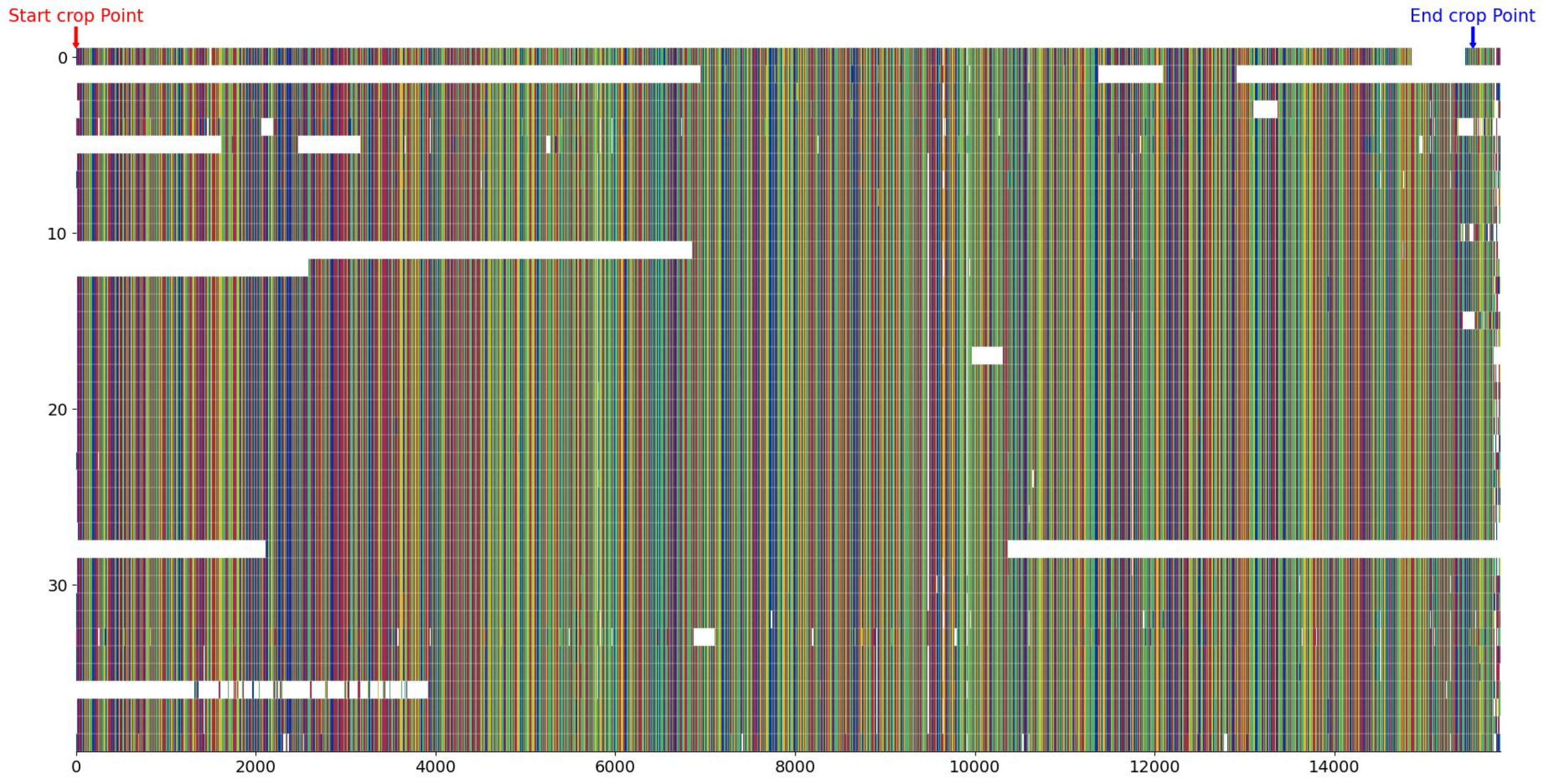
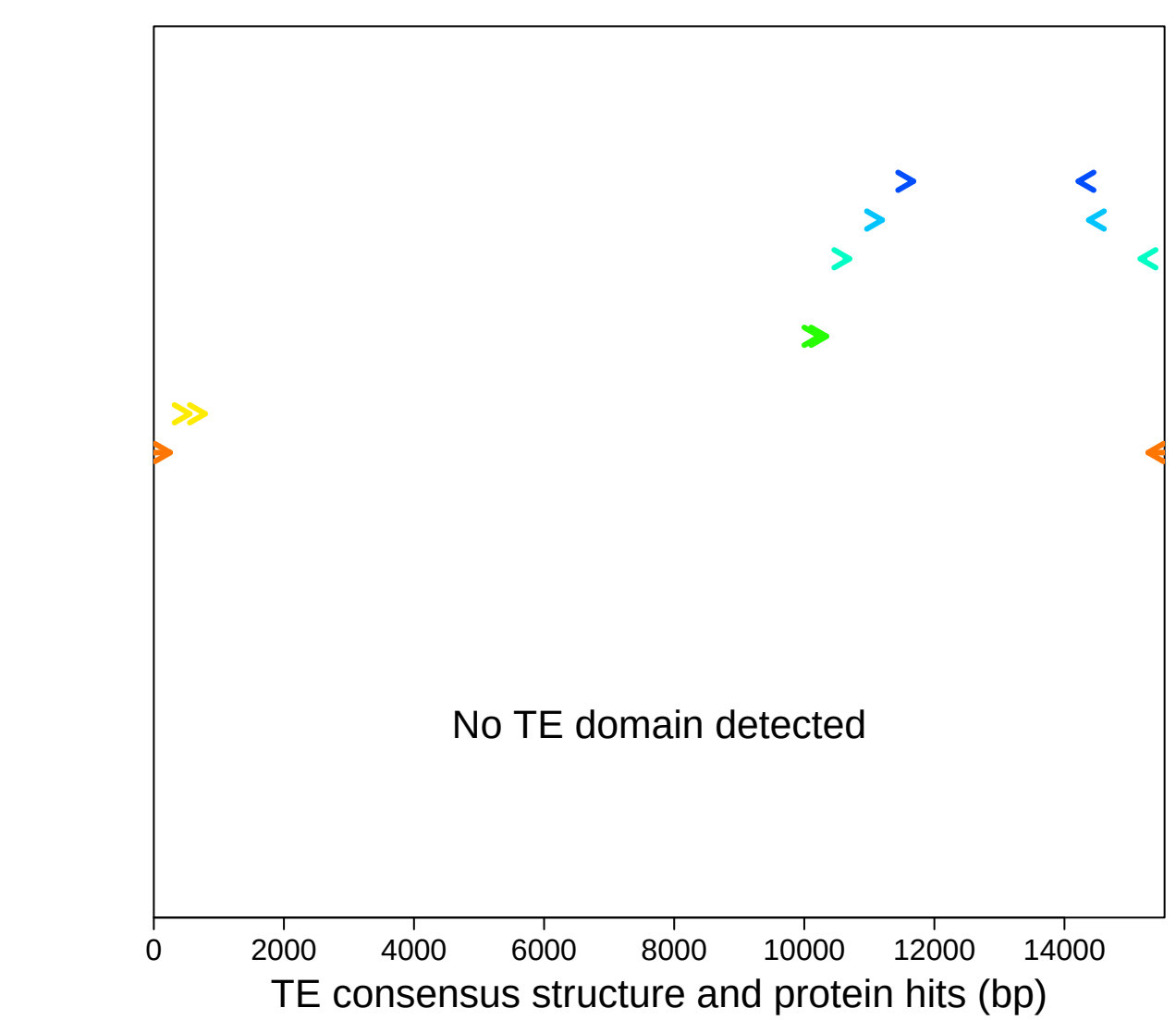
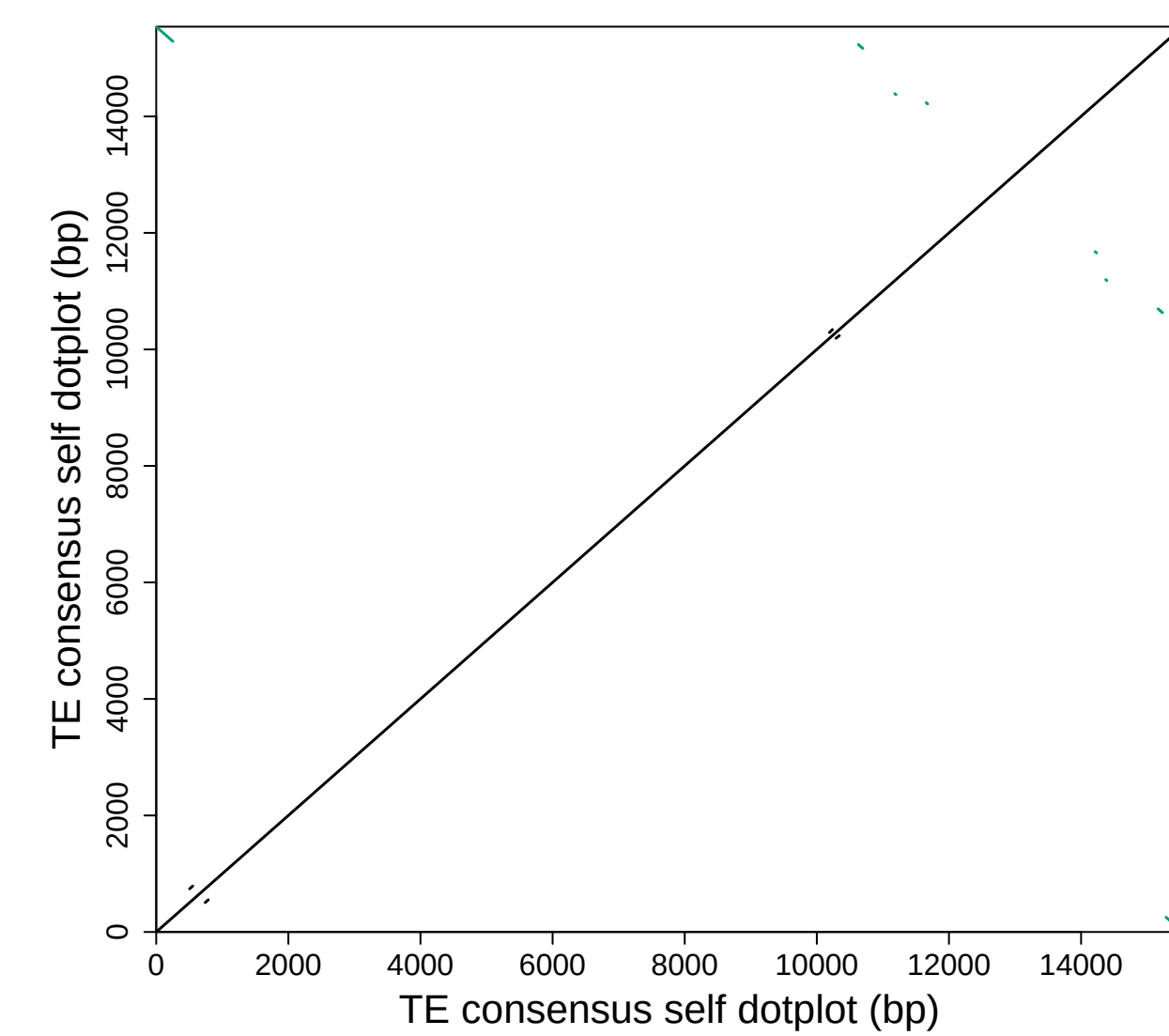
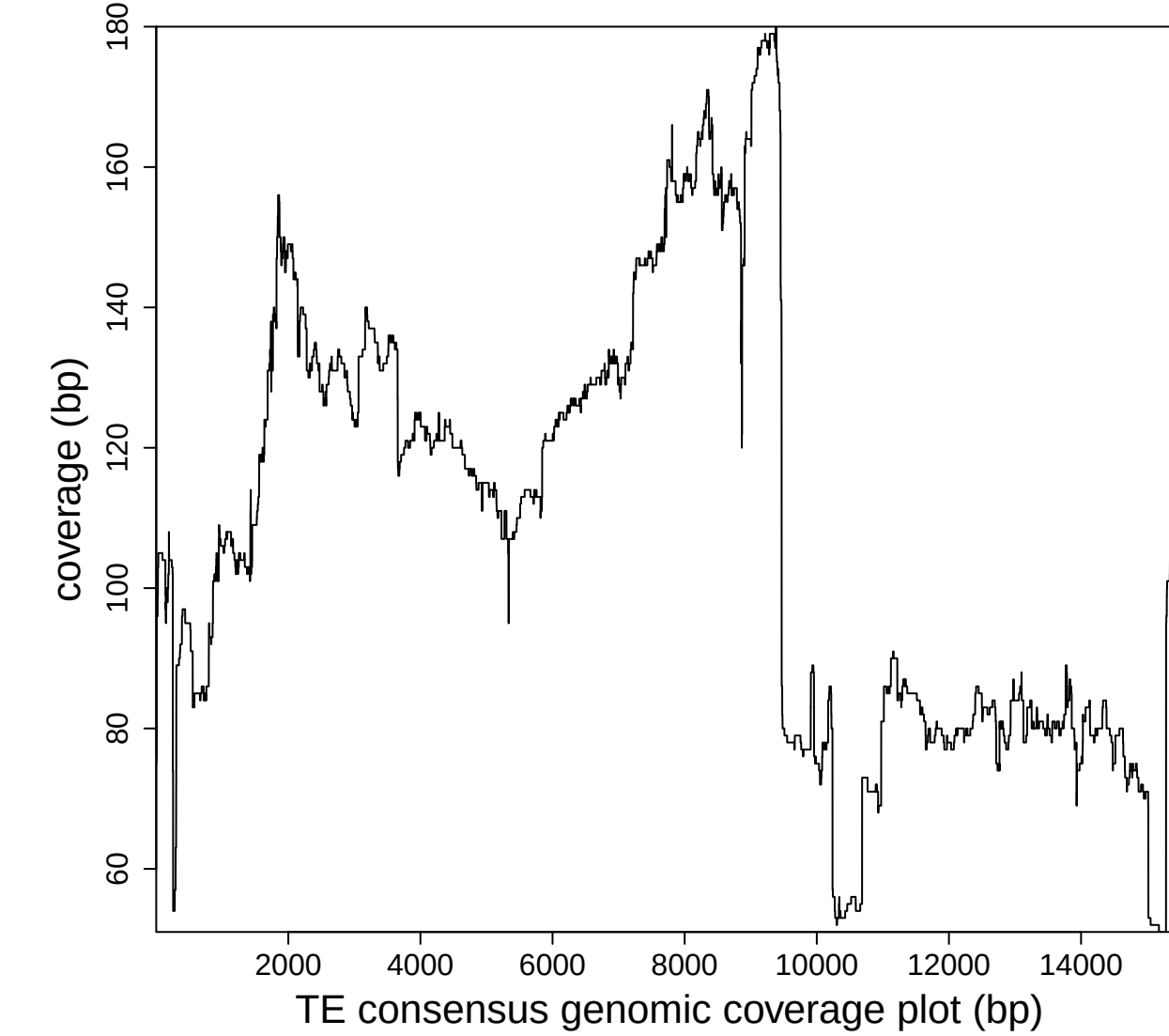
