

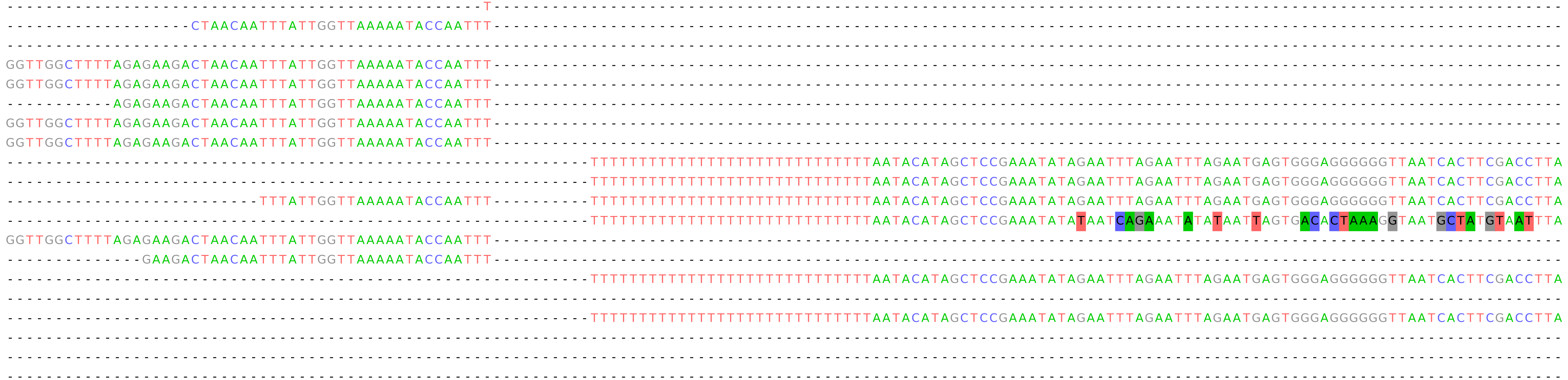
Start crop Point

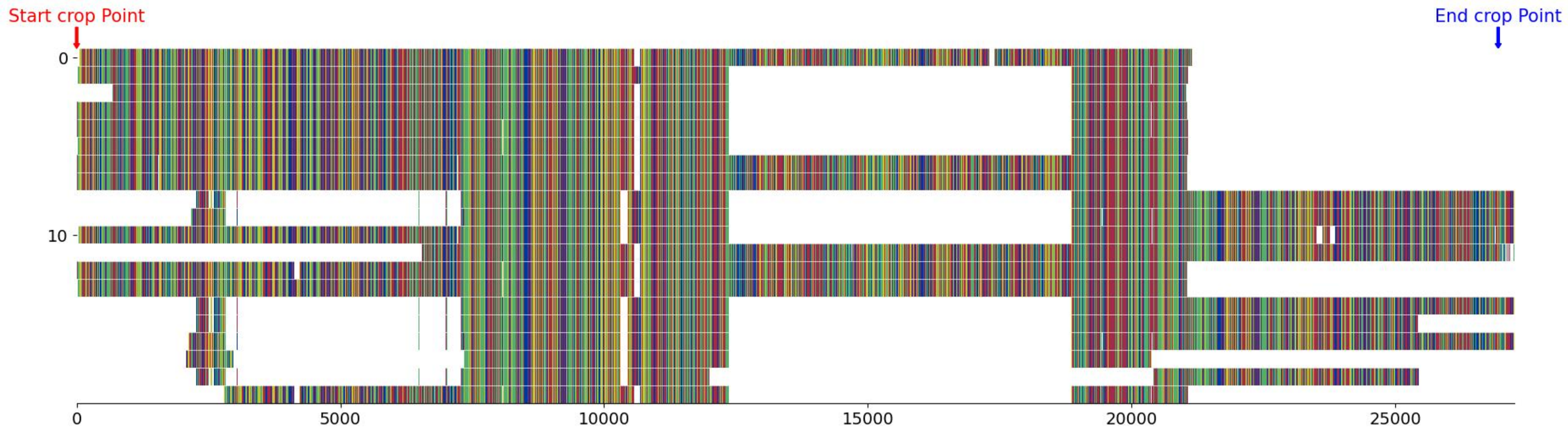


End crop Point



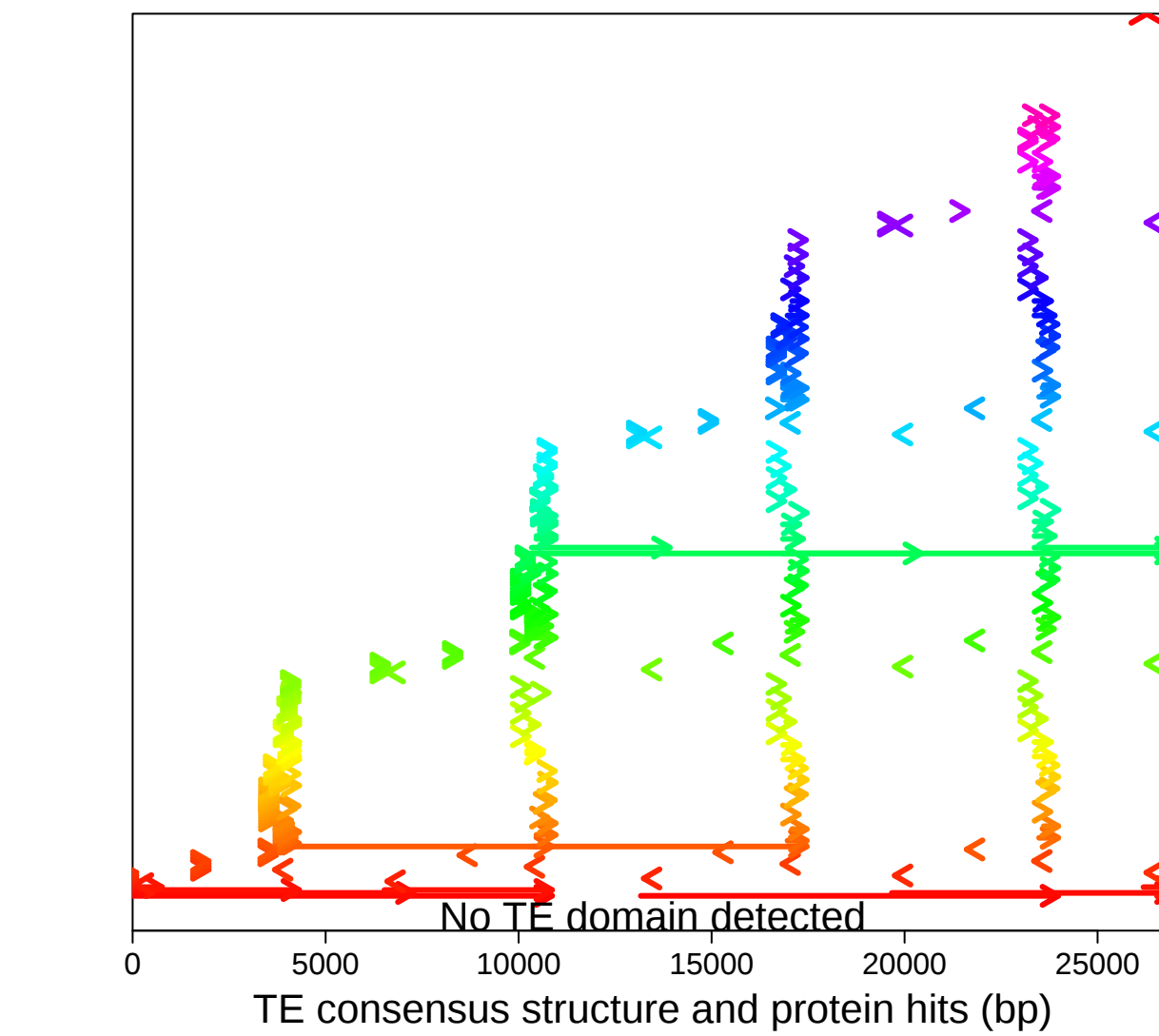
