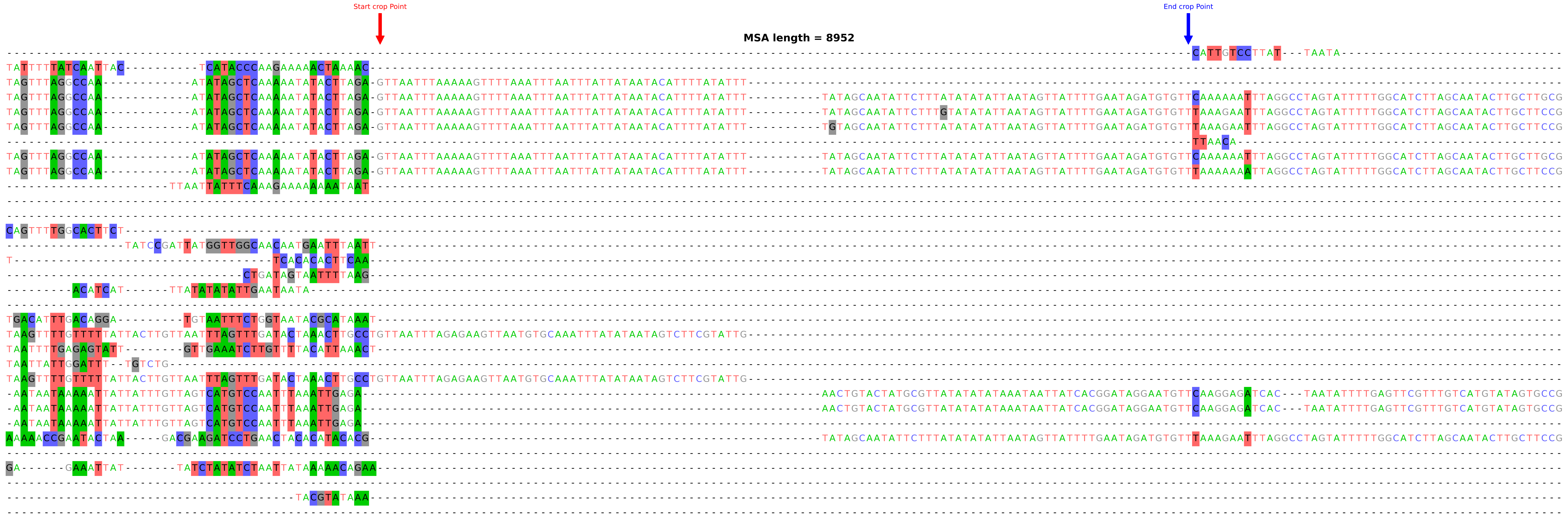
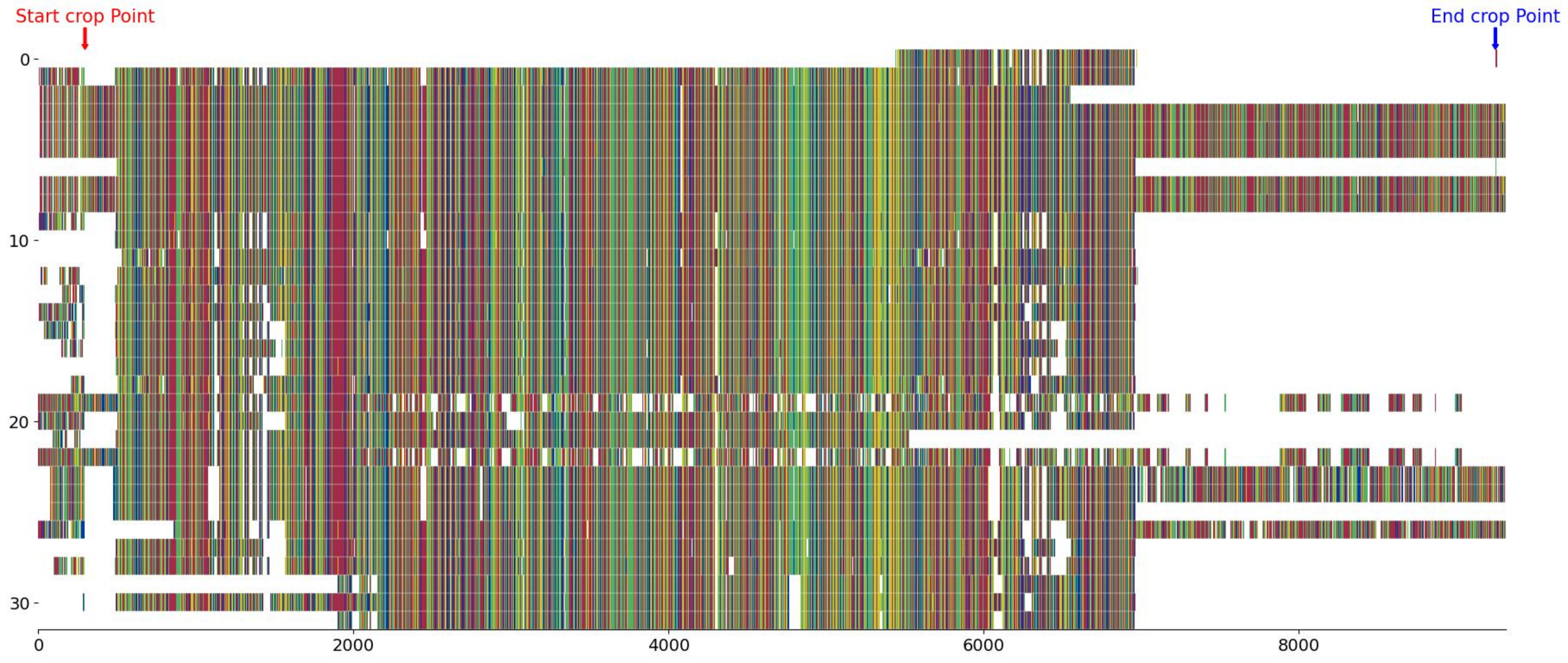
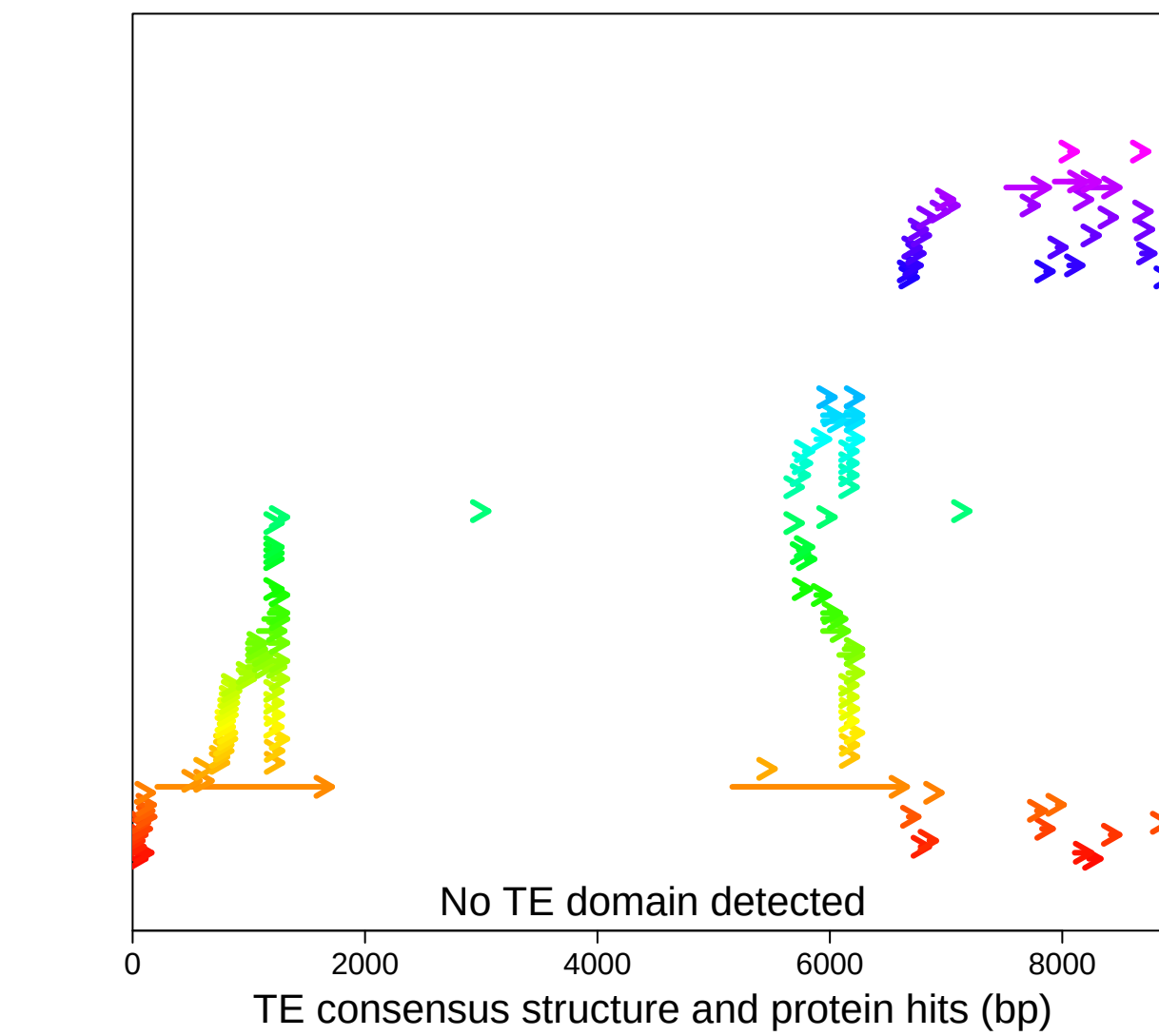
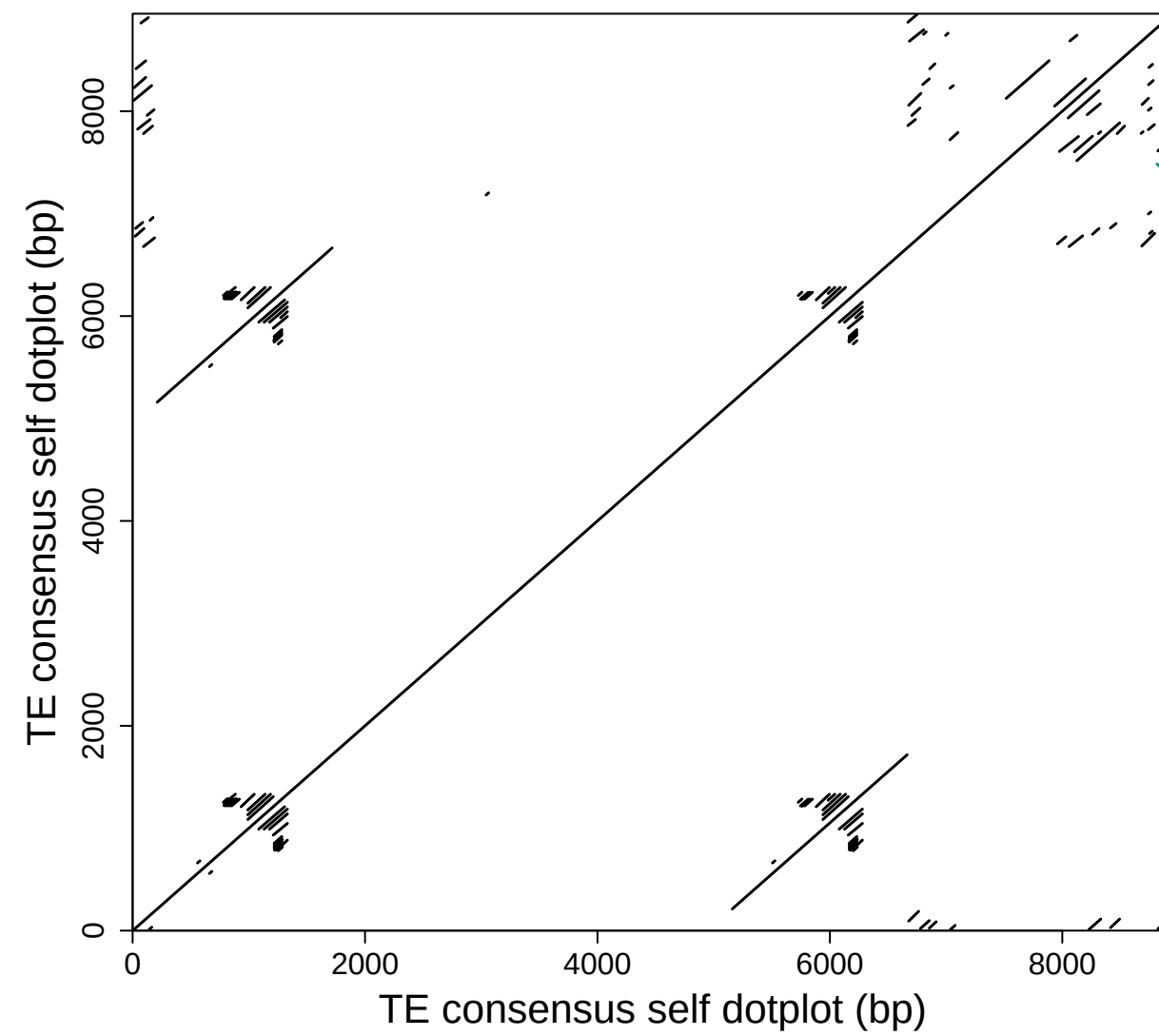
