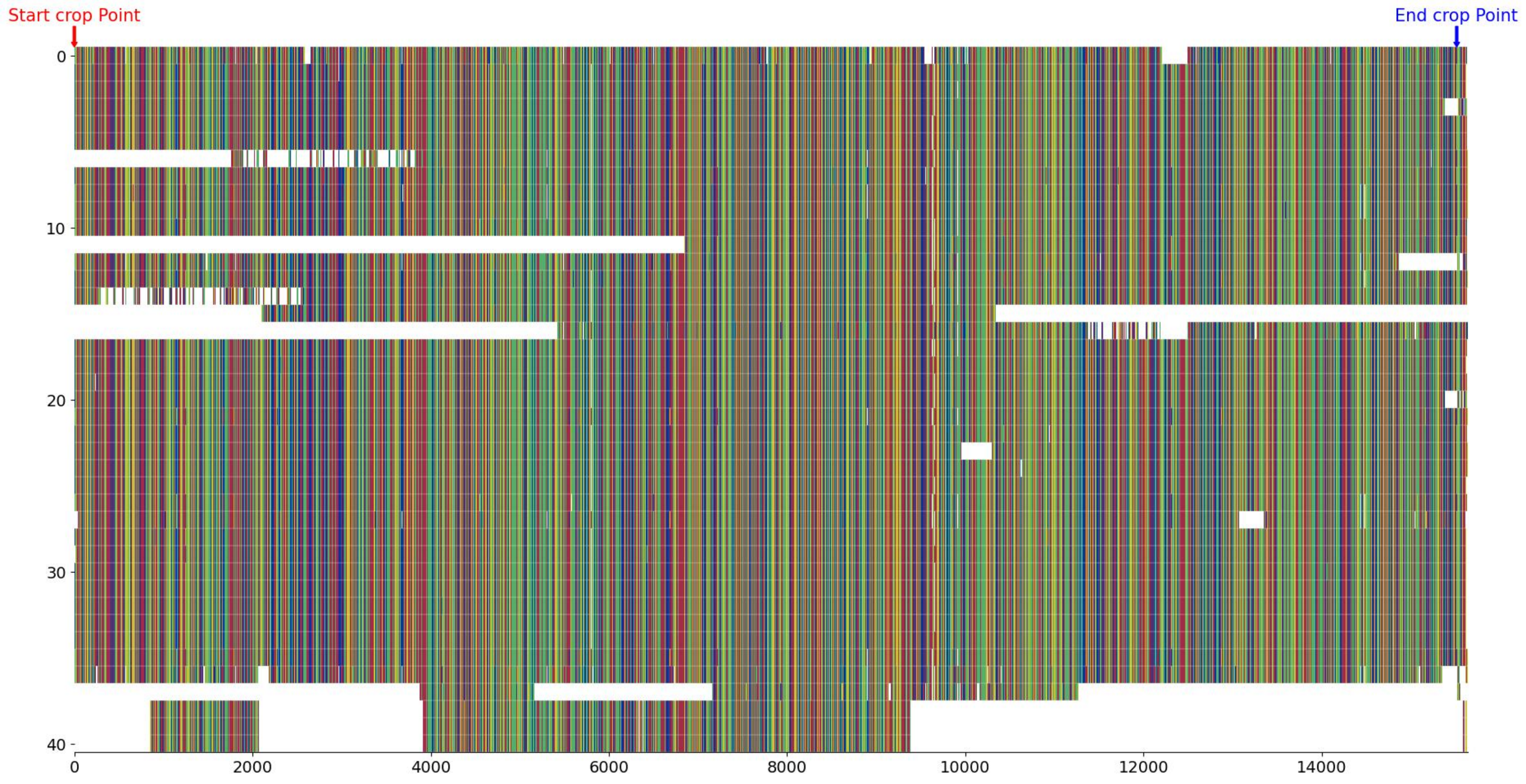
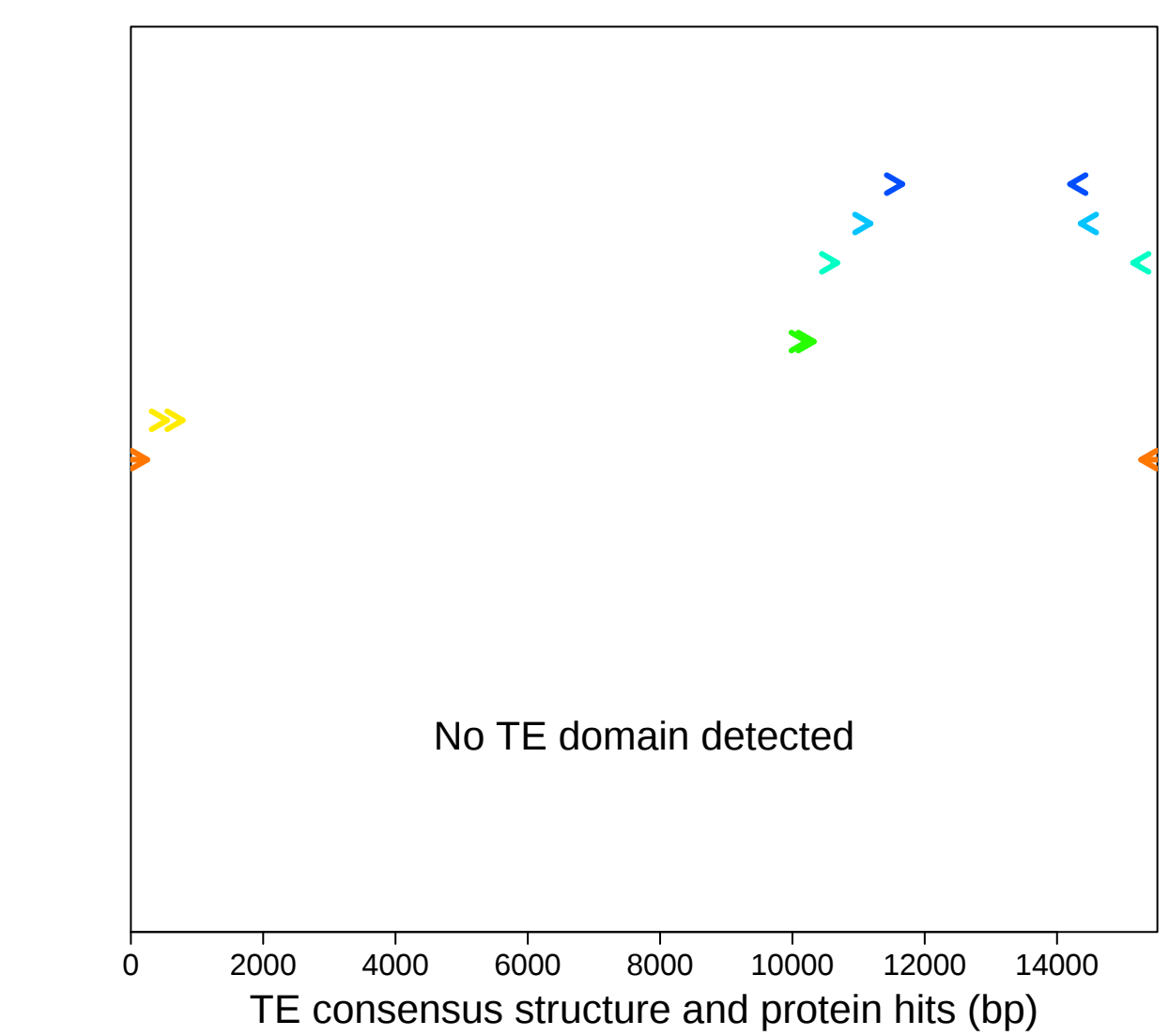
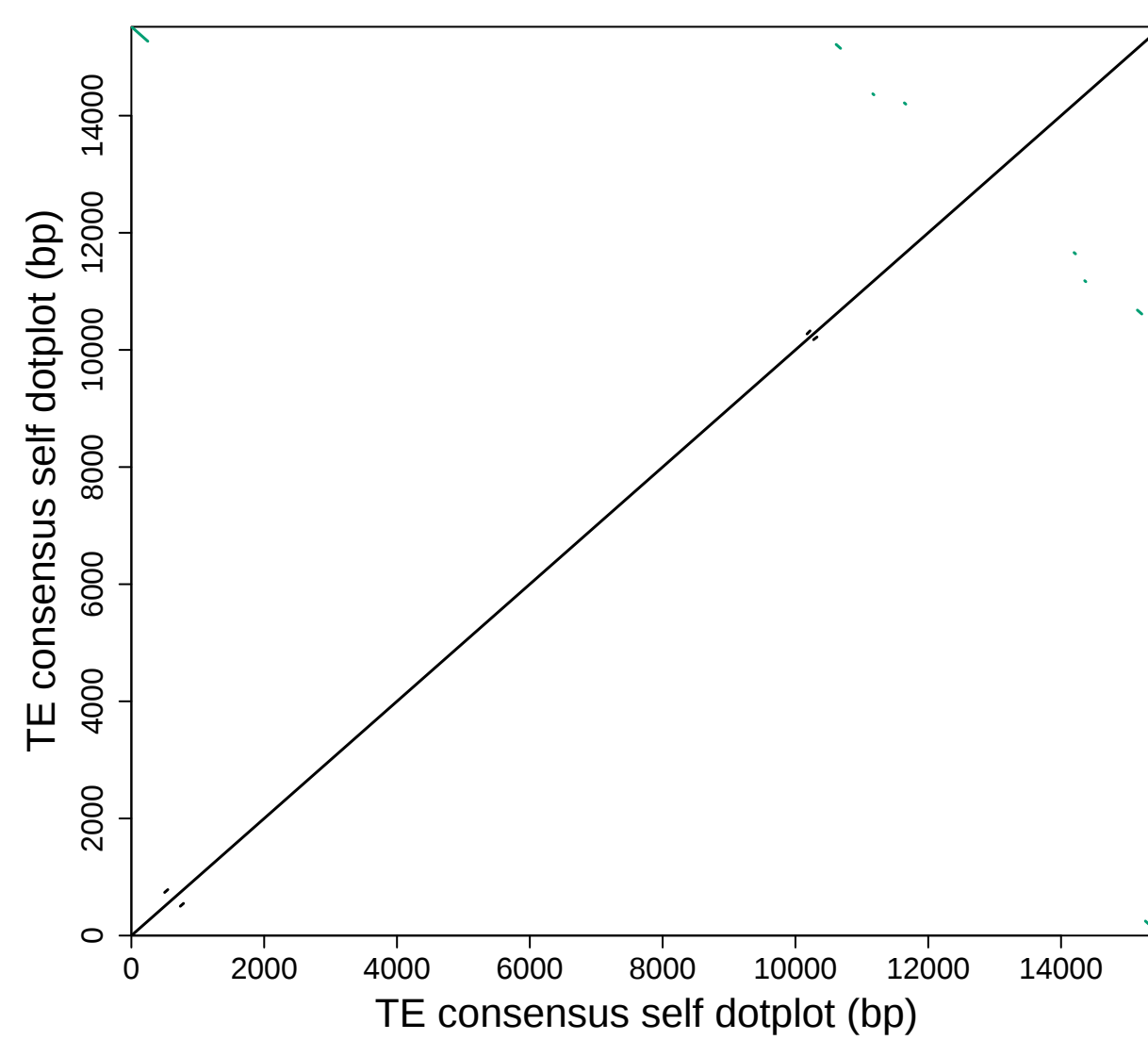
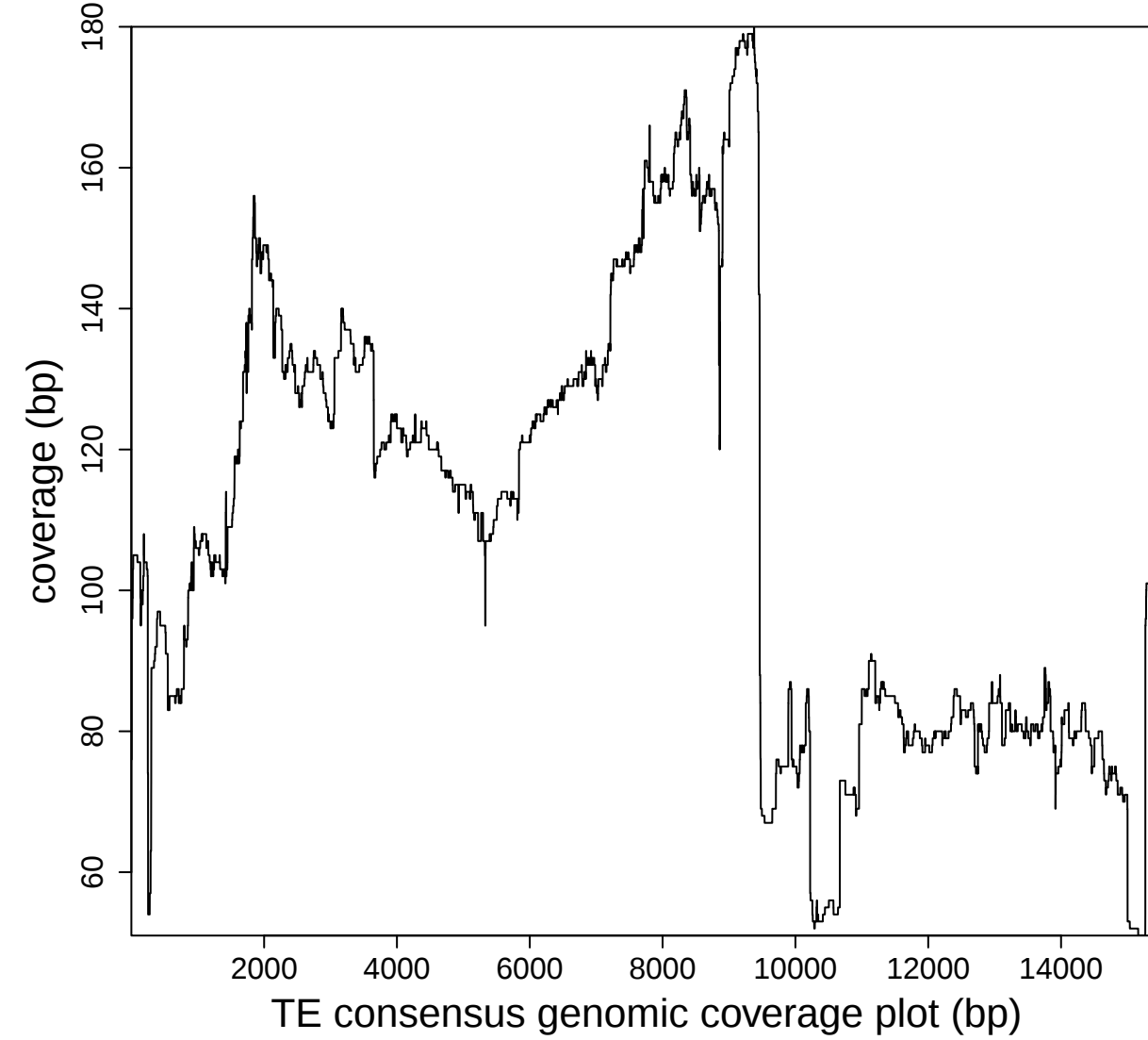