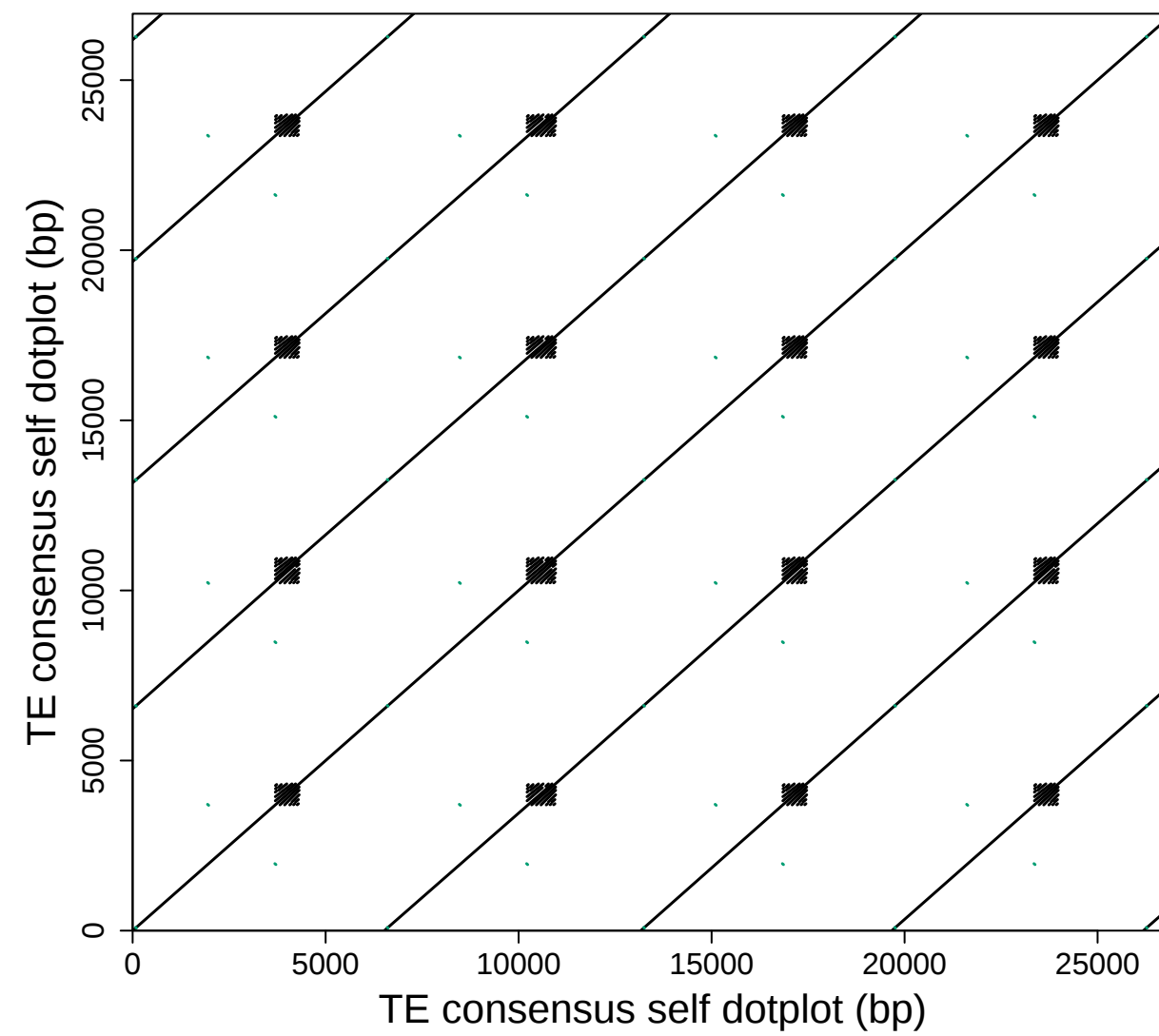
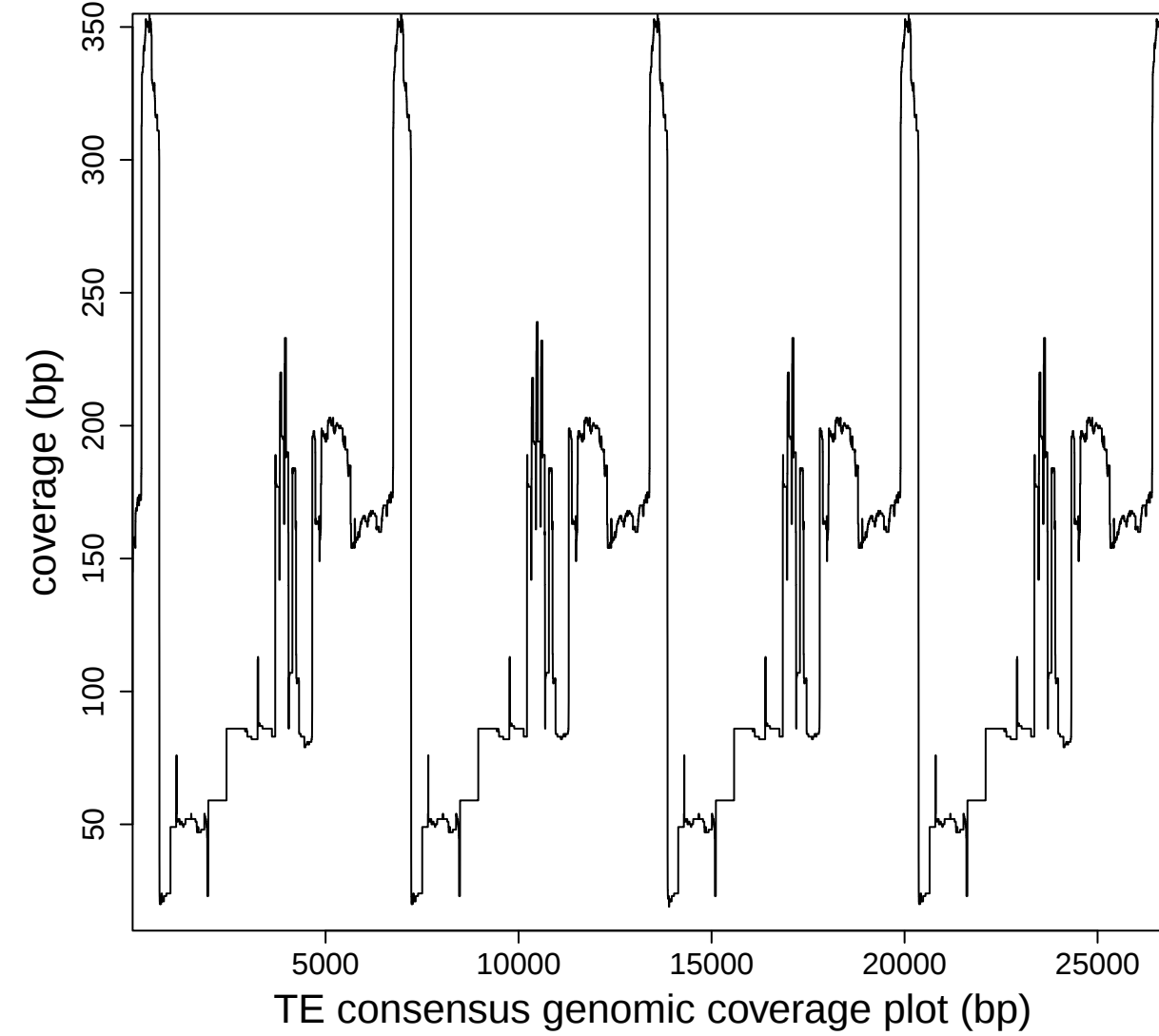
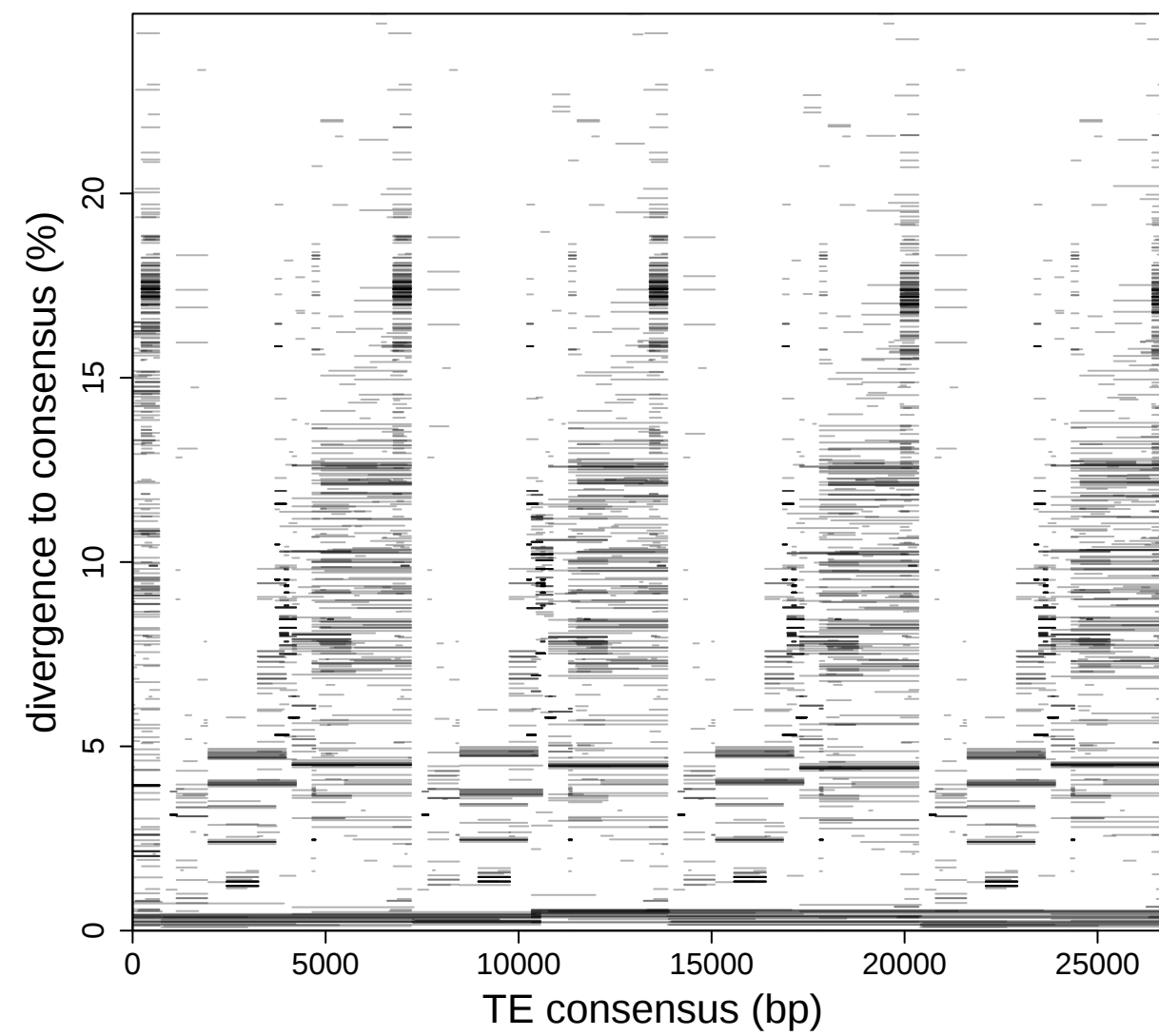
MSA length = 26952





