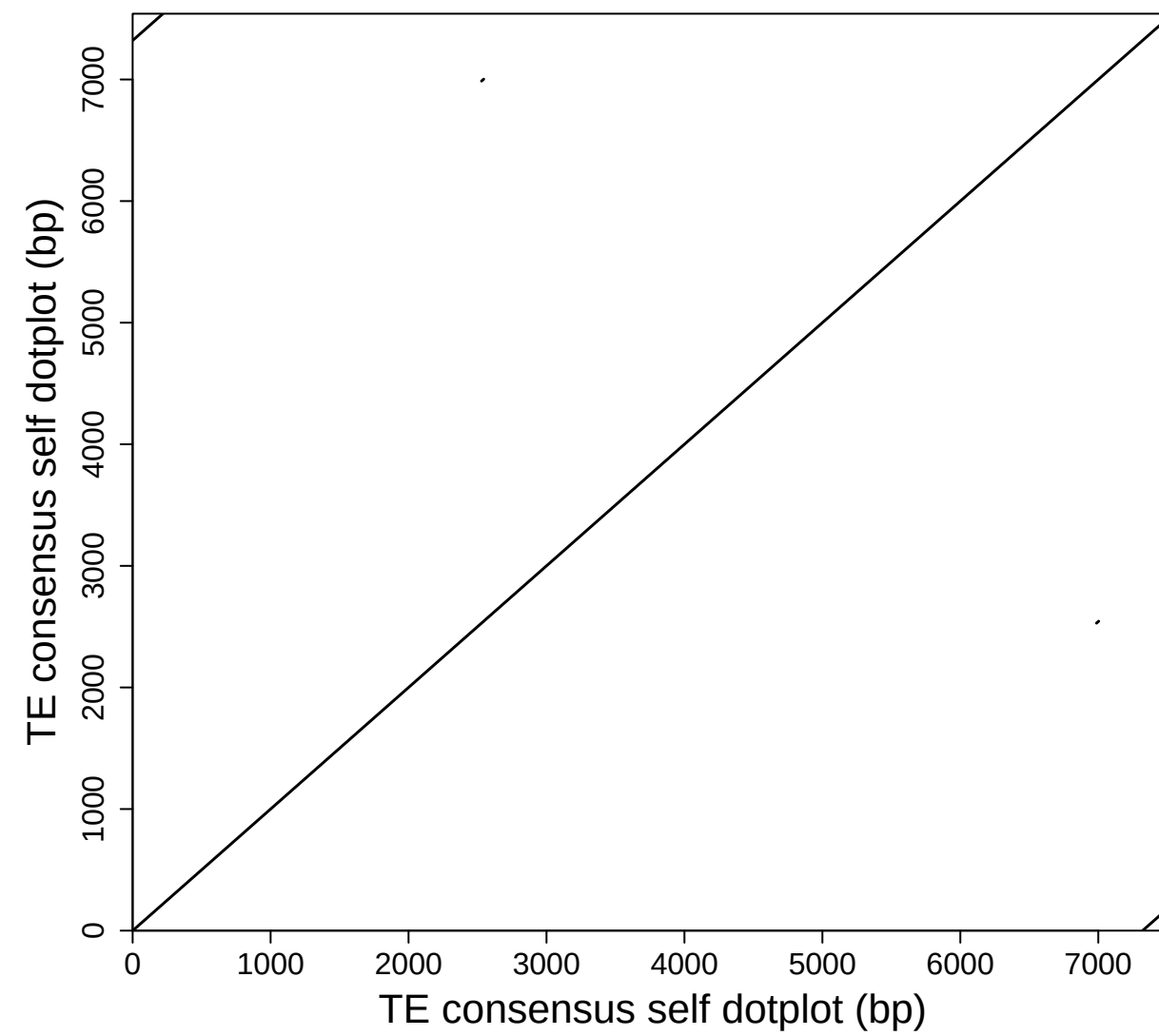
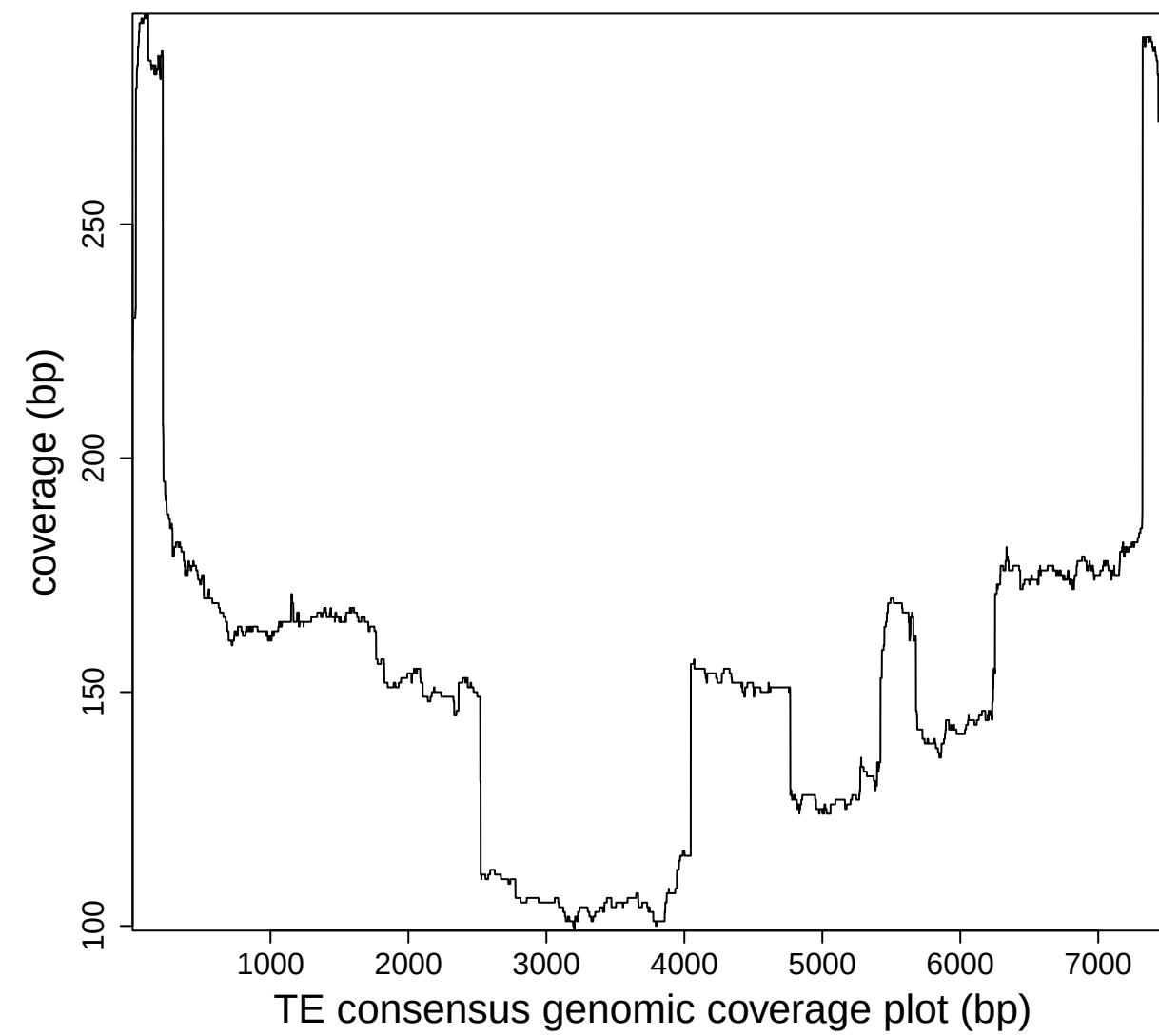
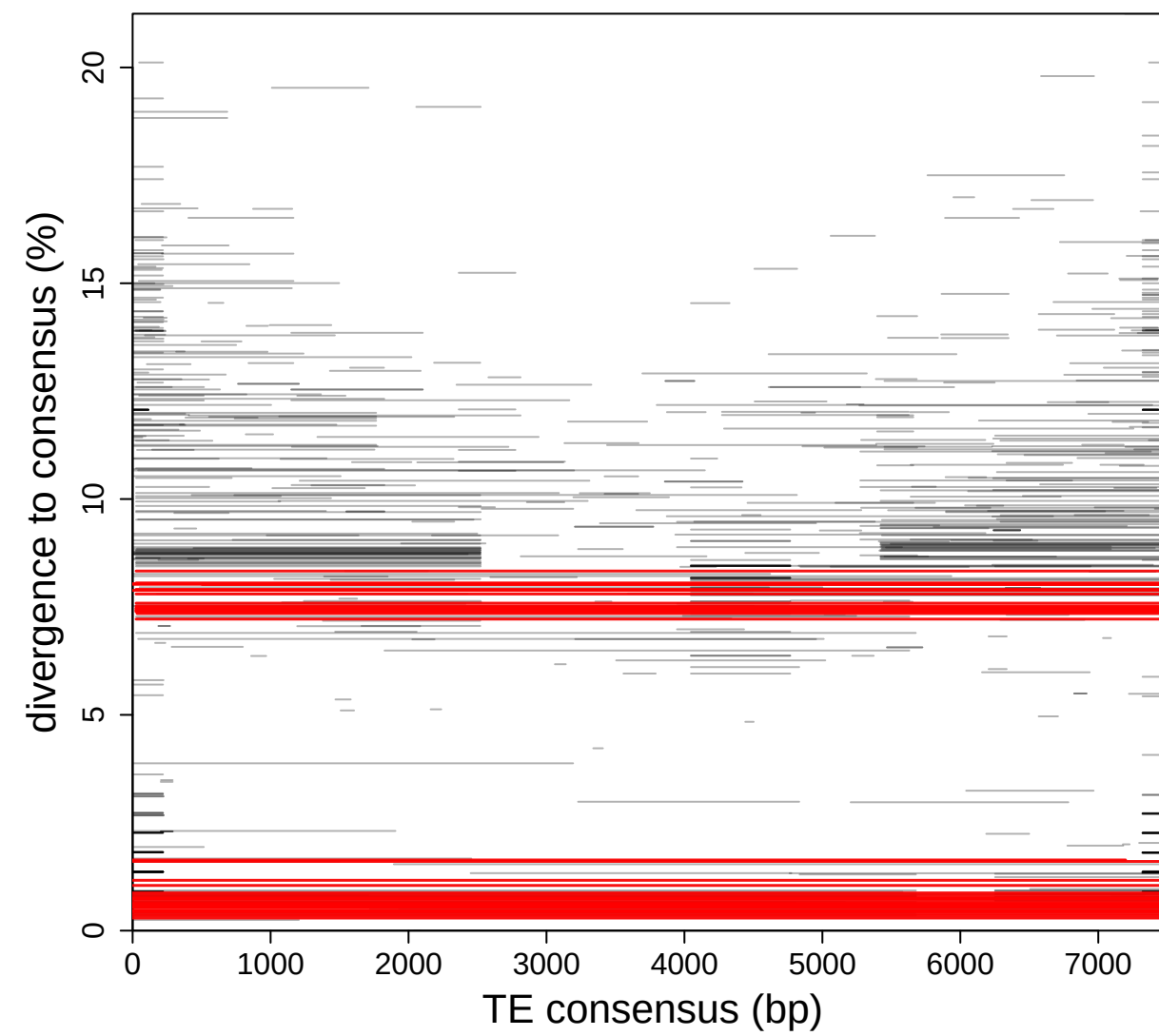
End crop Point