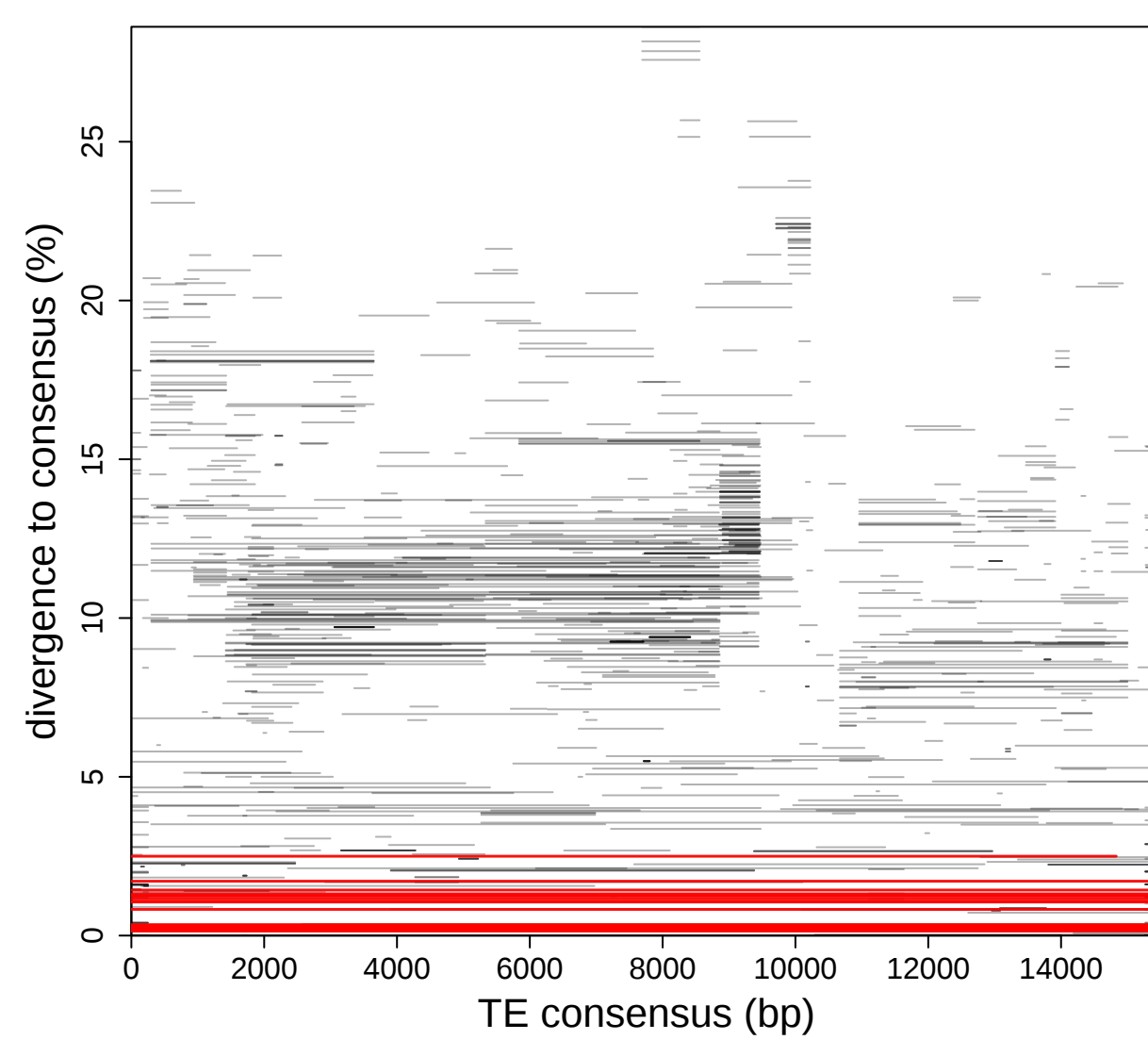


[illegible]



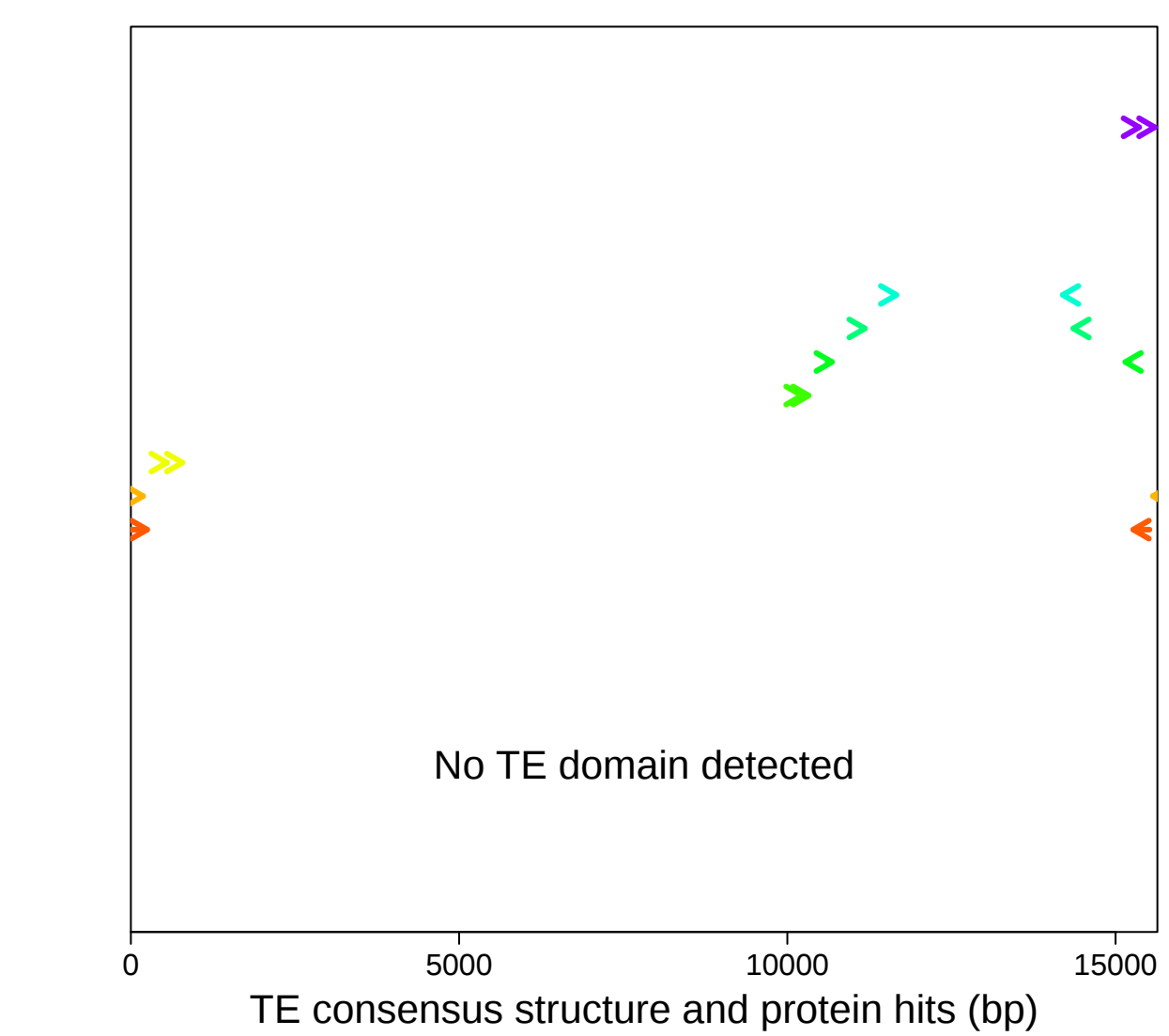
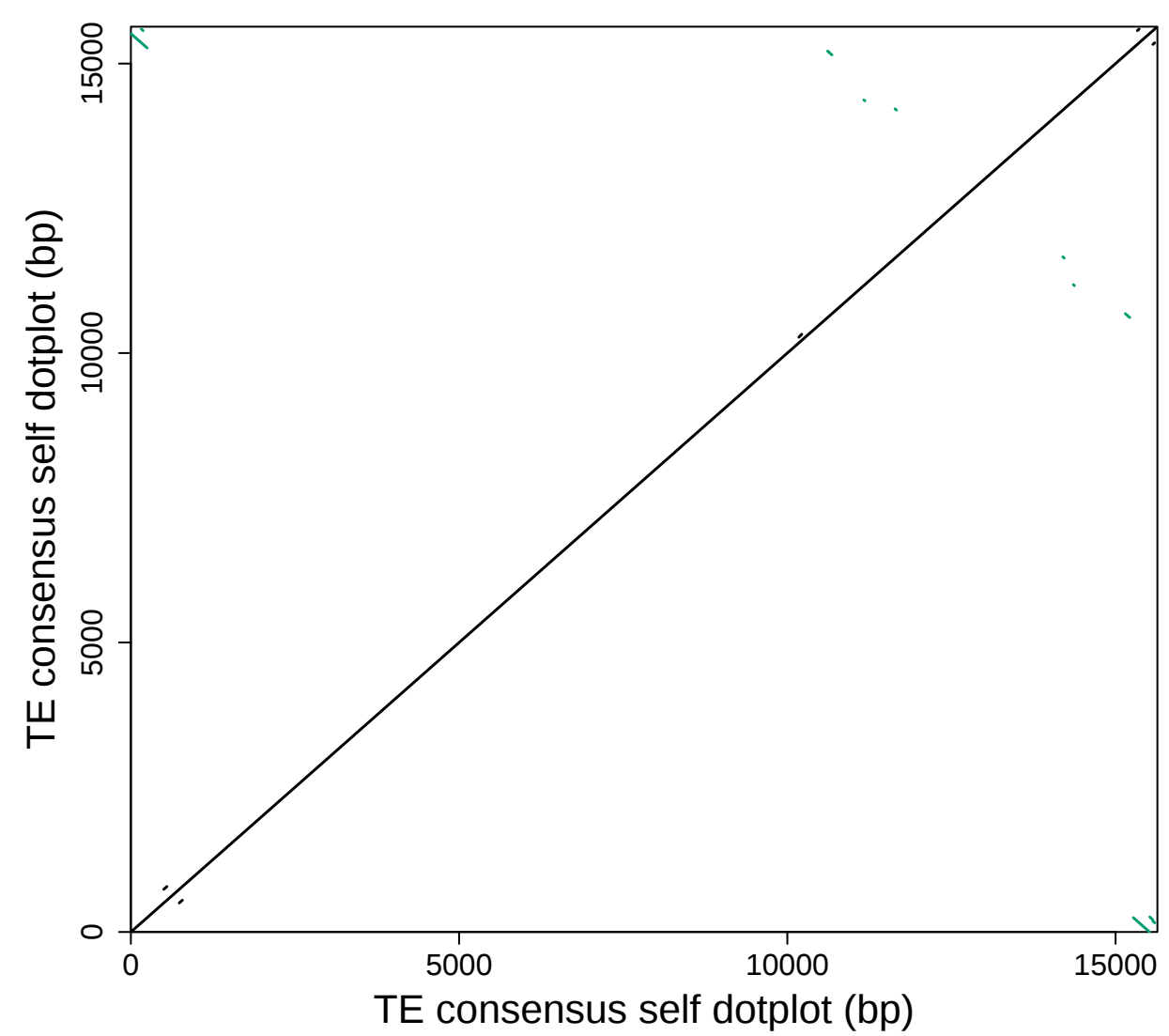