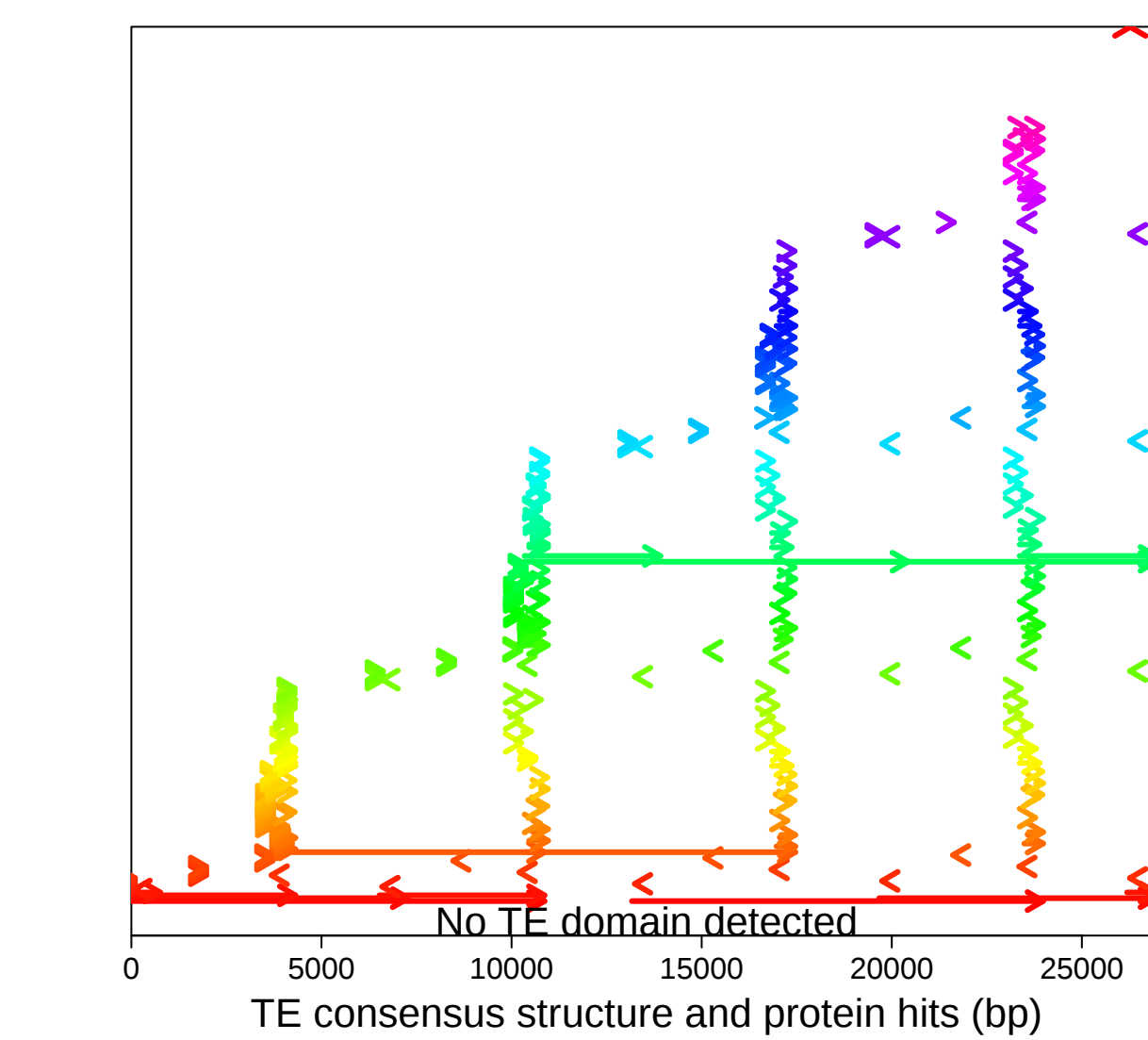
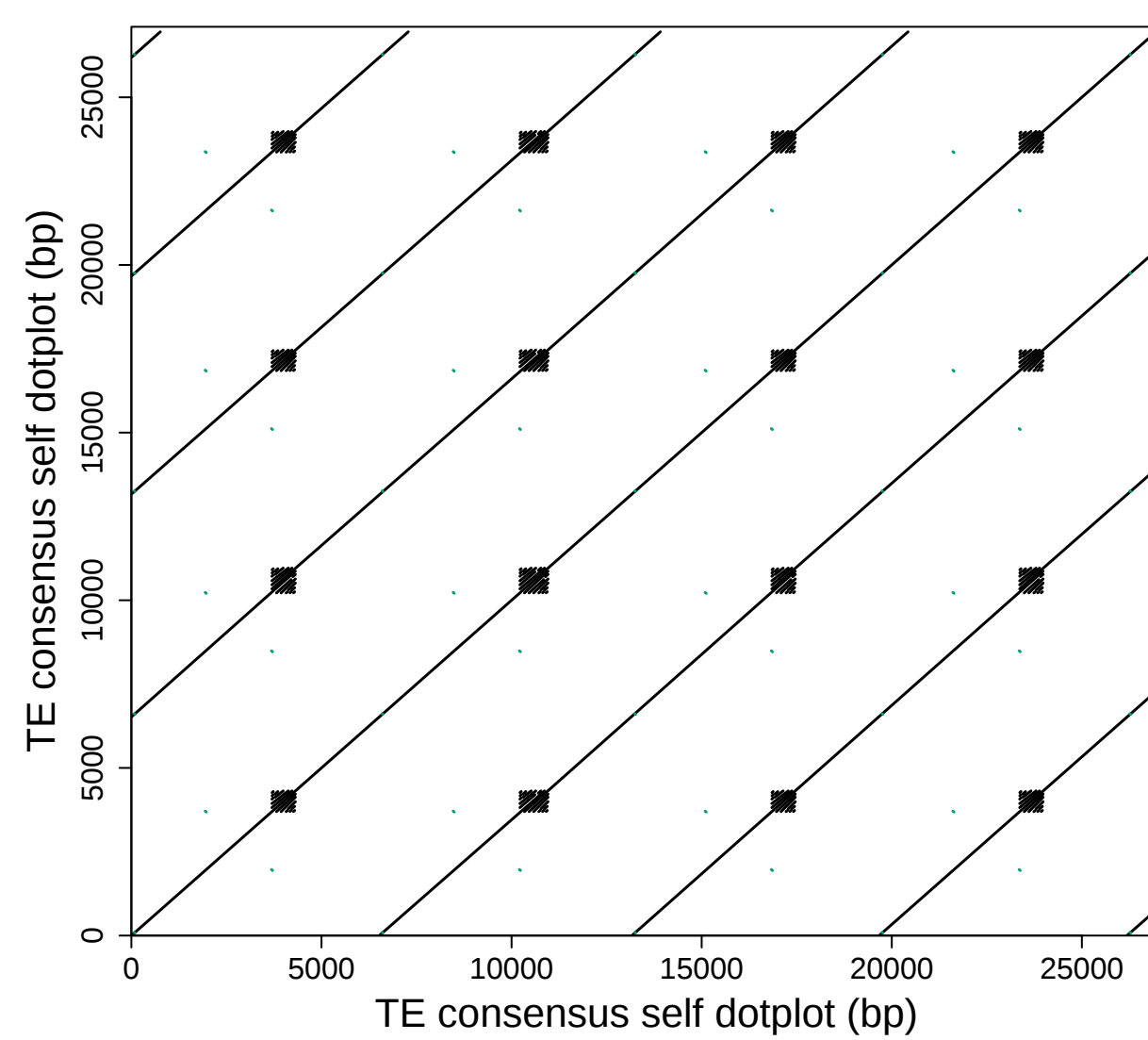
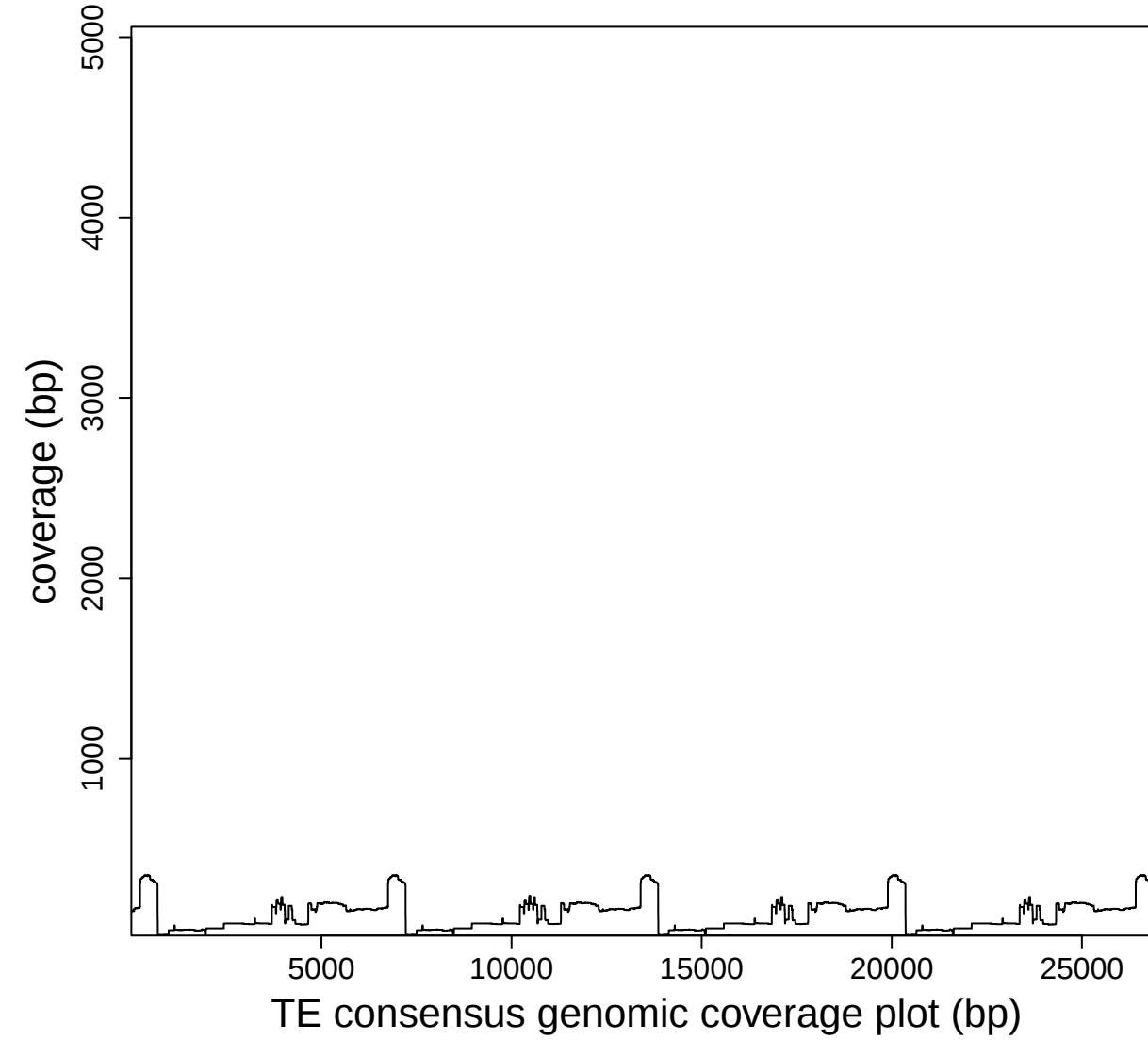
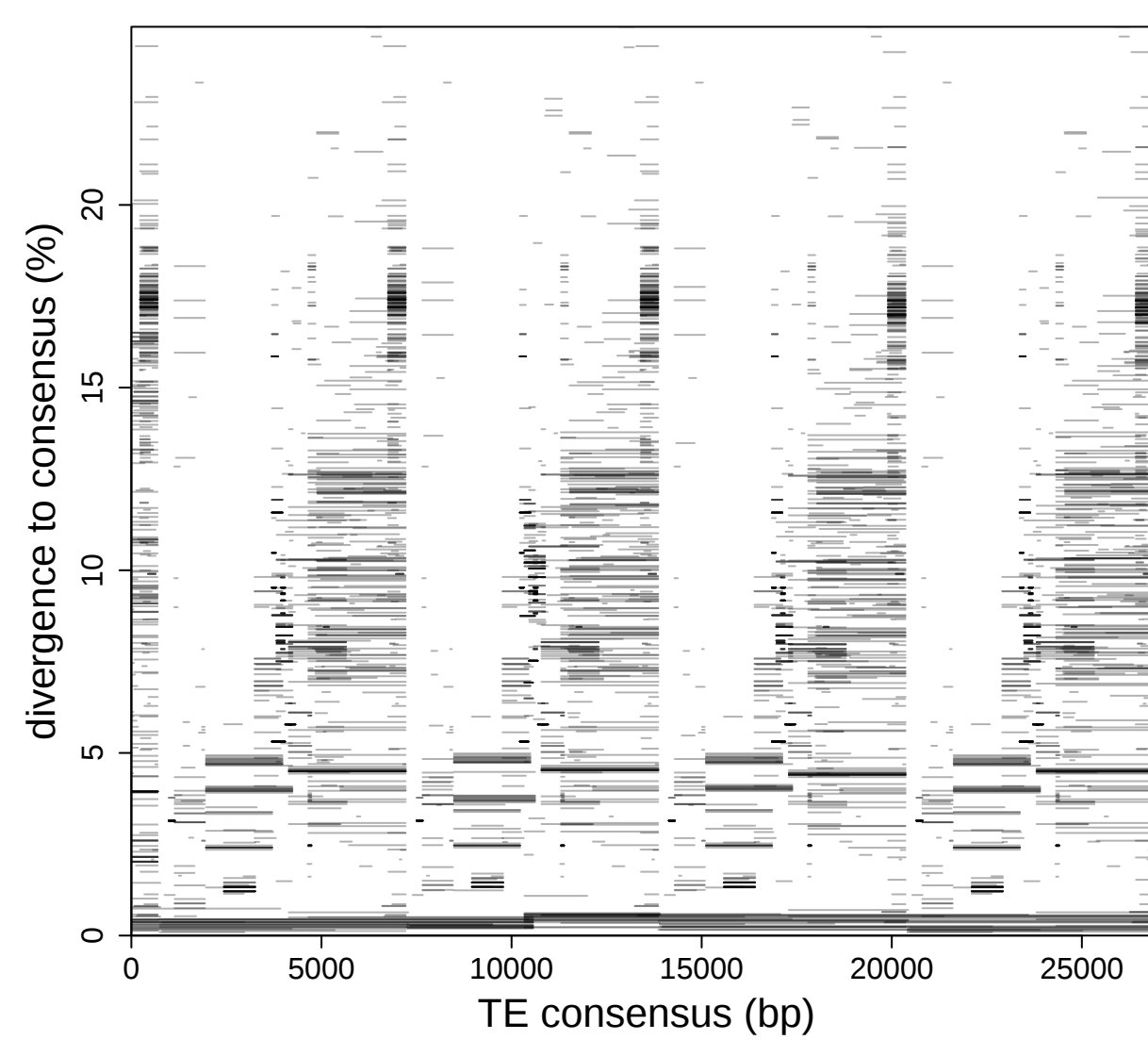
bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.fa_c
size: 26952bp; fragments: 4548; full length: 0 (>=24256.8bp)

After TEtrimmer 26952 bp

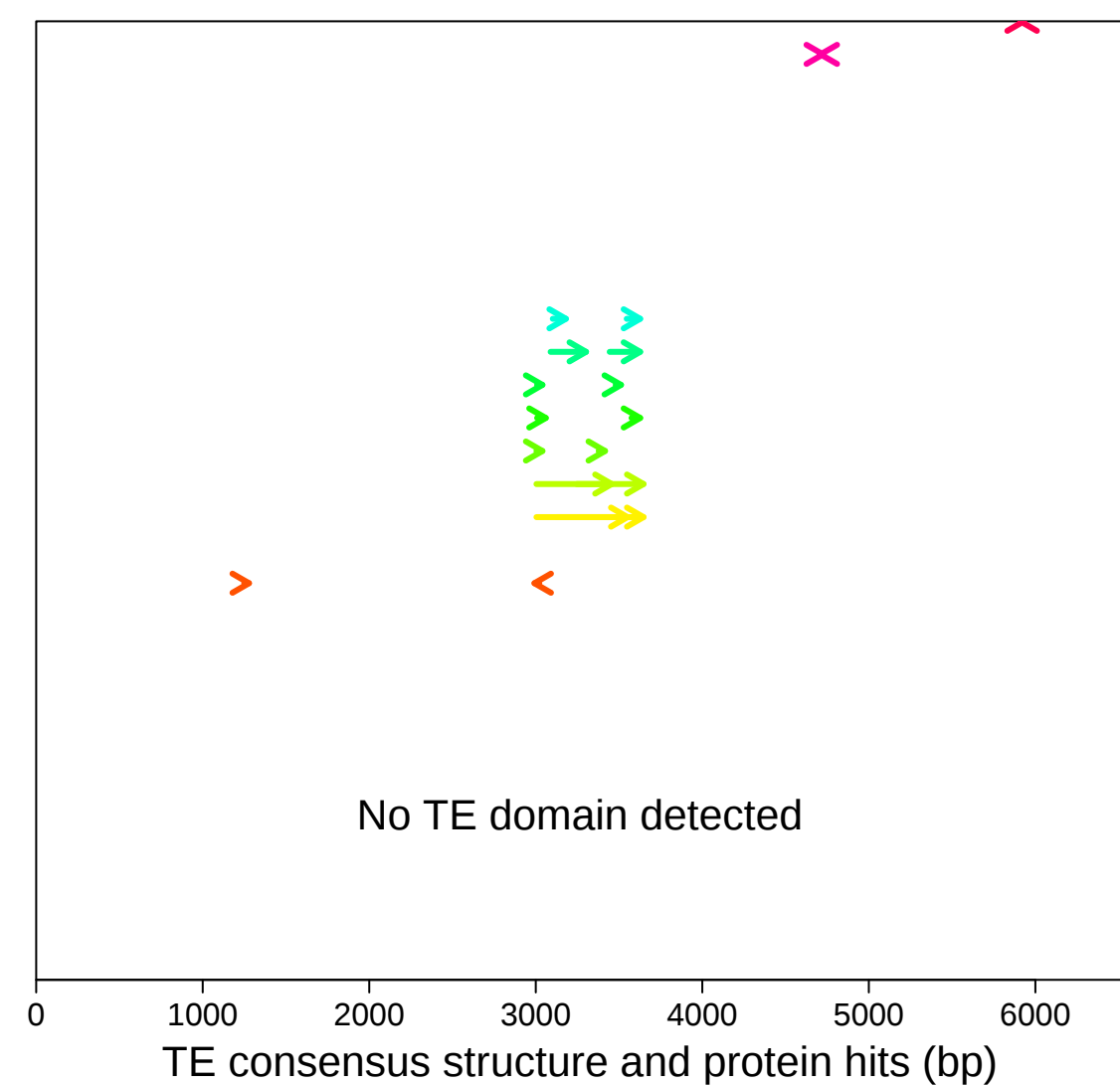
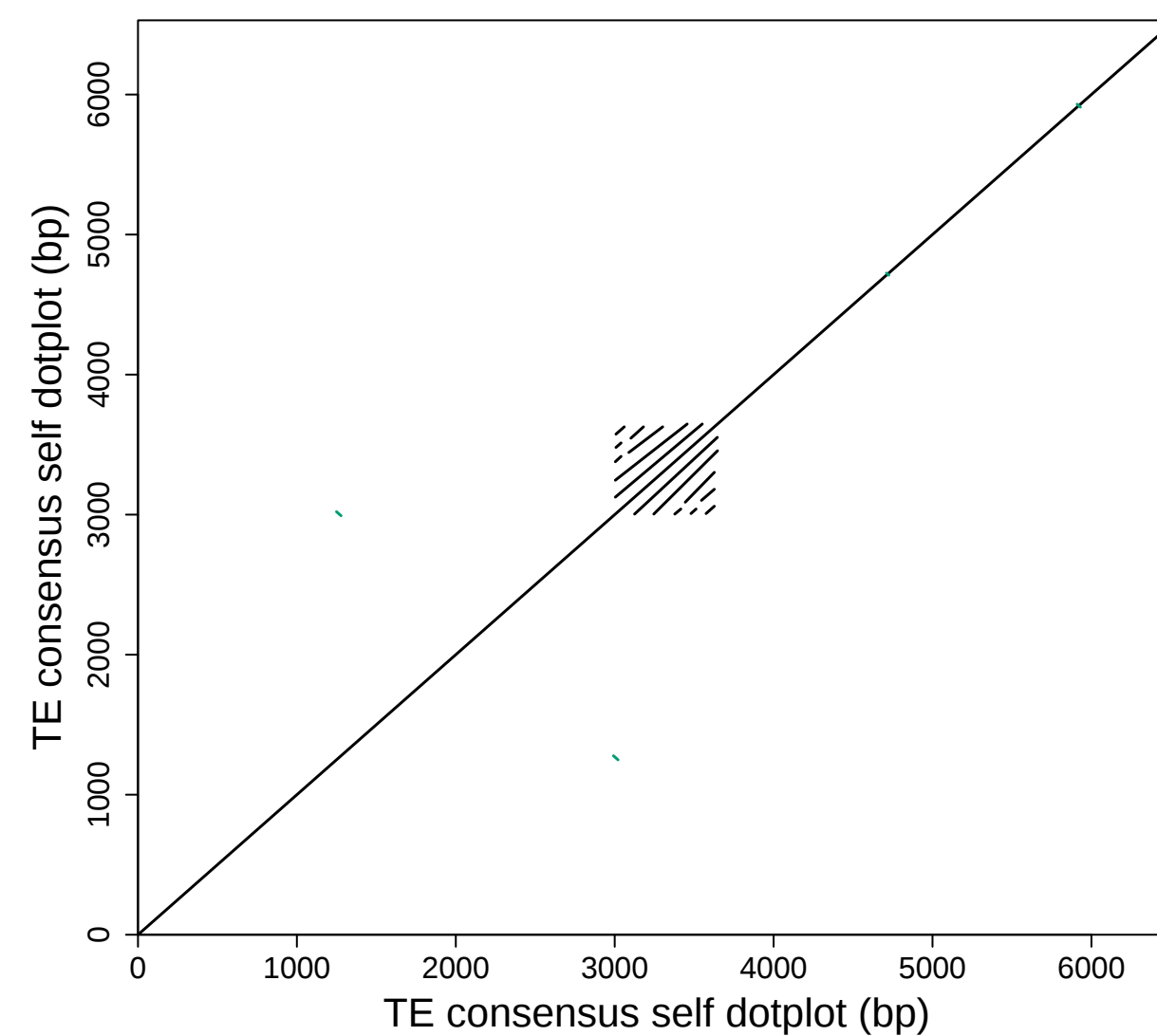
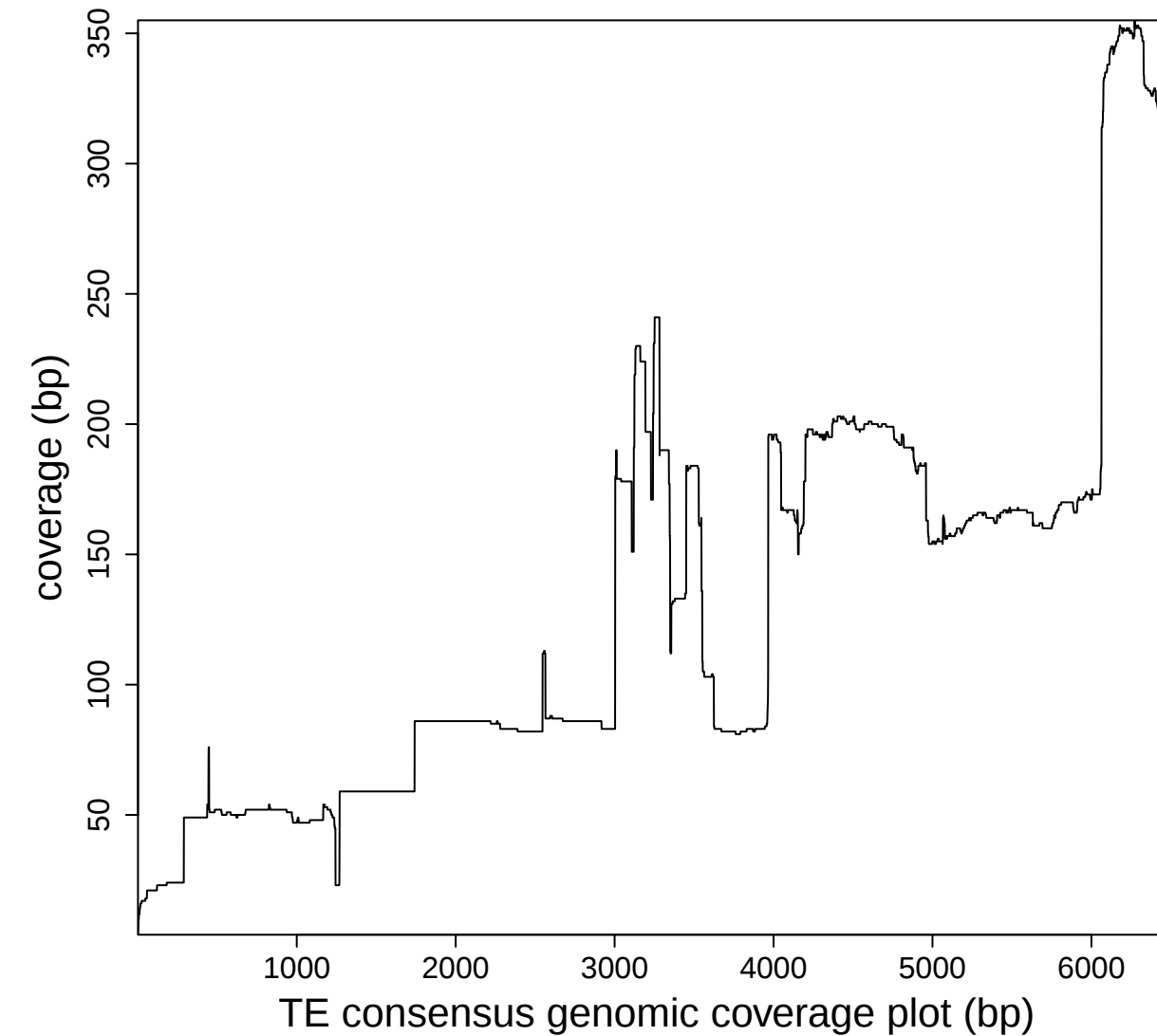
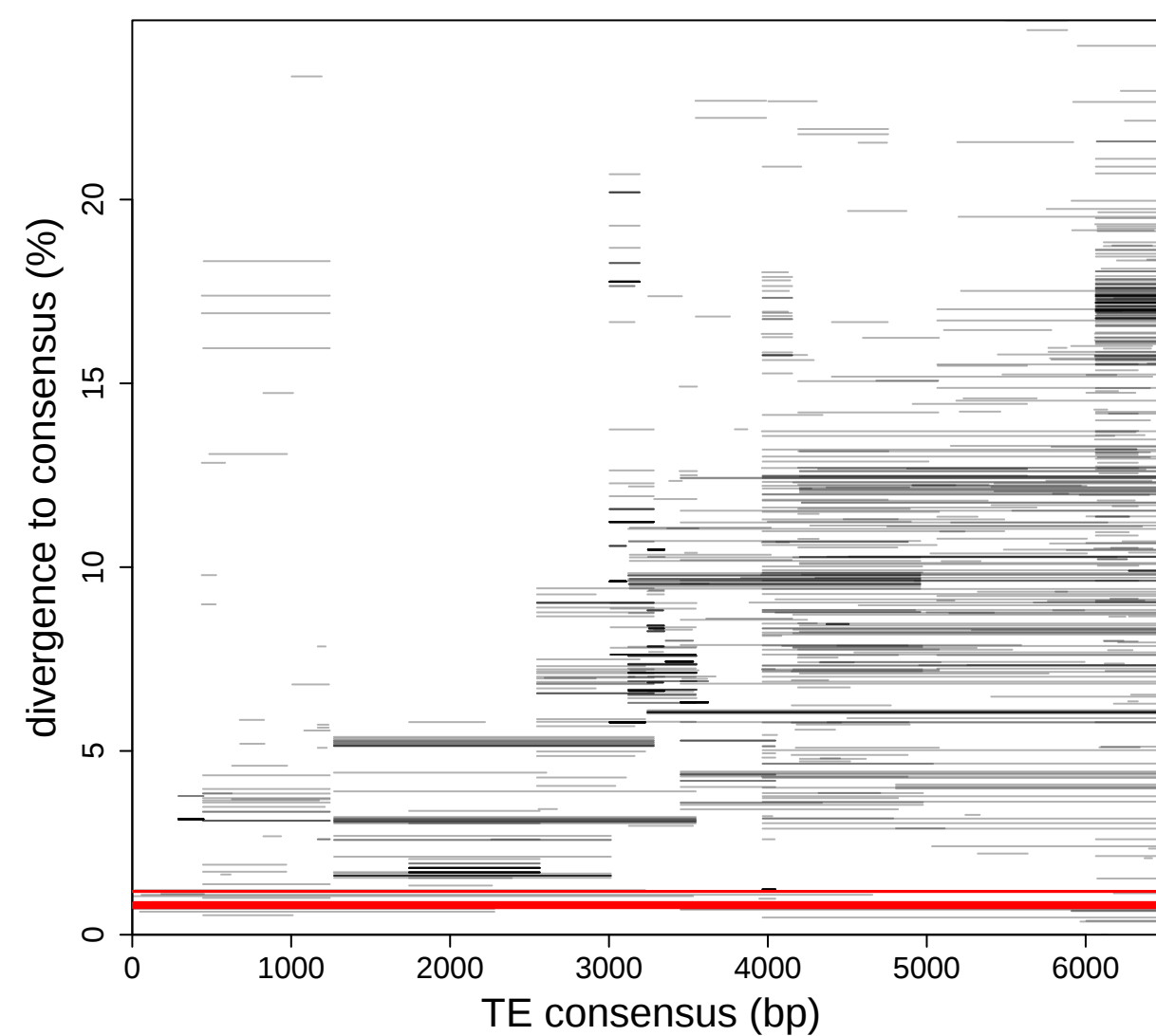