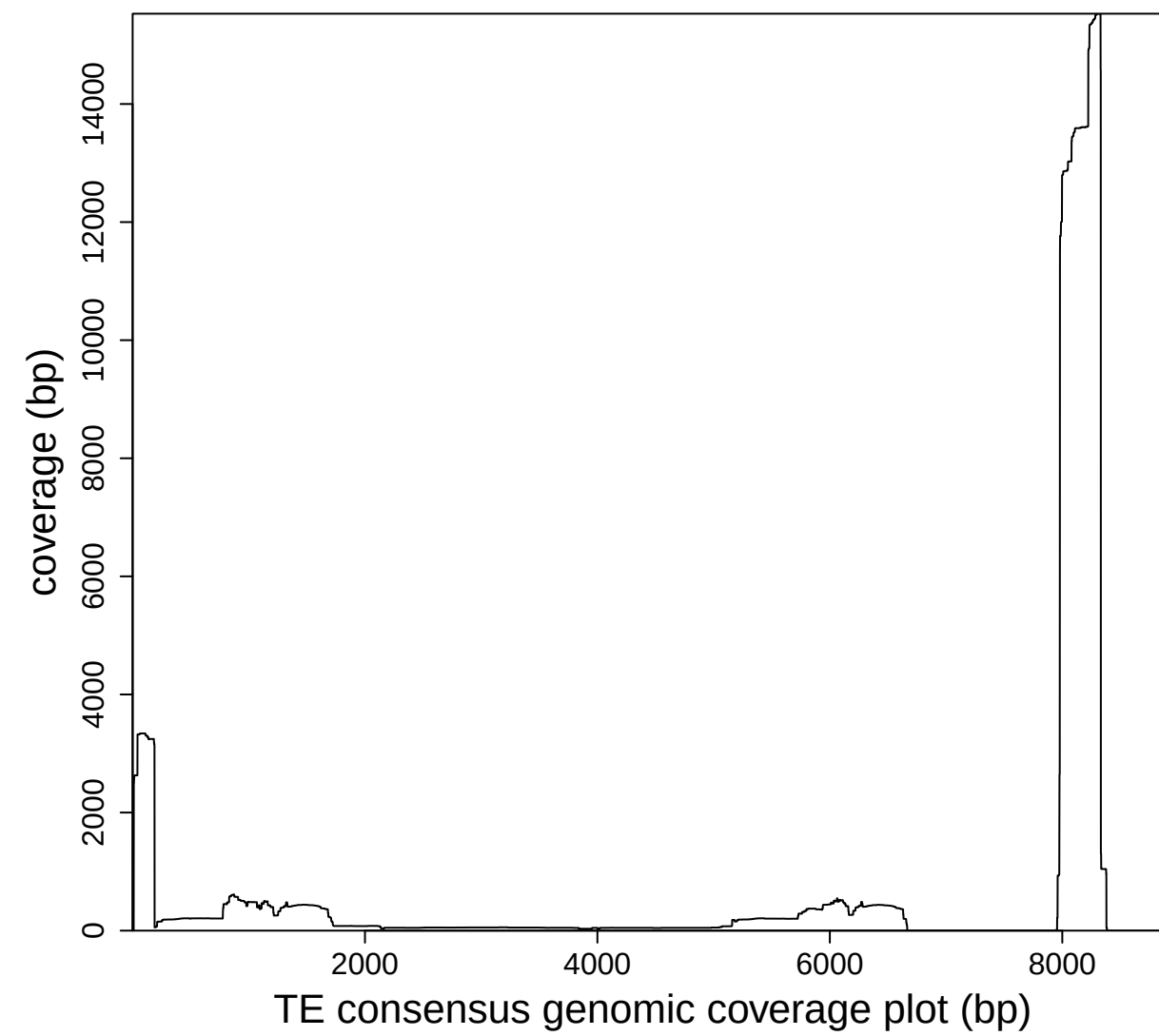
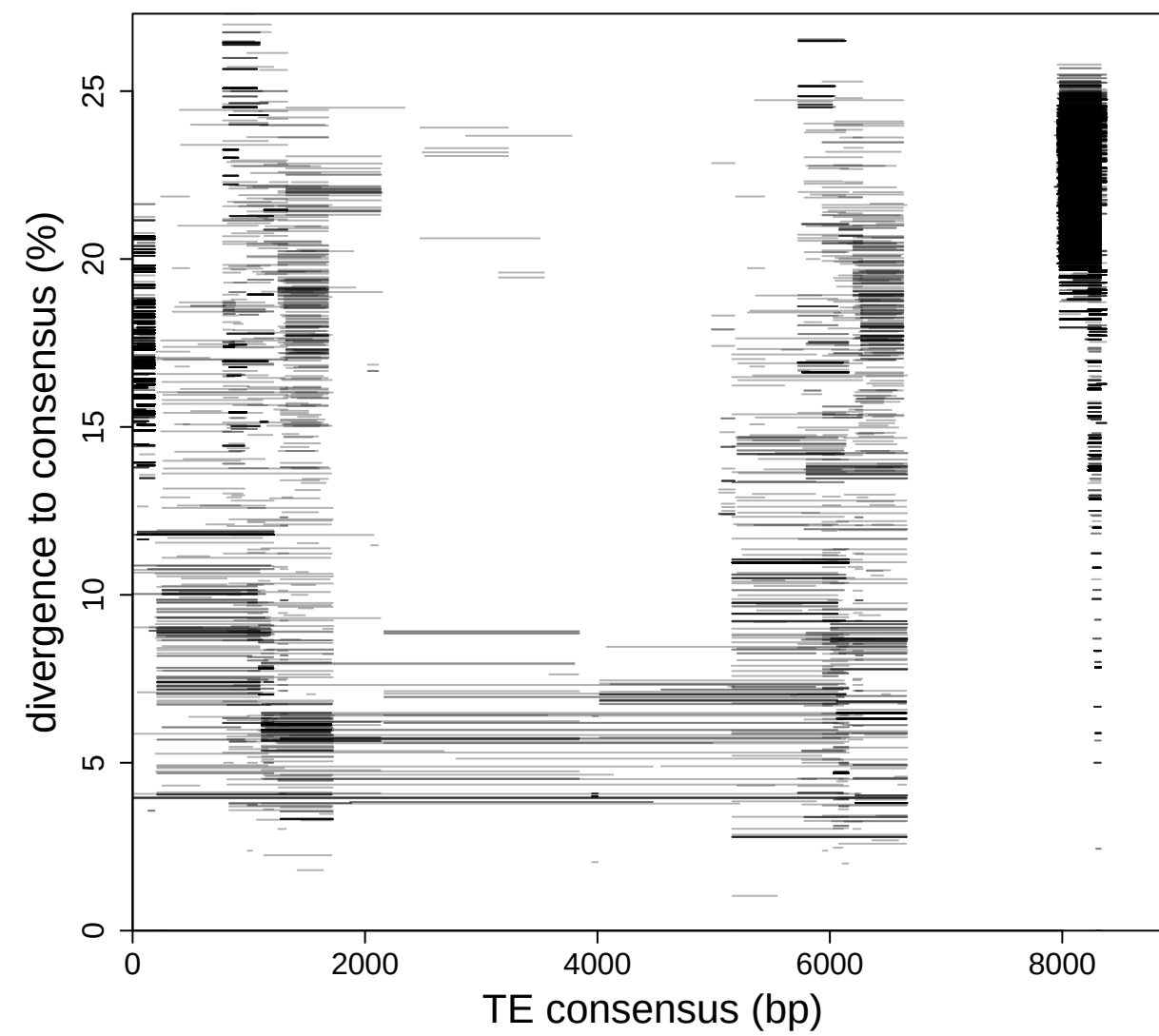






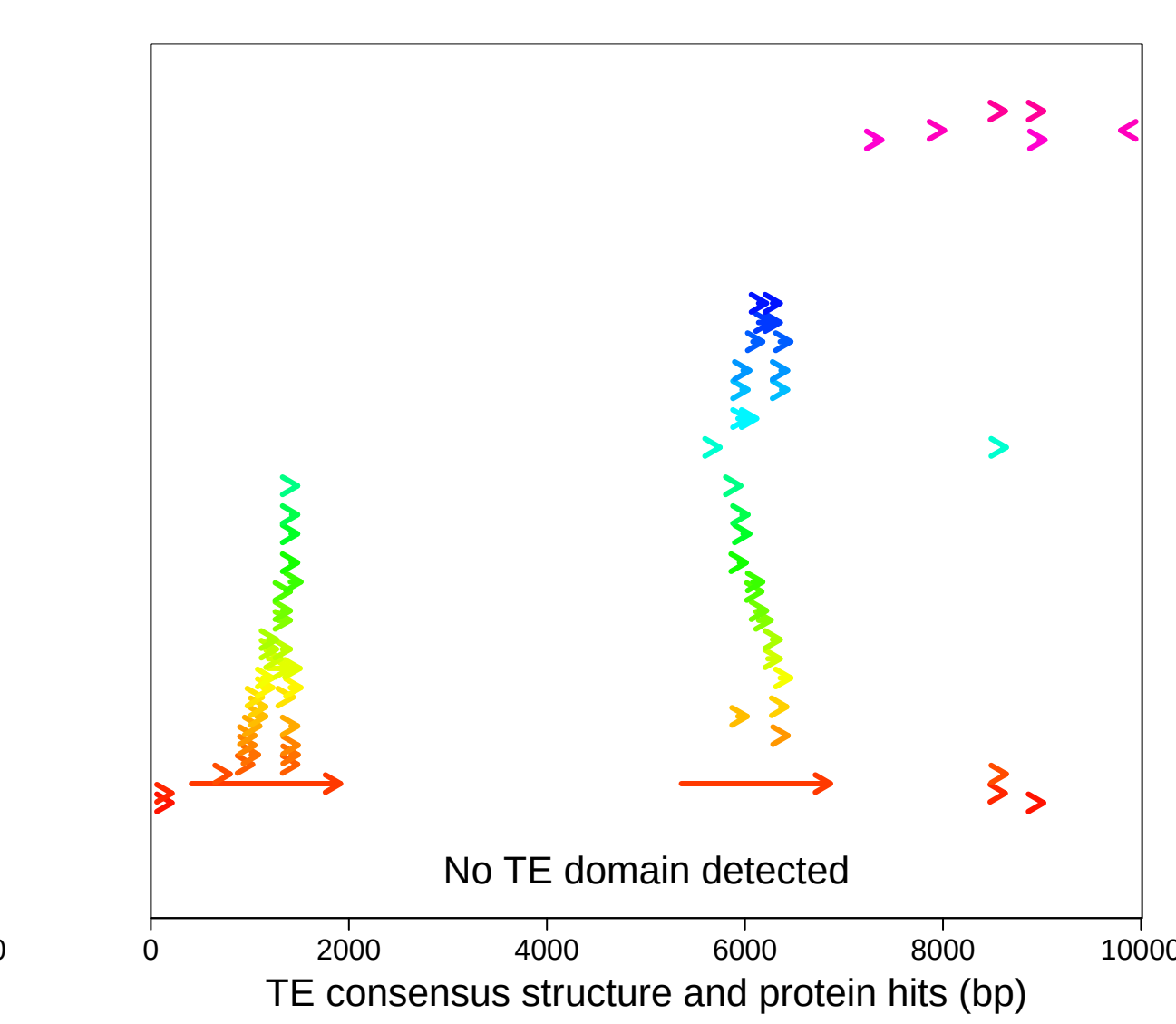
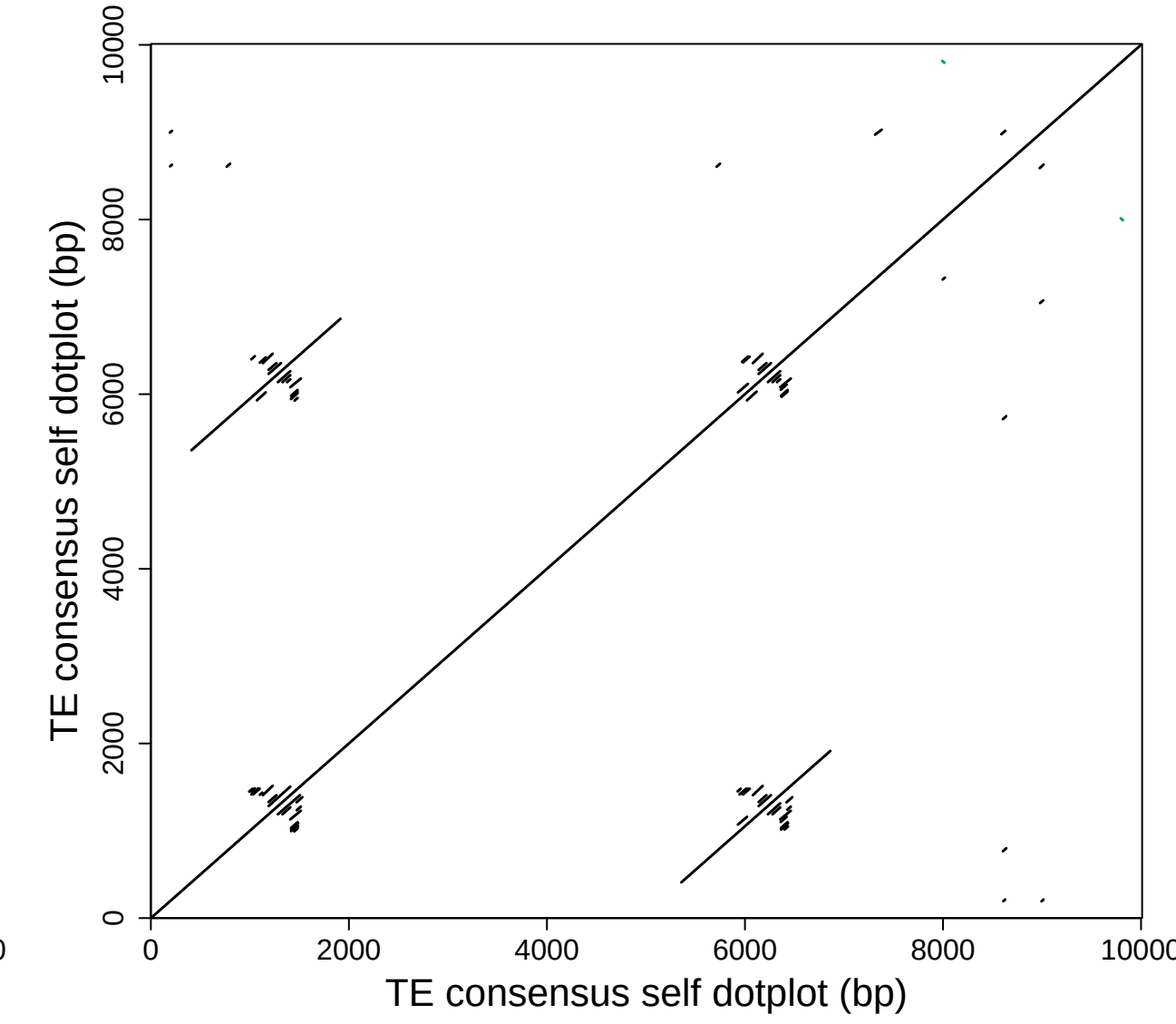