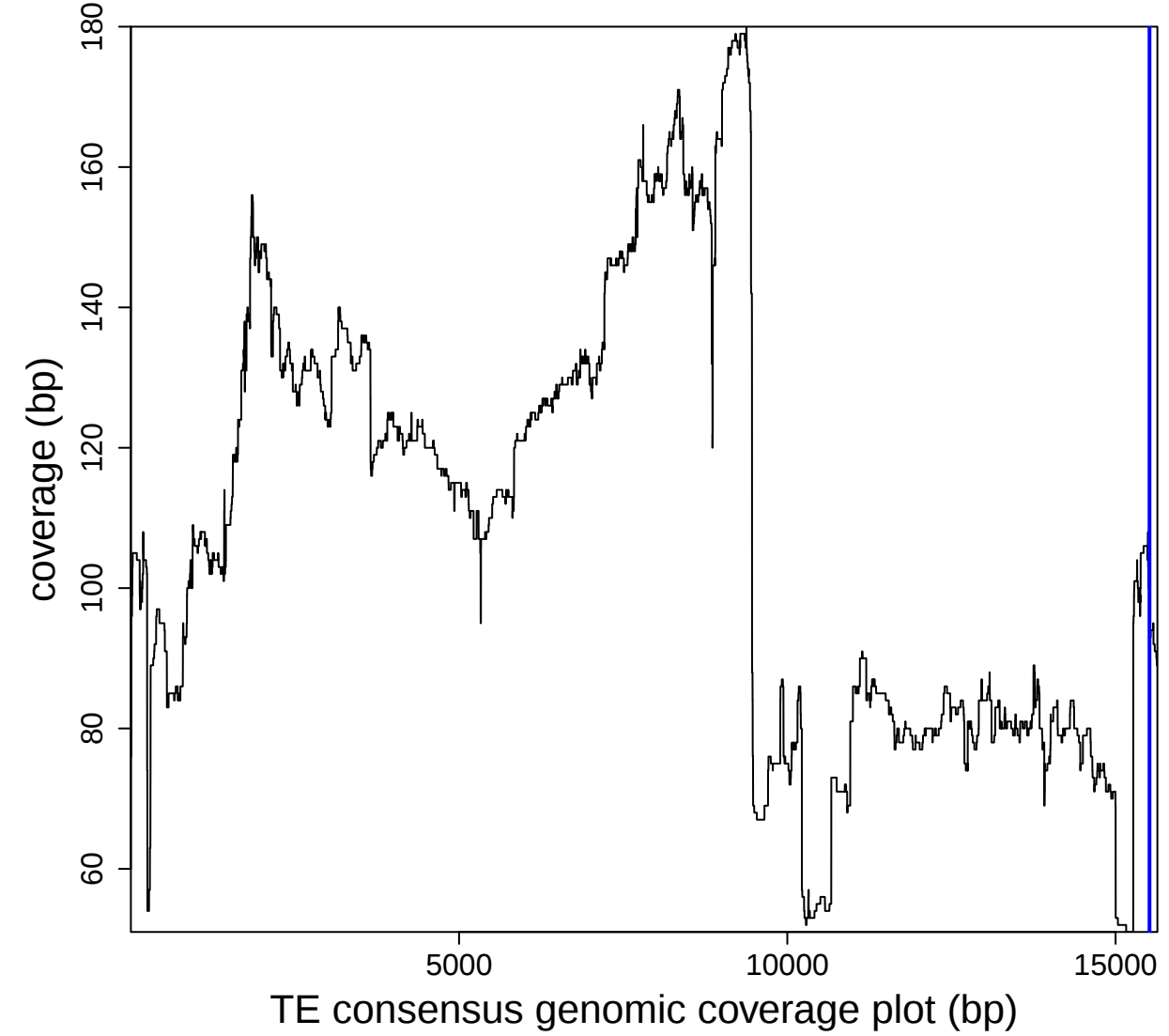
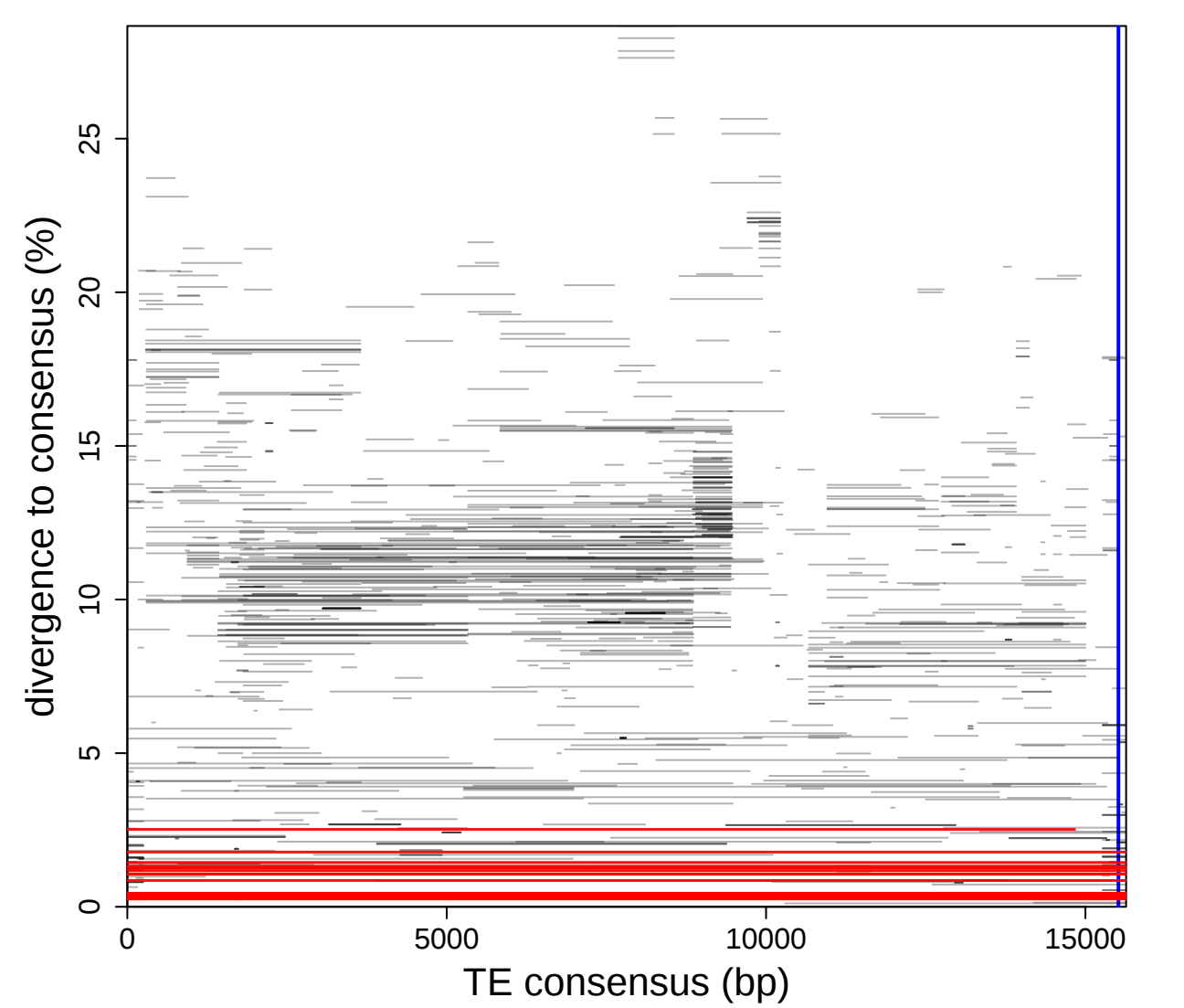
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 15517bp; fragments: 