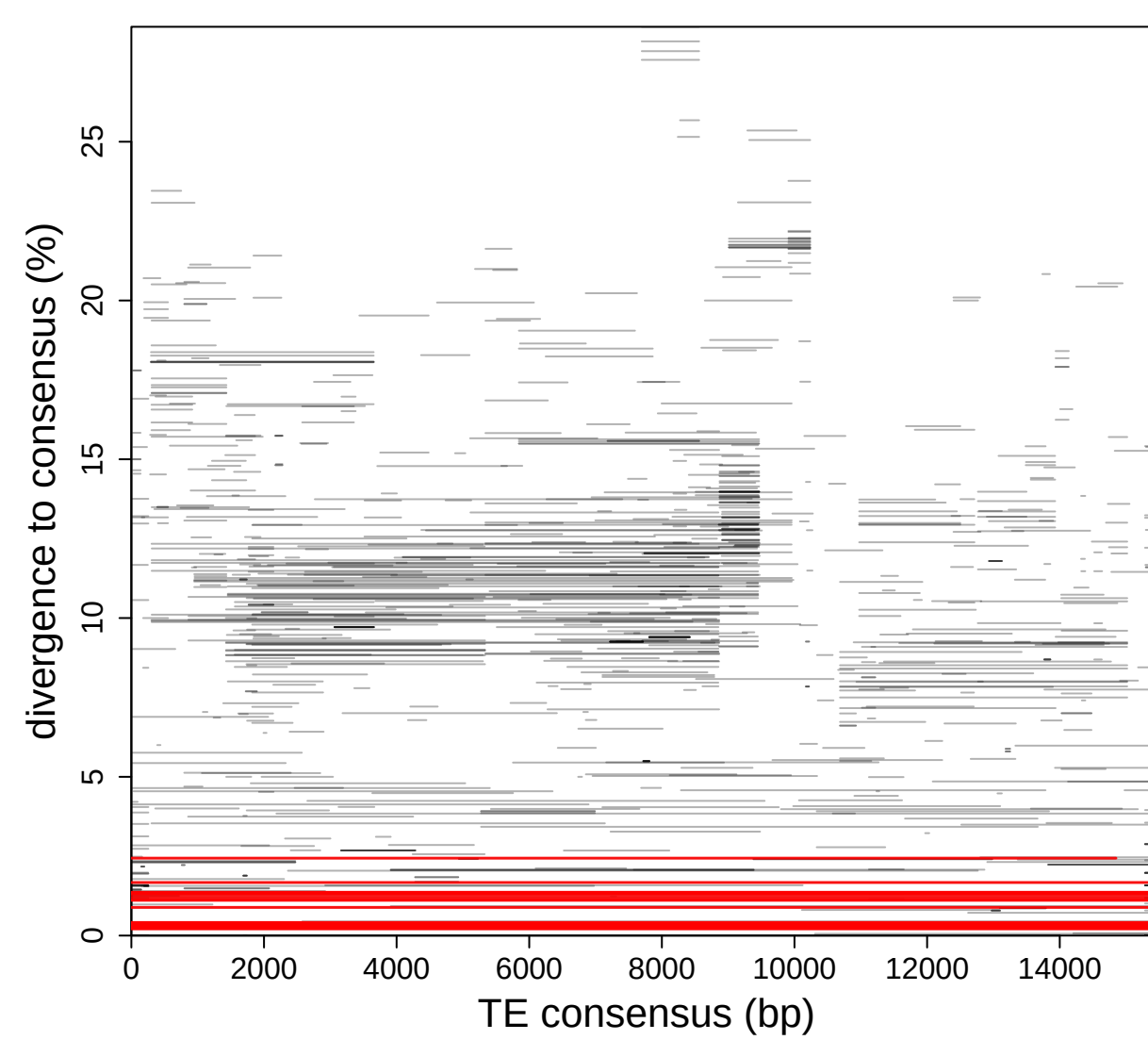


[illegible]



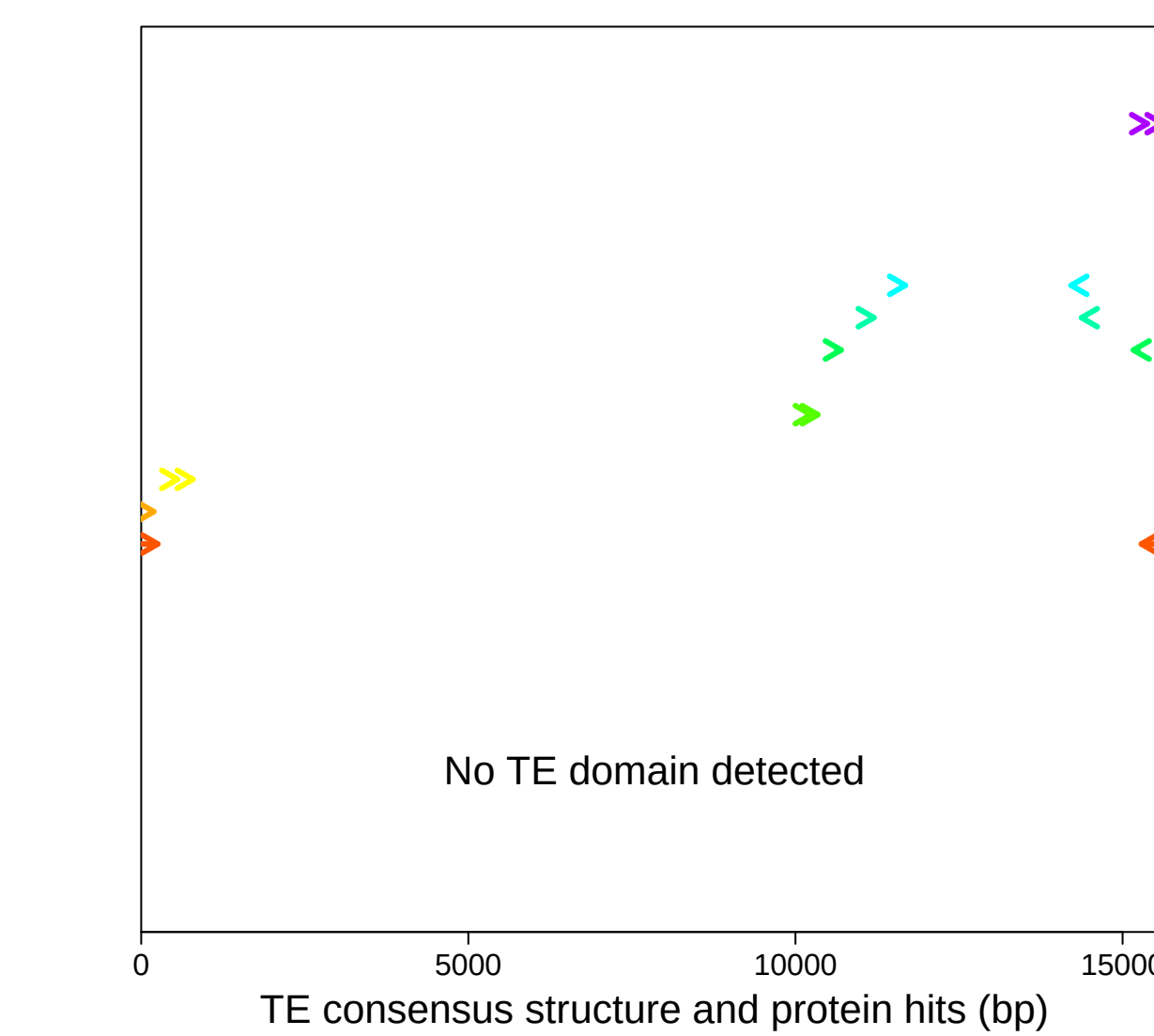
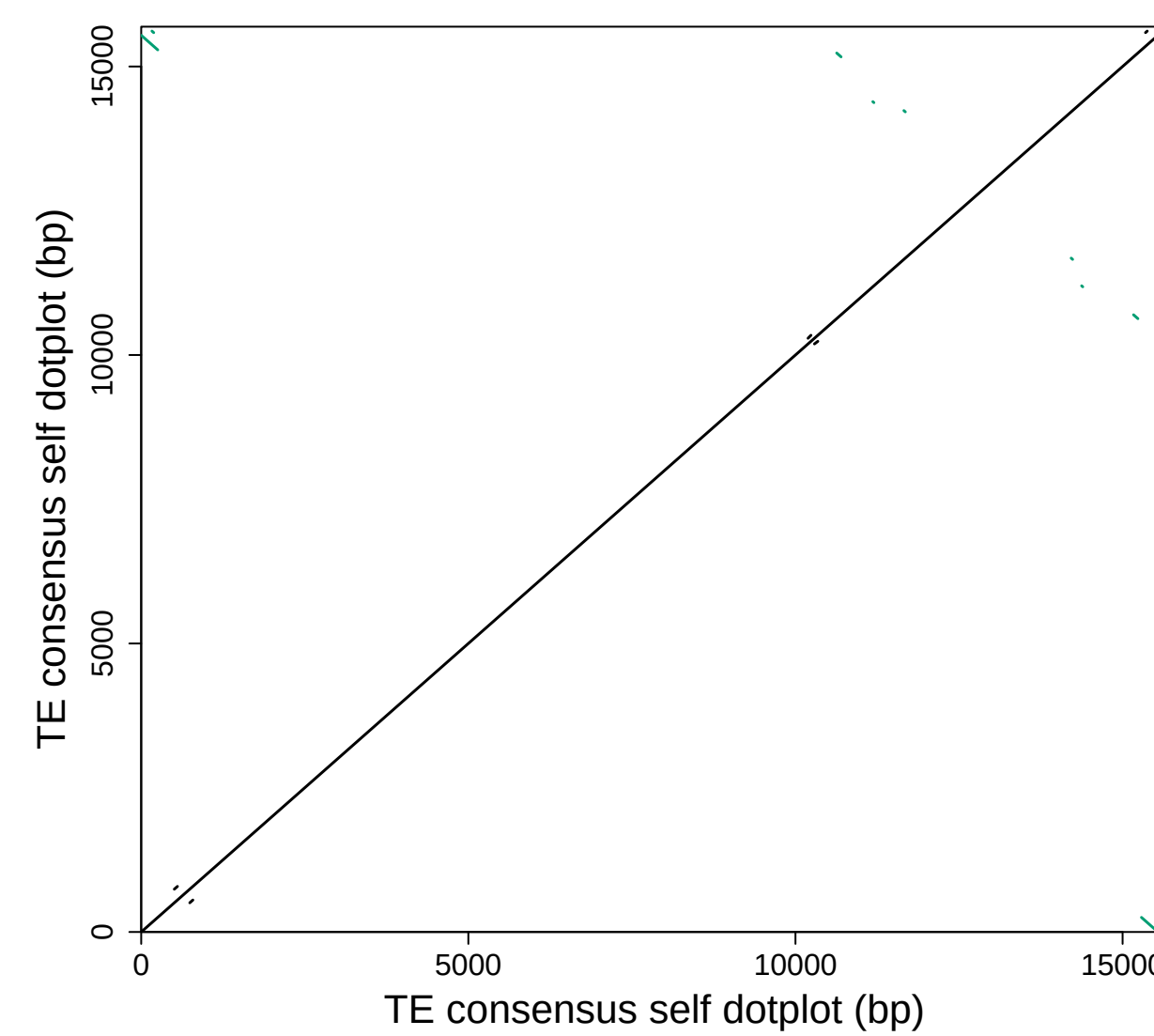