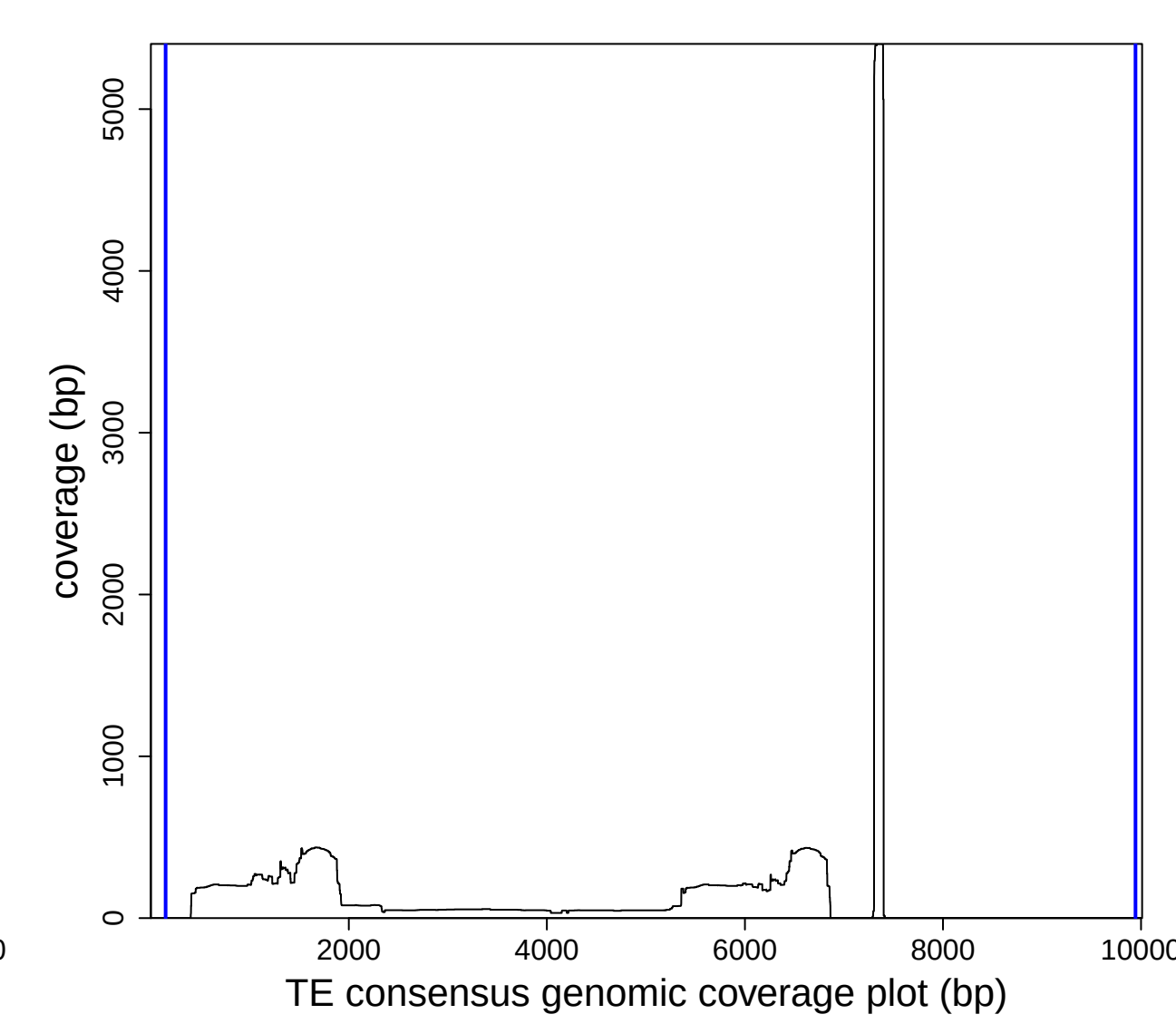
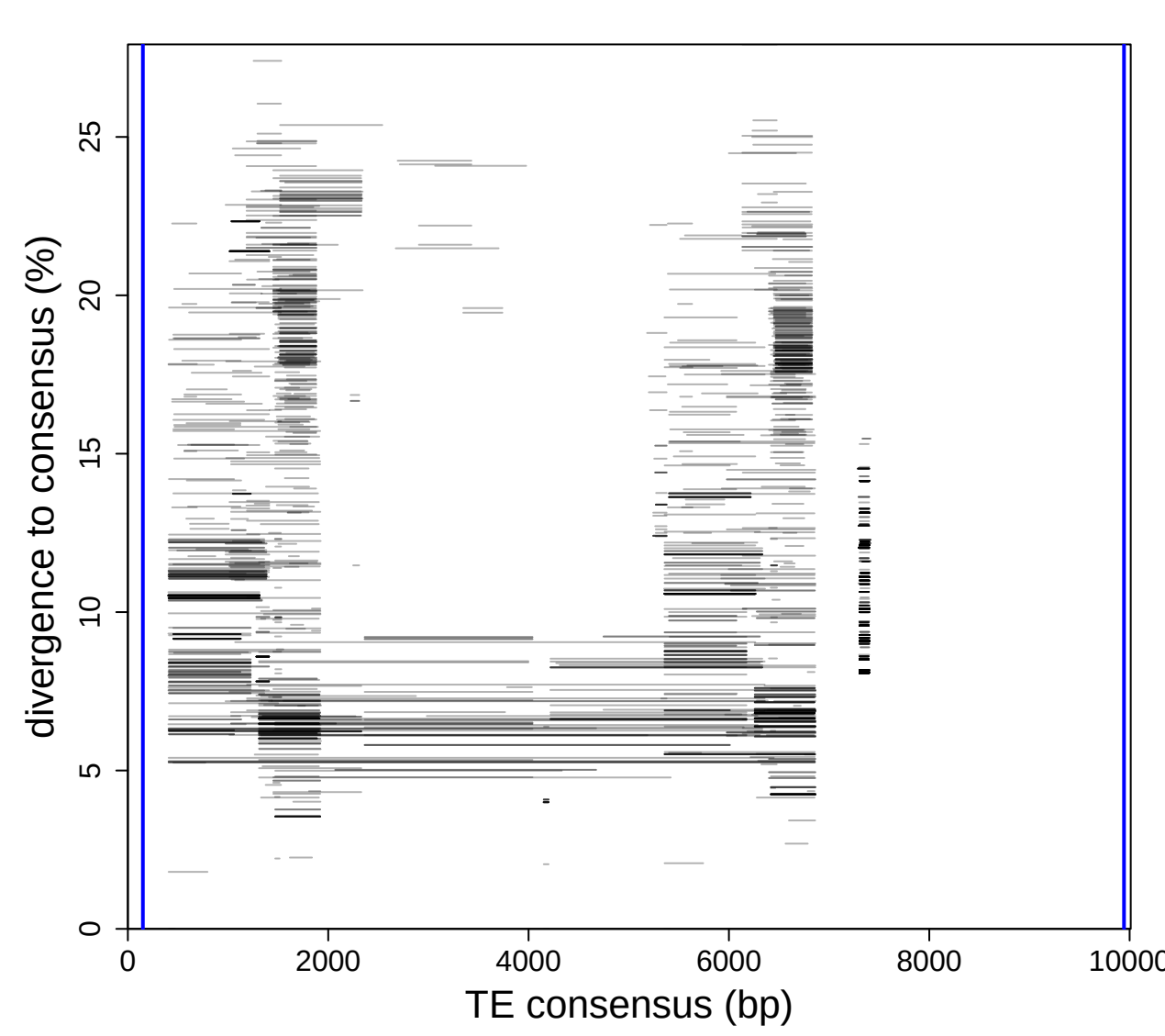
a.b.bed_uf.bed_g_2.bed_fm_1.bed_0_0_bcln.fa_aln.fa_cl.fa_gs.f
size: 8952bp; fragments: 21031; full