1002; full length: 29 (>=13965.3bp)

After TEtrimmer 15517 bp



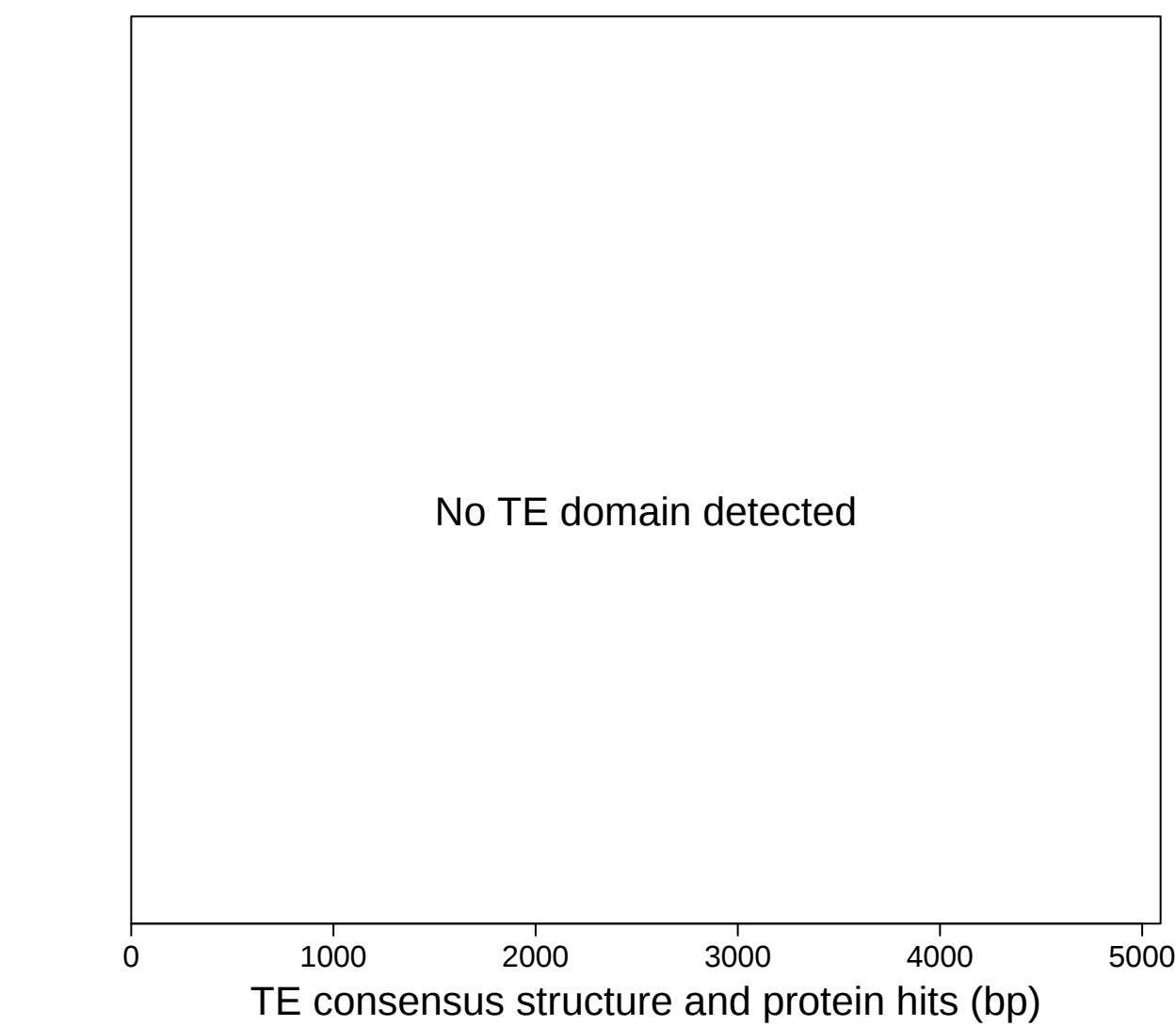
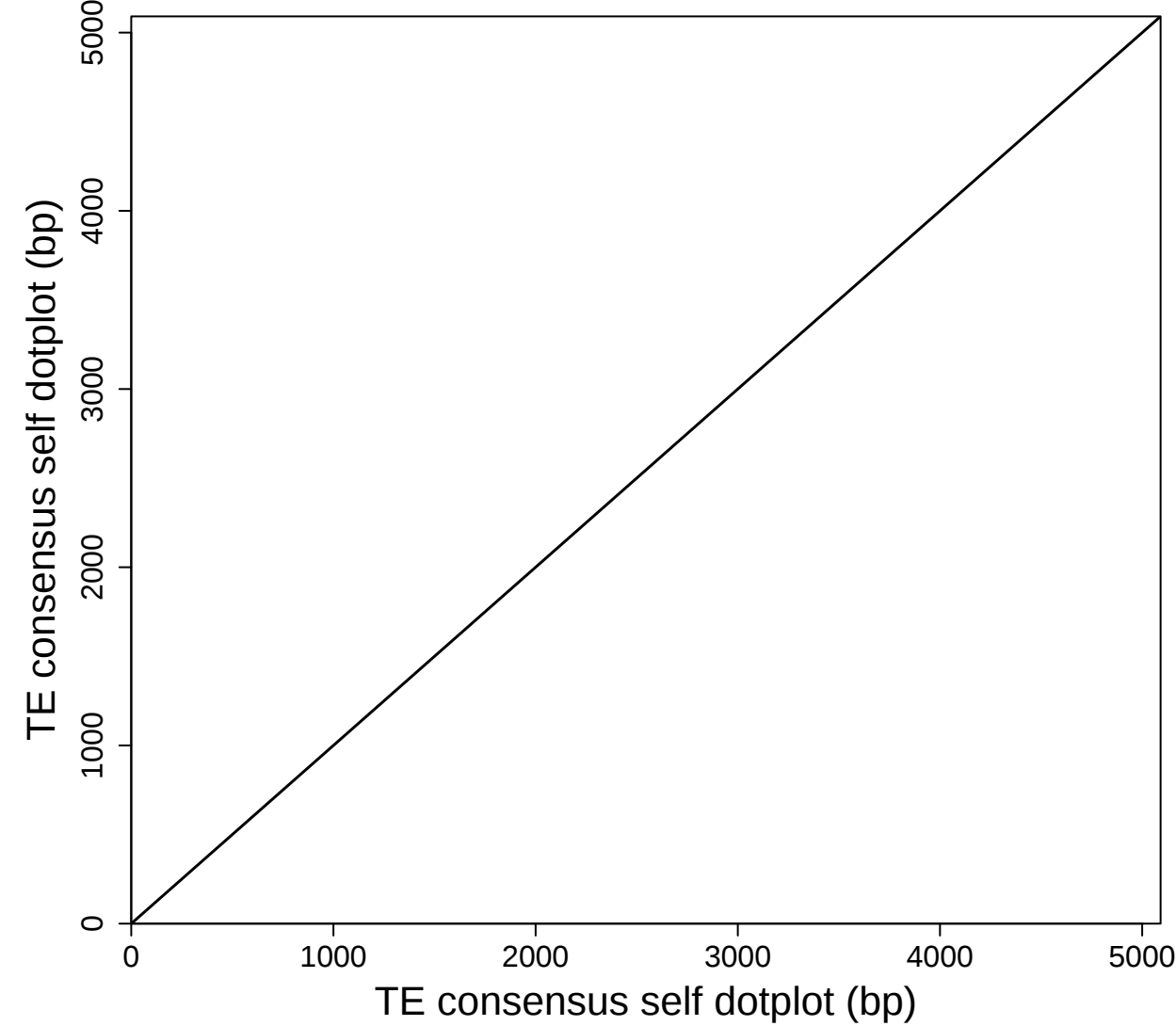