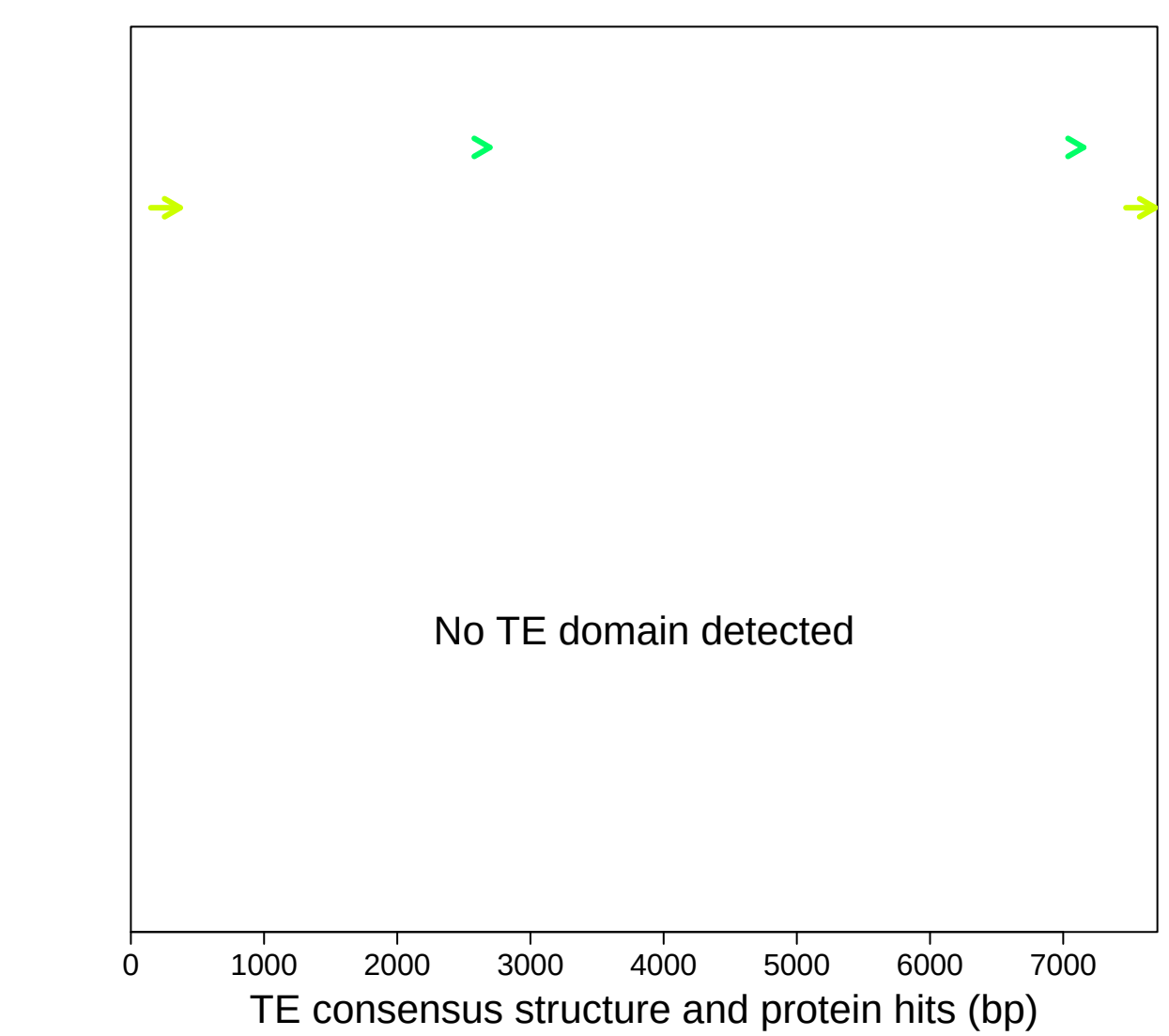
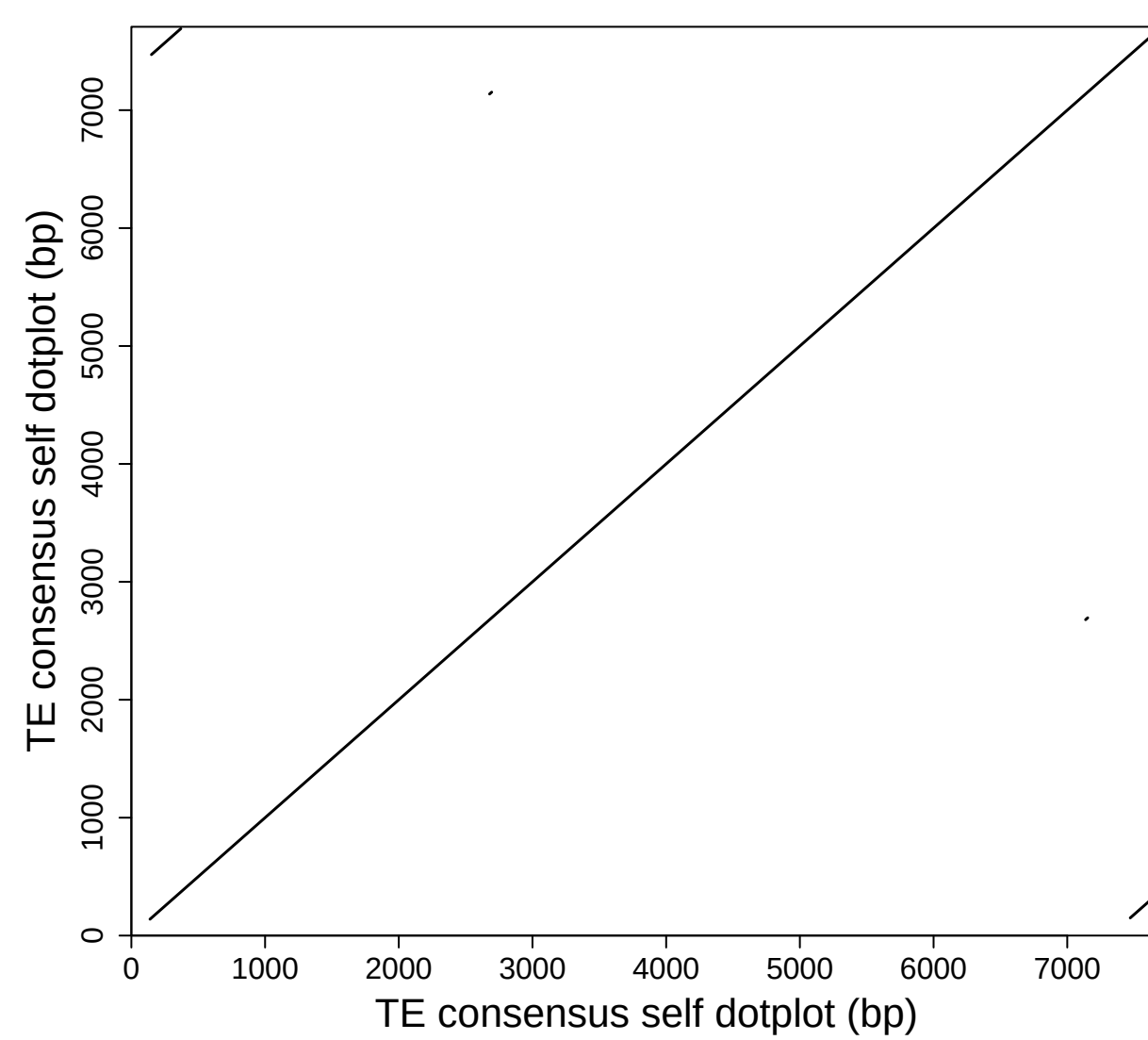
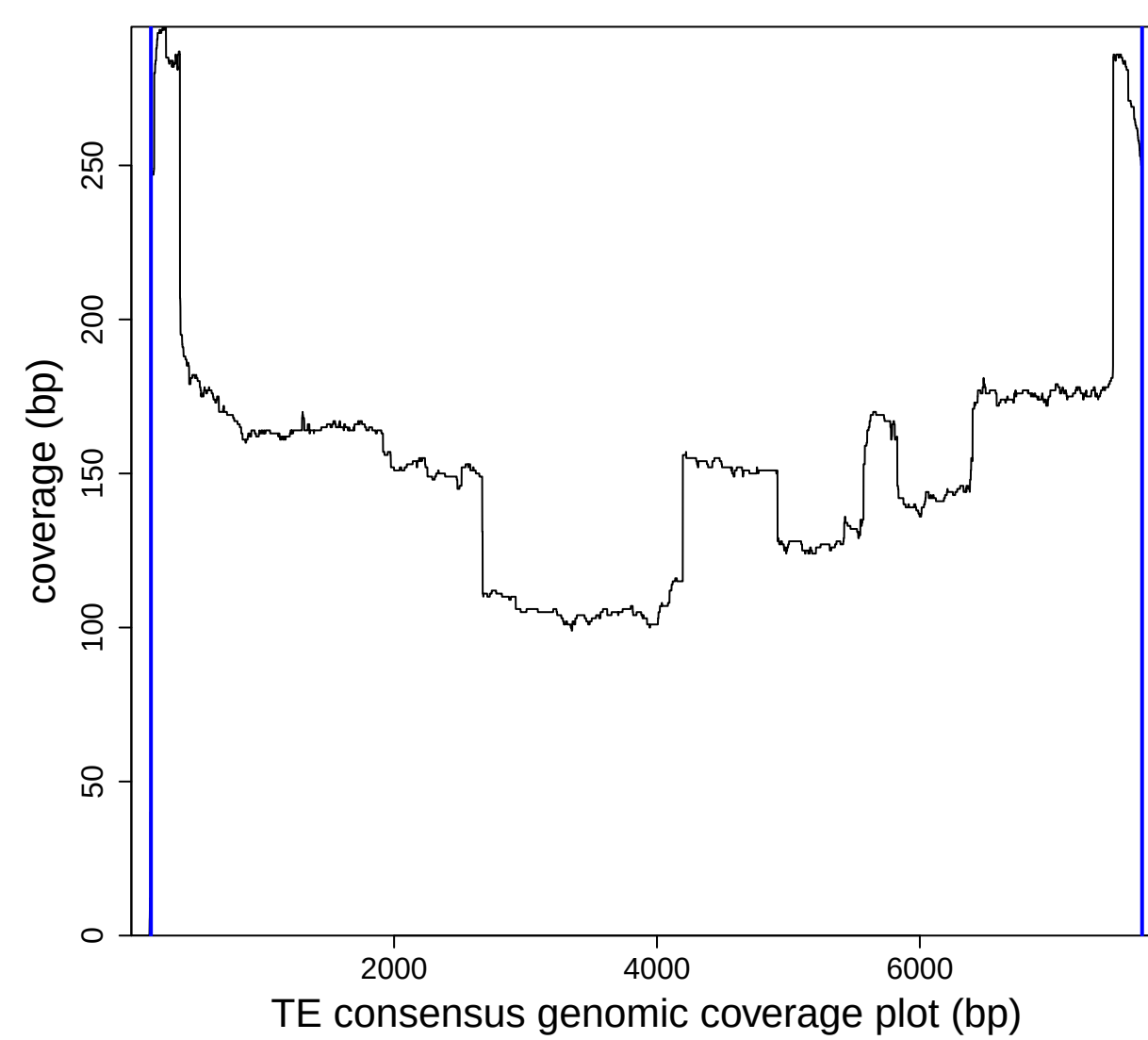