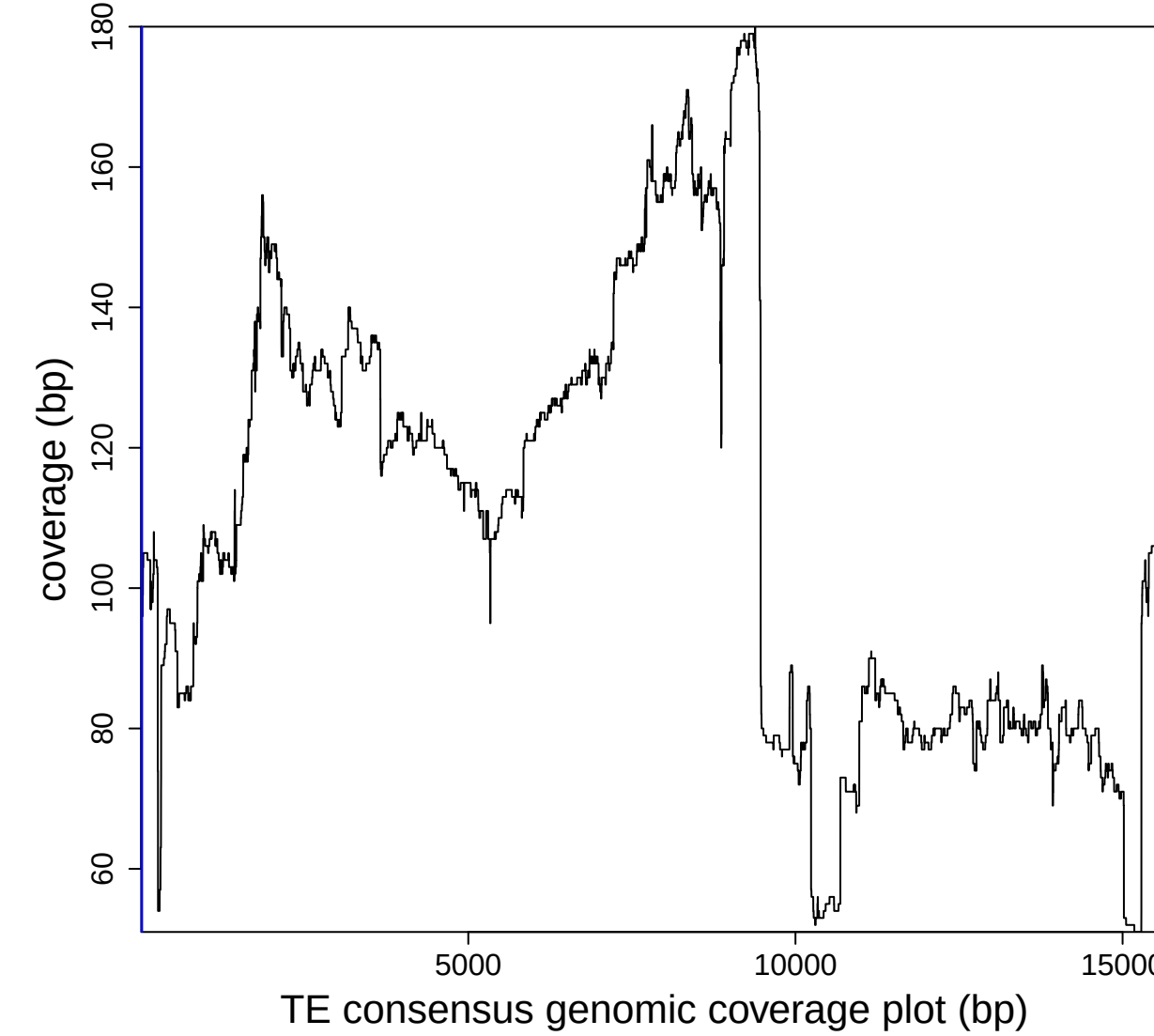
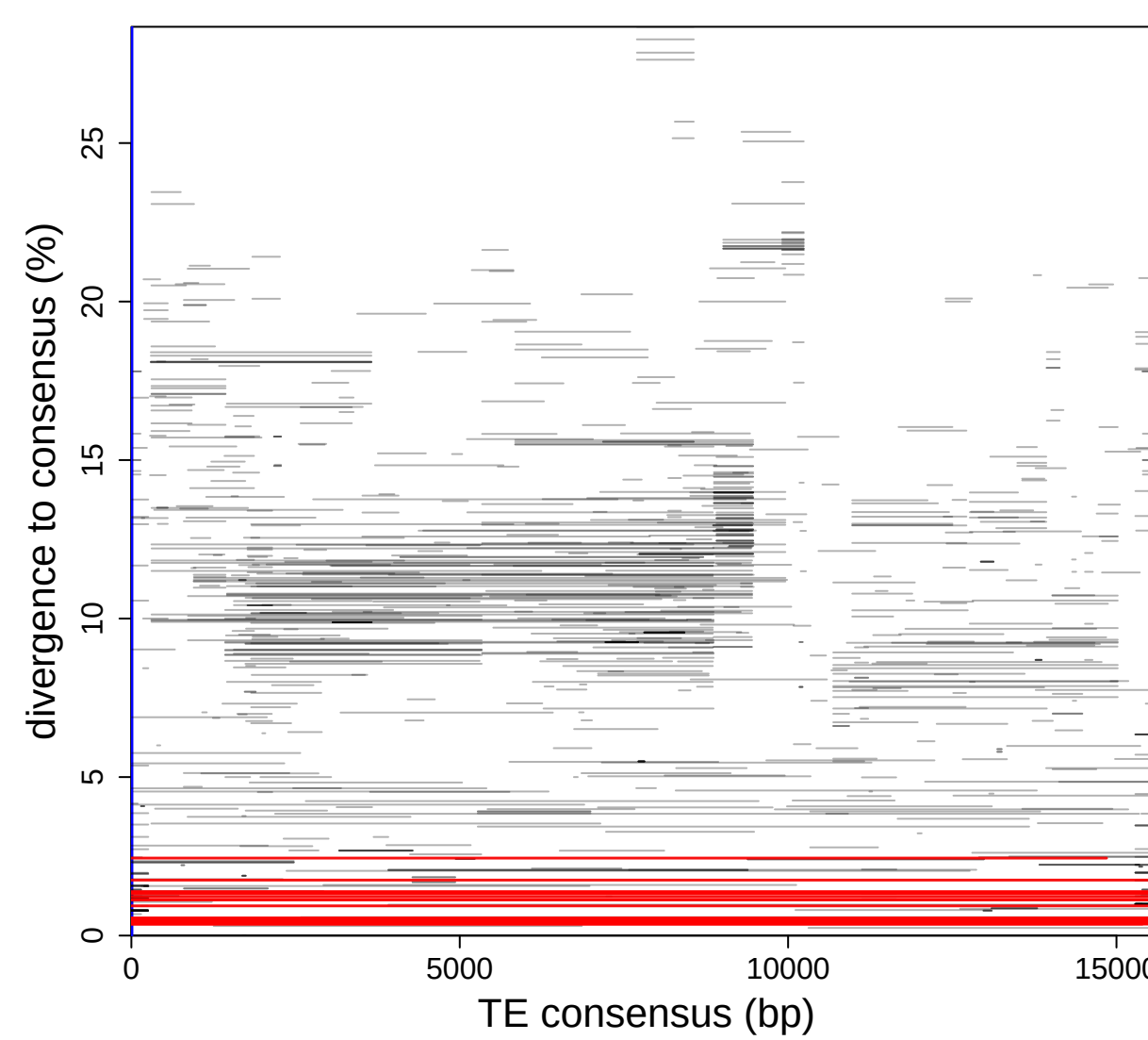
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 15539bp; fragments: 994; full length: 29 (≥ 13985.1 bp)