length: 0 (>=8056.8bp)

After TEtrimmer 8952 bp



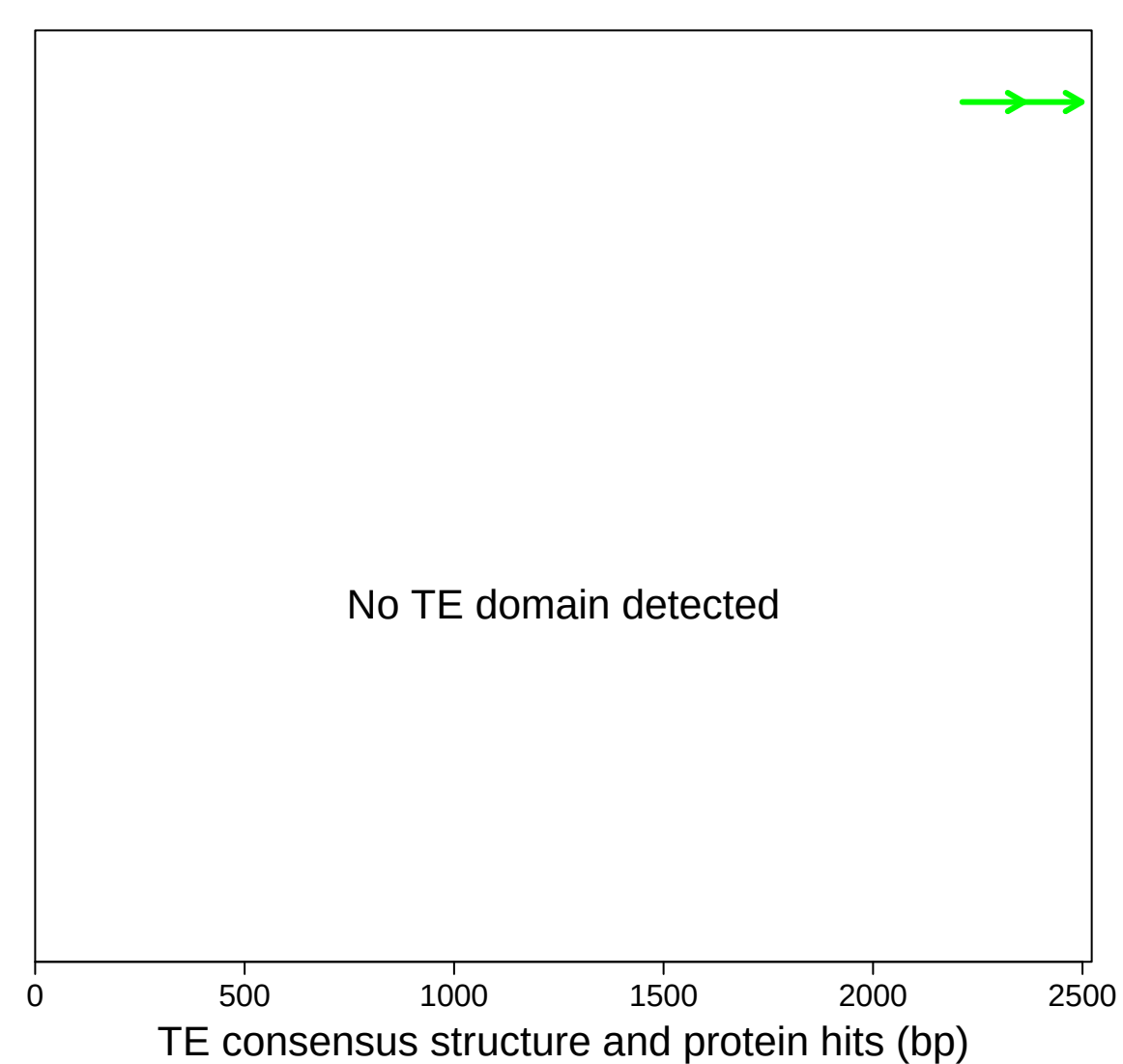
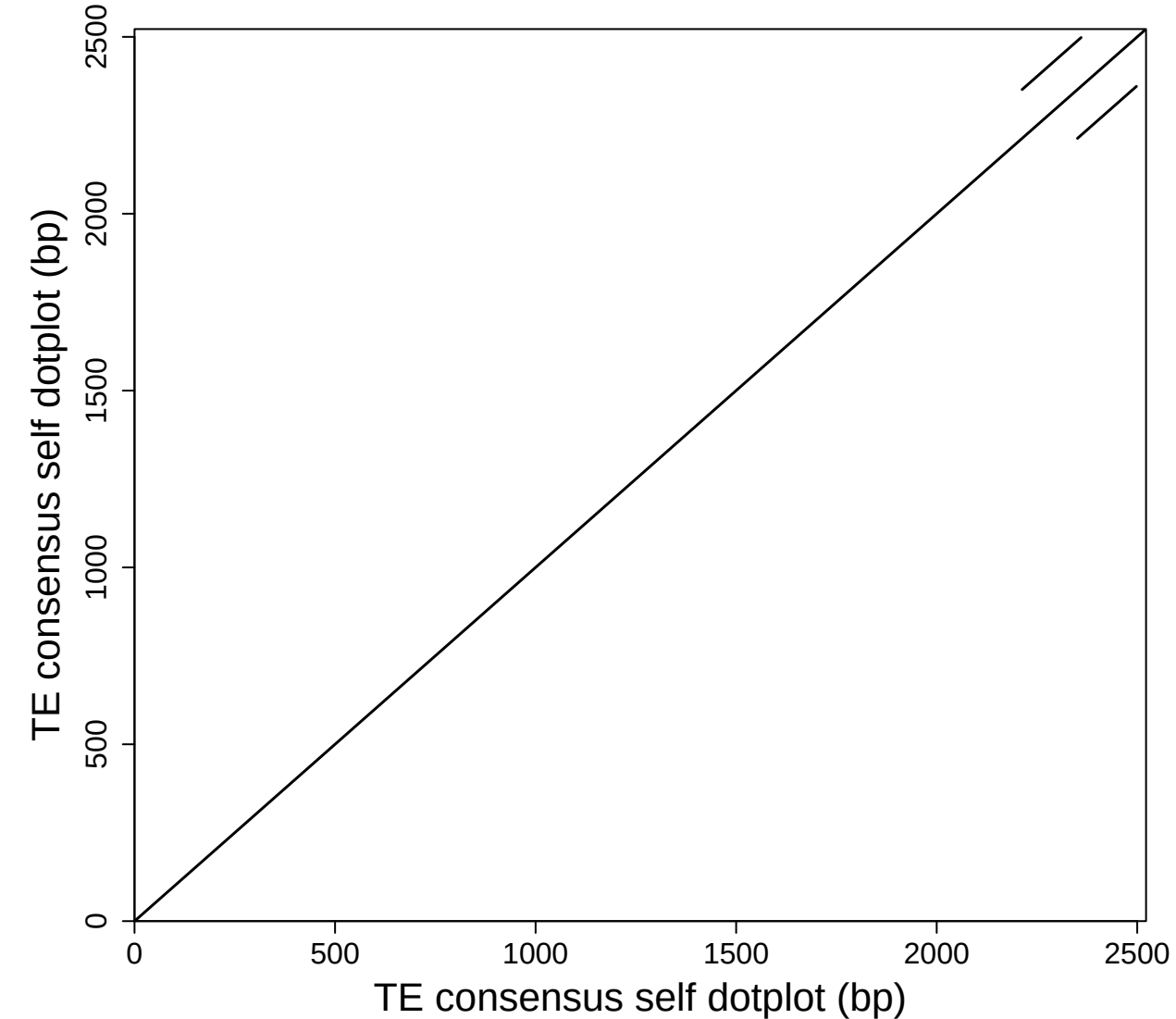
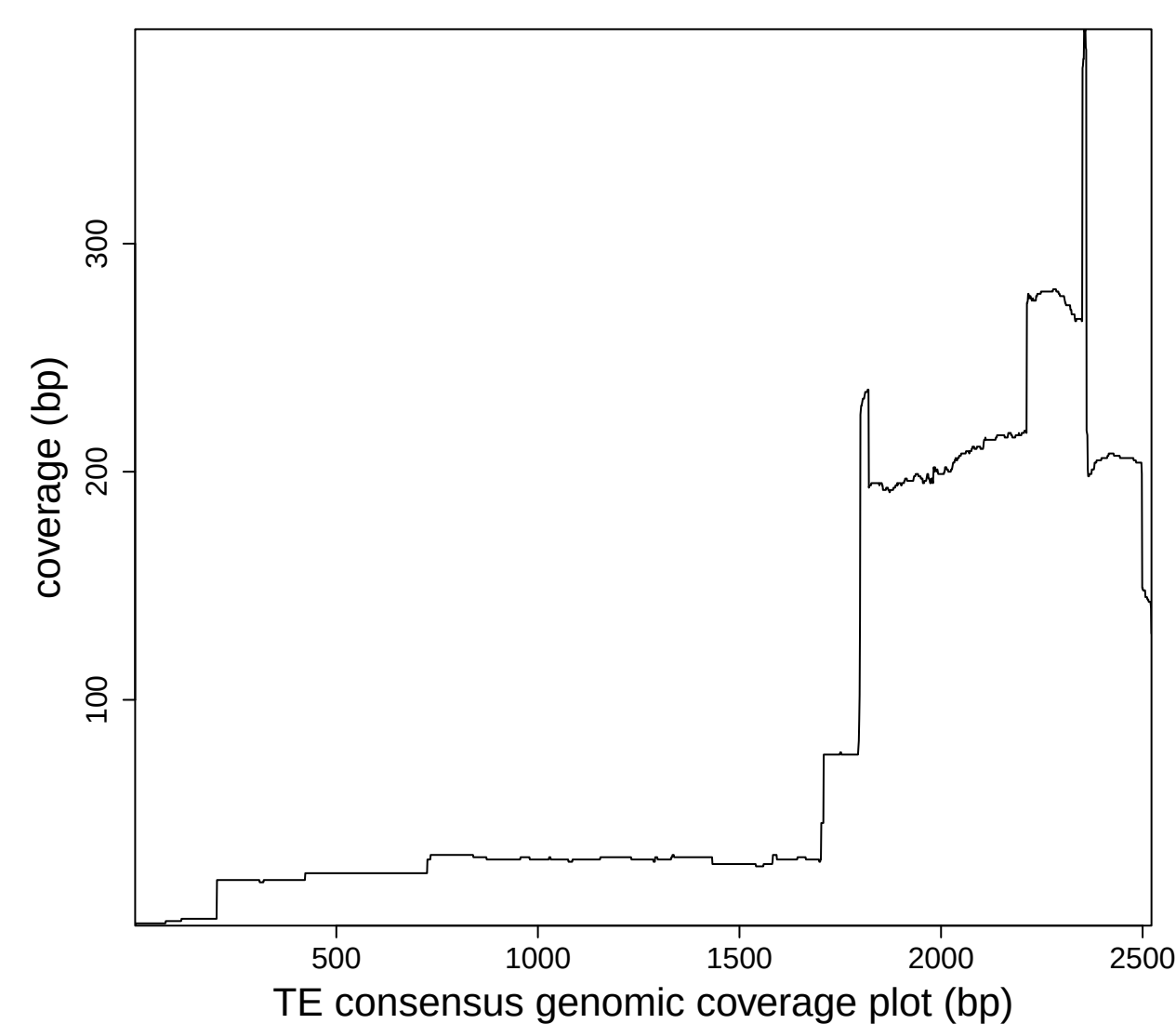