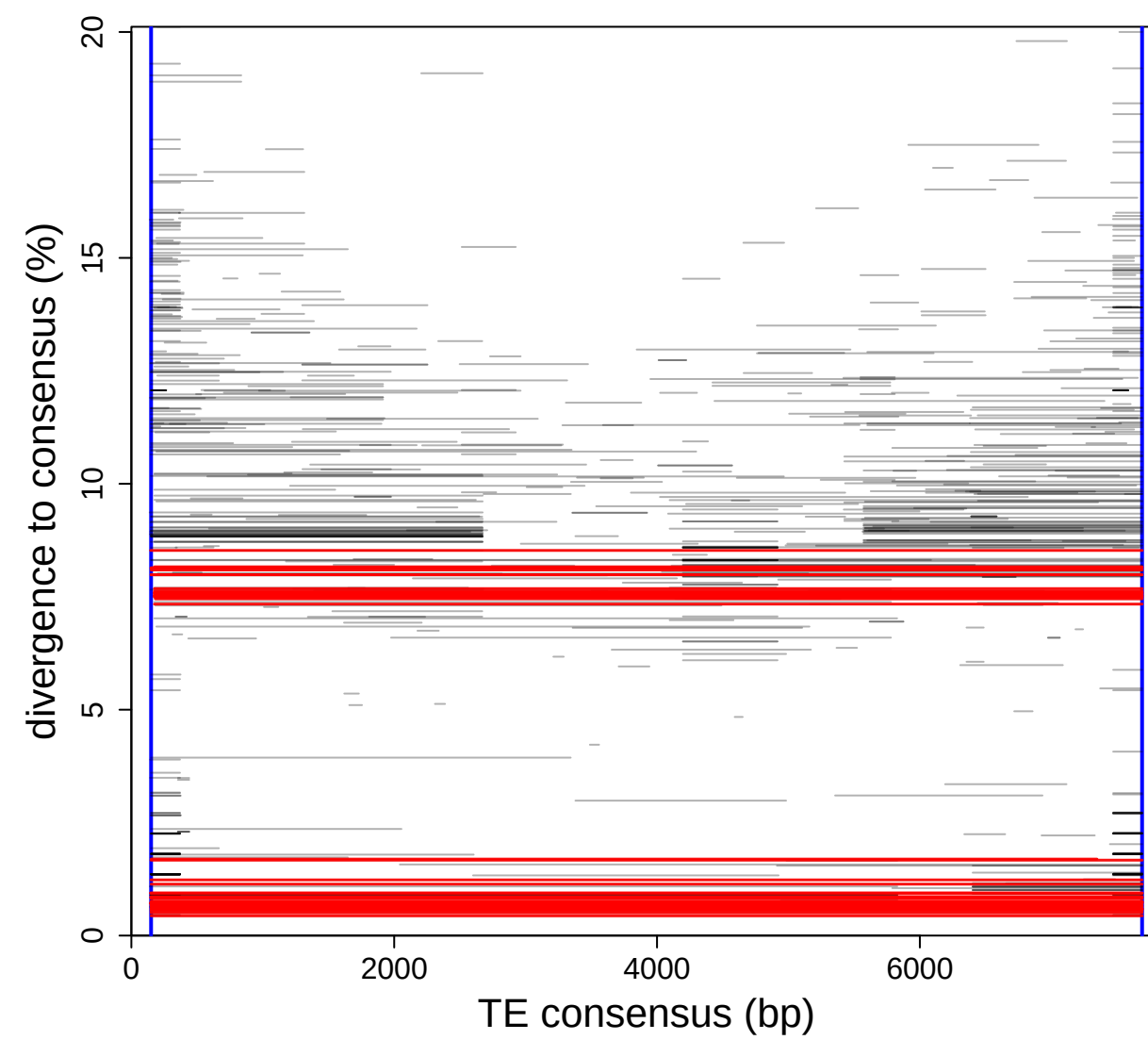
bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 7541bp; fragments: 804; full length: 51 (>=6786.9bp)