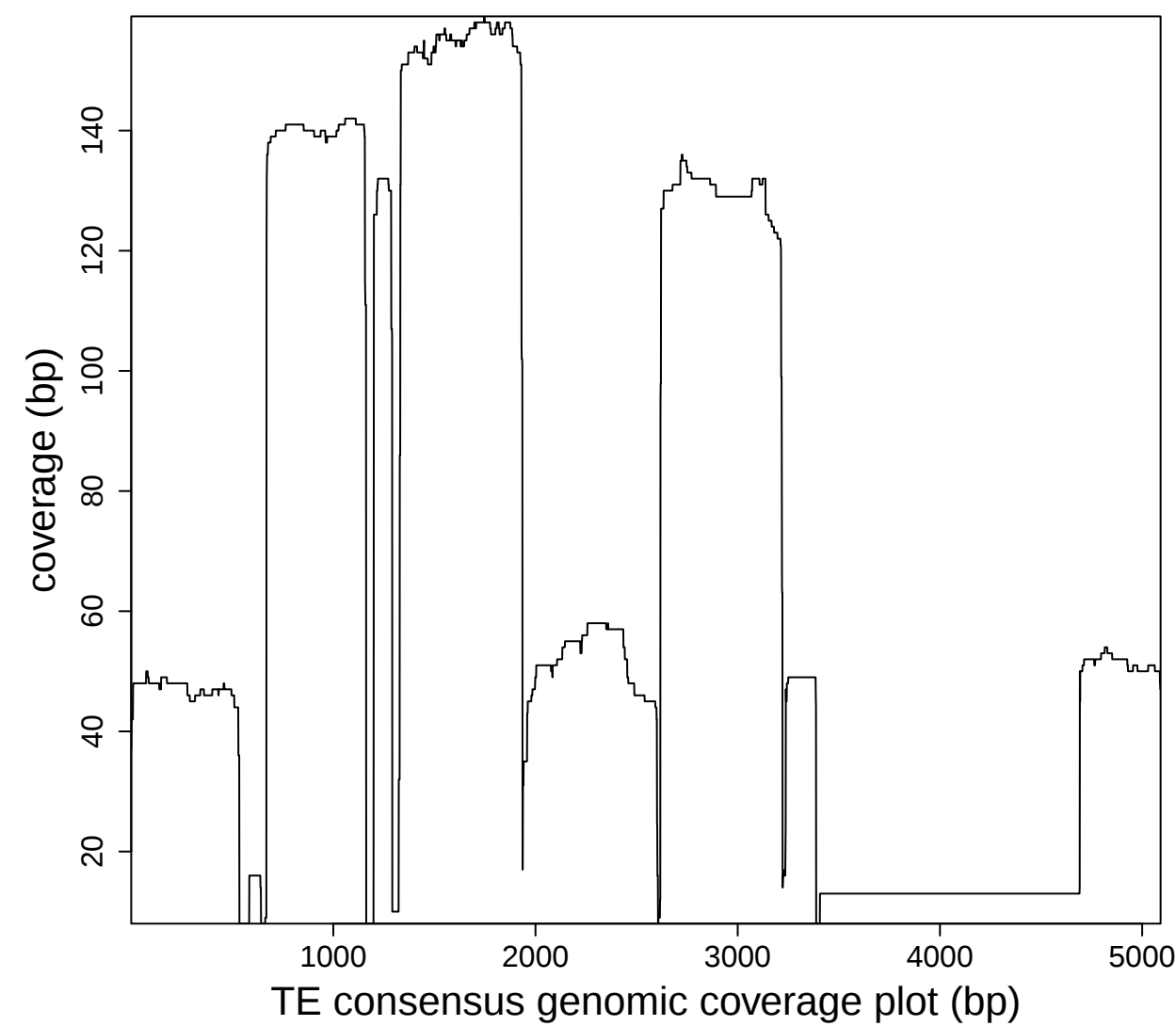
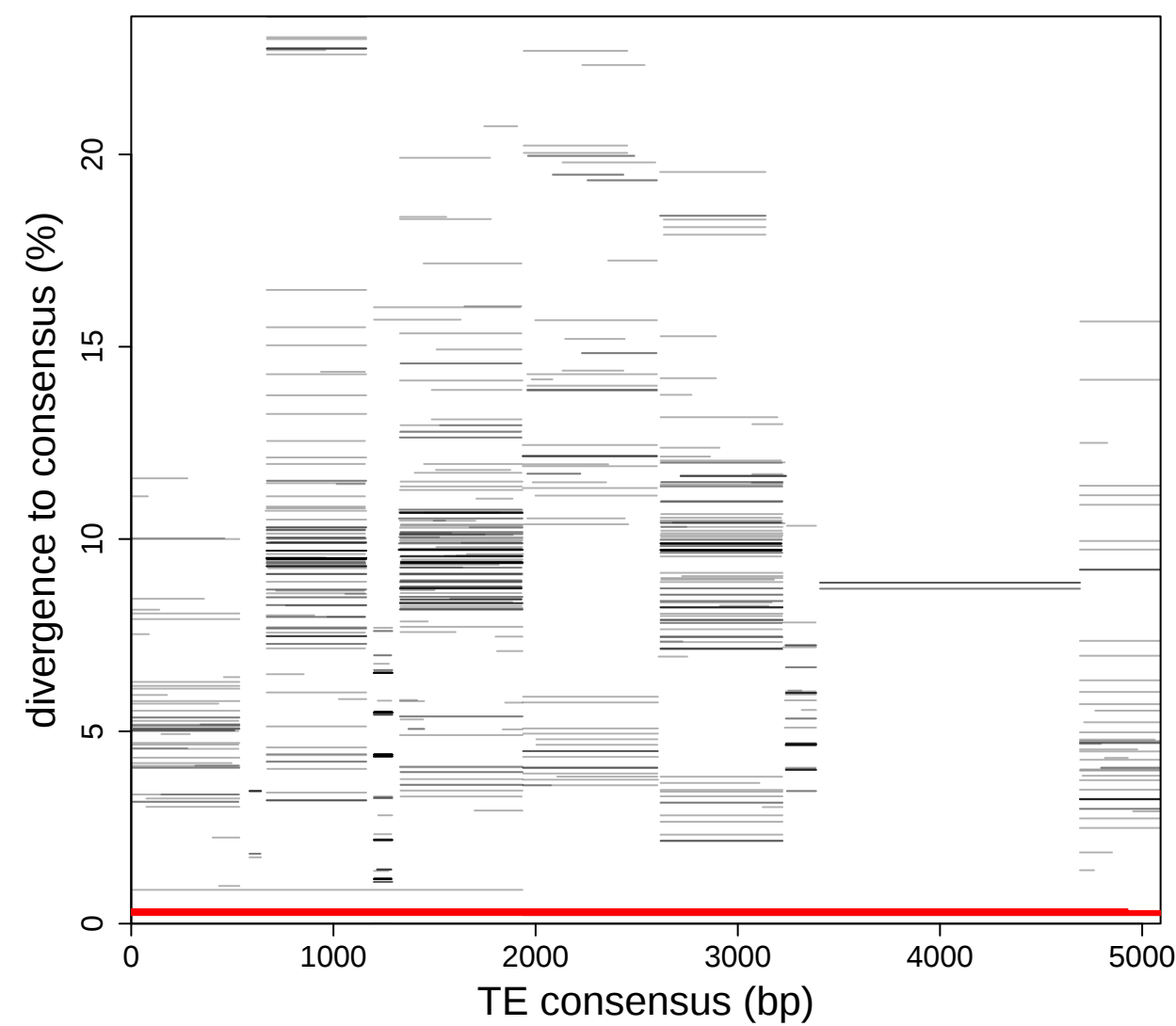
ly_7362.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.