ly_9115.fasta.b.bed_uf.bed_g_1.bed_fm_1.bed_0_0_bcln.fa_aln.f
size: 27102bp; fragments: 9822; full length: 0 (>=24391.8bp)

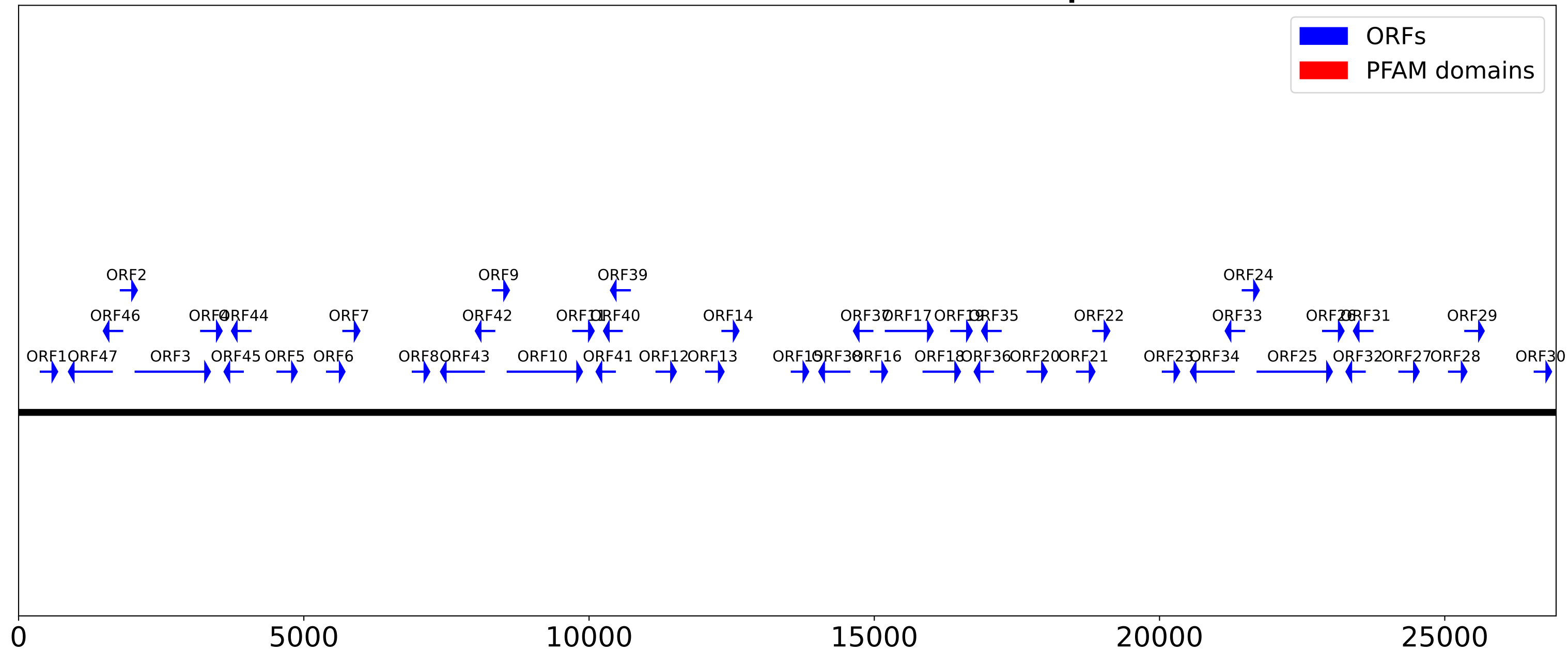