After TEtrimmer 15539 bp



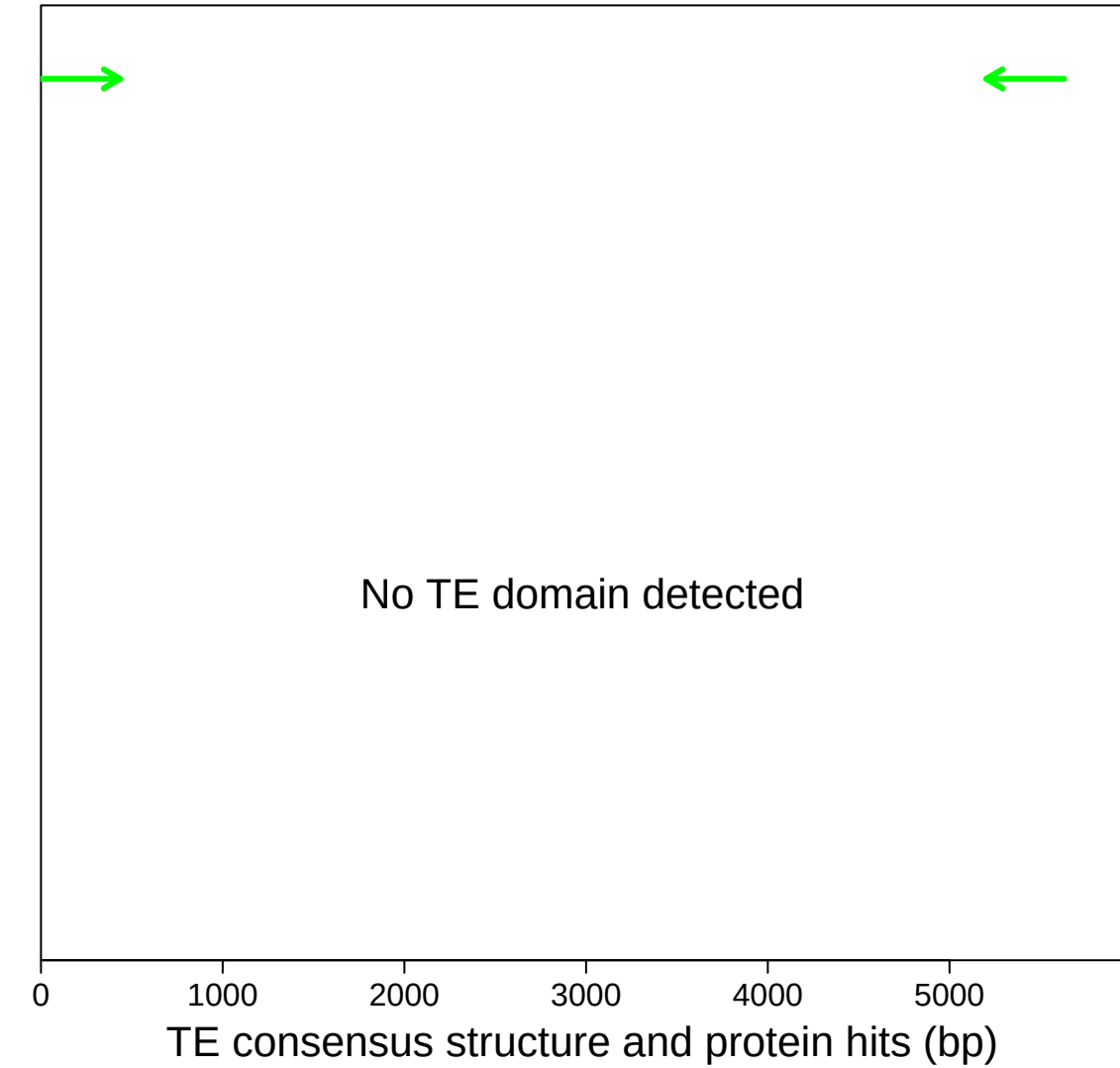
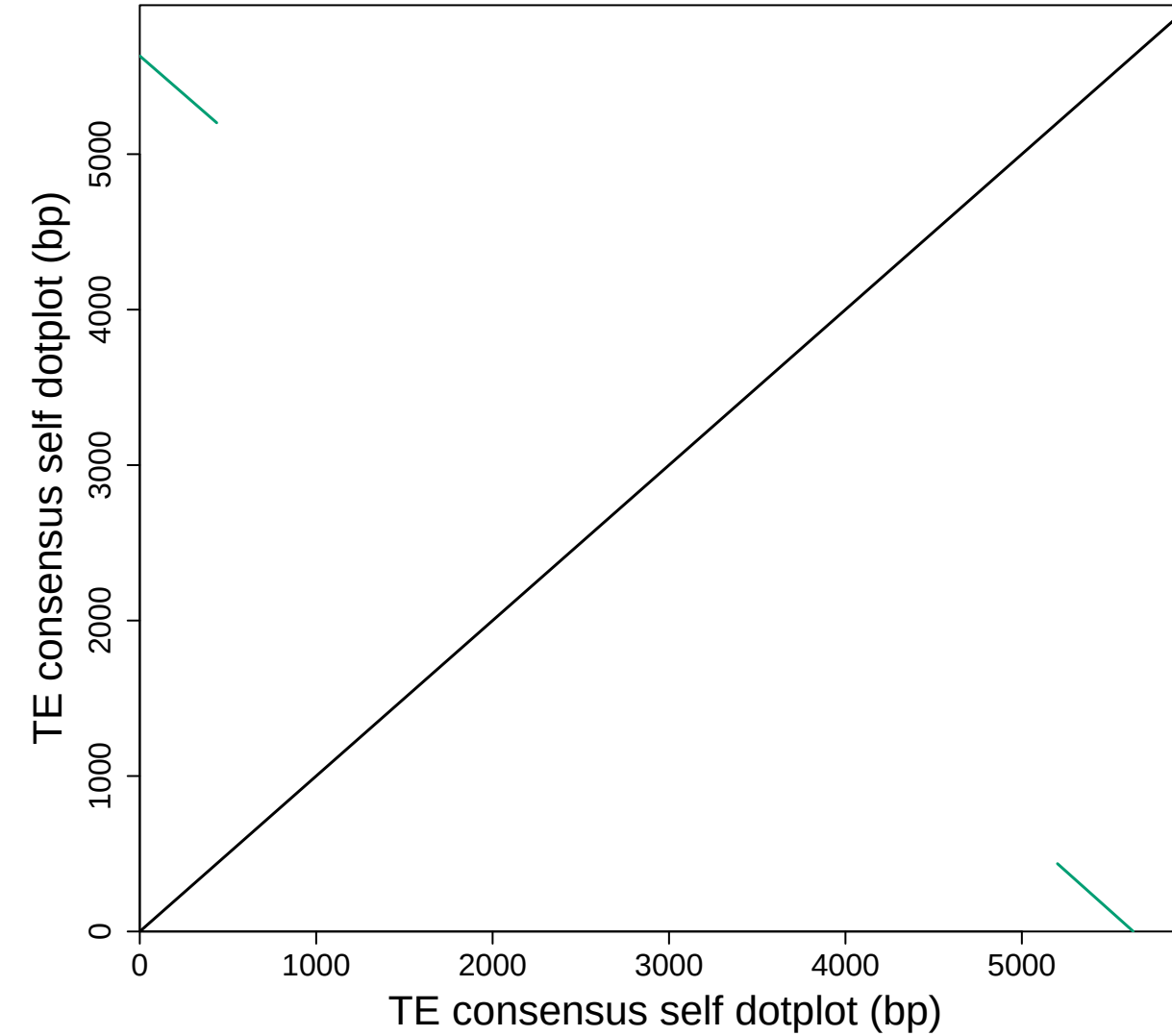
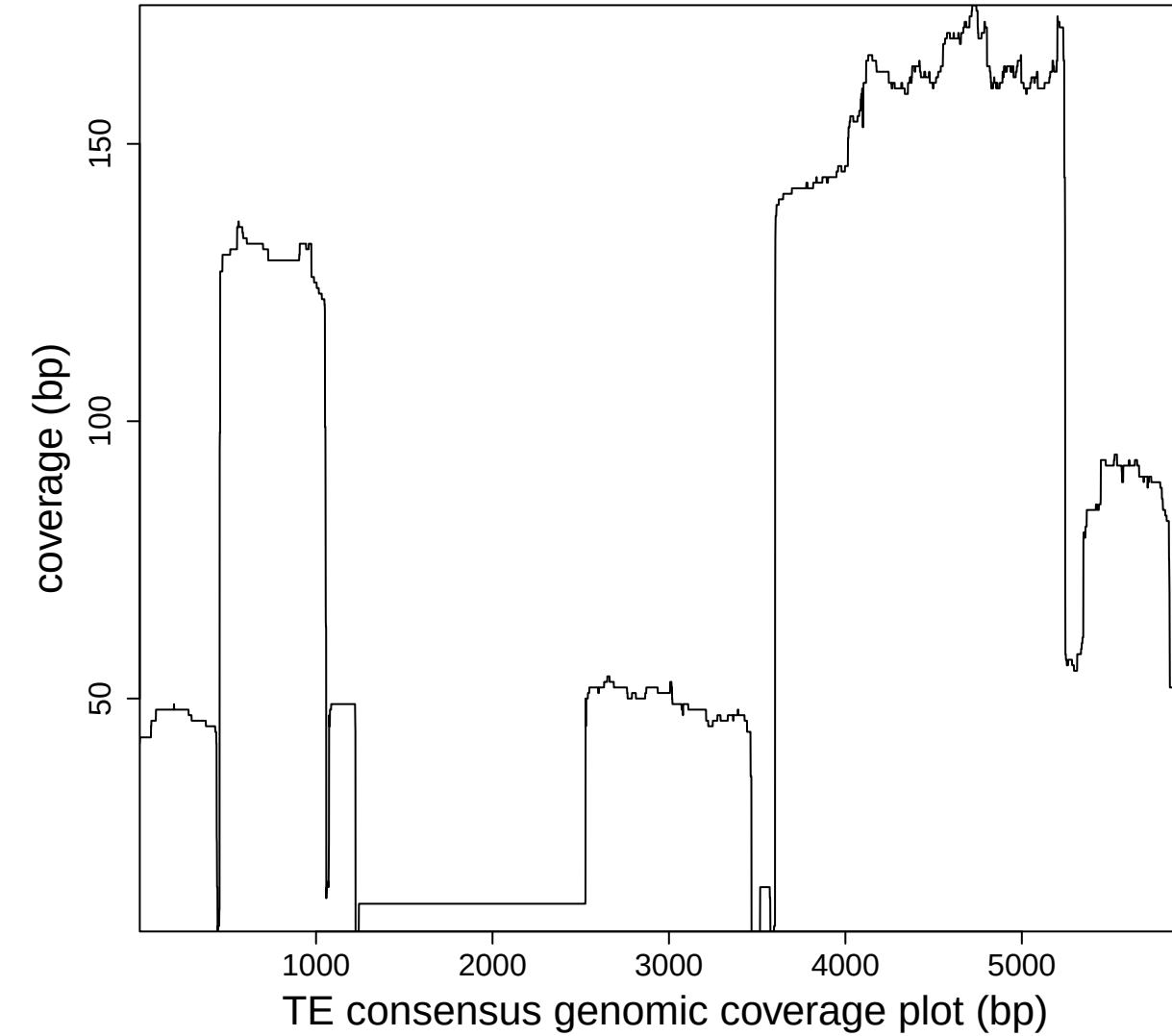