size: 15637bp; fragments: 1015; full length: 29 (>=14073.3bp)

After TEtrimmer Extended plot Blue lines are boundaries

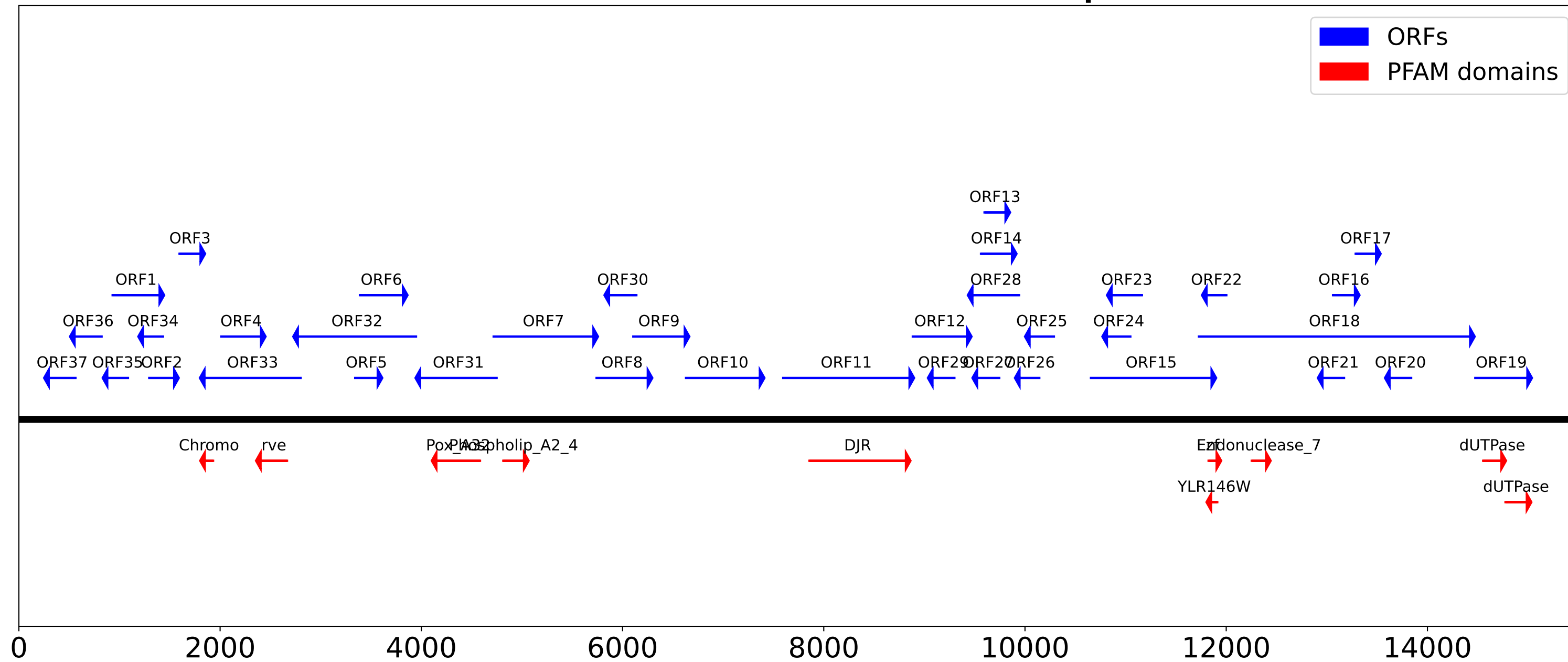