After TEtrimmer Extended plot Blue lines are boundaries



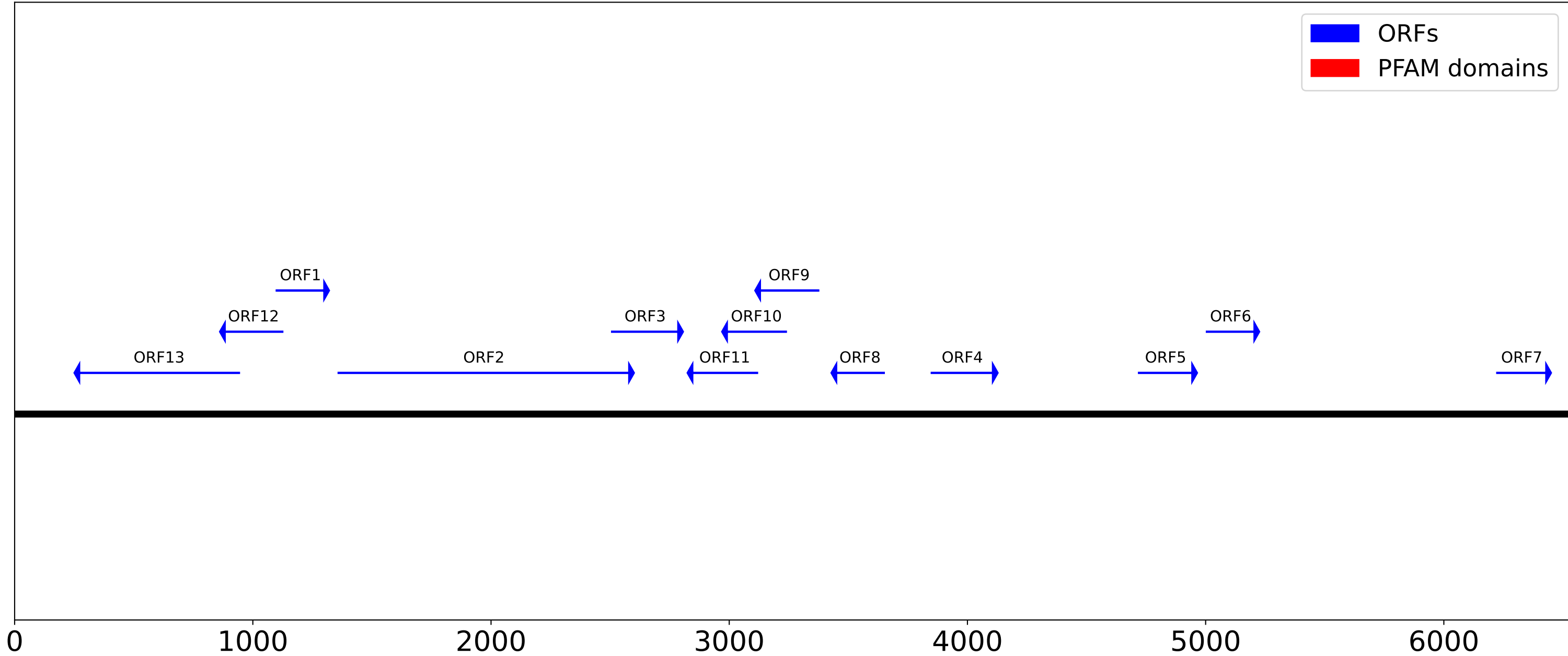
TE: rnd_5_family_9115
size: 6530bp; fragments: 1010; full length: 18 (≥ 5877 bp)