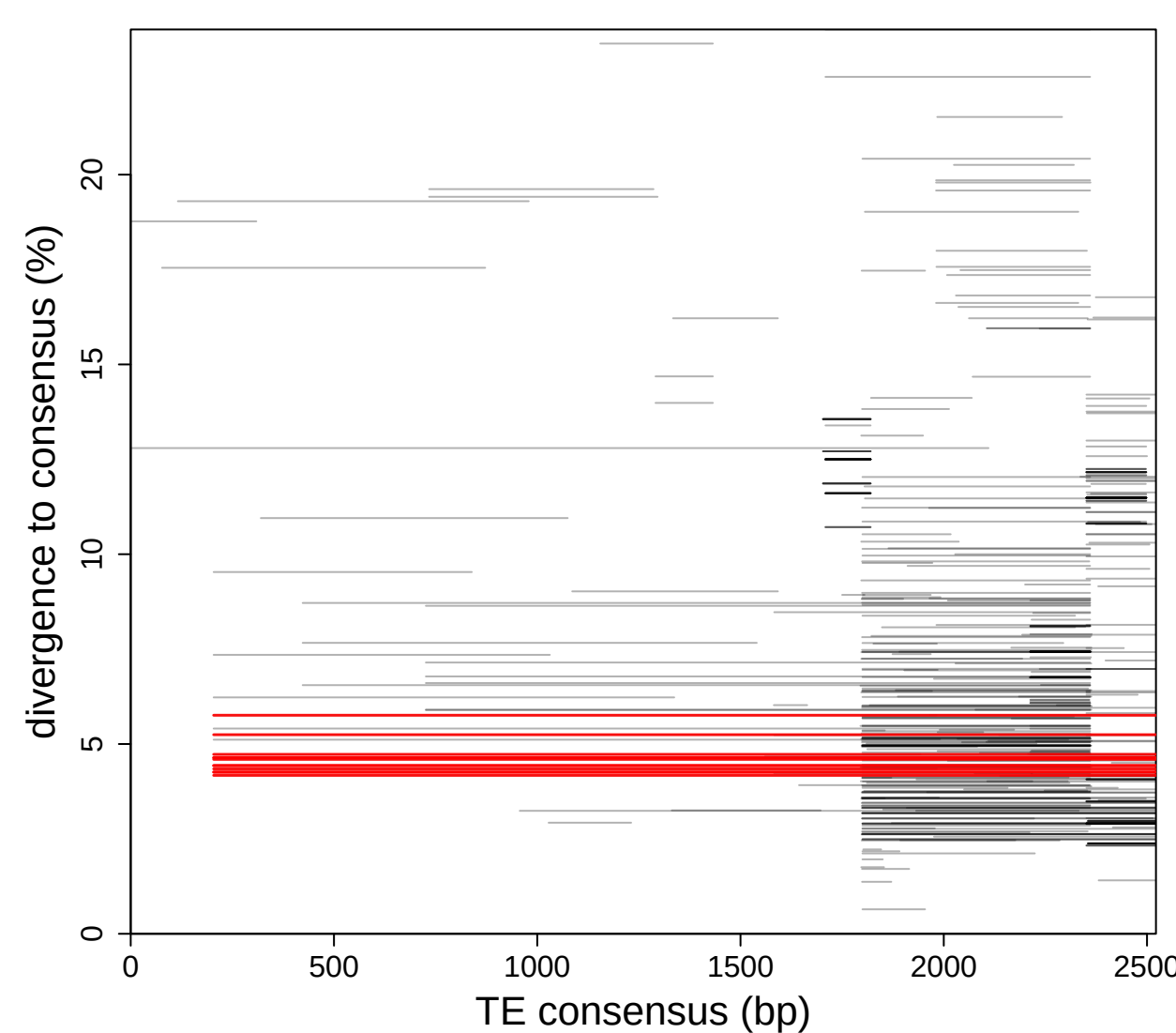
7.fasta.b.bed uf.bed g_2.bed fm_1.bed 0 0 bcln.fa aln.fa cl.f
size: 10011bp; fragments: 6866; full length: 0 (>=9009.9bp)