TE: rnd_5_family_7362
size: 5091bp; fragments: 781; full length: 7 (≥ 4581.9 bp)

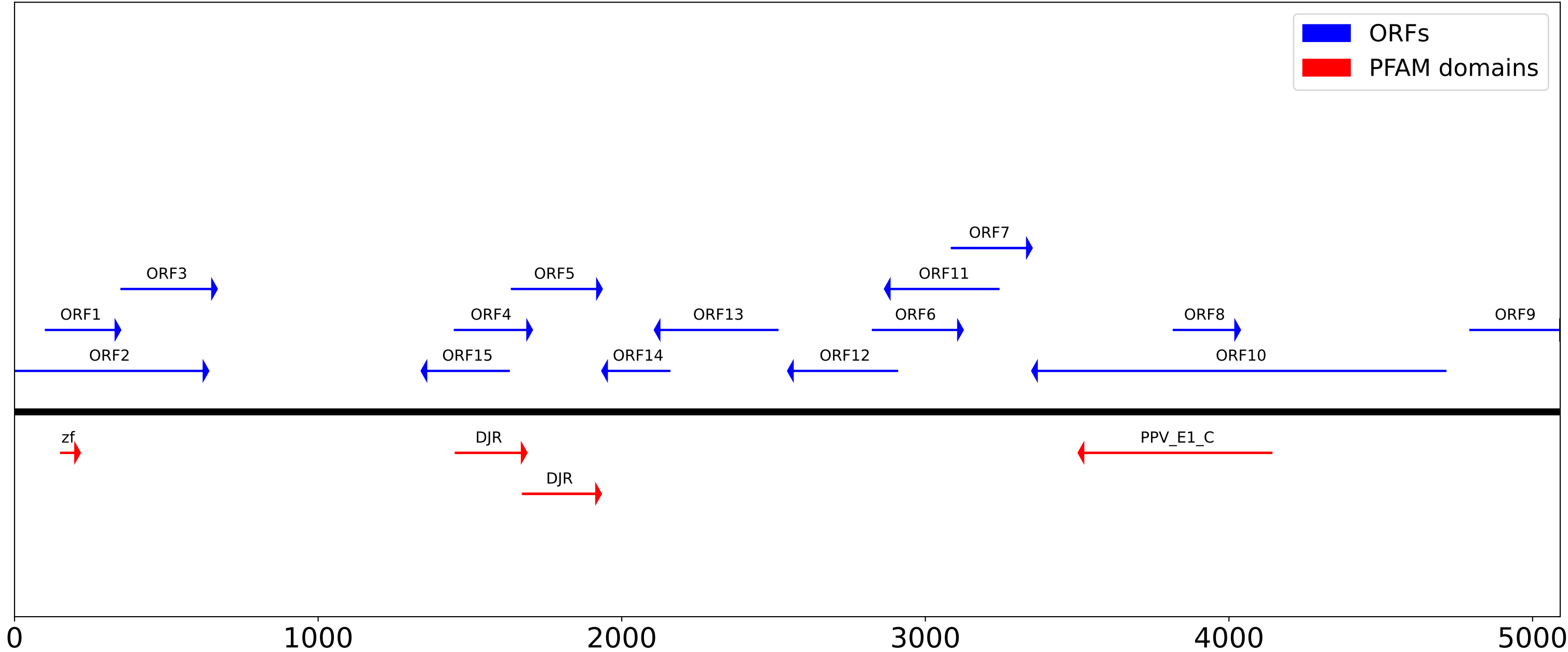
Before TEtrimmer 5091 bp