After TEtrimmer 7541 bp



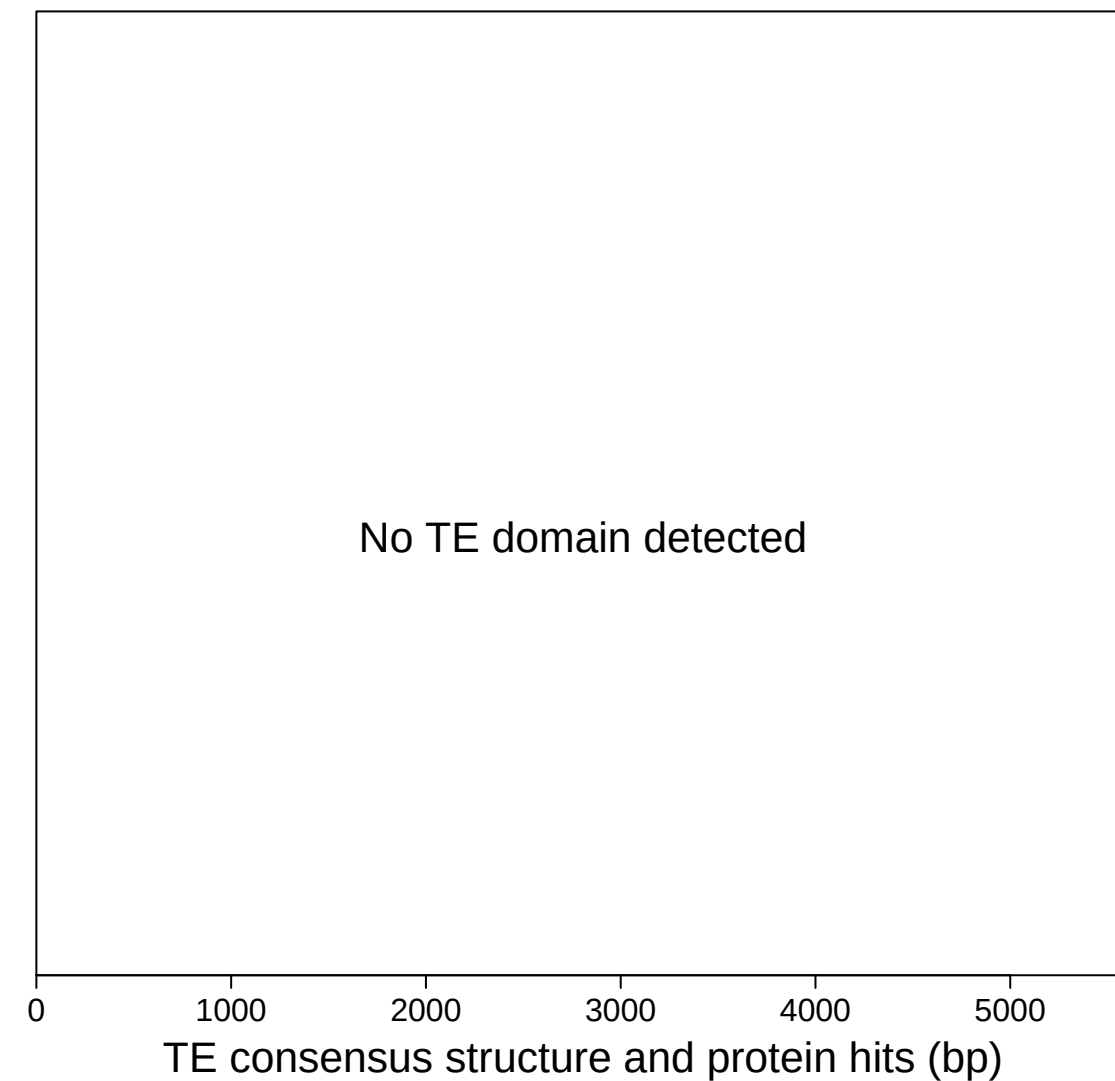
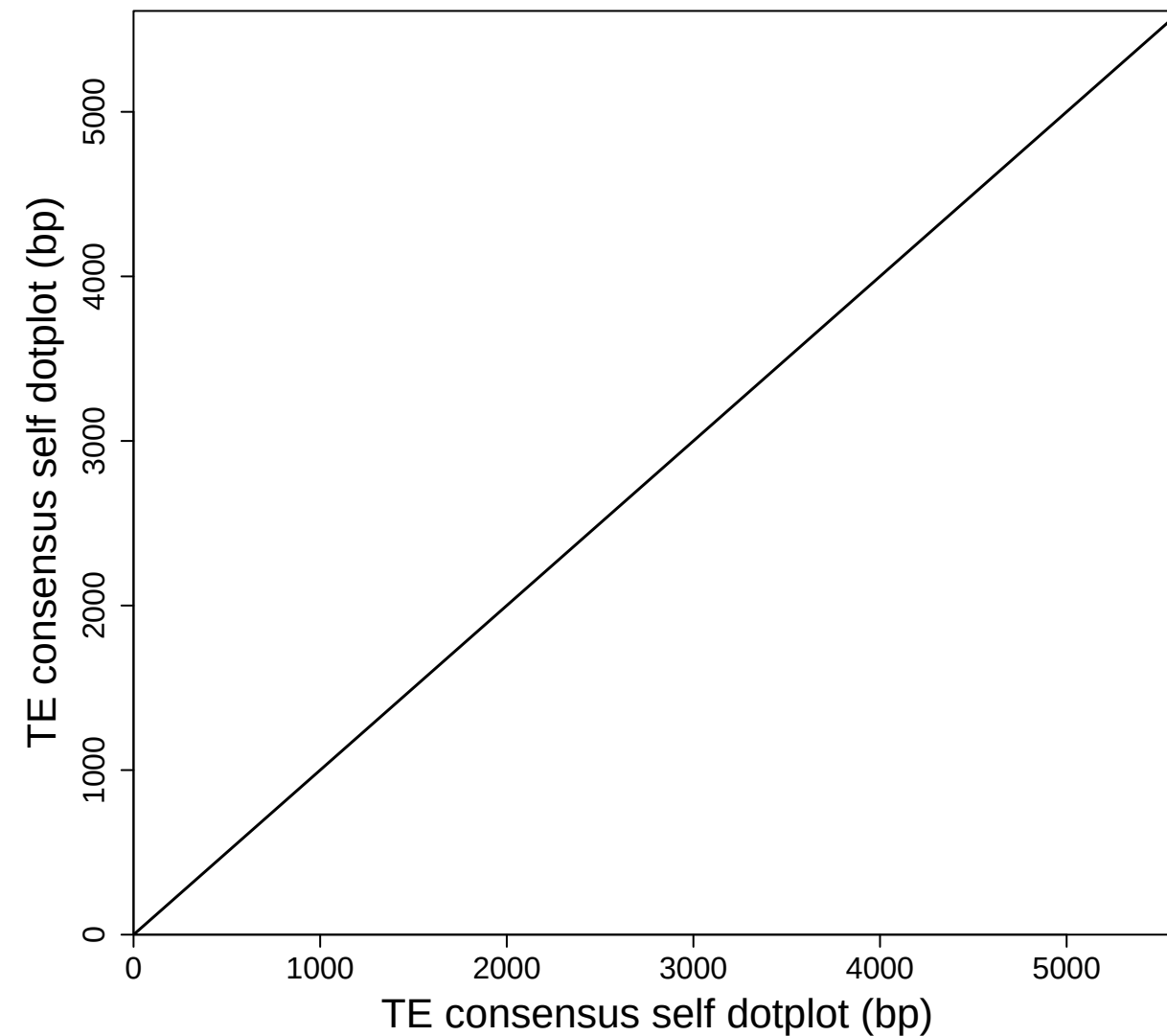