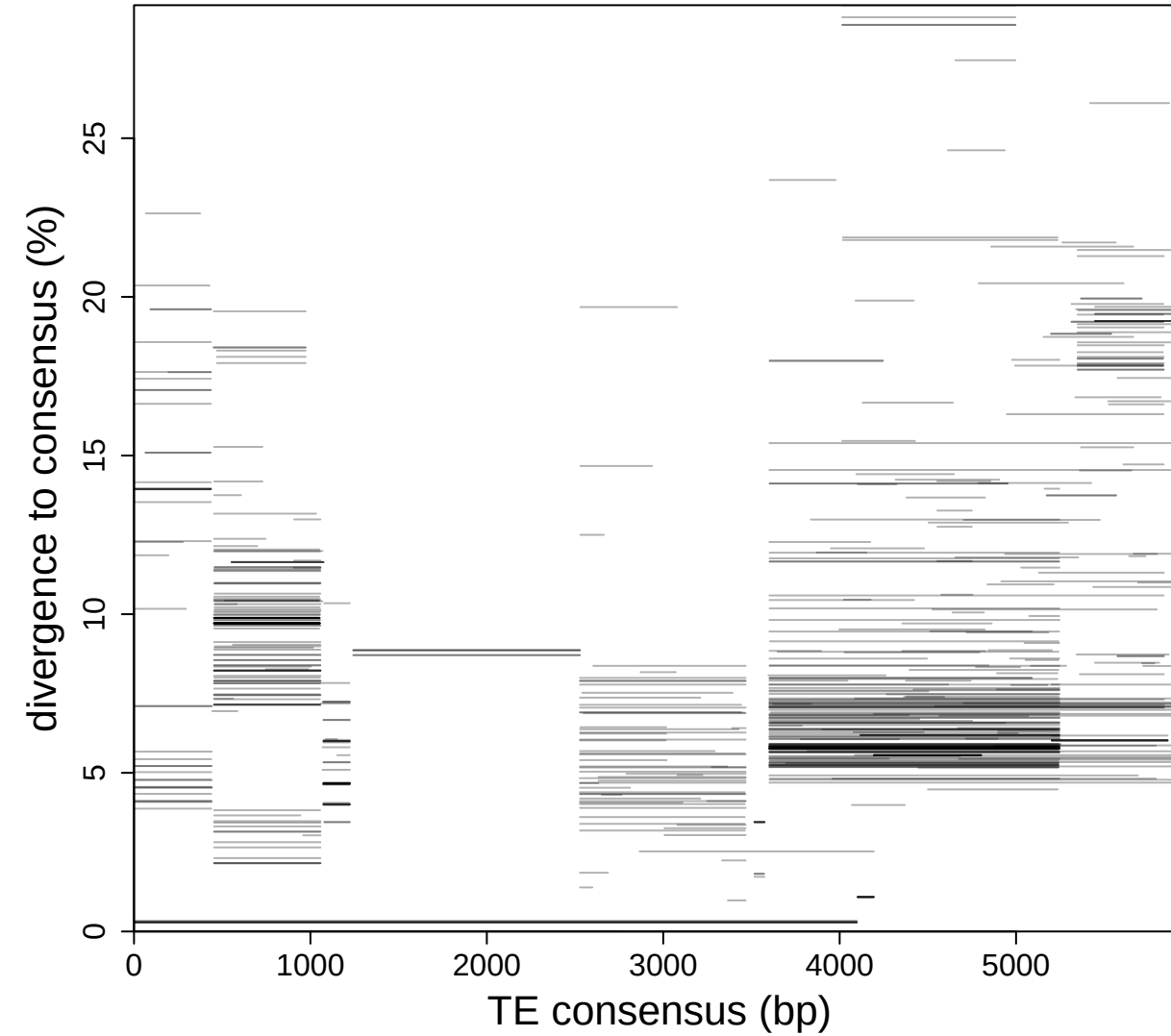
ly_3973.fasta.b.bed uf.bed g 1.bed fm 1.bed 0 0 bcln.fa aln.f
size: 15690bp; fragments: 1006; full length: 29 (>=14121bp)