After TEtrimmer ORF and PFAM domain plot



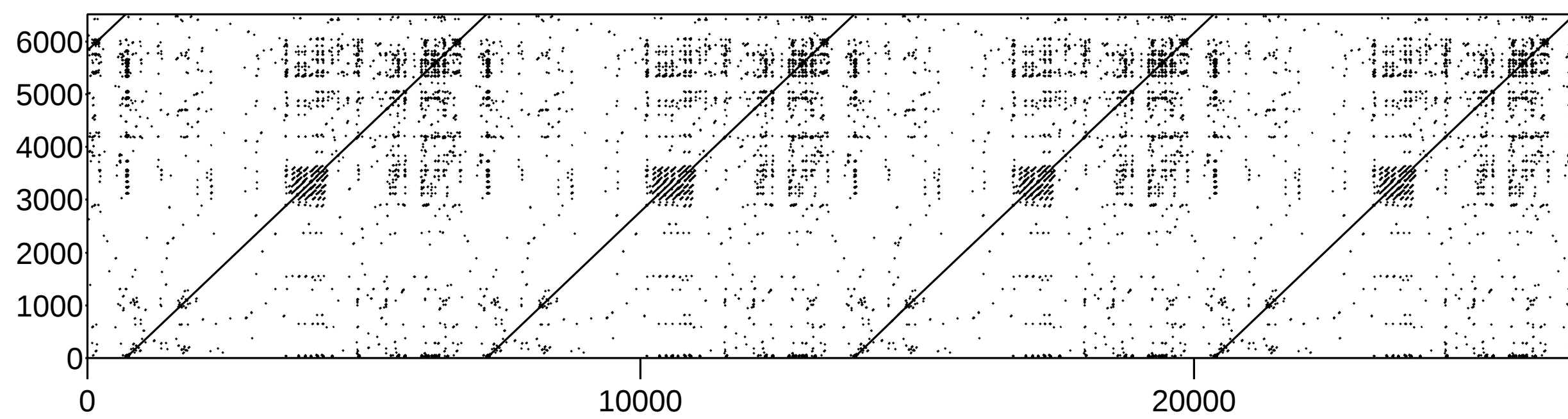
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)