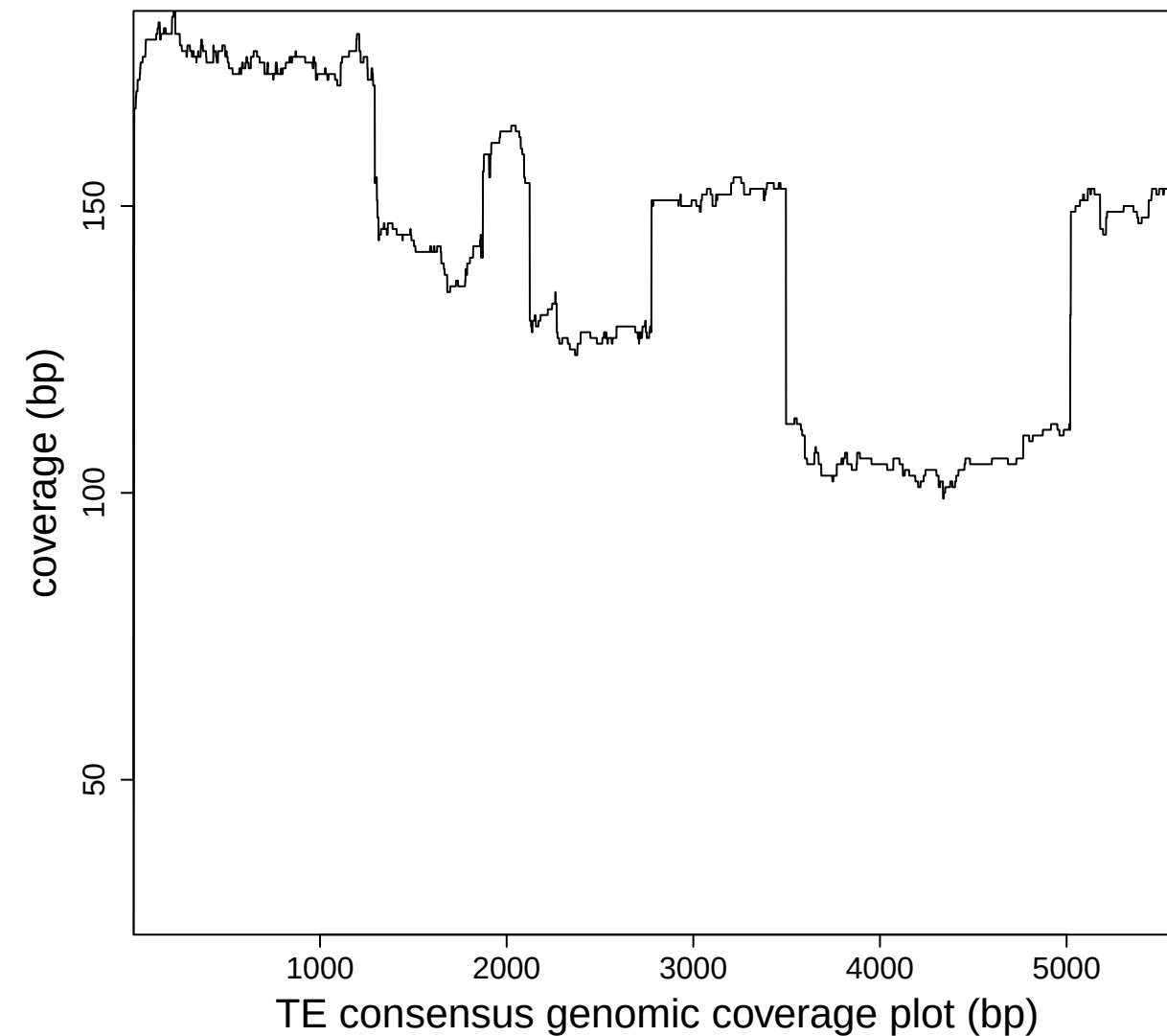
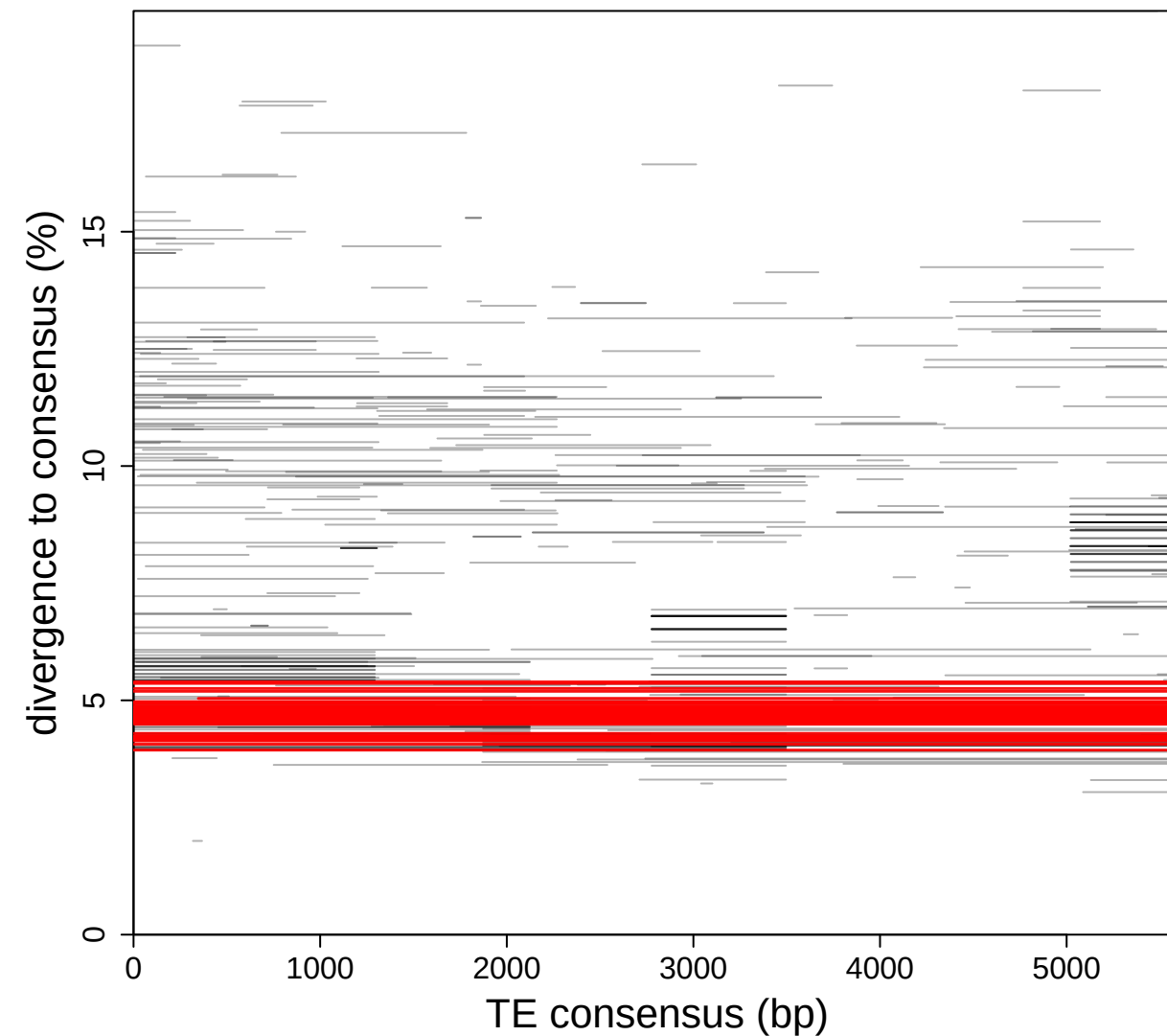
2267.fasta.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa
size: 7841bp; fragments: 798; full length: 50 (≥ 7056.9 bp)

After TEtrimmer Extended plot Blue lines are boundaries

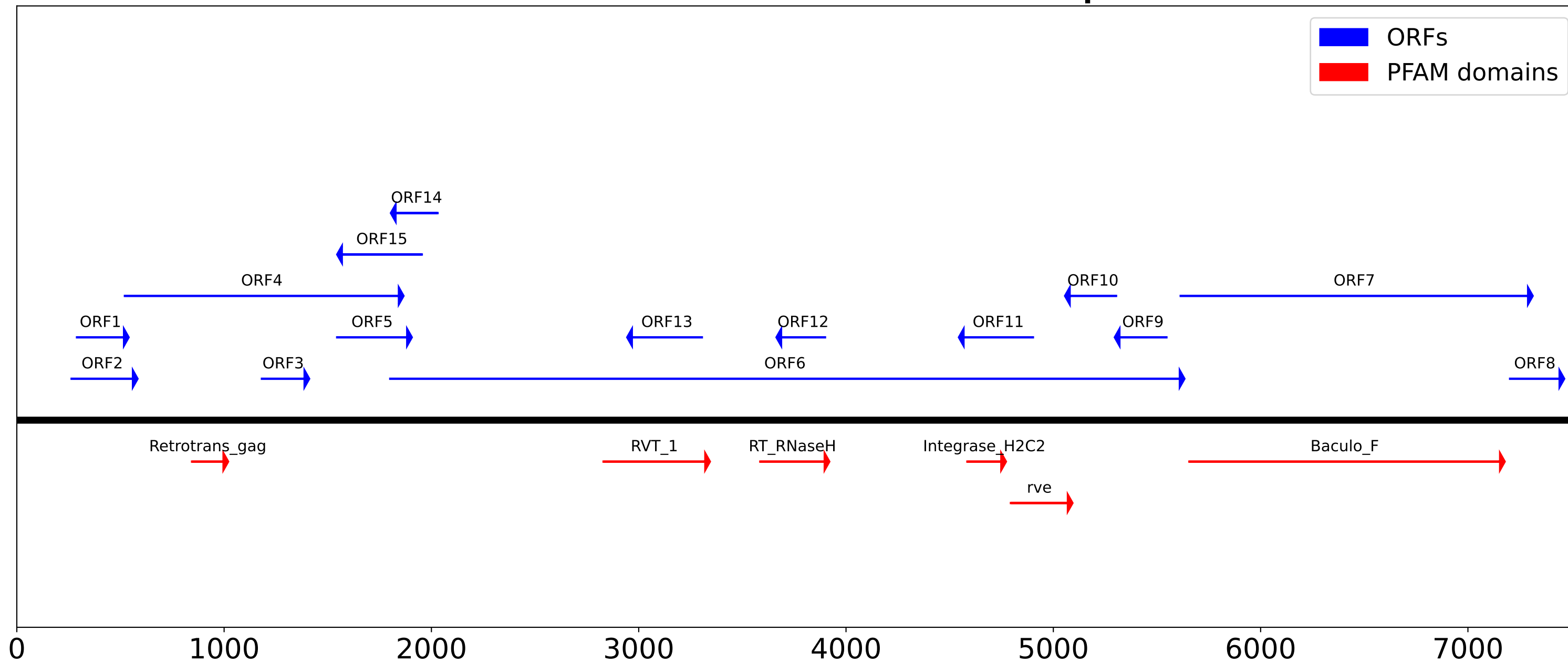