After TEtrimmer Extended plot Blue lines are boundaries

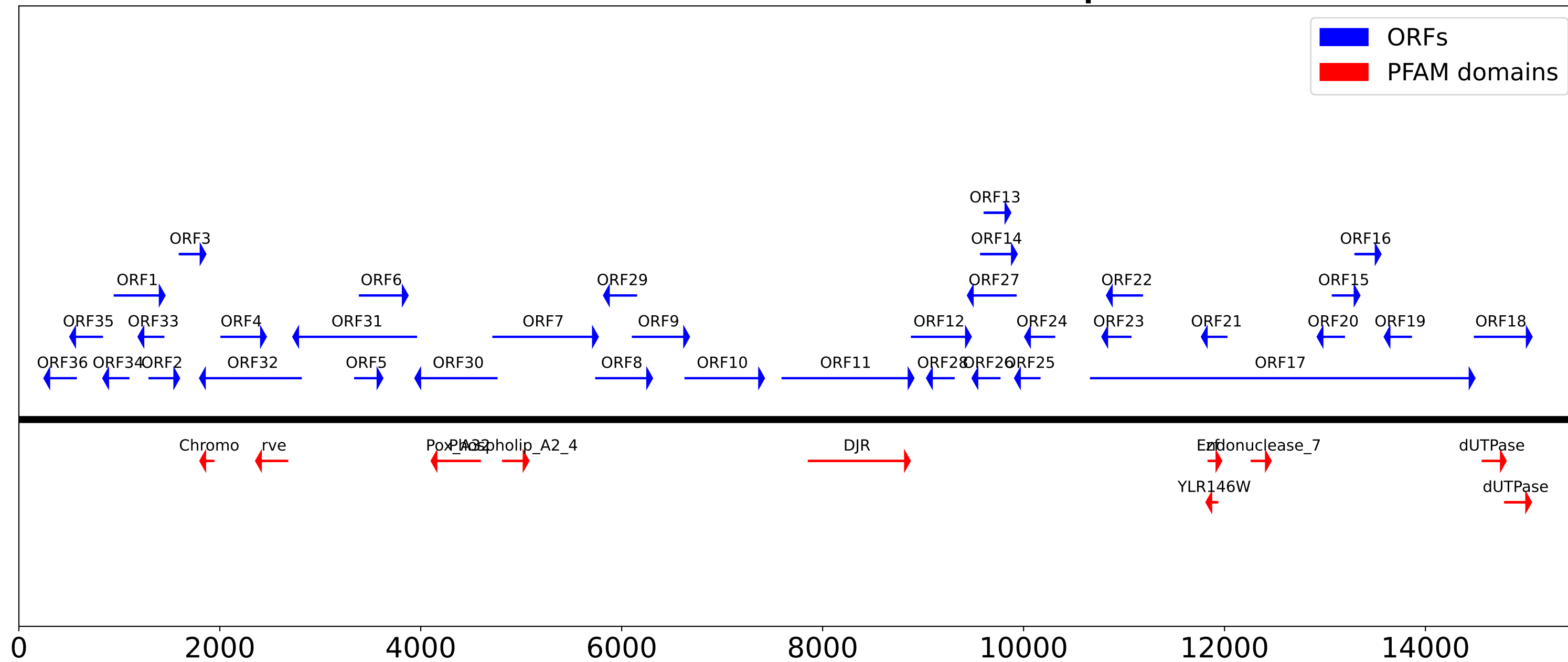


TE: rnd_5_family_3973