After TEtrimmer ORF and PFAM domain plot



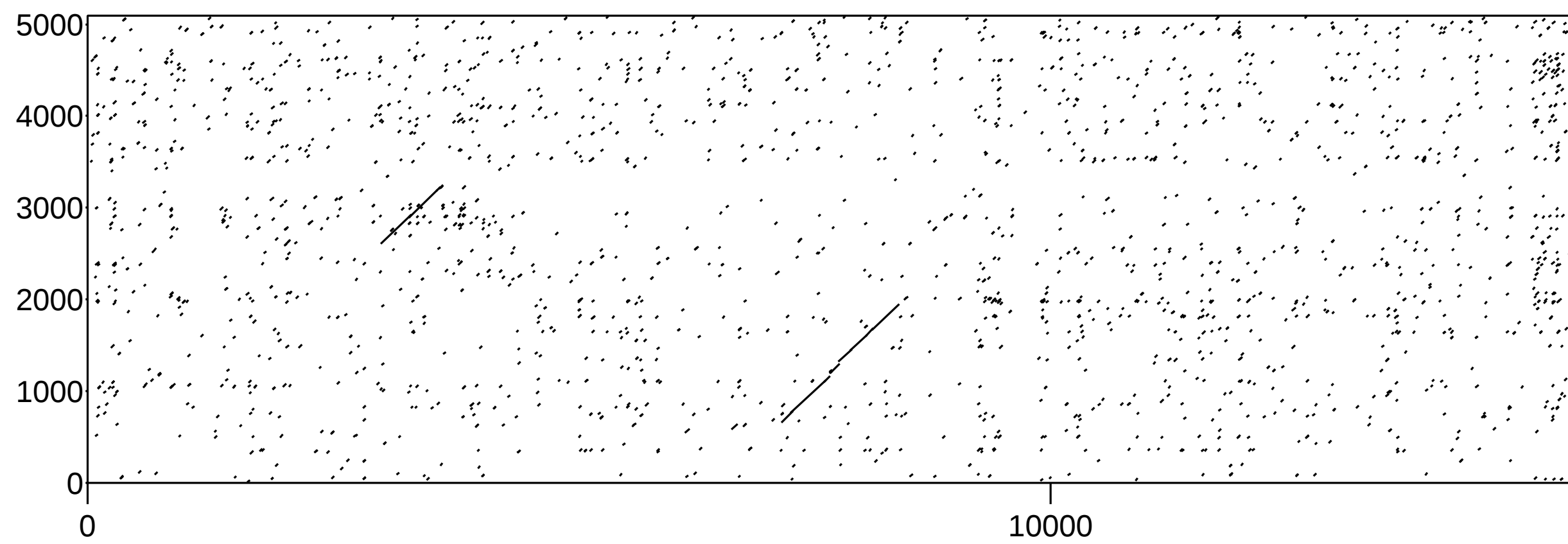
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)