TE: rnd_5_family_2267
size: 5613bp; fragments: 466; full length: 56 (≥ 5051.7 bp)

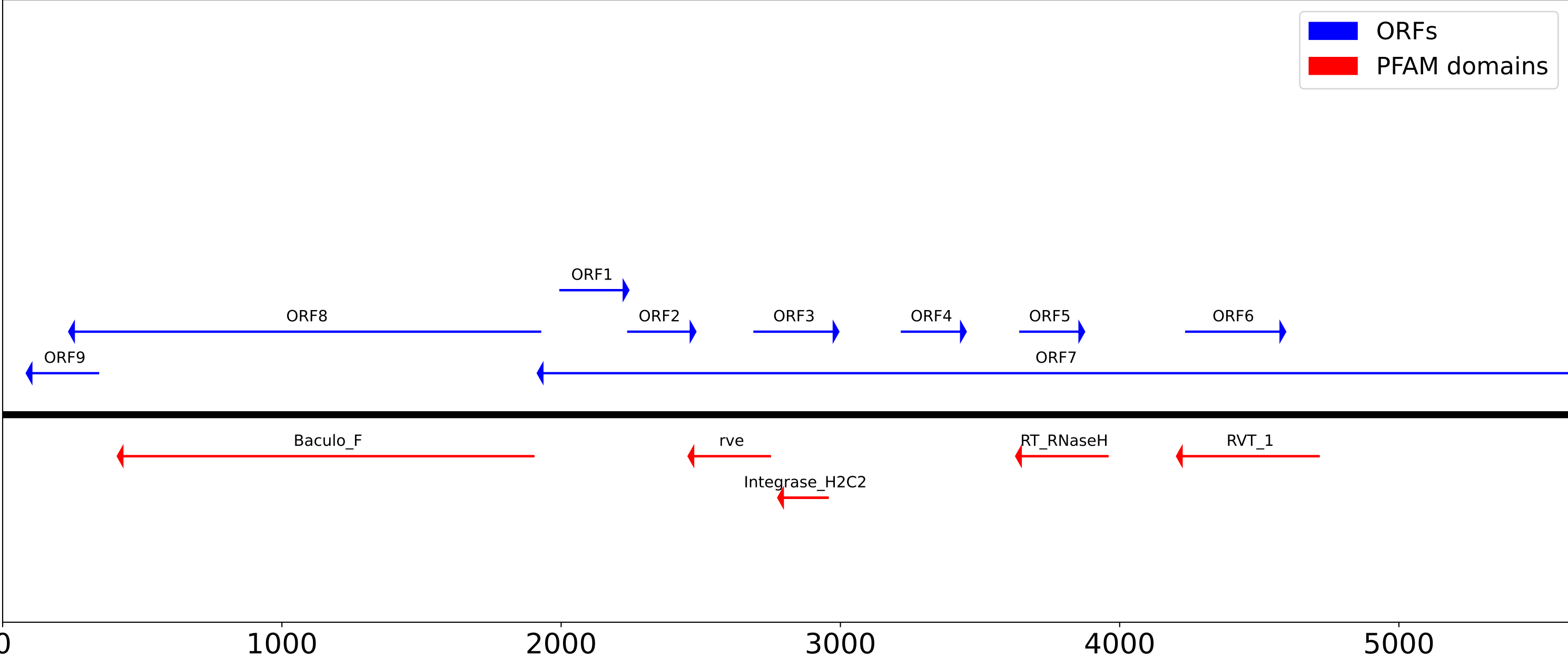
Before TEtrimmer 5613 bp



After TEtrimmer ORF and PFAM domain plot



Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