size: 5958bp; fragments: 598; full length: 0 (≥ 5362.2 bp)

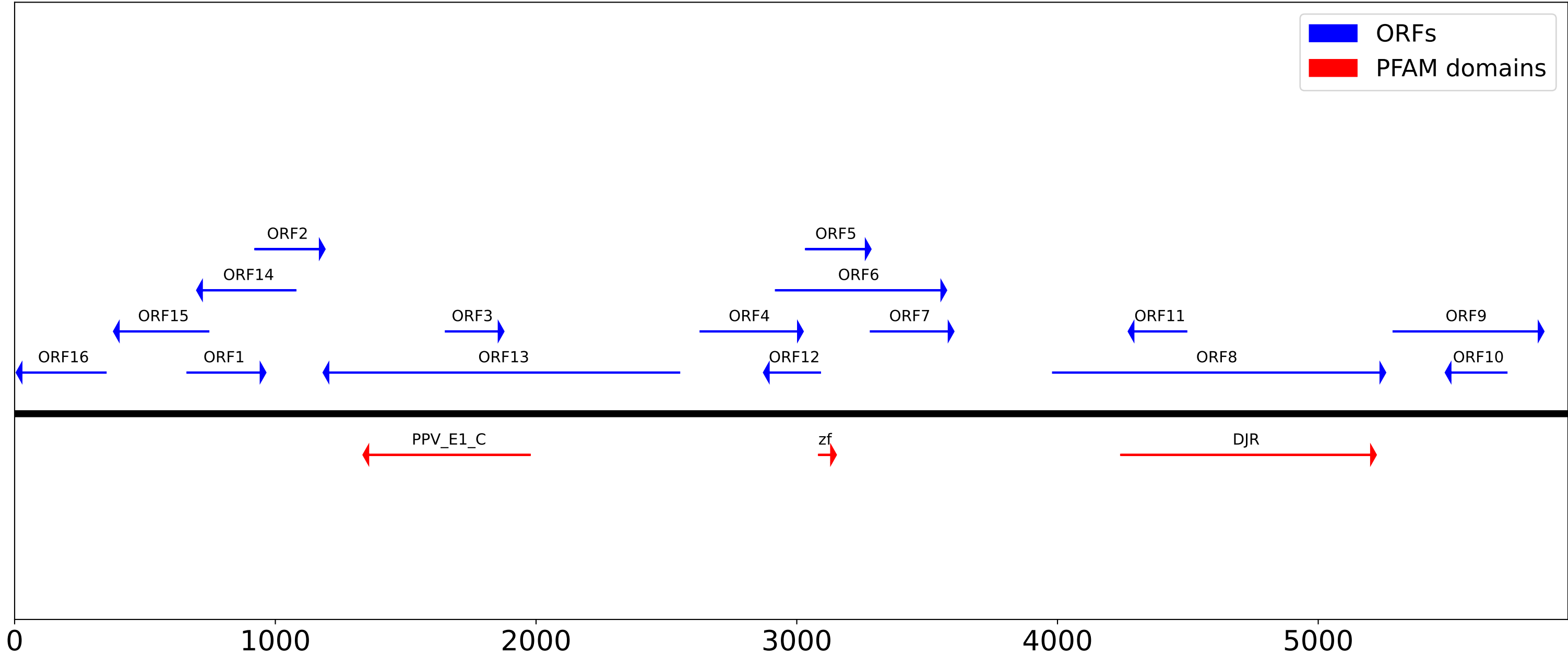
Before TEtrimmer 5958 bp



After TEtrimmer ORF and PFAM domain plot