After TEtrimmer Extended plot Blue lines are boundaries

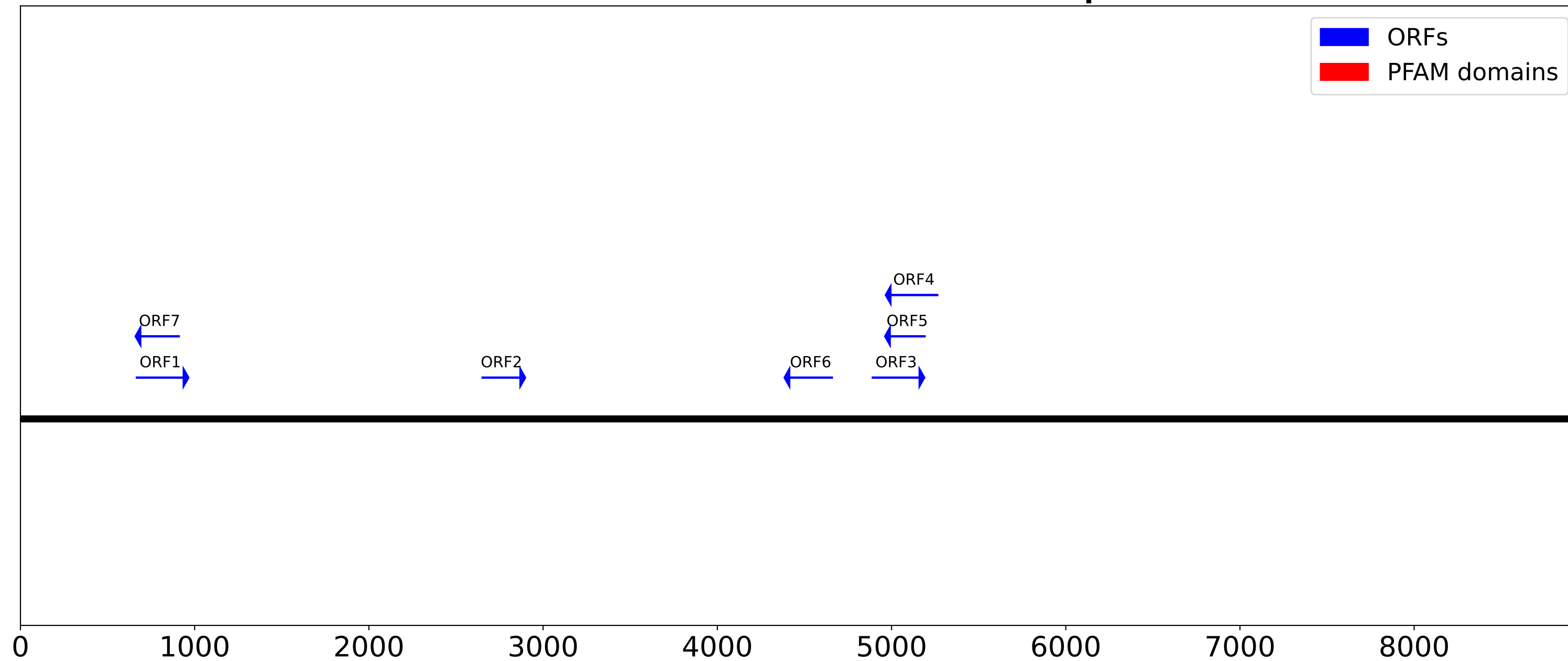


TE: rnd_5_family_6907
size: 2522bp; fragments: 535; full length: 12 (≥ 2269.8 bp)