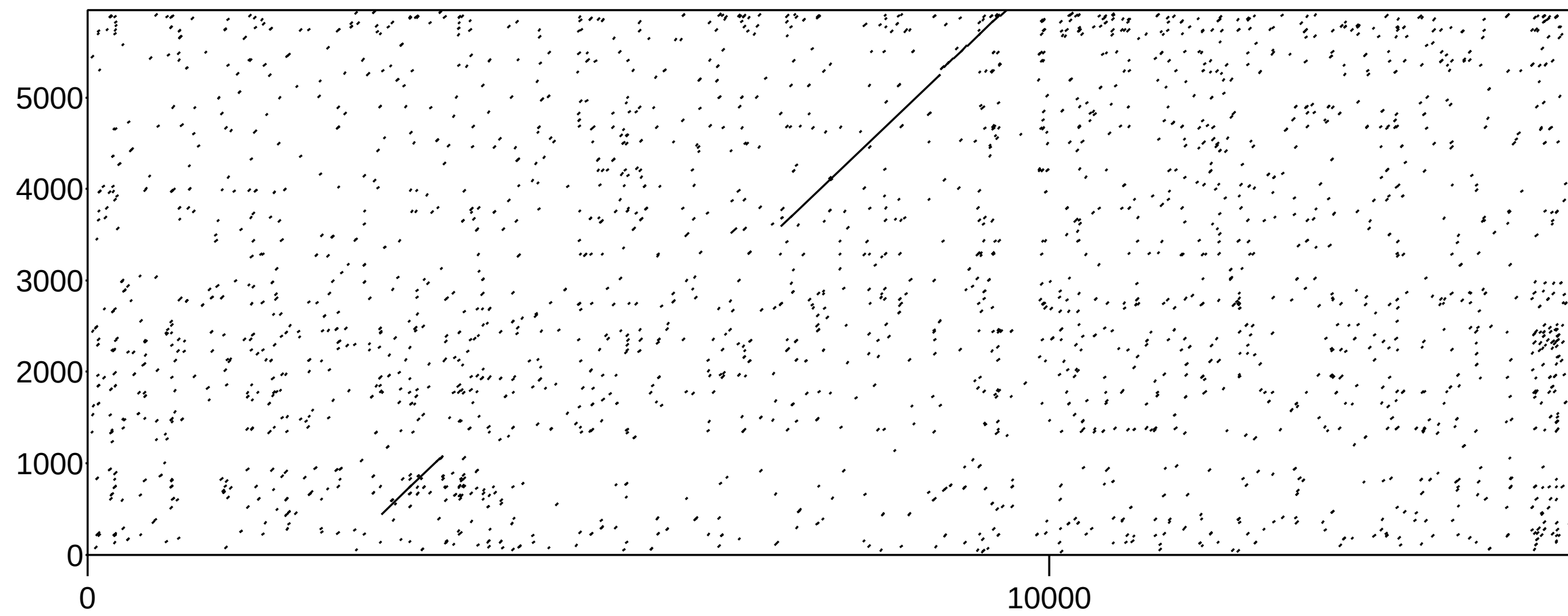
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

TE consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)