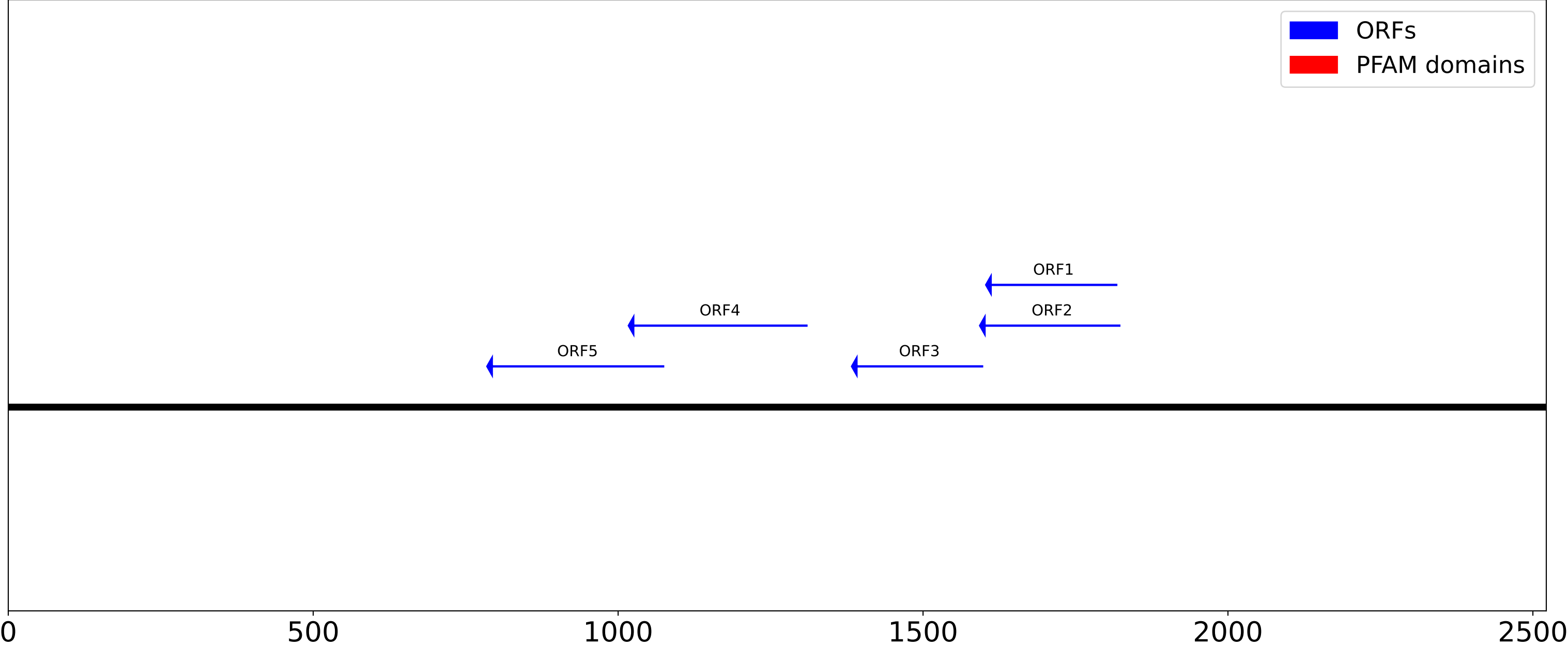
Before TEtrimmer 2522 bp



After TEtrimmer ORF and PFAM domain plot



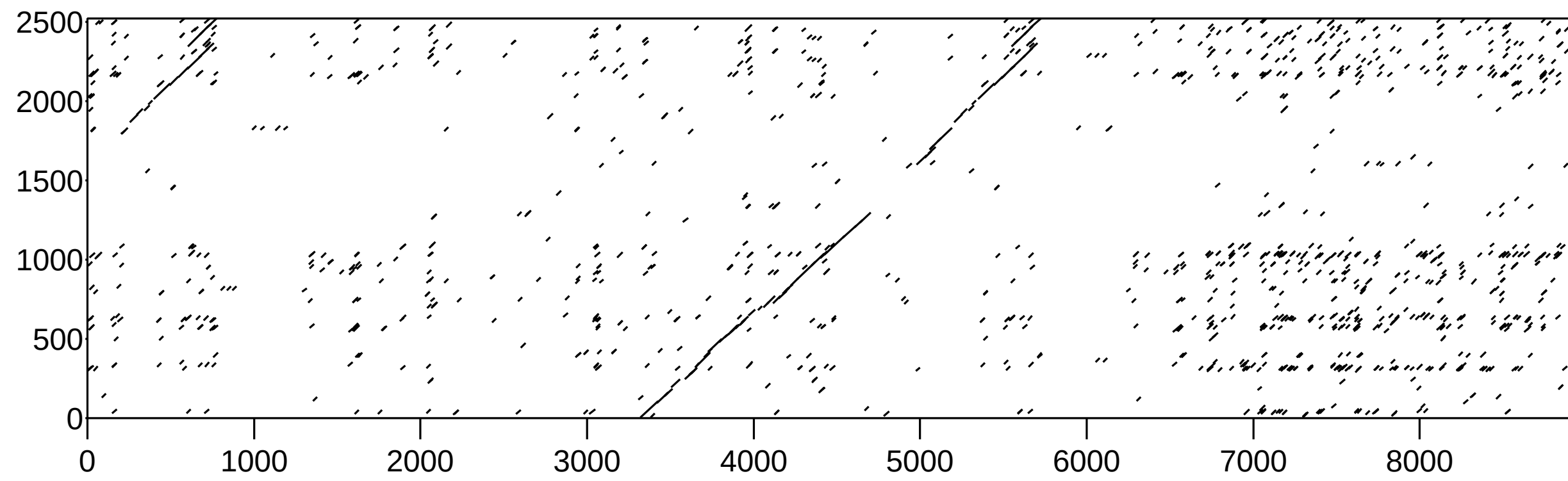
Before TEtrimmer ORF and PFAM domain plot



Dotplot

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

consensus before TEtrimmer (bp)



TE consensus after TEtrimmer (bp)